

# UC Riverside

## UC Riverside Electronic Theses and Dissertations

### Title

Systematic Research on the Millipede Assassin Bugs (Hemiptera: Reduviidae: Ectrichodiinae)

### Permalink

<https://escholarship.org/uc/item/01p4r9sc>

### Author

Forthman, Michael

### Publication Date

2016

### Copyright Information

This work is made available under the terms of a Creative Commons Attribution License, available at <https://creativecommons.org/licenses/by/4.0/>

Peer reviewed|Thesis/dissertation



UNIVERSITY OF CALIFORNIA  
RIVERSIDE

Systematic Research on the Millipede Assassin Bugs (Hemiptera: Reduviidae: Ectrichodiinae)

A Dissertation submitted in partial satisfaction  
of the requirements for the degree of

Doctor of Philosophy

in

Entomology

by

Michael Forthman

March 2016

Dissertation Committee:

Dr. Christiane Weirauch, Chairperson

Dr. John Heraty

Dr. Bradley Mullens

Dr. Mark Springer

Copyright by  
Michael Forthman  
2016

The Dissertation of Michael Forthman is approved:

---

---

---

---

Committee Chairperson

University of California, Riverside

## Acknowledgments

Attaining a doctorate in science is by no means an easy feat. I could not have achieved this dream had it not been for the support, encouragement, and love from so many people during my graduate career. First and foremost, I want to express my gratitude to my dissertation advisor, Christiane Weirauch, who offered substantial guidance with my research projects and professional development with grant proposals, scientific writing, presentations, and organizational involvement. I also want to thank all the members of my dissertation committee (John Heraty, Mark Springer, and Bradley Mullens), all of whom provided valuable support and guidance on my dissertation, as well as amazing encouragement, throughout my time at UCR. I also thank a good friend and collaborator, Dominik Chłond, who contributed to a massive monograph of Madagascan Ectrichodiinae.

There are many colleagues from the Weirauch and Heraty labs that I consider as family. All of them have provided a lot of perspective on grad school, project design, data analysis, the plethora of phylogenetic programs in existence, and life in general: Austin Baker, Lily Berniker, Roger Burks, Dimitri Forero, Sarah Frankenberg, Eric Gordon, John Hash, Judith Herreid, Rochelle Hoey-Chamberlain, Wei Song Hwang, Alexandr Knyshov, Sunghoon Jung, Stephanie Leon, Paul Masonick, Jason Mottern, Elizabeth Murray, Ryan Perry, Guanyang Zhang, and Junxia Zhang.

I would like to extend my appreciation to the following individuals who donated or loaned specimens for the purpose of this dissertation: Jean-Michel Berenger, Dominik Chłond, Eliane de Coninck, Jason Cryan, David Grimaldi, Eric Guilbert, Tom Henry, Michael Ivie, John Longino, Rudolf Meier, Joshua Oyelade, Norm Penny, Dávid Rédei, Ruth Salas, Toby Schuh, Michael Sharkey, David Wagner, Mick Webb, and the students and instructors of the 2010 Organization

for Tropical Studies field-course in Costa Rica. I want to thank Claudia Alvarez, Nick Duncan, Rochelle Hoey-Chamberlain, and Sarah Frankenberg for assisting me with specimen curation for a ridiculously large amount of material. Rochelle Hoey-Chamberlain also willingly assisted with some molecular benchwork for the last chapter. I extend my appreciation to Petra Sierwald for providing references on millipede taxonomy and to Wanzhi Cai for verifying the identifications of some Asian Ectrichodiinae. Ted Garland, Jr. was very helpful with running PLogReg.m.

I am lucky to have a very close, loving, and supportive group of friends and family. Unfortunately, many of my family members that supported my decision to get a Ph.D. in entomology are no longer here to see my achievements. Regardless, I owe everything I have and everything I am to living and deceased family members and all my friends. In particular, my late mother, Carol Forthman, always taught me to love who I am, pursue what I love, and always be humble and compassionate along the way. She sacrificed a lot in life for my education and career goals. I am also very appreciative of my great-uncle, Thomas Bruce, who passed away 11 days before my defense after a long fight with cancer. He was very encouraging and financially supportive in my endeavor to become a professional academic. Lastly, I must thank two very important people in my life, John Nelson and Steve Weber for their love, support in life, and sacrifices so that I could pursue my love of entomology and systematics. Both also put in a lot of effort on the final aspects of my dissertation: Steve literally made thousands of pie charts for the last chapter, and John helped to curate specimens.

This dissertation was supported by the National Science Foundation PEET DEB 093333853 grant (awarded to Christiane Weirauch); Entomological Society of America Systematic, Evolution, and Biodiversity Travel Award; Earle C. Anthony Graduate Student Travel Award; Graduate Dean's Dissertation Research Grant; Department of Entomology at the University of California – Riverside; and the University of California – Riverside Graduate Division

Dissertation Year Program Award. Some specimens loaned for this research were acquired under NSF DEB 0542864 (awarded to Michael Sharkey and Brian Brown), NSF BSI 0072713 (awarded to Brian Fisher and Charles Griswold), and NSF DEB 1354739 (awarded to John Longino).

The text of this dissertation, in part, is a modified reprint of the material as is appears in Forthman & Weirauch (2012; *European Journal of Entomology* 109: 147–153), Forthman et al. (in press; *Bulletin of the American Museum of Natural History*), and Forthman & Weirauch (in press; *Molecular Phylogenetics and Evolution*). The co-author Christiane Weirauch directed and supervised the research that forms the basis of this dissertation. The co-author Dominik Chłond contributed to the taxonomic descriptions for some of the new species, including images and drawings.

## **Dedication**

This dissertation is dedicated to  
all of my family, even those who rest eternally.

## ABSTRACT OF THE DISSERTATION

Systematic Research on the Millipede Assassin Bugs (Hemiptera: Reduviidae: Ectrichodiinae)

by

Michael Forthman

Doctor of Philosophy, Graduate Program in Entomology  
University of California, Riverside, March 2016  
Dr. Christiane Weirauch, Chairperson

The predominantly circumtropical millipede assassin bugs (Ectrichodiinae) comprise the fifth largest subfamily of Reduviidae with 736 described species in 121 genera. Ectrichodiinae appear to be specialized millipede predators and stand out among other Reduviidae in the high proportions of aposematically colored and sexually dimorphic species. These phenomena make Ectrichodiinae an attractive group for ecological, behavioral, and evolutionary research, but is hindered by a lack of thorough taxonomic treatments and phylogenetic hypotheses. This dissertation improves our knowledge of Ectrichodiinae life history, diversity, and phylogeny by: 1) reviewing Ectrichodiinae predatory behaviors and supplementing this with field observations; 2) revising the Madagascan fauna; 3) investigating historical biogeographic patterns of Madagascan Ectrichodiinae; and 4) constructing the most comprehensive Ectrichodiinae+ Tribelocephalinae phylogeny to study the evolution of aposematic coloration and sexual dimorphism. (1) About 2% of Ectrichodiinae species are documented to prey on juliform and, to a lesser extent, polydesmid millipedes. The majority of observations suggest that immatures typically exhibit communal predation, while adults usually engage in solitary predation. (2) The taxonomic monograph includes description of 63 new species and three new genera of endemic Madagascan Ectrichodiinae, as well as redescriptions and synonymies of previously described



taxa. (3) Madagascan Ectrichodiinae do not form a monophyletic group, with lineages exhibiting close relationships with Oriental or Afrotropical taxa. Ectrichodiinae colonized Madagascar in the last 68 my via transoceanic dispersal at least twice from the Oriental region and once from the Afrotropical region; a single out-of-Madagascar dispersal event to the Afrotropical region is recovered in some analyses. Dispersal between these regions were likely facilitated by islands and/or ocean currents. (4) Ectrichodiinae are not monophyletic with respect to Tribelocephalinae. Aposematic coloration evolved from a cryptic ancestor early in the evolutionary history of Ectrichodiinae+Tribelocephalinae, with subsequent reversals and regains. A positive association between the prevalence of aposematic coloration and body length is recovered when phylogenetic logistic regressions are conducted on ultrametric trees, but is not supported when phylograms are used. Extreme sexual dimorphism is shown to evolve multiple times from a more limited condition. Based on phylogenetic results, a new Ectrichodiinae classification is proposed.

## Table of Contents

Introduction .....	1
References .....	5
Chapter 1: Toxic associations: natural history observations and review of the predatory behaviors of millipede assassin bugs (Hemiptera: Reduviidae: Ectrichodiinae)	
Abstract .....	8
Introduction .....	8
Literature review and analysis of online image material .....	10
Field and laboratory procedures .....	11
Ectrichodiinae prey specificity .....	12
Point of stylet insertion .....	13
Communal predatory strategies .....	14
Conclusion .....	15
References .....	16
Figures .....	19
Tables .....	23
Chapter 2: Taxonomic monograph of the endemic millipede assassin bug fauna of Madagascar (Hemiptera: Reduviidae: Ectrichodiinae)	
Abstract .....	26
Introduction .....	27
Material and methods .....	30
Results of the molecular association of males and females .....	36
Taxonomy .....	37

Identification key to Ectrichodiinae genera of Madagascar .....	37
<i>Gibbosella</i> Chłond .....	39
Identification key to the males of species of <i>Gibbosella</i> .....	43
Identification key to the females of species of <i>Gibbosella</i> .....	45
<i>Gibbosella andasibe</i> , new species .....	46
<i>Gibbosella betampona</i> , new species .....	48
<i>Gibbosella brunalvus</i> , new species .....	50
<i>Gibbosella conisimilis</i> , new species .....	53
<i>Gibbosella elongata</i> Chłond .....	55
<i>Gibbosella fulva</i> , new species .....	58
<i>Gibbosella mantella</i> , new species .....	59
<i>Gibbosella megafrons</i> , new species .....	61
<i>Gibbosella mirabilis</i> Chłond .....	63
<i>Gibbosella nitida</i> , new species .....	64
<i>Gibbosella notoconica</i> , new species .....	65
<i>Gibbosella pallidacorium</i> , new species .....	68
<i>Gibbosella pallidalata</i> , new species .....	70
<i>Gibbosella planiscutum</i> , new species .....	73
<i>Gibbosella quadocris</i> , new species .....	77
<i>Gibbosella vangocris</i> , new species .....	80
<i>Glymmatophora</i> Stål .....	84
Identification key to the males and females of Madagascan species of	
<i>Glymmatophora</i> .....	85
<i>Glymmatophora (Glymmatophora) carolae</i> , new species .....	85

<i>Glymmatophora (Cyclosandalus) crassipes</i> Horváth .....	88
<i>Maraenaspis</i> Karsch .....	91
<i>Maraenaspis bidens</i> (Reuter) .....	92
<i>Marojejycoris</i> , new genus .....	95
Identification key to the males of species of <i>Marojejycoris</i> .....	97
<i>Marojejycoris auranticorium</i> , new species .....	98
<i>Marojejycoris brevifrons</i> , new species .....	100
<i>Marojejycoris francais</i> , new species .....	102
<i>Marojejycoris notadichroa</i> , new species .....	103
<i>Marojejycoris ranomafana</i> , new species .....	105
<i>Tanindrazanus</i> , new genus .....	107
Identification key to the males of species of <i>Tanindrazanus</i> .....	109
<i>Tanindrazanus amboasaricus</i> , new species .....	112
<i>Tanindrazanus andohahela</i> , new species .....	113
<i>Tanindrazanus anjozorobeus</i> , new species .....	115
<i>Tanindrazanus antananarivo</i> , new species .....	117
<i>Tanindrazanus bemaraha</i> , new species .....	119
<i>Tanindrazanus brunneus</i> , new species .....	120
<i>Tanindrazanus hannajagodae</i> , new species .....	122
<i>Tanindrazanus harinhali</i> , new species .....	123
<i>Tanindrazanus irwini</i> , new species .....	128
<i>Tanindrazanus joffrevillus</i> , new species .....	131
<i>Tanindrazanus kathrynae</i> , new species .....	132
<i>Tanindrazanus mahafaly</i> , new species .....	135

<i>Tanindrazanus marginatus</i> , new species .....	137
<i>Tanindrazanus marojejy</i> , new species .....	143
<i>Tanindrazanus nigripes</i> , new species .....	145
<i>Tanindrazanus notatus</i> , new species .....	147
<i>Tanindrazanus simulans</i> , new species .....	149
<i>Tanindrazanus tenebricus</i> , new species .....	150
<i>Tanindrazanus varicolor</i> , new species .....	152
<i>Tanindrazanus vohiparara</i> , new species .....	155
<i>Toliarus</i> , new genus .....	157
Identification key to the males of species of <i>Toliarus</i> .....	159
<i>Toliarus karinae</i> , new species .....	159
<i>Toliarus trichrous</i> , new species .....	160
<i>Toxopus</i> Bergroth .....	165
Identification key to the males of species of <i>Toxopus</i> .....	169
Identification key to the females of species of <i>Toxopus</i> .....	173
<i>Toxopus ambohitantely</i> , new species .....	173
<i>Toxopus ampitavananima</i> , new species .....	176
<i>Toxopus antsiranana</i> , new species .....	179
<i>Toxopus basalis</i> , new species .....	182
<i>Toxopus brucei</i> , new species .....	184
<i>Toxopus farafangana</i> , new species .....	187
<i>Toxopus fisheri</i> , new species .....	190
<i>Toxopus griswoldi</i> , new species .....	192
<i>Toxopus insignis</i> , new species .....	196

<i>Toxopus italaviana</i> , new species .....	199
<i>Toxopus melobrunneus</i> , new species .....	201
<i>Toxopus miandritsara</i> , new species .....	203
<i>Toxopus namoroka</i> , new species .....	205
<i>Toxopus pallidus</i> , new species .....	207
<i>Toxopus parkeri</i> , new species .....	209
<i>Toxopus politus</i> Bergroth .....	212
<i>Toxopus signoretii</i> (Reuter), new combination .....	212
<i>Toxopus simulans</i> , new species .....	215
<i>Toxopus steineri</i> , new species .....	217
<i>Toxopus tibialis</i> , new species .....	218
<i>Toxopus toamasina</i> , new species .....	221
<i>Toxopus toliara</i> , new species .....	223
<i>Toxopus vazimba</i> , new species .....	229
Conclusion .....	230
References .....	232
Figures .....	238
Maps .....	286
Tables .....	312

### Chapter 3: Phylogenetics and biogeography of the endemic Madagascan millipede assassin bugs

(Hemiptera: Reduviidae: Ectrichiinae)

Abstract .....	328
Introduction .....	329
Material and methods .....	332

Results .....	339
Discussion .....	344
Conclusion .....	350
References .....	352
Figures .....	360
Tables .....	368
Supplementary material .....	378
Chapter 4: Millipede assassin bugs (Heteroptera: Reduviidae: Ectrichodiinae) show off: evolution of aposematic coloration and extreme sexual dimorphism	
Abstract .....	471
Introduction .....	472
Material and methods .....	478
Results .....	487
Discussion .....	493
Taxonomy .....	498
Conclusion .....	507
References .....	510
Figures .....	518
Tables .....	530
Supplementary material .....	546
Conclusion .....	675

## List of Figures

Figure 1.1. Internet images used for analysis of Ectrichodiinae predatory behavior .....	19
Figure 1.2. Field and laboratory images of <i>Rhiginia cinctiventris</i> predation on millipedes at La Selva Biological Station, Costa Rica .....	21
Figure 2.1. Dorsal habitus of <i>Gibbosella</i> species, <i>Glymmatophora carolae</i> , and <i>Glymmatophora crassipes</i> .....	238
Figure 2.2. Dorsal habitus of <i>Glymmatophora crassipes</i> , <i>Maraenaspis bidens</i> , <i>Marojejycoris</i> species, and <i>Tanindrazanus</i> species ( <i>T. amboasaricus</i> – <i>T. kathrynae</i> ) .....	240
Figure 2.3. Dorsal habitus of <i>Tanindrazanus</i> species ( <i>T. mahafaly</i> – <i>T. vohiparara</i> ), <i>Toliarus</i> species, and <i>Toxopus</i> species ( <i>T. ambohitantely</i> – <i>T. griswoldi</i> [male]) .....	242
Figure 2.4. Dorsal habitus of <i>Toxopus</i> species ( <i>T. griswoldi</i> [female] – <i>T. vazimba</i> ) .....	244
Figure 2.5. Lateral habitus of <i>Gibbosella</i> species, <i>Glymmatophora</i> species, <i>Maraenaspis bidens</i> , and <i>Marojejycoris</i> species ( <i>M. auranticorium</i> – <i>M. francais</i> ) .....	246
Figure 2.6. Lateral habitus of <i>Marojejycoris notadichroa</i> , <i>Marojejycoris francais</i> , and <i>Tanindrazanus</i> species ( <i>T. amboasaricus</i> – <i>T. varicolor</i> ) .....	248
Figure 2.7. Lateral habitus of <i>Tanindrazanus varicolor</i> , <i>Tanindrazanus vohiparara</i> , <i>Toliarus</i> species, and <i>Toxopus</i> species .....	250
Figure 2.8. Ventral habitus of <i>Gibbosella</i> species, <i>Glymmatophora</i> species, and <i>Maraenaspis bidens</i> .....	252
Figure 2.9. Ventral habitus of <i>Marojejycoris</i> species and <i>Tanindrazanus</i> species ( <i>T. amboasaricus</i> – <i>T. nigripes</i> ) .....	254
Figure 2.10. Ventral habitus of <i>Tanindrazanus</i> species ( <i>T. notatus</i> – <i>T. vohiparara</i> ), <i>Toliarus trichrous</i> , and <i>Toxopus</i> species ( <i>T. ambohitantely</i> – <i>T. melobrunneus</i> ) .....	256



Figure 2.11. Ventral habitus of <i>Toxopus</i> species ( <i>T. miandritsara</i> – <i>T. vazimba</i> ) .....	258
Figure 2.12. Dorsal and lateral habitus of holotypes of <i>Gibbosella elongata</i> , <i>Gi. mirabilis</i> , <i>Glymmatophora crassipes</i> , <i>Synavecoris dimorphus</i> , and <i>Toxopus signoretii</i> .....	260
Figure 2.13. Dorsal and lateral head structures .....	262
Figure 2.14. Lateral antennal morphology of <i>Toxopus insignis</i> .....	264
Figure 2.15. Dorsal, lateral, and ventral thoracic structures .....	266
Figure 2.16. Metathoracic gland evaporatorium and femoral armature .....	268
Figure 2.17. Wing morphology .....	270
Figure 2.18. Lateral and ventral abdominal morphology .....	272
Figure 2.19. Pygophore morphology in caudal view .....	274
Figure 2.20. Pygophore morphology in lateral view .....	276
Figure 2.21. Phallus morphology in dorsal view .....	278
Figure 2.22. Phallus morphology in lateral view .....	280
Figure 2.23. Female external and internal genitalic morphology .....	282
Figure 2.24. Dorsal habitus of the eight females that cannot be associated with male-based species using molecular, morphological, and geographic data .....	284
Figure 3.1. Examples and criteria of limited and extreme sexual dimorphism in Ectrichodiinae .....	360
Figure 3.2. Best tree based on maximum likelihood (ML) analysis of 56 taxa, four G-INS-I aligned gene partitions, and GTR+ $\Gamma$ +I model of sequence evolution .....	362
Figure 3.3. Strict consensus tree from 938 parsimonious trees from a TNT analysis of 110 taxa and a combined morphological and molecular dataset .....	364
Figure 3.4. Ancestral range reconstructions from DEC and DEC+j models for select nodes .....	366

Figure 4.1. Examples of limited and extreme sexual dimorphism in Ectrichodiinae species .....	518
Figure 4.2. ML best tree based on 156 taxa and a combined morphological and molecular dataset .....	520
Figure 4.3. BayesTraits Multistate ML ancestral state reconstruction of color character 198 on the ML phylogram for Tribelocephalinae and Ectrichodiinae taxa .....	522
Figure 4.4. BayesTraits MultiState ML reconstruction of ancestral sexual dimorphism states on the ML phylogram for Tribelocephalinae and Ectrichodiinae taxa .....	524
Figure 4.5. Combined morphological and molecular ML phylogram showing new classification of Ectrichodiinae and unambiguous character optimizations .....	526

## List of Maps

Map 2.1. Localities of <i>Gibbosella brunalvus</i> , <i>G. conisimilis</i> , <i>G. megafrons</i> , <i>G. mirabilis</i> , <i>G. pallidacorium</i> , and <i>G. pallidalata</i> .....	286
Map 2.2. Localities of <i>Gibbosella andasibe</i> , <i>G. elongata</i> , <i>G. fulva</i> , <i>G. nitida</i> , <i>G. notoconica</i> and <i>G. quadocris</i> .....	288
Map 2.3. Localities of <i>Gibbosella betampona</i> , <i>G. mantella</i> , <i>G. planiscutum</i> , and <i>G. vangocris</i> .....	290
Map 2.4. Localities of <i>Glymmatophora</i> species and <i>Maraenaspis bidens</i> .....	292
Map 2.5. Localities of <i>Marojejycoris</i> species .....	294
Map 2.6. Localities of <i>Tanindrazanus antananarivo</i> , <i>T. hannajagodae</i> , <i>T. harinhali</i> , <i>T. joffrevillus</i> , <i>T. marojejy</i> , and <i>T. notatus</i> .....	296
Map 2.7. Localities of <i>Tanindrazanus andohahela</i> , <i>T. bemaraha</i> , <i>T. irwini</i> , <i>T. kathrynae</i> , <i>T. mahafaly</i> , and <i>T. varicolor</i> .....	298
Map 2.8. Localities of <i>Tanindrazanus brunneus</i> , <i>T. marginatus</i> , <i>T. nigripes</i> , <i>T. tenebricus</i> , and <i>T. vohiparara</i> .....	300
Map 2.9. Localities of <i>Tanindrazanus amboasaricus</i> , <i>Ta. anjzorobeus</i> , and <i>Toliarus (To.)</i> species .....	302
Map 2.10. Localities of <i>Toxopus antsiranana</i> , <i>T. griswoldi</i> , <i>T. miandritsara</i> , <i>T. toamasina</i> , <i>T. toliara</i> , and <i>T. vazimba</i> .....	304
Map 2.11. Localities of <i>Toxopus ambohitantely</i> , <i>T. basalis</i> , <i>T. farafangana</i> , <i>T. melobrunneus</i> , <i>T. parkeri</i> , and <i>T. tibialis</i> .....	306
Map 2.12. Localities of <i>Toxopus ampitavananima</i> , <i>T. brucei</i> , <i>T. fisheri</i> , <i>T. italaviana</i> , <i>T. pallidus</i> , and <i>T. steineri</i> .....	308

Map 2.13. Localities of *Toxopus insignis*, *T. namoroka*, *T. politus*, *T. signoretii*, and *T. simulans*

..... 310

## List of Tables

Table 1.1. Data extracted from published literature and Internet images for analysis of Ectrichodiinae predatory behavior .....	23
Table 2.1. Measurements (in millimeters) .....	312
Table 2.2. GenBank accession numbers of COI sequences .....	326
Table 3.1. Taxon sampling, unique specimen identifier (USI) codes, ethanol molecular specimen collection numbers, depositories, and GenBank accession numbers for sequenced specimens .....	368
Table 3.2. Morphological characters and character state codings .....	371
Table 4.1. Morphological characters and character state codings .....	530
Table 4.2. Taxon sampling, USI codes, ethanol molecular specimen collection numbers, depositories, and GenBank accession numbers for sequenced specimens .....	540
Table 4.3. Phylogenetic logistic regression parameter estimates for the effects of log-transformed male body length on color character 198 in 133 Ectrichodiinae and Tribelocephalinae species .....	545

## List of Supplementary Material

Supplementary material 3.1. Morphological character matrix in nexus format for cladistic analysis .....	378
Supplementary material 3.2. Molecular character matrix in nexus format for cladistic analysis .....	386
Supplementary material 3.3. Unambiguous character optimizations on the strict consensus tree from parsimony analysis of 110 taxa and a combined morphological and molecular dataset .....	450
Supplementary material 3.4. Fossil information and calibration nodes, means, standard deviations or sigma values, and offsets (in real space) .....	454
Supplementary material 3.5. DEC and DEC+j time-stratified dispersal rate matrix .....	455
Supplementary material 3.6. Equal and implied weights molecular parsimony results for 56 taxa .....	456
Supplementary material 3.7. Best tree based on maximum likelihood (ML) analysis of 110 taxa, gene partitions and morphology .....	458
Supplementary material 3.8. Divergence time estimates based on a BEAST analysis using the ML best tree .....	460
Supplementary material 3.9. Ancestral range reconstructions from the DIVA model for select nodes .....	462
Supplementary material 3.10. Discussion of morphological support for major lineages recovered in the combined morphological and molecular phylogenetic analysis .....	464
Supplementary material 4.1. Twenty-eight female-based terminal taxa, with sources of male morphological data .....	546

Supplementary material 4.2. Morphological character matrix in nexus format for cladistic analysis .....	547
Supplementary material 4.3. Polymerase chain reaction protocols for each targeted gene region .....	565
Supplementary material 4.4. Molecular character matrix in nexus format for cladistic analysis .....	566
Supplementary material 4.5. Molecular parsimony, ML, and Bayesian phylogenetic analyses of 59 taxa .....	651
Supplementary material 4.6. Strict consensus tree from 269 parsimonious trees based on TNT analysis of 156 taxa and morphological data .....	653
Supplementary Material 4.7. Morphological ML and Bayesian phylogenetic analyses of 156 taxa .....	655
Supplementary material 4.8. Combined morphological parsimony phylogenetic analyses of 156 taxa using equal and implied weights .....	657
Supplementary material 4.9. Combined morphological parsimony phylogenetic analyses of 156 taxa using implied weights .....	659
Supplementary Material 4.10. Bayesian Markov Chain Monte Carlo (MCMC) majority-rule consensus tree based on of 156 taxa .....	661
Supplementary material 4.11. Parsimony ancestral state reconstruction on the ML phylogram for Tribelocephalinae and Ectrichodiinae taxa and color character 198 .....	663
Supplementary material 4.12. Figshare data repository link to BayesTraits MultiState ML and MCMC ancestral state reconstruction output for color character 198 on ML phylogram and ultrametric trees .....	665

Supplementary material 4.13. Parsimony ancestral state reconstruction on the ML phylogram (Fig. 4.2) for Tribelocephalinae and Ectrichodiinae taxa (tree converted to cladogram for visual) and color character 199 .....	666
Supplementary material 4.14. BayesTraits MultiState ML ancestral state reconstruction of color character 199 on the ML phylogram for Tribelocephalinae and Ectrichodiinae taxa .....	668
Supplementary material 4.15. Figshare data repository link to BayesTraits MultiState ML and MCMC ancestral state reconstruction output for color character 199 on ML phylogram and ultrametric trees .....	670
Supplementary material 4.16. Phylogenetic logistic regression parameter estimates for the effects of log-transformed male body length on coloration in 133 Ectrichodiinae and Tribelocephalinae species using color character 199 .....	671
Supplementary material 4.17. Parsimony ancestral sexual dimorphism reconstruction on the ML phylogram for Tribelocephalinae and Ectrichodiinae taxa .....	672
Supplementary material 4.18. Figshare data repository link to BayesTraits MultiState ML and MCMC ancestral state reconstruction output for sexual dimorphism character 200 on ML phylogram and ultrametric trees .....	674



## **Introduction**

Assassin bugs, or Reduviidae, are the largest hemimetabolous clade of predatory insects (>7,000 spp.) and the second most speciose family of Heteroptera (Hemiptera). While most species are generalist predators, there is a striking range of prey specialization in several subfamilies (e.g., termite-feeding Salyavatinae [Gordon & Weirauch, 2016]; araneophagic Emesinae [Wignall & Taylor, 2011]). Reduviidae also include the notorious vertebrate-blood feeding Triatominae that vector Chagas disease (Lent & Wygodzinsky, 1979). Species in some clades have evolved morphological adaptations for prey-capture, e.g., sticky-trap predation (Harpactorinae; Weirauch, 2006; Zhang & Weirauch, 2013), subchelate and chelate forelegs (Phymatinae; Weirauch et al., 2011), and tibial attachment structures in several subfamilies (Weirauch, 2007). A number of studies have advanced our understanding of Reduviidae phylogenetics (e.g., Weirauch, 2008; Weirauch & Munro, 2009; Hwang & Weirauch, 2012; Zhang et al., 2016), but the relationships and monophyly of several subfamilies remain uncertain.

The predominantly circumtropical millipede assassin bugs, or Ectrichodiinae, comprise the fifth largest Reduviidae subfamily (736 described species, 121 genera; Gil-Santana, 2015; Forthman et al., in press and references within). Ectrichodiinae are the most evolutionarily successful, in terms of extant diversity, clade of millipede specialists, a group of prey organisms few arthropods exploit (Forthman & Weirauch, 2012). They further stand out among other Reduviidae in the high proportions of species with aposematic coloration and metallic features. A number of species also exhibit drastic sexual dimorphism, which poses significant difficulty in associating sexes when using morphology alone. These phenomena make Ectrichodiinae an attractive group for ecological, behavioral, and evolutionary research, which is hindered by a lack of thorough taxonomic treatments and phylogenetic hypotheses. Prior to this dissertation, relationships between genera had not been cladistically tested beyond a genus-level

phylogeny of New World taxa (Dougherty, 1995). Faunistic checklists and descriptions of taxa based on few specimens of a single sex predominate the taxonomic literature, which is characterized by a lack of illustrations, identification keys, and synthetic revisions. Few Ectrichodiinae have been incorporated in recent Reduviidae phylogenetic analyses based on morphological or molecular datasets (Weirauch, 2008; Weirauch & Munro, 2009; Weirauch, 2010; Hwang & Weirauch, 2012; Zhang et al., 2016). Based on these studies, Ectrichodiinae monophyly is not firmly established and remains to be investigated using morphological and molecular data for a larger sample of taxa. My dissertation research integrates natural history, revisionary taxonomy, and morphological and molecular phylogenetics to improve our knowledge of Reduviidae phylogenetics, specifically the subfamily Ectrichodiinae.

**Chapter 1: Predatory behaviors of Ectrichodiinae:** Several species are known to feed on chemically defended millipedes (e.g., Green, 1925; Haridass & Ananthakrishnan, 1980; Haridass, 1985). However, details on the ectrichodiine-millipede relationship are scarce, and documentation is mostly based on anecdotal evidence. To investigate predatory behaviors in Ectrichodiinae, natural history observations in the field and laboratory are made and supplemented with published literature and online images to investigate prey specificity, communal predation, and insertion of mouthparts into millipedes.

**Chapter 2: Taxonomic monograph of the Madagascan Ectrichodiinae:** Madagascar is one of the world's premier biodiversity hotspots (Myers et al., 2000), and the island's terrestrial arthropods are in danger of extinction due to anthropogenic influences (Sussman et al., 1996; Myers et al., 2000). An impediment to identifying areas for conservation is the incomplete knowledge of the island's pattern of species richness, turnover, and endemism (Schatz, 2002; Smith et al., 2005). The California Academy of Sciences' Terrestrial Arthropod Inventory of Madagascar Project addressed the need for taxonomic exploration of the arthropod fauna and has

yielded >2,000 Ectrichodiinae specimens, revealing a dramatic undescribed species-level diversity. Thus, the Madagascan Ectrichodiinae is documented and revised.

**Chapter 3: Biogeography of the Madagascan Ectrichodiinae:** Despite Madagascar's proximity to Africa, a large proportion of the known biota is closely related to Oriental lineages (Vences, 2004; Yoder & Nowak, 2006). Although Gondwanan vicariance is thought to be responsible for the presence of some Madagascan lineages, most colonized the island via transoceanic dispersal after the island separated from Africa (~160 mya) and India (80–90 mya) (Vences 2004; Yoder & Nowak, 2006). A recent divergence dating analysis of Reduviidae suggests that Ectrichodiinae arose after the Madagascar-India split (Hwang & Weirauch, 2012). Thus, dispersal is likely responsible for the endemic Madagascan Ectrichodiinae fauna, but the origin and timing of colonization remains unknown. Morphological and molecular phylogenies are constructed and used for divergence dating and biogeographic analyses to investigate geographic and oceanic factors that may have facilitated long-distance dispersal between Madagascar and other regions.

**Evolution of aposematic coloration and sexual dimorphism:** A few comparative phylogenetic studies support a positive correlation between aposematic coloration and body size (e.g., Hagman & Forsman, 2003; Rudh, 2013), while others support a negative or no correlation (Nilsson & Forsman, 2003; Cheney et al., 2014). Few insect phylogenetic studies investigating the relationship between these two traits are restricted to the Lepidoptera (Tullberg & Hunter, 1996; Nilsson & Forsman, 2003), but larval gregariousness may confound results (Nilsson & Forsman, 2003). Ectrichodiinae exhibit a diversity of color patterns across all body sizes, are generally solitary, and are thus an excellent model for investigating the coloration-body size relationship. Some species also exhibit drastic sexual dimorphism with major changes in female morphology, and this remains to be explored in a phylogenetic context. A comprehensive

phylogeny is constructed for Ectrichodiinae and Tribelocephalinae using morphology and molecular data, ancestral state reconstructions on coloration and sexual dimorphism are performed, and a phylogenetic comparative method is employed to test for a coloration-body size association. Phylogenetic results are used to revise the Ectrichodiinae classification.

## References

- Cheney, K.L., Cortesi, F., How., M.J., Wilson, N.G., Blomberg, S.P., Winters, A.E., Umanzör, S., Marshall, N.J. 2014. Conspicuous visual signals do not coevolve with increased body size in marine sea slugs. *Journal of Evolutionary Biology* 27: 676–687.
- Dougherty, V. 1995. A review of the New World Ectrichodiinae genera (Hemiptera: Reduviidae). *Transactions of the American Entomological Society* 121: 173–225.
- Forthman, M., Weirauch, C. 2012. Toxic associations: a review of the predatory behaviors of millipede assassin bugs (Hemiptera: Reduviidae: Ectrichodiinae). *European Journal of Entomology* 109: 147–153.
- Forthman, M., Chłond, D., Weirauch, C. (in press). Taxonomic monograph of the endemic millipede assassin bug fauna of Madagascar (Hemiptera: Reduviidae: Ectrichodiinae). *Bulletin of the American Museum of Natural History*.
- Gil-Santana, H.R. 2015. First record of the genus *Pseudopothea* from South America, with description of a new species from Brazil (Hemiptera: Heteroptera: Reduviidae: Ectrichodiinae). *Zootaxa* 3904: 541–552.
- Gordon, E.R.L., Weirauch, C. 2016. Efficient capture of natural history data reveals prey conservatism of cryptic termite predators. *Molecular Phylogenetics and Evolution* 94: 65–73.
- Green, W.E. 1925. The President's address. *Proceedings of the Entomological Society of London* 1924: clxi–ccii.
- Hagman, M., Forsman, A. 2003. Correlated evolution of conspicuous coloration and body size in poison frogs (Dendrobatidae). *Evolution* 57: 2904–2910.
- Haridass, E.T. 1985. Feeding and ovipositional behavior in some reduviids (Insecta – Heteroptera). *Proceedings of the Indian Academy of Sciences, Animal Sciences* 94: 239–247.
- Haridass, E.T., Ananthkrishnan, T.N. 1980. Models for the predatory behavior of some reduviids from southern India (Insecta – Heteroptera – Reduviidae). *Proceedings of the Indian Academy of Sciences, Animal Sciences* 89: 387–402.
- Hwang, W.S., Weirauch, C. 2012. Evolutionary history of assassin bugs: insights from divergence dating and ancestral state reconstruction. *PLoS ONE* 7: e45523.
- Lent, H., Wygodzinsky, P. 1979. Revision of the Triatominae (Hemiptera, Reduviidae), and their significance as vectors of Chagas' disease. *Bulletin of the American Museum of Natural History* 163: 125–520.
- Myers, N., Mittermeier, R.A., Mittermeier, C.G., da Fonseca, G.A.B., Kent, J. 2000. Biodiversity hotspots for conservation priorities. *Nature* 403: 853–858.

- Nilsson M., Forsman, A. 2003. Evolution of conspicuous colouration, body size and gregariousness: a comparative analysis of lepidopteran larvae. *Evolutionary Ecology* 17: 51–66.
- Rudh, A. 2013. Loss of conspicuous coloration has co-evolved with decreased body size in populations of poison dart frogs. *Evolutionary Ecology* 27: 755–767.
- Schatz, G.E. 2002. Taxonomy and herbaria in service of plant conservation: lessons from Madagascar's endemic families. *Annals of the Missouri Botanical Garden* 89: 145–152.
- Smith, M.A., Fisher, B.L., Hebert, P.D.N. 2005. DNA barcoding for effective biodiversity assessment of a hyperdiverse arthropod group: the ants of Madagascar. *Philosophical Transactions of the Royal Society B* 360: 1825–1834.
- Sussman, R.W., Green, G.M., Sussman, L.K. 1996. The use of satellite imagery and anthropology to assess the causes of deforestation in Madagascar. In Sponsel, L.E., Headland, T.N., & Bailey, R.C. (eds.): *Tropical Deforestation: The Human Dimension*. Columbia University Press, New York, pp. 296–315.
- Tullberg, B.S., Hunter, A.F. 1996. Evolution of larval gregariousness in relation to repellent defences and warning coloration in tree-feeding Macrolepidoptera: a phylogenetic analysis based on independent contrasts. *Biological Journal of the Linnean Society* 57: 253–276.
- Vences, M. 2004. Origin of Madagascar's extant fauna: a perspective from amphibians, reptiles and other non-flying vertebrates. *Italian Journal of Zoology* 71(S2): 217–228.
- Weirauch, C. 2006. Observations on the sticky trap predator *Zelus luridus* Stål (Heteroptera, Reduviidae, Harpactorinae), with the description of a novel gland associated with the female genitalia. *Denisia* 19: 1169–1180.
- Weirauch, C. 2007. Hairy attachment structures in Reduviidae (Cimicomorpha, Heteroptera), with observations on the fossula spongiosa in some other Cimicomorpha. *Zoologischer Anzeiger - A Journal of Comparative Zoology* 246: 155–175.
- Weirauch, C. 2008. Cladistic analysis of Reduviidae (Heteroptera: Cimicomorpha) based on morphological characters. *Systematic Entomology* 33: 229–274.
- Weirauch, C. 2010. *Tribelocodia ashei*, new genus and new species of Reduviidae (Insecta: Hemiptera), has implications on character evolution in Ectrichodiinae and Tribelocephalinae. *Insect Systematics and Evolution* 41: 103–122.
- Weirauch, C., Munro, J.B. 2009. Molecular phylogeny of the assassin bugs (Hemiptera: Reduviidae), based on mitochondrial and nuclear ribosomal genes. *Molecular Phylogenetics and Evolution* 53: 287–299.
- Weirauch, C., Forero, D., Jacobs, D.H. 2011. On the evolution of raptorial legs – an insect example (Hemiptera: Reduviidae: Phymatinae). *Cladistics* 27: 138–149.

- Wignall, A.E., Taylor, P.W. 2011. Assassin bug uses aggressive mimicry to lure spider prey. *Proceedings of the Royal Society of London B* 278: 1427–1433.
- Yoder, A.D., Nowak, M.D. 2006. Has vicariance or dispersal been the predominant biogeographic force in Madagascar? Only time will tell. *Annual Review of Ecology, Evolution, and Systematics* 37: 405–431.
- Zhang, G., Weirauch, C. 2013. Sticky predators: a comparative study of sticky glands in harpactorine assassin bugs (Insecta: Hemiptera: Reduviidae). *Acta Zoologica* 94: 1–10.
- Zhang, J., Gordon, E., Forthman, M., Hwang, W.S., Walden, K., Swanson, D., Johnson, K.P., Meier, R., Weirauch, C. 2016. Evolution of the assassin's arms: insights from a phylogeny of combined transcriptomic and ribosomal DNA data (Heteroptera: Reduvidae). *Scientific Reports* 6: 22177.

## **Chapter 1: Toxic associations: natural history observations and review of the predatory behaviors of millipede assassin bugs (Hemiptera: Reduviidae: Ectrichodiinae)**

### **Abstract**

Ectrichodiinae (Hemiptera: Reduviidae), the millipede assassin bugs, are a speciose group (>660 species) of assassin bugs that appear to be specialist predators on Diplopoda, or millipedes. Apparently capable of coping with the noxious defensive compounds produced by many millipedes, Ectrichodiinae are engaged in a predator-prey relationship with millipedes realized only by few other arthropods. Unfortunately, feeding behaviors of Ectrichodiinae are inadequately documented, rendering this exciting phenomenon largely inaccessible. We here present a literature review on ectrichodiine prey selection and feeding behaviors, with supplemental original observations on *Rhiginia cinctiventris* (Stål), 1872, in Costa Rica. Thirteen species in 12 genera have been observed to feed on millipedes. The majority of diplopod prey species were reported from the orders Spirostreptida and Spirobolida, whereas Polydesmida are rarely attacked. Ectrichodiinae insert their stylets at the millipede's intersegmental membranes on the ventral and ventrolateral trunk area or between the head and collum. Communal predation was observed among conspecific nymphs, among groups of nymphs with a conspecific adult, and more rarely among adults. Immature ectrichodiines were rarely observed to engage in solitary predation. Observations on *R. cinctiventris* indicate that this species preys on spirobolid and polydesmid millipedes and are in agreement with behaviors described for other Ectrichodiinae.

### **Introduction**

Millipedes (Diplopoda) are a diverse group of arthropods that include 16 orders represented by approximately 12,000 species in 145 families (Sierwald & Bond, 2007). In 11 of these orders,



millipedes are protected from predators by chemical defenses produced in glands that vary in number among orders and are located laterally or mid-dorsally in the diplosegments (Eisner et al., 1978; Hopkin & Read, 1992). The major defense components of these secretions seem to exhibit a phylogenetic pattern: Julida, Spirostreptida, and Spirobolida (Superorder Juliformia) produce benzoquinones; Callipodida and Chordeumatida produce phenols; Glomerida and Polyzoniida produce alkaloids and quinazolinones; Polydesmida produce cyanogenic compounds; and Platydesmida, Siphonocryptida, and Siphonophorida produce terpenoids (Eisner et al., 1978; Hopkin & Read, 1992; Sierwald & Bond, 2007). Despite such defenses, many vertebrates prey on millipedes (Sierwald & Bond, 2007). In addition, some invertebrates are specialized predators or parasitoids on juliform millipedes (Banks, 1911; Picard, 1930; Lawrence, 1984; Eisner et al., 1998; Dejean et al., 2001; Brunke et al., 2009; Larsen et al., 2009). None of these groups of millipede specialists are particularly speciose. This is dramatically different in the Ectrichodiinae, the fifth largest subfamily of Reduviidae (Hemiptera: Heteroptera), or assassin bugs, that appears to be specialized on millipedes (Green, 1925; Haridass & Ananthkrishnan, 1980; Haridass, 1985).

The circumtropical millipede assassin bugs comprise >660 species in 123 genera (Maldonado, 1990; Dougherty, 1995; Carpintero & Maldonado, 1996; Weirauch et al., 2009; Chłond, 2010). Species have been collected while feeding on millipedes, but details on the predator-prey relationships, including prey specificity and point of mouthpart insertion are largely undocumented. Thus, it is unknown if Ectrichodiinae target the millipede's nervous system to maximize the impact of their toxic saliva. It is also unknown if Ectrichodiinae target the body regions that possess defensive glands to sequester their own defensive compounds, or if they avoid the millipedes' defense glands. In addition, predation in Ectrichodiinae may be a communal activity (Haridass & Ananthkrishnan, 1980; Haridass, 1985) rather than a solitary behavior as

usually observed in Reduviidae (Readio, 1927; McMahan, 1983; Li et al., 2010), but details on this unusual subsocial behavior are scarce.

Here, we compile published data and evaluate images available on the Internet for prey specificity, point of stylet insertion, and communal predation. In addition, data for *Rhiginia cinctiventris* (Stål), 1872, are presented based on observations in the field and laboratory at one site in Costa Rica; feeding observations have never been reported for this genus before. Literature and original data were analyzed in response to three questions: (1) Does the literature data suggest a pattern of Ectrichodiinae prey preference for species in particular millipede orders? (2) Assuming that Ectrichodiinae either target or circumvent the millipede's defense glands, where do they insert their mandibular and maxillary stylets to inject toxic saliva and digestive enzymes? (3) Are communal predation strategies commonly observed in nymphs and adults?

### **Literature Review and Analysis of Online Image Material**

Publications and online images containing information on Ectrichodiinae feeding behavior were evaluated for information regarding ectrichodiine and millipede taxa, ectrichodiine developmental stage, stylet insertion, and communal predation (Table 1.1). Even though some references identified millipedes to genus or species (Cachan, 1952; Miller, 1953; Haridass & Ananthkrishnan, 1980; Lawrence, 1984; Giliomee, 1985; Ambrose, 1999), the validity of these names have changed over the years and identification without voucher material is tentative. Thus, We focus on establishing a pattern based on the prey order level instead. Victor Cheah (<http://www.clubsnap.com/forums/threads/861603-Assassin-feed-on-millipede>; accessed 20 Jun. 2011), Jonker Fourie (<http://fireflyafrica.blogspot.com/2009/01/shongololo-millipede.html>; accessed 13 Jun. 2011), Helcio Gil-Santana (<http://www.nadiplochilo.com/fpseudonannolenidae.html>; accessed 18 Aug. 2011), Joce Gordon (<http://www>.

marlothparkhonoraryrangers.co.za/?page\_id=976; accessed 18 Oct. 2011), Zichen Wang (<http://www.flickr.com/photos/biozcw/5247453267/>; accessed 18 Aug. 2011), Eddy Lee (<http://www.flickr.com/photos/lonesomecrow/5750042813/>; accessed 18 Aug. 2011), Ted MacRae (<http://beetlesinthebush.wordpress.com/2009/02/18/millipede-assassin-bug/>; accessed 18 Aug. 2011), Alan Weaving (<http://imagepro.photography.com/weaving>; accessed 18 Aug. 2011), and Arend van de Wetering (<http://www.whatsthatbug.com/2008/08/22/millipede-assassin-bugs-from-south-africa/>; accessed 18 Oct. 2011) took images used in the analysis and gave permission to publish the data and images (Fig. 1.1). Two images did not specify the photographer or provide contact information; these images were found on the following websites: <http://www.flickr.com/photos/12639178@N07/5631578778/> (accessed 18 Oct. 2011) and <http://www.whatsthatbug.com/2008/08/22/millipedeassassin-bugs-from-south-africa/> (accessed 18 Oct. 2011).

### **Field and Laboratory Procedures**

Specimens of *Rhiginia cinctiventris* were collected at La Selva Biological Research Station, Costa Rica (10.430862°N 84.006467°W) from August 9–15, 2010. Ectrichodiinae and millipedes were placed individually into petri dishes or larger containers with moist tissue. Immature ectrichodiines were determined to be the same species as adults based on coloration and proximity to adults. Sex and developmental stages were determined prior to feeding observations. Individual specimens or groups of individuals were exposed to millipedes representing the orders Spirobolida and Polydesmida, resulting in 25 trials. For each trial, specimens were removed after 20 min if the ectrichodiine(s) did not approach, grip, or probe the millipede. If individuals interacted with the millipede, observations continued until the individual(s) had ceased these interactions for more than 15 min or until after feeding. All ectrichodiine specimens and one

representative of each millipede species were preserved in 100% ethanol, barcode labeled, databased, and deposited at the Entomological Research Museum at the University of California, Riverside (UCR).

### **Ectrichodiinae Prey Specificity**

Representatives of 12 Ectrichodiinae genera have been documented in the literature to feed on millipedes in 42 separate observations (Table 1.1). Seventeen observations involved millipedes in the order Spirostreptida, six in Spirobolida, and one in Polydesmida. Six additional observations included millipedes that could only be identified to the superorder Juliformia, based on identifications made by Rowland Shelley (some online images) and identification keys (Loomis, 1968; Sierwald, [http://archive.fieldmuseum.org/millipeet/milli\\_key.html](http://archive.fieldmuseum.org/millipeet/milli_key.html); accessed 18 Oct. 2011). Two of the 25 laboratory observations and one field observation made by the senior author indicate that *Rhiginia cinctiventris* preys on Spirobolida and Polydesmida. Although feeding was not observed directly in the field, observation of an adult female and two late instar (3–5) nymphs with swollen abdomens in close proximity to a dead spirobolid millipede (Fig. 1.2A) indicate a recent meal.

The observation that only three of the 16 millipede orders have been reported as prey of Ectrichodiinae may be, in part, due to a lack of data or may be based on the fact that these orders have high species diversity (Sierwald & Bond, 2007). However, this observation may also hint to a potential prey preference for the benzoquinone-producing Juliformia over cyanogenic Polydesmida. The relative toxicity of benzoquinones and cyanogenic compounds to ectrichodiines is unknown. Based on material safety data sheets, benzoquinone has an LD<sub>50</sub> of 25 mg/kg (mouse; oral), whereas hydrogen cyanide has an LD<sub>50</sub> of 3.7 mg/kg (mouse; oral). The degree of toxicity between the two compounds may be relatively similar for ectrichodiines.

### **Point of Stylet Insertion**

From the literature, only 18 observations made reference to the millipede's body region attacked by ectrichodiines, all of which were on the trunk. Fourteen of these stated that the ventral or ventrolateral area was targeted (Figs. 1.1C, D, F, J), with 8 of these attacks at intersegmental membranes (Fig. 1.1I). Four less specific observations by Miller (1953) indicated that ectrichodiines targeted trunk intersegmental membranes, and Lawrence (1984) mentioned intersegmental membranes without reference to body region. Haridass (1985) stated that ectrichodiines also insert stylets at the head-collum intersegmental membrane. In our two laboratory trials, specimens of *R. cinctiventris* inserted stylets at the head-collum intersegmental membrane and/or at intersegmental membranes along the ventral, ventrolateral, and/or dorsal trunk (Figs. 1.2B). Overall, the data suggest Ectrichodiinae primarily target the millipede's ventral or ventrolateral trunk, with some exceptions.

There are several possible explanations for this behavior. These areas may be more penetrable than others, making a more suitable entry point for the typically smooth, unarmored stylets (Weirauch, 2003, 2008). Unfortunately, little is known about diplopod cuticular ultrastructure and chemical composition to further support this claim (Subramoniam, 1974; Walker & Crawford, 1980; Ansenne et al., 1990). Alternatively, Ectrichodiinae may initially target the millipede's ventral ganglia or the supra-esophageal ganglion to expose the nervous system to toxic salivary secretions. The millipede's ventral or head-collum region may be selected to avoid inserting stylets into laterally or dorsally located defensive glands along the trunk during millipede immobilization and consumption. It remains to be shown if, once immobilized, the millipede's defense gland reservoirs are targeted by ectrichodiines to sequester toxic secondary metabolites. At least half of the genera of Ectrichodiinae exhibit aposematic coloration to a larger extent than seen in most other reduviid subfamilies. Although ectrichodiines

are presumably well protected from vertebrate predation due to their generally powerful bites and defensive secretions (Louis, 1974; Ambrose, 1999), a millipede diet may provide ectrichodiines with additional defensive benefits in the form of toxic secondary metabolites. Secondary defense metabolites have been investigated in other insects, such as aposematically colored Lepidoptera that sequester many chemicals from their host plants and biochemically alter them for defense purposes (Bowers, 1990; Nishida, 2002). Ectrichodiinae may potentially employ a similar strategy by targeting the defense gland reservoirs.

### **Communal Predatory Strategies**

Evidence for communal predation comprises 13 observations. Of these, two involved groups of conspecific nymphs in the presence of a conspecific adult, whereas eight involved groups of conspecific nymphs. Conspecific adults engaging in communal behaviors have only been documented in *Cleptria cinctiventris* Stål, 1855 (Lawrence, 1984). Evidence of immatures engaging in solitary predation has only been documented for a *Haematorrhophus* species (Green, 1925). Communal predation was observed in only one of our laboratory trials with *R. cinctiventris* (Fig. 1.2C): early instar (1–2) nymphs, in the presence of an adult female and two later instar nymphs, frequently attempted predation on a millipede without immobilizing it. Overall, the data may indicate that communal, conspecific feeding among immatures or immatures and adults is the rule but rather an exception among adults. One potential explanation is that the salivary secretions produced by a single nymph may not be sufficiently toxic to immobilize and kill a millipede. Thus, the attack of multiple individuals may be necessary to increase the toxic effects, especially to subdue large-bodied prey items.

## **Conclusion**

The predator-prey association between Ectrichodiinae and millipedes was first documented almost a century ago (Green, 1925), but has since received little attention. Judging from published data, it is obvious that the feeding behavior of Ectrichodiinae is still poorly understood. About 2% of ectrichodiine species have been documented to prey on millipedes, leaving ample room to further test the claim that Ectrichodiinae, as a group, specialize on millipedes. Based on our analysis, Ectrichodiinae may be exclusive millipede specialists, with a potential preference for the benzoquinone-producing Juliformia. Millipedes were usually not identified below the order level in the literature and online images. Thus, future investigations should attempt accurate genus or species determination of millipedes. Furthermore, the millipede's ventral trunk area and intersegmental membranes, including between the head and collum, are typically targeted. Whether these areas are attacked in relation to the location of the millipede nervous system or defense glands remains to be investigated. Lastly, immature and adult Ectrichodiinae exhibit both solitary and communal predation. The majority of observations suggest that immatures typically exhibit communal predation, whereas adults usually predate solitarily. Given that these conclusions are general and preliminary, it is imperative that future studies provide detailed documentation regarding these aspects of ectrichodiine predation. Other relevant investigations that will contribute to understanding this phenomenon include examining the chemical composition of ectrichodiine salivary secretions and the ability of ectrichodiines to sequester secondary metabolites from diplopods for defense.

## References

- Ambrose, D.P. 1999. *Assassin Bugs*. Science Publishers, Enfield, NH.
- Ansenne, A., Compère, P., Goffinet, G. 1990. Ultrastructural organization and chemical composition of the mineralized cuticle of *Glomeris marginata* (Myriapoda, Diplopoda). In Minelli A. (ed.): *Proceedings of the 7th International Congress of Myriapodology*. Brill, Leiden, pp. 125–134.
- Banks, N. 1911. A curious habit of one of our phorid flies. *Proceedings of the Entomological Society of Washington* 13: 212–214.
- Bowers, M.D. 1990. Recycling plant natural products for insect defense. In Evans D.L. & Schmidt J.O. (eds): *Insect Defenses: Adaptive Mechanisms and Strategies of Prey and Predators*. SUNY Press, Albany, NY, pp. 353–386.
- Brunke, A.J., Bahlai, C.A., Sears, M.K., Hallett, R.H. 2009. Generalist predators (Coleoptera: Carabidae, Staphylinidae) associated with millipede populations in sweet potato and carrot fields and implications for millipede management. *Environmental Entomology* 38: 1106–1116.
- Cachan, P. 1952. Etude de la prédation chez les Réduvides de la région Éthiopienne. 1. La prédation en groupe chez *Ectrichodia gigas* H-Sch. *Physiologia Comparata et Oecologia* 2: 378–385.
- Carpintero, D.J., Maldonado, J. 1996. Diagnostic characters and key to the genera of American Ectrichodiinae (Heteroptera, Reduviidae). *Caribbean Journal of Science* 32: 125–141.
- Chłond, D. 2010. A new, remarkable genus and two new species of Ectrichodiinae (Hemiptera: Heteroptera: Reduviidae) from Madagascar. *Zootaxa* 2522: 61–68.
- Dejean, A., Suzzoni, J.P., Schatz, B. 2001. Behavioral adaptations of an African ponerine ant in the capture of millipedes. *Behaviour* 138: 981–996.
- Dougherty, V. 1995. A review of the New World Ectrichodiinae genera (Hemiptera: Reduviidae). *Transactions of the American Entomological Society* 121: 173–225.
- Eisner, T., Alsop, D., Hicks, K., Meinwald, J. 1978. Defensive secretions of millipedes. In Bettini, S. (ed.): *Arthropod Venoms*. Springer, Berlin, pp. 41–72.
- Eisner, T., Eisner, M., Attygalle, A.B., Deyrup, M., Meinwald, J. 1998. Rendering the inedible edible: circumvention of a millipede's chemical defense by a predaceous beetle larva (Phengodidae). *Proceedings of the National Academy of Sciences of the United States of America* 95: 1108–1113.
- Giliomee, J.H. 1985. A millipede assassinated. *African Wildlife* 39: 149.



- Green, W.E. 1925. The President's address. *Proceedings of the Entomological Society of London* 1924: clxi–ccii.
- Haridass, E.T. 1978. *Biological and Ethological Studies on some South Indian Reduviids (Hemiptera – Reduviidae)*. Ph.D. Thesis, University of Madras, Madras.
- Haridass, E.T. 1985. Feeding and ovipositional behavior in some reduviids (Insecta – Heteroptera). *Proceedings of the Indian Academy of Sciences, Animal Sciences* 94: 239–247.
- Haridass, E.T., Ananthkrishnan, T.N. 1980. Models for the predatory behavior of some reduviids from southern India (Insecta – Heteroptera – Reduviidae). *Proceedings of the Indian Academy of Sciences, Animal Sciences* 89: 387–402.
- Hopkin, S.P., Read H.J. 1992. *The Biology of Millipedes*. Oxford University Press, Oxford, UK.
- Larsen, T.H., Lopera, A., Forsyth, A., Génier, F. 2009. From coprophagy to predation: a dung beetle that kills millipedes. *Biology Letters* 5: 152–155.
- Lawrence, R.F. 1984. *The Centipedes and Millipedes of Southern Africa: a Guide*. A.A. Balkema, Cape Town.
- Li, H., Zhao, G.Y., Cao, L.M., Xu, K., Cai, W.Z. 2010. Taxonomic and bionomic notes on the white spot assassin bug *Platyeris biguttatus* (Linnaeus) (Hemiptera: Reduviidae: Reduviinae). *Zootaxa* 2644: 47–56.
- Livingstone, D., Ambrose, D.P. 1984. Adaptive modifications of the Reduviidae of the scrub jungles and semi-arid zones of the Palghat Gap, India – an evolutionary approach. *The Journal of the Bombay Natural History Society* 81: 583–595.
- Loomis, H.F. 1968. A checklist of the millipedes of Mexico and Central America. *Bulletin of the United States National Museum* 266: 1–137.
- Louis, D. 1974. Biology of Reduviidae of cocoa farms in Ghana. *American Midland Naturalist* 91: 68–89.
- Maldonado, J. 1990. Systematic catalogue of the Reduviidae of the world (Insecta: Heteroptera). *Caribbean Journal of Science*, Special Edition: 1–694.
- McGavin, G.C. 1993. *Bugs of the World*. Facts on File, New York, NY.
- McMahan, E.A. 1983. Adaptations, feeding preferences, and biometrics of a termite-baiting assassin bug (Hemiptera: Reduviidae). *Annals of the Entomological Society of America* 76: 483–486.
- Miller, N.C.E. 1953. Notes on the biology of the Reduviidae of southern Rhodesia. *Transactions of the Zoological Society of London* 27: 541–672.
- Miller, N.C.E. 1971. *The Biology of the Heteroptera*. E.W. Classey, Hampton.

- Nishida, R. 2002. Sequestration of defensive substances from plants by Lepidoptera. *Annual Review of Entomology* 47: 57–92.
- Paulian, R. 1948. Sur quelques insectes guano-bies de la Coted'Ivoire. *Publications du Muséum National d'Histoire Naturelle* 10: 63–68.
- Picard, F. 1930. Sur le parasitisme d'un Phoride (*Megaselia cuspidata* Schmitz) aux dépens d'un Myriapode. *Séance du 25 Février 1930*: 180–183.
- Readio, P.A. 1927. Studies on the biology of the Reduviidae of America north of Mexico. *Kansas University Science Bulletin* 17: 1–291.
- Sahayaraj, K. 2006. Ecological adaptive features of hunter reduviids (Insecta: Heteroptera: Reduviidae Latreille 1807) and their biological control potential. In Gupta V .K. & Verma A.K. (eds): *Perspectives in Animal Ecology and Reproduction. Vol. 3*. Daya Publishing House, Delhi, pp. 22–48.
- Sierwald, P., Bond, J.E. 2007. Current status of the myriapod class Diplopoda (millipedes): taxonomic diversity and phylogeny. *Annual Review of Entomology* 52: 401–420.
- Subramoniam, T. 1974. Histochemical study on cuticle of a millipede *Spirostreptus asthenes* (Diplopoda: Myriapoda). *Acta Histochemica* 51: 200–204.
- Walker, L.J., Crawford, C.S. 1980. Integumental ultrastructure of the desert millipede, *Orthoporus ornatus* (Girard) (Diplopoda: Spirostreptidae). *International Journal of Insect Morphology and Embryology* 9: 231–249.
- Weirauch, C. 2003. *Ein Beitrag zum phylogenetischen System der Reduviidae (Heteroptera)*. Ph.D. Dissertation. Fachbereich Biologie/Chemie/Pharmazie. Freie Universität Berlin.
- Weirauch, C. 2008. Cladistic analysis of Reduviidae (Heteroptera: Cimicomorpha) based on morphological characters. *Systematic Entomology* 33: 229–274.
- Weirauch, C., Rabitsch, W., Redei, D. 2009. *Austrokatanga*, gen. nov., new genus of Ectrichiinae (Hemiptera: Heteroptera: Reduviidae) from Australia. *Zootaxa* 2094: 1–15

Figure 1.1. Internet images used for analysis of Ectrichodiinae predatory behavior. A. An adult *Brontostoma trux* feeding on a spirostreptid millipede (Gil-Santana). Millipede identified by R. Shelley and ectrichodiine identified by H. Gil-Santana. B. *Ectrichodia* sp. nymphs communally feed on a spirostreptid millipede (Fourie). C. *Ectrichodia* sp. nymphs feeding on the venter of a millipede (Weaving). D. An adult *Neozirta eidmanni* feeding on a juliform millipede carcass (Wang). W. Cai identified the ectrichodiine. E. *Ectrichodia* sp. nymphs communally feed on a juliform millipede (van de Wetering). F. An adult and three immature *Ectrichodia crux* share a juliform millipede carcass (MacRae). T. MacRae identified the ectrichodiine. G. An adult *Ectrichodia crux* consumes a juliform millipede (<http://www.flickr.com>). H. *Ectrichodia crux* nymphs communally feed on a spirostreptid millipede (<http://www.whatsthatbug.com>). R. Shelley identified the millipede. I. An adult *Schottus* sp. feeds on a spirobolid millipede (Cheah). J. An adult *Schottus* sp. feeds on a juliform millipede (Lee).



©H. Gil-Santana



©J. Fourie



©A. van de Wetering



©A. Weaving



©J. Wang



©T. MacRae



©www.flickr.com



©www.whatsthatbug.com



©V. Cheah



©E. Lee

Figure 1.2. Field and laboratory images of *Rhiginia cinctiventris* predation on millipedes at La Selva Biological Station, Costa Rica. A. Field image of an adult and two immature *Rhiginia cinctiventris* near a spirobolid millipede carcass at La Selva Biological Station, Costa Rica. B. An adult female feeds on a small polydesmid millipede. C. An adult female, two early instar nymphs, and two later instar nymphs in the presence of a spirobolid millipede.





Table 1.1. Data extracted from published literature and Internet images for analysis of Ectrichodiinae predatory behavior.

<b>Ectrichodiinae taxon</b>	<b>Stage</b>	<b>Prey millipede taxa</b>	<b>Stylet insertion</b>	<b>Communal predation</b>	<b>Reference</b>
<i>Brontostoma notatum</i> (Stål), 1859	Adult	Diplopoda		Solitary	McGavin (1993)
<i>Brontostoma trux</i> (Stål), 1859	Adult	Spirostreptida		Solitary	Gil-Santana (image)
<i>Cleptria cinciventris</i> Stål, 1855	Adults	Spirostreptida	Intersegmental membrane	Conspecific adults	Lawrence (1984)
<i>Ectrichodia</i> spp.		Spirostreptida	Intersegmental membrane on trunk		Miller (1953)
<i>Ectrichodia</i> spp.		Spirostreptida			Miller (1971)
<i>Ectrichodia</i> sp.	Nymphs	Juliformia		Conspecific nymphs	van de Wetering (image)
<i>Ectrichodia</i> sp.	Nymphs	Spirostreptida		Conspecific nymphs	Fourie (image)
<i>Ectrichodia</i> sp.	Nymphs	Diplopoda	Ventral trunk	Conspecific nymphs	Weaving (image)
<i>Ectrichodia barbicornis</i> (Fabricius), 1775		Diplopoda			Louis (1974)
<i>Ectrichodia crux</i> (Thunberg), 1783	Nymphs	Juliformia		Conspecific nymphs	Gordon (image)
<i>Ectrichodia crux</i> (Thunberg), 1783		Spirostreptida	Intersegmental membrane on ventral trunk	Solitary	Gilimee (1985)
<i>Ectrichodia crux</i> (Thunberg), 1783		Diplopoda	Intersegmental membrane on ventral trunk	Solitary	Gilimee (1985)
<i>Ectrichodia crux</i> (Thunberg), 1783	Adult & Nymphs	Juliformia	Ventral trunk	Conspecific nymphs with adult	MacRae (image)
<i>Ectrichodia crux</i> (Thunberg), 1783	Adults	Spirostreptida	Intersegmental membrane on ventral trunk	Solitary	Miller (1953)
<i>Ectrichodia crux</i> (Thunberg), 1783		Diplopoda			Paulian (1948)

Table 1.1. (Continued).

<b>Ectrichodiinae taxon</b>	<b>Stage</b>	<b>Prey millipede taxa</b>	<b>Stylet insertion</b>	<b>Communal predation</b>	<b>Reference</b>
<i>Ectrichodia crux</i> (Thunberg), 1783	Adult	Juliformia		Solitary	<a href="http://www.flickr.com">http://www.flickr.com</a> (image)
<i>Ectrichodia crux</i> (Thunberg), 1783	Nymphs	Spirostreptida		Conspecific nymphs	<a href="http://www.whatsthatbug.com">http://www.whatsthatbug.com</a> (image)
<i>Ectrichodia gigas</i> (Herrich-Schaeffer), 1848	Adult & Nymphs	Spirostreptida	Intersegmental membrane on ventral trunk	Conspecific nymphs; nymphs with adult	Cachan (1952)
<i>Ectrychotes pilicornis</i> (Fabricius), 1794		Diplopoda			Haridass (1978)
<i>Glymmatophora</i> spp.		Spirostreptida	Intersegmental membrane on trunk		Miller (1953)
<i>Glymmatophora</i> sp.	Nymphs	Spirostreptida	Intersegmental membrane	Conspecific nymphs	Lawrence (1984)
<i>Guionius nigripennis</i> (Fabricius), 1794		Diplopoda			Haridass (1978)
<i>Haematorrhophus</i> sp.		Diplopoda			Livingstone & Ambrose (1984)
<i>Haematorrhophus</i> sp.		Spirobolida			Ambrose (1999)
<i>Haematorrhophus</i> sp.		Spirobolida			Ambrose (1999)
<i>Haematorrhophus</i> sp.		Spirobolida			Ambrose (1999)
<i>Haematorrhophus</i> sp.	Nymph	Diplopoda	Ventral posterior trunk	Solitary	Green (1925)
<i>Haematorrhophus linnaei</i> (Stål), 1859	Adult	Diplopoda			Green (1925)
<i>Haematorrhophus linnaei</i> (Stål), 1859	Adult	Diplopoda	Ventral posterior trunk		Sahayaraj (2006)
<i>Haematorrhophus nigroviolaceus</i> (Reuter), 1873	Adult & Nymphs	Spirobolida	Intersegmental membrane on ventral posterior trunk	“Nymphs as well as adults”	Haridass & Ananthakrishnan (1980)
<i>Haematorrhophus nigroviolaceus</i> (Reuter), 1873	Adult & Nymphs	Spirobolida	Intersegmental membrane on ventral posterior trunk	“Nymphs as well as adults”	Haridass & Ananthakrishnan (1980)



Table 1.1. (Continued).

<b>Ectrichodiniinae taxon</b>	<b>Stage</b>	<b>Prey millipede taxa</b>	<b>Stylet insertion</b>	<b>Communal predation</b>	<b>Reference</b>
<i>Haematorrhophus nigroviolaceus</i> (Reuter), 1873	Adult & Nymphs	Polydesmida	Intersegmental membrane on ventral posterior trunk	“Nymphs as well as adults”	Haridass & Ananthakrishnan (1980)
<i>Haematorrhophus nigroviolaceus</i> (Reuter), 1873		Spirostreptida			Haridass & Ananthakrishnan (1980)
<i>Maraenaspis</i> spp.		Spirostreptida			Miller (1953)
<i>Maraenaspis</i> spp.		Spirostreptida			Miller (1971)
<i>Maraenaspis problematica</i> (Gerstaecker), 1892	Nymphs	Spirostreptida	Intersegmental membrane on trunk	Conspecific nymphs	Miller (1953)
<i>Neozirta eidmanni</i> (Taeuber), 1930	Adult	Juliformia	Ventral posterior trunk	Solitary	Wang (image)
<i>Nularda nobilitata</i> Stål, 1859		Diplopoda			Louis (1974)
<i>Scadra</i> spp.		Spirostreptida	Intersegmental membrane on trunk		Miller (1953)
<i>Scadra</i> spp.		Spirostreptida			Miller (1971)
<i>Schottus</i> sp.	Adult	Spirobolida	Intersegmental membrane on ventrolateral trunk	Solitary	Cheah (image)
<i>Schottus</i> sp.	Adult	Juliformia	Ventral or ventrolateral trunk	Solitary	Lee (image)

## **Chapter 2: Taxonomic monograph of the endemic millipede assassin bug fauna of Madagascar (Hemiptera: Reduviidae: Ectrichodiinae)**

### **Abstract**

Madagascar is one of the world's most recognized biodiversity hotspots and has a diverse assassin bug (Reduviidae) fauna that is still incompletely known, especially for the Ectrichodiinae, or millipede assassin bugs. Ectrichodiinae are a speciose (673 described species, 118 genera), worldwide group of assassin bugs that is most diverse in the Old and New World tropics, but so far only six genera and 10 species have been described from Madagascar. Based on examination of 1,981 ectrichodiine specimens mostly from the California Academy of Sciences, as well as a few other collections, dramatic undescribed species-level diversity is revealed. The island's Ectrichodiinae fauna is taxonomically revised with the description of three new genera (*Marojejycoris*, *Tanindrazanus*, and *Toliarus*) and 63 new species: *Gibbosella andasibe*, *Gibbosella betampona*, *Gibbosella brunalvus*, *Gibbosella conisimilis*, *Gibbosella fulva*, *Gibbosella mantella*, *Gibbosella megafrons*, *Gibbosella nitida*, *Gibbosella notoconica*, *Gibbosella pallidacorium*, *Gibbosella pallidalata*, *Gibbosella planiscutum*, *Gibbosella quadocris*, *Gibbosella vangocris*, *Glymmatophora (Glymmatophora) carolae*, *Marojejycoris auranticorium*, *Marojejycoris brevifrons*, *Marojejycoris francais*, *Marojejycoris notadichroa*, *Marojejycoris ranomafana*, *Tanindrazanus amboasarius*, *Tanindrazanus andohahela*, *Tanindrazanus anjzorobeus*, *Tanindrazanus antananarivo*, *Tanindrazanus bemaraha*, *Tanindrazanus brunneus*, *Tanindrazanus hannajagoda*, *Tanindrazanus harinhali*, *Tanindrazanus irwini*, *Tanindrazanus joffrevillus*, *Tanindrazanus kathrynae*, *Tanindrazanus mahafaly*, *Tanindrazanus marginatus*, *Tanindrazanus marojejy*, *Tanindrazanus nigripes*, *Tanindrazanus notatus*, *Tanindrazanus simulans*, *Tanindrazanus tenebricus*, *Tanindrazanus varicolor*, *Tanindrazanus vohiparara*,

*Toliarus karinae*, *Toliarus trichrous*, *Toxopus ambohitantely*, *Toxopus ampitavananima*, *Toxopus antsiranana*, *Toxopus basalis*, *Toxopus brucei*, *Toxopus farafangana*, *Toxopus fisheri*, *Toxopus griswoldi*, *Toxopus insignis*, *Toxopus italaviana*, *Toxopus melobrunneus*, *Toxopus miandritsara*, *Toxopus namoroka*, *Toxopus pallidus*, *Toxopus parkeri*, *Toxopus simulans*, *Toxopus steineri*, *Toxopus tibialis*, *Toxopus toamasina*, *Toxopus toliara*, and *Toxopus vazimba*. *Toxopus* Bergroth, 1905, is redescribed, with the first description of males in the genus, and *Cleptria signoretii* Reuter, 1887, is transferred to *Toxopus*. *Gibbosella* Chłond, 2010, is also redescribed and includes description of males for the first time. *Gibbosella elongata* Chłond, 2010, *Glymmatophora crassipes* Horváth, 1914, and *Maraenaspis bidens* (Reuter), 1887, are redescribed, with the first description of males for each species. Habitus images, documentation of male and female genitalic features, identification keys, and distribution maps are provided. Males and females are associated based on morphology, geographic information, and, when available, molecular data. The newly discovered species diversity constitutes a sixfold increase over the previously documented millipede assassin bug fauna in Madagascar.

## **Introduction**

Madagascar, often referred to as “the island continent” due to its wide range of ecosystems and size (Scales, 2014), is one of the most biologically diverse places in the world. More than 13,000 species of plants, 900 of vertebrates, and 5,800 of invertebrates are known to occur on the island (Goodman & Benstead, 2005; Phillipson et al., 2006), although, for invertebrates at least, these numbers are probably gross underestimates of true diversity. Among noninvasive species, about 50% of birds, 95% of reptiles, 100% of amphibians and mammals, 86% of invertebrates, and more than 90% of vascular plants are endemic to the approximately 600,000 km<sup>2</sup> island (Goodman & Benstead, 2005; Yoder & Nowak, 2006; Phillipson et al., 2006; Buerki et al., 2013).

Such diversity and endemism is extraordinary when compared to islands of relatively similar size, e.g., Sumatra (~473,500 km<sup>2</sup>: 201 mammals, 4% endemic; 580 birds, 3% endemic) (Whitten et al., 2000) or Borneo (~748,000 km<sup>2</sup>: 1,500 vertebrates, 30% endemic) (Quek, 2009). The remarkable Madagascan biodiversity is under threat from a variety of anthropogenic influences, such as introduction of invasive species (Myers et al., 2000) and poaching of endangered species (Rakotomanana et al., 2013), although the greatest threat comes from habitat loss and fragmentation (Myers et al., 2000). Given the immense threat to its largely endemic biota, Madagascar has been deemed the world's "hottest" biodiversity hotspot and its flora and fauna is of critical conservation priority (Myers et al., 2000; Ganzhorn et al., 2001; Scales, 2014).

Nevertheless, knowledge of the island's patterns of species richness, turnover, and endemism is still incomplete (Schatz, 2002; Smith et al., 2005). This gap impedes the ability to identify and prioritize critical areas of endemism for conservation efforts (Schatz, 2002; Smith et al., 2005). Thus, taxonomic surveys of the Madagascan biota are severely needed. The California Academy of Sciences' (CAS) Terrestrial Arthropod Inventory of Madagascar Project (2000–2009) was conducted to address the need for taxonomic exploration of the arthropod fauna. This project has yielded a plethora of arthropod specimens for many taxonomic studies (e.g., Mugarib & Azevedo, 2010; Krishnankutty & Dietrich, 2011; Álvarez-Padilla et al., 2012), and has yielded more than 3,000 specimens of Reduviidae, or assassin bugs.

The morphologically and ecologically diverse Reduviidae is the second largest family of true bugs (Hemiptera: Heteroptera) that comprises 7,000 described species worldwide (Putshkov & Putshkov, 1986–1989; Maldonado, 1990; Weirauch et al., 2014). Prior to the turn of the 21st century, 253 species were known to occur on Madagascar (Maldonado, 1990), ~85% of which were described by André Villiers in 30 papers published between 1948–1979 (e.g., Villiers, 1948; Villiers, 1968a; Villiers, 1979; see Maldonado, 1990, for other references). A number of small

taxonomic studies (i.e., no monographs) have since focused on Madagascan assassin bugs and the described diversity has increased by 24 species (Labina & Kerzhner, 2001; Weirauch, 2008a; Chłond, 2010a, 2010b, 2010c, 2011a, 2011b, 2014; Chłond & Junkiert, 2010, 2011; Hwang & Weirauch, 2010; Zhang & Weirauch, 2011; Chłond & Guilbert, 2012; Chłond & Bañař, 2013). Of these 277 species, only 10 currently classified in six genera belong to the fifth largest reduviid subfamily Ectrichodiinae, the millipede assassin bugs (673 species, 118 genera) (Carpintero & Maldonado, 1990, 1991; Maldonado, 1990, 1995; Dougherty, 1995; Murugan & Livingstone, 1995; Kerzhner & Günther, 2003; Gil-Santana & Costa, 2005; Gil-Santana et al., 2004; Gil-Santana et al., 2005; Gil-Santana & Baena, 2009; Weirauch et al., 2009; Chłond, 2010a; Rédei & Tsai, 2012; Rédei et al., 2012; Gil-Santana et al., 2013; Gil-Santana, 2014, 2015). All described Madagascan ectrichodiine species and three of the six genera (*Distirogaster* Horváth, 1914; *Gibbosella* Chłond, 2010; *Toxopus* Bergroth, 1905) are endemic to the island. The 1,819 ectrichodiine specimens collected by CAS's taxonomic inventory, together with 162 specimens loaned from various European and American museums and colleagues, has provided a unique opportunity to examine, document, and revise the dramatic undescribed species-level ectrichodiine diversity in Madagascar. We here describe three new genera and 63 new species, revise the existing genera *Toxopus* and *Gibbosella* and describe the males of both genera for the first time. We also transfer *Cleptria signoretii* Reuter, 1887, to *Toxopus*. The species *Gibbosella elongata* Chłond, 2010, *Glymmatophora crassipes* Horváth, 1914, and *Maraenaspis bidens* (Reuter), 1887, are redescribed, including, for the first time, descriptions of males for each species. The genus *Distirogaster* will be treated in a separate revision (Chłond et al., in prep.) due to the limited CAS material, the abundant material from European collections available to the second author (Chłond; taking the lead on the revision of *Distirogaster*), and timely finalization of the monograph; however, the taxon is included in the key to Madagascan genera and several

specimens are sequenced for molecular data. Our taxonomic approach is largely focused on male specimens, since ~90% of the material collected by CAS are males (likely due to collecting methods). Sexual dimorphism, where males are typically macropterous and females are apterous, is widespread among Ectrichodiinae and poses a significant problem for associating male and female specimens. Wherever feasible, males are associated with females (and immatures) based on a combination of morphology and geographic information and, where successfully amplified, molecular data. Cladistic methods are utilized in formulating generic concepts and informing diagnostic features of Madagascan genera and species, but results of these analyses are being published in a companion paper (Forthman & Weirauch, in press).

## **Material and methods**

### *Specimens and databasing*

A total of 1,981 specimens were examined for this taxonomic revision, of which 1,661 are males, 48 females, and 272 immatures; five additional specimens classified as *Distirogaster* were included in the molecular approach to associating dimorphic sexes. The skewed abundance of males compared to females and immatures is likely a result of collecting techniques employed by the CAS taxonomic survey: a significant proportion of the material (~80%) was collected in Malaise traps that typically favor the capture of winged insects, and thus in the case of Ectrichodiinae, male specimens. Leaf litter sifting, pitfall traps, light traps, and hand collecting resulted in a small sample (~12% of total material indicating collecting method from CAS material) of apterous female and immature specimens. Only 10 immatures were collected in Malaise traps.

Unique specimen identifier (USI) labels were affixed to specimens. All USI labels are comprised of the prefix UCR\_ENT followed by an eight-digit number. USI codes are provided in

the material examined section of each species description, but the prefix (UCR\_ENT) is omitted to save space. Specimen information was databased using the American Museum of Natural History's Planetary Biodiversity Inventory (PBI) Arthropod Easy Capture Software database (<http://www.research.amnh.org/pbi/locality/index.php>) and are available through the Heteroptera Species Pages (<http://research.amnh.org/pbi/heteropterasespeciespage/>). Specimens with locality data including GPS coordinates were databased verbatim. For some older specimens deposited in European collections, locality records were georeferenced using Google Earth v7.1.2.2041 and GeoLocate (<http://www.museum.tulane.edu/geolocate/>) to determine geographic coordinates and elevation; locality data from these specimens were databased using current geopolitical names. In several cases, locality records on older specimens could not be successfully georeferenced.

#### *Morphological methods*

External morphology and genitalic characters were examined using Nikon NiU (University of Silesia [SU]; Chłond), Nikon SMZ1000 (University of California-Riverside [UCR]; Forthman and Weirauch), Nikon SMZ1500 (SU and UCR), and Olympus SZH 10 (SU) dissecting microscopes. For males, genitalia (abdominal segment 8, pygophore, and phallus) were dissected from the body, cleared in heated 10% potassium hydroxide (KOH) for 3–8 minutes, washed in distilled water and 100% ethanol (EtOH), examined in glycerol, and permanently stored in genitalic capsules pinned to the specimen. For females, genitalia were dissected from the body using a similar protocol with the exception that the entire abdomen was detached from the thorax, heated in KOH initially, opened to expose internal structures, and subsequently cleared in heated KOH for 5–10 minutes. Female internal genitalia were permanently stored in genitalic capsules pinned to the specimen, while external abdominal structures were mounted to card stock pinned to the specimen. Genitalic structures were stained with Chlorazol Black E in 70% EtOH solution

to provide contrast to membranous areas. Morphological characters were coded in the Descriptive Language for Taxonomy (DELTA) program. Natural language taxonomic descriptions were subsequently generated.

#### *Imaging, distribution maps, and measurements*

Dorsal (Figs. 2.1–2.4), lateral (Figs. 2.5–2.7), and ventral (Figs. 2.8–2.11) habitus images were produced for representatives of species and types of some described species (except *Glymmatophora crassipes* provided by Dávid Rédei) (Fig. 2.12) using Microptics-USA or Leica Z16 APO imaging systems LAS v4.3 (UCR) or a Nikon D300 on Fomei CS-920 copy stand (SU). Images of select morphological features (head: Fig. 2.13; antenna: Fig. 2.14; thorax: Fig. 2.15; metathoracic gland evaporatorium and leg armature: Fig. 2.16; wings: Fig. 2.17; abdomen: Fig. 2.18; male and female genitalia: Fig. 2.19–23) were produced using a Leica Z16 APO imaging system. For genitalic images, genitalia were mounted on top of hand sanitizer and immersed in 70% EtOH. All images were stacked using LAS v4.3 and Zerene Stacker v1.04 (UCR). Distribution maps were created using SimpleMappr (<http://www.simplemappr.net>) based on GPS coordinates (Maps 2.1–2.13). Measurements (in mm) were made using an Olympus SZX9 stereoscopic microscope with a micrometer (SU) and a Lomo MBC-10 stereoscopic microscope with a two-axes movable stage and two digital micrometers connected to a Microcode II RS-232 readout (UCR) (Table 2.1). Because the hemelytron is often distorted in mounted specimens, body length is reported as the length from the clypeal apex to the posterior margin of the abdomen for the holotype in species descriptions, followed by a range where multiple specimens are available. In cases where the hemelytron obscured the posterior abdominal margin in dorsal view, a light source was positioned under the specimen and illuminated the outline of the abdomen under the hemelytron in dorsal view.



### *Identification keys*

An identification key is provided for genera found in Madagascar. Keys to species are given for each genus with more than one Madagascan species known. For genera having more than one extremely sexually dimorphic species, separate male and female identification keys are given. No specimens were examined for *Toxopus politus* Bergroth, 1905, which is known only from the female holotype, which may be lost (Jansson & Coscarón, 1989; Larry Huldén, personal commun.). This species is included in the key to females of this genus, but diagnostic features are based on the original description.

### *Molecular data and pairwise genetic distance analysis*

To associate males and females, sequencing and pairwise genetic distance analysis of part of the mitochondrial cytochrome oxidase I (COI) gene was performed following Zhang & Weirauch (2011) for 61 male specimens representing 39 species (including representatives of *Distirogaster*) and 10 female specimens representing 10 putative female-based species; sequence-quality specimens were not available for 35 male- and 17 putative female-based species. A hind leg was removed from each specimen for DNA extraction using QIAGEN DNeasy Blood and Tissue Kit. Amplification of partial COI was performed using COI primers C1-J-2183 (5'-CAACATTTATTTGATTTTTTGG-3', forward) (Simon et al., 1994) and C1-N-2609 (5'-CGAATACTGCTCCTATTGATA-3'' reverse) (Damgaard et al., 2000), GE Healthcare Life Sciences PuReTaq Ready-To-Go PCR Beads, and a Fisher Scientific Thermocycler with the following settings: denaturation 94° C (30 s), annealing 48° C (30 s), and extension 72° C (45 s) for 35 cycles, with an initial denaturation at 94° C (2 min) and a final extension at 72° C (7 min). PCR products were cleaned using Bioline SureClean. Sequencing was performed on an Applied Biosystems 3730xl DNA Sequencer at UCR's Institute for Integrative Genome Biology.

Sequences were assembled and beginning and ending nucleotides with unresolved chromatograms were deleted in Sequencher v4.8. Sequences are available at GenBank under the accession numbers listed in Table 2.2. Sequence alignment was performed with the online version of MAFFT (Kato et al., 2005; Kato et al., 2008; <http://mafft.cbrc.jp/alignment/server/index.html>) using the E-INS-i algorithm, which produced the shortest aligned dataset without internal gaps. TaxonDNA/SpeciesIdentifier 1.5alpha10 (Meier et al., 2006) was used to compute uncorrected pairwise genetic distances. The smallest distance between female and male sequences were determined and compared to intra- and interspecific distances. Following Zhang & Weirauch (2011), putative matches were then investigated further to assess morphological similarities and geographic proximity (DeSalle et al., 2005).

#### *Terminology, specimen depositories, and abbreviations*

Terminology generally follows a subset of terms used by Dougherty (1995), Weirauch (2008b), and Forero & Weirauch (2012). Terminology for wing venation follows Hill (2014) and Weirauch (2008b), although homology concepts across Paraneoptera are currently being reviewed (Dávid Rédei, personal commun.). Abbreviations used in plates and/or text (capitalized abbreviations used within generic and species descriptions with the exception of standard wing venation abbreviations): **1A**, first anal vein; **aa**, articulatory apparatus (basal plate and basal plate extension); **ad**, aedeagus; **afsp**, anterior femoral subapical protuberance; **alp**, anterolateral pronotal projection; **ana**, antennal articulation; **anf**, antennifer; **aoc**, anteocular area; **ap**, antennal pseudoarticulation; **apl**, anterior pronotal lobe; **as**, antennal shield; **asp8**, abdominal spiracle 8; **bc**, bursa copulatrix; **bcl**, bursa copulatrix lateral lobe; **BFLA**, basiflagellomere; **bp**, basal plate; **BPE**, basal plate extension; **ca**, clypeal apex; **cl**, collar; **cly**, clypeus; **co**, corium; **cp**, corial

pterostigma; **Cu**, cubitus; **DFLA**, distiflagellomere; **dl**, dorsal laterotergite; **dlp**, dorsal laterotergite protuberance; **dpes**, dorsal phallothecal sclerite-endosomal struts fusion; **dps**, dorsal phallothecal sclerite; **en**, endosoma; **es**, endosomal struts; **exM**, extension of M beyond M+Cu distal junction; **fmp**, femoral medial protuberance; **ft**, femoral tubercle; **gl**, gula; **is**, interocular sulcus; **L2**, labial segment II (first visible segment); **L3**, labial segment III (second visible segment); **L4**, labial segment IV (third visible segment); **lb**, labrum; **M**, media; **MGE**, metathoracic gland evaporatorium; **mms**, transverse suture between meso- and metasterna; **mo**, median oviduct; **mpp**, medial pygophore process; **mss**, mesosternum; **mtc**, metacoxa; **mts**, metasternum; **mxp**, maxillary plate; **nk**, neck; **oc**, ocelli; **ot**, ocellar tubercle; **p**, pedicel; **pa**, paramere; **pap**, trochanter and/or femoral papillae; **pc**, sternal paramedian carinae; **pcd**, postclypeal depression; **pfsp**, posterior femoral subapical protuberance; **pltf**, lateral furrow of posterior pronotal lobe; **pls**, pronotal longitudinal sulcus; **poc**, postocular; **ppl**, posterior pronotal lobe; **pts**, pronotal transverse suture; **R**, radius; **s**, scape; **S2**, sternite 2; **S3**, sternite 3; **sc**, scutellum; **sis**, sternal intersegmental suture; **sl**, synthlipsis; **sld**, sternal medial longitudinal depression; **T8**, tergite 8; **T9**, tergite 9; **T10**, tergite 10; **V1**, valvula 1; **V3**, valvula 3; **vap**, ventral anterior process of endosomal struts; **vfl**, valvifer 1; **vpp**, ventral posterior process of endosomal struts.

Abbreviations for depositories: **AMNH**, American Museum of Natural History, New York; **BMNH**, Natural History Museum, London, United Kingdom; **CAS**, California Academy of Sciences, California; **HNHM**, Hungarian Natural History Museum, Budapest, Hungary; **MMBC**, Moravské Museum, Brno, Czech Republic; **MNHN**, Muséum National d'Histoire Naturelle, Paris, France; **MRAC**, Musée Royal de l'Afrique Centrale, Tervuren, Belgium; **MTEC**, Montana State University, Montana; **NMW**, Naturhistorisches Museum, Vienna, Austria; **SU**, Department of Zoology, University of Silesia, Poland; **TLMF**, Tiroler Landesmuseum Ferdinandeum,

Innsbruck, Austria; **UCR**, University of California, Riverside Entomological Research Museum, California; **USNM**, National Museum of Natural History, Washington D.C.

### **Results of the molecular association of males and females**

Pairwise genetic distances of COI between intraspecific males were less than 3.75%. Between interspecific males within a genus or between genera, COI pairwise genetic distances were greater than 7.10%. Of the 10 females that were sequenced for COI, two had pairwise genetic distances with two male-based species that were less than 0.80% (*Toxopus griswoldi*: 0.76% [Figs. 2.3, 2.4, 2.7, 2.10]; an undescribed species of *Distirogaster*: 0.75%). These two results were corroborated by examination of geographic distance between male and female specimens (*Toxopus griswoldi*: same locality; *Distirogaster* sp.: 2.28 km), as well as morphology (see taxonomic descriptions and discussions). For the remaining eight females (Fig. 2.24) sequenced, female-to-male pairwise distances exceeded 9.5%, and therefore, COI sequences did not offer a way of associating these females with conspecific males. In addition, these females were collected at localities that were near those of males belonging to several species, such that geographic proximity could not be used as a line of evidence. Generic and species diagnoses based on males heavily rely on structural differences on the head, thorax, and wings, among other features. However, all eight females have drastically reduced morphological features (e.g., head structures, scutellum, wings, etc.), and three of these females could not be unambiguously associated with any of the male-based genera treated in this monograph. The remaining five females could be associated with the genus *Gibbosella* based on morphology (see generic discussion) but could not be unambiguously associated to any male-based species within this genus.

## Taxonomy

### Identification key to Ectrichodiinae genera of Madagascar

1. Abdominal spiracles circular; body small (<10 mm); dull colored .....2
  - Abdominal spiracles ovoid to elliptical; body medium to large (≥10 mm); color variable .....3
2. Male: Metathoracic gland evaporatorium (MGE) not visible in lateral view (Fig. 2.16B, C); scutellar processes dorsally directed in lateral view (except in *Gibbosella planiscutum*) (Fig. 2.15C); hemelytron with distal part of M and Cu fused basally (Fig. 2.17A); dorsal laterotergites with posterior protuberances (Fig. 2.18A). Female: apterous; 6- or 8-segmented antenna with short vestiture; dorsal laterotergites with posterior protuberances reduced to absent. Bicolored, dark brown and pale in both sexes ..... *Gibbosella* Chłond
  - Males only known. MGE visible in lateral view (Fig. 2.16A); scutellar processes horizontally directed (i.e., parallel to dorsal abdominal surface) in lateral view (Fig. 2.15D); hemelytron with distal part of M and Cu separate basally (Fig. 2.17B, C); dorsal laterotergites unarmed (Fig. 2.18B); coloration generally uniform, orange-brown ..... *Marojejycoris*, new genus
3. Distiflagellomere (DFLA) (Fig. 2.14A, C) divided into four pseudosegments; antenna thus appearing 8-segmented; very pronounced anterolateral protuberances on anterior pronotal lobe (Fig. 2.15A); metallic coloration .....4
  - DFLA divided into two or three pseudosegments; antenna thus appearing 6- or 7-segmented; anterolateral protuberances on anterior pronotal lobe much smaller or absent; coloration not metallic .....5
4. Dorsal laterotergites with posterior protuberances (Fig. 2.18A); abdominal sternites with paramedian, longitudinal carinae and shallow medial longitudinal depression (Fig. 2.18C). Male: macropterous; subquadrate head in lateral view; relatively slender abdomen. Female: apterous; more ovate head shape; abdomen wider ..... *Distirogaster* Horváth

- Dorsal laterotergites without posterior protuberances; abdominal sternites without paramedian longitudinal carinae and shallow medial longitudinal depression; head ovoid (Fig. 2.13E); abdomen wide; males and females of the only Madagascan species apterous  
.....*Maraenaspis* Karsch
- 5. Forefemur with anterior subapical and medial protuberances and mid and hind femora with anterior and posterior subapical and medial protuberances in both sexes (Fig. 2.16E); body glabrous except on antenna and tibiae; DFLA divided into two pseudosegments, thus, antenna appearing 6-segmented ..... *Glymmatophora* Stål
- Femora unarmed in males, with small papillae (Fig. 2.16D) in females; body with dense vestiture; DFLA divided into two or three pseudosegments, thus, antenna appearing 6- or 7-segmented .....6
- 6. Males only known; meso- and metasterna completely divided by a transverse suture (Fig. 2.15F); dorsal laterotergite II not distinctly expanded; head subtriangular in lateral view (Fig. 2.13G–I).....7
- Male: meso- and metasterna not completely divided by a transverse suture (Fig. 2.15E); dorsal laterotergite II distinctly expanded; head short and subquadrate (Fig. 2.13K) to subovate or elongate cylindrical (Fig. 2.13J). Female: apterous; shiny blackish coloration, sometimes with pale yellow markings; 7-segmented antenna with short vestiture; fore and mid trochanters with small papillae (Fig. 2.16D) ..... *Toxopus* Bergroth
- 7. DFLA divided into three pseudosegments; antenna thus appearing 7-segmented (except 6-segmented in some specimens of *Tanindrazanus harinhali*); posterior pronotal lobe smooth or transversely striated ..... *Tanindrazanus*, new genus
- DFLA divided into two pseudosegments; antenna thus appearing 6-segmented; posterior pronotal lobe punctate.....*Toliarus*, new genus

*Gibbosella* Chłond, 2010

Figs. 2.1, 2.5, 2.8, 2.12, 2.13C, 2.15C, 2.16D, 2.17A, 2.18A, 2.19A, 2.19B, 2.20A–C, 2.21A, 2.21B, 2.22A, 2.23A, 2.23B, 2.24; Maps 2.1–2.3

*Gibbosella* Chłond, 2010: 62.

TYPE SPECIES: *Gibbosella mirabilis* Chłond, 2010

REVISED DIAGNOSIS: Males recognized by the small body size, shallow anteromedial depression on the ventral head surface, large antennal shield that does not conceal the antennal insertion in lateral view, depressed postclypeus, 8-segmented antenna (6-segmented in *G. pallidalata*), dorsally oriented long and slender scutellar processes (short, stout, and horizontally oriented in *G. planiscutum*), basally fused distal part of M and Cu, dorsal laterotergites transversely bicolored pale yellow and brown and with posterolateral protuberances, and apex of abdomen with very long setae. Females recognized by the apterous condition, reduced pale markings, head about as wide as anterior margin of pronotum, 6- or 8-segmented antenna with short vestiture, scape (Fig. 2.14A) longer than distance between anterior margin of eye and apex of head, anterior pronotal lobe much longer than posterior lobe, scutellar processes reduced and dorsally directed, and dorsal laterotergites with posterior protuberances reduced or absent. Males are similar to males of the monotypic Afrotropical genus *Synavecoris* Villiers, 1968b (Fig. 2.12), but the presence of the antennal shield, lack of femoral armature, separation of the proximal part of M and Cu (Fig. 2.17A), and armature of the dorsal laterotergites differentiate males of *Gibbosella* from *Synavecoris*. and armature of the dorsal laterotergites differentiate males of *Gibbosella* from *Synavecoris*. Due to reduced morphological features in both genera, females of *Gibbosella* appear very similar to females of *Synavecoris* but may be distinguished by the apterous condition, dorsally directed scutellar processes, and the pronotum lacking lateral carinae.

REDESCRIPTION: **MALE:** Macropterous, small body size. COLORATION: Pale to dark brown color patterns. VESTITURE: Sparse to dense, semierect to erect, long pale to brown setae on head, thorax, legs, corium of hemelytra, and abdomen; setae longer on apex of abdomen compared to rest of vestiture on body; setae on tibiae stouter and denser near apex.

STRUCTURE: HEAD (Fig. 2.13C): Circular, ovoid, or cylindrical; shorter than pronotum; ventrally with shallow anteromedial depression; clypeal apex not dorsally elevated relative to labrum; maxillary plate not reaching dorsal clypeal surface (Fig. 2.13E, I); postocular (hind eye margin to posterior constriction; Fig. 2.13J) broad in dorsal view (Fig. 2.13B) ocelli present, separated by less than diameter of ocellus; distinct constriction between postocular and neck; antennal shield not concealing antennal insertion in lateral view; scape surpassing clypeal apex; pedicel (Fig. 2.14A) slightly curved; flagellum subdivided into BFLA (Fig. 2.14A, B) and DFLA (Fig. 2.14A, C); BFLA divided into two pseudosegments; DFLA divided into two or four pseudosegments; antenna thus appearing 6- or 8-segmented. THORAX (Fig. 2.15C): Pronotum wider than long, anterior margin distinctly concave, smooth, collar distinct (Fig. 2.15B); anterior pronotal lobe shorter than posterior lobe, more than half as wide as posterior lobe; pronotal longitudinal furrow reaching anterior but not posterior margin of pronotum, foveate posteriorly (Fig. 2.15B); pronotal transverse furrow distinct; lateral depressions on posterior pronotal lobe distinct; scutellum with two moderately separated apical processes, directed dorsally in most species (horizontally directed in *G. planiscutum*), disc medially depressed; MGE with shallow meshlike cuticle that does not extend dorsally in lateral view (Fig. 2.16B, C); fossula spongiosa on foretibia, absent on mid tibia in some species; hemelytron (Fig. 2.17A) with corium restricted to areas adjacent to basal wing veins, with pterostigmalike appearance on anterodistal margin; proximal parts of M and Cu veins separate; distal parts of M and Cu fused basally; distal part of M extending beyond apical junction of



M+Cu. ABDOMEN (Fig. 2.18A): Dorsal laterotergites II-VI with posterolateral protuberances; sternal intersegmental sutures (Fig. 2.18B, C) carinulate; spiracles circular; pygophore process directed dorsoposteriad, not surpassing posterior margin of pygophore; DPS apex rounded (Fig. 2.21A, B); endosomal struts with anterior ventral process (Fig. 2.22G); endosoma weakly to strongly sclerotized medially (Fig. 2.21A, B). **FEMALE:** Differs from males in the following characteristics: apterous; reduced pale markings; head about as wide as anterior pronotal margin; in some species, head ventrally with small tubercles; postclypeus not depressed; antennal shield not expanded; ocelli absent; eye small; antennal vestiture much shorter; scape longer than distance between anterior margin of eye and apex of head; collar not distinct; anterior pronotal lobe much longer than and as wide as or wider than posterior lobe (Fig. 2.15A), sometimes elevated dorsally; pronotal longitudinal furrow reduced to deep medial depression near posterior margin of anterior pronotal lobe (Fig. 2.15A) and present or obsolete on posterior lobe; lateral depressions on posterior pronotal lobe present or obsolete; scutellum processes weakly developed, dorsally directed; meso- and metasterna convex; in some species, fore- and mid trochanters and base of forefemur with ventral patches of small papillae (Fig. 2.16D) and femora ventrally with small tubercles (Fig. 2.16D); hind femur slightly curved in dorsal view; fossula spongiosa larger; dorsal laterotergites with posterior protuberances reduced or absent; external genitalia short, platelike (Fig. 2.23A, B).

**DISTRIBUTION:** Species are known from all provinces and occur in habitats between 9–1,600 m elevation. Macrohabitats are described as spiny forests, tropical forests, mixed tropical forests, low and high altitude rainforests, tropical dry forests, montane rainforest, sclerophyl forests, dwarf littoral forests, and secondary tropical forests.

**DISCUSSION:** Chłond (2010a) recently described this genus and two species, *G. mirabilis* and *G. elongata*, from single female representatives with no males or immatures known. One male

specimen from undetermined material showed some morphological features similar to the described females, e.g., an elongate cylindrical head, a slightly dorsally pronounced anterior pronotal lobe, dorsally directed scutellar processes, general color pattern, and body size. Given the morphological similarities, we assign this male specimen to the genus and to the species *G. elongata* (see species redescription for discussion on species-level assignment). Examination of other undetermined male specimens revealed similar morphology to the previously mentioned male: general coloration, body size, dorsally directed scutellar processes, slightly to very distinctly dorsally pronounced anterior pronotal lobe, and wing venation pattern, among many other features. Given the morphological similarities between these males and the male of *G. elongata*, we assign these males to *Gibbosella* despite the fact that head shapes in these males are not elongated but rather spherical or short and ovoid.

Nineteen female specimens have been identified as representatives of this genus based on morphological features mentioned in the generic redescription, e.g., small body sizes, antennal segmentation, slightly dorsally projecting scutellar processes, and dark coloration. Despite availability of morphological, molecular, and geographic data, we were unable to associate 16 of them with male-based species. The remaining three female specimens were associated with two male-based species (*G. brunalvus* and *G. planiscutum*) using morphology and geographic data (see species discussions).

There is scutellar and antennal variation among species in this genus: the short, stout scutellar processes in *G. planiscutum* are horizontally oriented (Fig. 2.15D) and the antenna 6-segmented in *G. pallidalata*. As previously mentioned, head shape is also variable among *Gibbosella* species, ranging from spherical or ovoid in most species to elongate and cylindrical in two previously described species.

This genus is very similar to *Synavecoris* by the small size, general dull coloration, ventral head depression restricted to the anteocular (apex of clypeus to anterior eye margin; Fig. 2.13K) region, moderately separated scutellar processes, shallowly depressed MGE that does not extend dorsally in lateral view, basally fused M and Cu in the distal part of the hemelytron, and circular spiracles, but is distinguished by the characters mentioned in the diagnosis. Maldonado (1990) incorrectly listed 1953 as the year *Synavecoris* was described and indicated that the description was based on “nymphs, probably of *Ectrichodia*.” Villiers (1968b) described *Synavecoris* based on one macropterous adult male (Fig. 2.12) and two micropterous adult females; nymphs of *Synavecoris* were not described by Villiers. The small size of *Synavecoris*, relatively slender legs, pronotal structure, and wing venation patterns distinguish *Synavecoris* from *Ectrichodia* Lepeletier & Serville, 1825, and thus, we considered *Synavecoris* a valid genus.

*Identification key to the males of species of Gibbosella*

- 1. Scutellar processes slender, long, and dorsally directed (Fig. 2.15C) .....2
- Scutellar processes short, stout, and horizontally directed (Fig. 2.15D) .....  
.....*planiscutum*, new species
- 2. Head circular or ovoid in lateral view, not distinctly elongate and cylindrical .....3
- Head very elongate, cylindrical in lateral view ..... *elongata* Chłond
- 3. DFLA divided into four pseudosegments; antenna thus appearing 8-segmented.....4
- DFLA divided into two pseudosegments; antenna thus appearing 6-segmented.....  
.....*pallidalata*, new species
- 4. Head as long as wide.....5
- Head longer than wide.....9

5. Head ovoid to subpentagonal; labial segment II longer than III; anterior pronotal lobe not laterally carinate.....	6
- Head nearly circular in lateral view (Fig. 2.13C); labial segments II and III subequal in length; anterior pronotal lobe slightly carinate laterally .....	<i>pallidacorium</i> , new species
6. Synthlipsis about two times width of eye .....	7
- Synthlipsis less than two times width of eye .....	8
7. Head subpentagonal; eye not reaching dorsal and ventral head surfaces; fossula spongiosa absent on mid tibia; body length <7 mm .....	<i>nitida</i> , new species
- Head ovoid; eye almost reaching dorsal and ventral head surfaces; fossula spongiosa present on mid tibia; body length >7 mm.....	<i>fulva</i> , new species
8. Synthlipsis about width of eye; fossula spongiosa absent on mid tibia .....	<i>betampona</i> , new species
.....	
- Synthlipsis less than width of eye; fossula spongiosa present on mid tibia .....	<i>mantella</i> , new species
.....	
9. Antecular region shorter than postocular.....	10
- Antecular region as long as postocular (Fig. 2.13C).....	11
10. Postclypeus not medially depressed (Fig. 2.13A); synthlipsis width about 2.5 times width of eye; sternal intersegmental sutures carinulate between II–IV .....	<i>andasibe</i> , new species
- Postclypeus medially depressed (Fig. 2.13B); synthlipsis width about 3.5 times width of eye; sternal intersegmental sutures carinulate between II–VI.....	<i>megafrons</i> , new species
11. Ventral margin of labial segment III convex (Fig. 2.13C); anterior pronotal lobe not dorsally projecting (Fig. 2.15D); mesosternum with medial and lateral depressions.....	12

- Ventral margin of labial segment III straight (Fig. 2.13G–I, K); anterior pronotal lobe dorsally projecting above pronotal disc (Fig. 2.15C); mesosternum with one large, shallow medial depression ..... 14
- 12. Postclypeus distinctly depressed (Fig. 2.13B); synthlipsis width 1.5 times width of eye; pronotal transverse suture complete, not divided by paramedian ridges; vestiture sparse ..... 13
- Postclypeus not distinctly depressed (Fig. 2.13A); synthlipsis width two times width of eye; pronotal transverse suture incomplete, divided by paramedian ridges (Fig. 2.15B); vestiture dense ..... *brunalvus*, new species
- 13. Labial segment II longer than III; abdominal sternites convex, keellike; median pygophore process spadelike in caudal view (Fig. 2.19B) ..... *vangocris*, new species
- Labial segment II and III subequal in length; abdominal sternites with shallow medial longitudinal depression (Fig. 2.18C); median pygophore process subquadrate in caudal view (Fig. 2.19A) ..... *quadocris*, new species
- 14. Synthlipsis width about 1.5 times width of eye; eye not reaching dorsal and ventral head surfaces; sternal intersegmental sutures carinulate between II and III and laterally between III–VI ..... *conosimilis*, new species
- Synthlipsis width about width of eye; eye reaching dorsal and ventral head surfaces; sternal intersegmental sutures carinulate between II–IV and laterally between IV–VI ..... *notoconica*, new species

*Identification key to the females of species of Gibbosella*

- 1. Head elongate, cylindrical (Fig. 2.12); anterior pronotal lobe strongly elevated dorsally in lateral view; body length >8 mm ..... 2

- Head shorter, ovate (Fig. 2.13F); anterior pronotal lobe slightly to moderately elevated dorsally in lateral view; body length <8 mm .....3
- 2. Scutellar processes convergent; hind tibia with dark apical half..... *mirabilis* Chłond
- Scutellar processes not convergent; hind tibia yellowish with dark annulus in basal part.....  
..... *elongata* Chłond
- 3. Synthlipsis width three times width of eye; generally brown to dark brown with scutellar apical projections whitish to pale ..... *planiscutum*, new species
- Synthlipsis width two times width of eye; generally brown to dark brown with scutellar apical projections brown to dark brown ..... *brunalvus*, new species

*Gibbosella andasibe*, new species

Figs. 2.1, 2.5, 2.8; Map 2.2

DIAGNOSIS: Males recognized among other species in this genus by the very small body size, coloration, antecular region shorter than the postocular, postclypeus not depressed, synthlipsis width about 2.5 times the width of an eye, and fossula spongiosa absent on mid tibia. This species is similar to *G. megafrons*, from which it differs by the light brown anterior pronotal lobe, brown posterolateral margin of pronotum, pale posterior half of scutellum, the flat postclypeus, smaller synthlipsis width, lack of a fossula spongiosa on the mid tibia, and intersegmental sutures carinulate between sternites II–IV.

DESCRIPTION: **MALE:** Body length: 3.64 mm (holotype). COLORATION: Dark brown with antennal segments VII and VIII, posterolateral margin of pronotum, pleura dorsad of coxae, spot on dorsoposterior margin of mesopleuron, posterior half of scutellum, corium basally, coxae, trochanters, femora basally, laterotergites, except posterior half of VII, and sternites medially and anterolaterally pale. Wing membrane, remainder of antenna, legs, and corium brown. Labium,

neck dorsally, and anterior pronotal lobe light brown. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region shorter than postocular; gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus flat (Fig. 2.13A); synthlipsis about 2.5 times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel as long as scape; antenna 8-segmented; labium stout; labial segment III subequal to II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections slightly dorsally directed (Fig. 2.15C); meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); mesosternum with medial and paramedial longitudinal depressions; metasternum slightly medially longitudinally depressed posteriorly; legs slender; fossula spongiosa absent on mid tibia; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex; distal part of R absent. ABDOMEN: Apex medially notched; sternites medially convex, intersegmental sutures carinulate between II–IV; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate.

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Andasibe National Park, Madagascar.

DISTRIBUTION: The single known specimen was collected southeast of Andasibe National Park headquarters in the Toamasina province (Map 2.2).

HOLOTYPE: Male: **Madagascar: Toamasina:** 7 km SE of Andasibe National Park headquarters, 18.96266°S 48.45266°E, 1050 m, 07 Jan 2001–22 Jan 2001, M. Irwin, R. Harin'Hala (00044816) (CAS).

*Gibbosella betampona*, new species

Figs. 2.1, 2.5, 2.8; Map 2.3

DIAGNOSIS: Males recognized among other species in *Gibbosella* by the circular head shape in lateral view that is as long as it is wide in dorsal view, large ocelli, eyes reaching dorsal and ventral head margin, labial segment II longer than III, mesosternum with large medial depression, absence of the fossula spongiosa on the mid tibia, and BPE shorter than the basal plate. This species is similar to *G. mantella*, but the pale posterior margin of the pronotum and lateral spots on the meso- and metasterna, the brown corium, larger synthlipsis width, meso- and metasterna not completely separated by a distinct suture, absence of the fossula spongiosa on the mid tibia, and shorter BPE relative to the basal plate distinguish this species from *G. mantella*.

DESCRIPTION: **MALE:** Body length: 6.40 mm (holotype). COLORATION: Dark brown with pale labial segment IV, antennal segment V apically, antennal segments VI–VIII, lateral spots on meso- and metasterna, coxae, trochanters, forefemur basally, mid and hind femora basally and subapically, foretibia except medially, mid and hind tibiae, tarsi, anterior half of laterotergites, paramedian spots on sternites III–V, and lateral margin of sternite VII. Remainder of labium, corium, and basal wing veins brown. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13C); as long as wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about width of eye; interocular



sulcus near hind margin of eye (Fig. 2.13B); ocelli large, located on distinct median tubercle (Fig. 2.13C); eye about half of head length, not reaching dorsal head surface, reaching ventral head surface; antenna inserted dorsally on head; pedicel about as long as scape; antenna 8-segmented; labium stout; labial segment III shorter than II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna not completely or weakly separated by distinct transverse suture (Fig. 2.15E); mesosternum with large medial depression; metasternum slightly medially longitudinally depressed; forefemur slightly incrassate; fossula spongiosa absent on mid tibia; tarsomeres I and II combined shorter than III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex medially notched; sternites flat, intersegmental sutures carinulate between II–IV and laterally between IV–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G); area of endosomal struts–DPS fusion ovate (Fig. 2.21A).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the locality in which the two known specimens were collected, Reserve Betampona, Madagascar.

DISTRIBUTION: Betampona Reserve in the Toamasina province (Map 2.3).

HOLOTYPE: Male: **Madagascar: Toamasina:** Reserve Betampona, Camp Vohitsivalana, 37.1 km 338 Toamasina, 17.88666°S 49.2025°E, 520 m, 01 Dec 2005–03 Dec 2005, Fisher et al. (00006143) (CAS).

PARATYPE: **Madagascar: Toamasina:** Reserve Betampona, Camp Vohitsivalana, 37.1 km  
338 Toamasina, 17.88666°S 49.2025°E, 520 m, 01 Dec 2005–03 Dec 2005, Fisher et al., 1♂  
(00007255) (CAS).

*Gibbosella brunalvus*, new species

Figs. 2.1, 2.5, 2.8, 2.23A; Map 2.1

DIAGNOSIS: Males are recognized among others in *Gibbosella* by a combination of the following characters: the coloration, dense vestiture, postclypeus not depressed, pronotal transverse suture divided by paramedian ridges, and scutellar processes dorsally directed. Females are recognized by the coloration and synthlipsis about two times the width of an eye. Males are very similar to *G. fulva*, from which *G. brunalvus* differs by the more abundant pale markings on the pronotum, scutellum, corium, and sternites, smaller body size, dense vestiture, head longer than wide, flat postclypeus, and divided pronotal transverse furrow, among several other features. Females are similar to *G. planiscutum* based on the features mention in the generic description, but are distinguished by the dark brown antennal segment VIII and scutellum, femora basally and tibiae apically and basally pale to light brown, and smaller synthlipsis width.

DESCRIPTION: **MALE:** Body length: 4.89 mm (holotype), 4.79–5.13 mm. **COLORATION:** Dark brown with pale color on posterolateral margin of pronotum, scutellar processes, basal half of corium, coxae ventrally, trochanters, femora and tibiae basally, tarsi, anterior half of laterotergites, and sternites anterolaterally pale. Antennal segment V apically and segments VI–VIII white. Dorsal head surface, except synthlipsis and ocellar tubercle, markings throughout anterior pronotal lobe, labium, and antenna brown. **VESTITURE:** Dense; other features as in generic description. **STRUCTURE: HEAD:** Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; antecular region as long as postocular (Fig. 2.13C); gula moderately swollen

ventrolaterally, not distinctly produced beyond ventral head margin (Fig. 2.13E, G–K); postclypeus flat (Fig. 2.13A); synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about as long as scape; antenna 8-segmented; labium stout; labial segment III shorter than II, ventrally convex. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); mesosternum with medial and paramedial longitudinal depressions; metasternum medially longitudinally depressed; forefemur incrassate; fossula spongiosa on mid tibia; tarsomeres I and II combined subequal to III; hemelytron reaching or surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex rounded; sternites convex, shallowly depressed between paramedian ridges on sternites II–V, intersegmental sutures carinulate between II–IV and laterally between IV–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G), fused anterior to fusion with DPS; area of endosomal struts–DPS fusion circular. **FEMALE:** Body length: 4.54 mm. Similar to males, but differ in the following characters: COLORATION: Dark brown with pale to light brown scape, pedicel, antennal segments III–IV and basal half of V, labium, femora except basal, and tibiae except apical and basal parts. Remaining antennal segments, coxae, trochanters, femora basally, tibiae apically and basally, and tarsi whitish to pale brown. STRUCTURE: In addition to characters mentioned in the generic description: HEAD: Synthlipsis about two times width of eye. THORAX:

Pronotal longitudinal furrow restricted to anterior lobe; lateral depressions on posterior pronotal lobe obsolete; forefemur with small ventral tubercles (Fig. 2.16D). ABDOMEN: Intersegmental sutures carinulate between sternites II–VI; external genitalia as in Fig. 2.23A; bursa copulatrix membranous, with lateral lobes (Fig. 2.23F); vermiform gland and lateral spermathecae damaged.

ETYMOLOGY: The species epithet is a noun in the nominative case and is named after the nearly uniform dark brown venter.

DISTRIBUTION: Tsingy de Bemaraha National Park in Mahajanga province (Map 2.1).

DISCUSSION: The coloration is entirely dark brown on the head and anterior pronotal lobe, pale to brown medial stripes on the abdominal sternites in some male specimens. A female specimen was associated with males based on morphology and the same collection event as one other male.

HOLOTYPE: Male: **Madagascar: Mahajanga:** Parc National Tsingy de Bemaraha, 3.4 km 93°E Bekopaka, Tombeau Vazimba, 19.14194°S 44.82805°E, 50 m, 06 Nov 2001–10 Nov 2001, Fisher et al. (00006317) (CAS).

PARATYPES: **Madagascar: Mahajanga:** Parc National Tsingy de Bemaraha, 3.4 km 93°E Bekopaka, Tombeau Vazimba, 19.14194°S 44.82805°E, 50 m, 06 Nov 2001–10 Nov 2001, Fisher et al., 3♂ (00006122, 00006318, 00099043) (UCR), 3♂ (00006319, 00006321, 00099044) (SU), 5♂ (00006322–00006326) (CAS), 2♂ (00006327, 00006328) (BMNH), 2♂ (00006329, 00006455) (USNM). Parc National Tsingy de Bemaraha, 10.6 km ESE 123° Antsalova, 19.70944°S 44.71806°E, 150 m, 16 Nov 2001–20 Nov 2001, Fisher et al., 1♂ (00045686), 1♀ (00048066) (CAS). **Unknown:** 2♂ (00007154, 00044944) (CAS).

*Gibbosella conisimilis*, new species

Figs. 2.1, 2.5, 2.8, 2.15C, 2.18A; Map 2.1

DIAGNOSIS: Males are recognized among other species in *Gibbosella* by the larger body size, synthlipsis width 1.5 times the width of an eye, labial segment III ventrally straight, conically elevated anterior pronotal paramedian lobes in lateral view, anterior pronotal lobe without anterolateral protuberances, continuous pronotal transverse suture, and mesosternum with large medial depression. This species is similar to *G. notoconica*, but the slightly smaller body size, pale apex of antennal segment V, dark brown pronotum (except anterior and posterolateral margins and disc) and mesosternum (except anterolaterally and posteriorly), larger synthlipsis width, interocular sulcus posterior to the hind margin of the eye, smaller ocelli, eye size relative to the head length and height, and intersegmental sutures carinulate between II and III and laterally between III–VI distinguish this species from *G. notoconica*. The conical anterior pronotal paramedian lobes are also not as distinct as in *G. notoconica*.

DESCRIPTION: **MALE:** Body length: 7.92 mm (holotype), 7.86–8.44 mm. COLORATION: Dark brown with pale dorsal head surface (except clypeal apex, mandibular plates, antennifers, anterior of interocular sulcus, interocular sulcus, and dorsal surface of ocellar tubercle), scape basally, antennal segment V apically, and segments VI–VIII, anterior and posterolateral margins of pronotum, pronotal disc, lateral longitudinal depressions of posterior pronotal lobe, stridulatory groove, mesosternum anterolaterally and posteriorly, metasternum, scutellum medially, corium basally, coxae, trochanters, femora basally and ventrally, mid and hind tibiae medially, tarsi, anterior margin of laterotergites, and sternites medially and anterolaterally. VESTITURE: Very sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with shallow, broad

medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about 1.5 times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about one-third longer than scape; antenna 8-segmented; labium slender; labial segment III subequal to II, ventrally straight (Fig. 2.13G, H, K). THORAX (Fig. 2.15C): Anterior pronotal lobe slightly conical in lateral view, without distinct anterolateral projections; pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed; meso- and metasterna not completely or weakly separated by distinct transverse suture (Fig. 2.15E); mesosternum with large medial depression; metasternum slightly medially longitudinally depressed; legs slender; fossula spongiosa on mid tibia; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN (Fig. 2.18A): Apex medially notched or sinuate; sternites medially convex, intersegmental sutures carinulate between II and III and laterally between III–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS, ventrally obscured by semiextended endosoma; area of endosomal struts–DPS fusion ovate (Fig. 2.21A).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the similar pronotal appearance of the pronotum in *G. notoconica*.

DISTRIBUTION: Marojejy Nature Reserve in Antsiranana province (Map 2.1).

DISCUSSION: Coloration is variable with the dark brown area around the ocellar tubercle and interocular sulcus variable in size. The anterior pronotal lobe may be dorsally pale, and the pronotal transverse sulcus entirely or nearly entirely pale.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 26 Sep 2005–04 Oct 2005, M. Irwin, R. Harin'Hala (00044850) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 15 Dec 2004–20 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00007113) (CAS), 1♂ (00045329) (SU); 25 Dec 2004–30 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00007260) (UCR); 10 Jan 2005–15 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00045259) (CAS); 25 Feb 2005–04 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00007066) (CAS); 25 Mar 2005–04 Apr 2005, M. Irwin, R. Harin'Hala, 1♂ (00045427) (CAS); 04 Apr 2005–16 Apr 2005, M. Irwin, R. Harin'Hala, 2♂ (00007079, 00007083) (CAS), 2♂ (00007150, 00007199) (BMNH); 18 May 2005–30 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00044819) (CAS); 30 May 2005–11 Jun 2005, M. Irwin, R. Harin'Hala, 1♂ (00045317) (SU); 11 Jun 2005–28 Jun 2005, M. Irwin, R. Harin'Hala, 1♂ (00045495) (UCR); 14 Oct 2005–22 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00007054) (AMNH). R.N.I. de Marojejy, 8.0 km NW Manantenina, 14.43667°S 49.775°E, 450 m, 05 Oct 1996–13 Oct 1996, E. Quinter and T. Nguyen, 1♂ (00078363) (AMNH). **Unknown:** 1♂ (00007064) (CAS).

*Gibbosella elongata*, Chłond, 2010

Figs. 2.1, 2.5, 2.12; Map 2.2

*Gibbosella elongata* Chłond, 2010: 64.

REVISED DIAGNOSIS: Males recognized among other species in this genus by the larger body size, synthlipsis width same as width of an eye, labial segment III ventrally convex, elevated anterior pronotal paramedian lobes in lateral view, anterior pronotal lobe with small anterolateral projections. Females are recognized by the cylindrical head, scape surpassing apex of the head,

and very gibbous anterior pronotal lobe with small anterolateral projections. This species can be easily distinguished from *G. mirabilis* by the enlarged, elongated, and much more elevated anterior pronotal lobe and lack of a depression in the posterior part of anterior pronotal lobe.

REDESCRIPTION: **MALE:** Body length: 9.90 mm. COLORATION: Dark brown with pale basal part of scape, coxa, trochanters, basal part of forefemur, basal and apical part of mid and hind femur, apical and middle part of mid tibiae, hind tibiae except small subapical ring, anterior part of laterotergites III-VII and middle part of abdominal tergites III-VII. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Elongate ovoid to cylindrical in lateral view; longer than wide in dorsal view; antecular region longer than postocular; gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with deep, narrow medial longitudinal depression to middle of interocular area; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli large sized, located on distinct median tubercle; eye about one-third of head length, reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel slightly longer than scape; labium stout; labial segment III shorter than II, ventrally convex. THORAX: Anterior pronotal lobe gibbous (Fig. 2.5) with small anterolateral projections; pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe striated transversely; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); forefemur incrassate; fossula spongiosa on mid tibia; tarsomeres I and II combined same length as III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Sternites medially convex, intersegmental sutures carinulate between II-IV; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE as long as basal plate; area of endosomal struts-DPS subquadrate. **FEMALE:** Body length: 9.40 mm (holotype). Similar to males, but differ in the following characters:



COLORATION: Shining, dark brown with pale labrum, DFLA, small spots on pronotum and thorax, coxae, trochanters, basal half of forefemur and two-thirds of mid and hind femora, as well as apical part of all femora, tibiae and anterior part of laterotergites. STRUCTURE: In addition to characters mentioned in the generic description: HEAD (damaged): Slightly convex gula.

THORAX: Anterior pronotal lobe gibbous with small anterolateral projections (Fig. 2.12); visible medial longitudinal sutures on both pronotal lobes; lateral depressions on posterior pronotal lobe striated transversely. ABDOMEN: Tergites laterally and laterotergites striated transversely; tergites posterior distinctly rounded, except tergite III; sternites with distinct visible fold near lateral margin; intersegmental sutures carinulate between sternites II-VII; valvifer I triangular with flat dorsal part and one long setae; valvula I small with long hairs placed on small fold visible in middle part; valvula III elongate, convolute in apical part with distinct dome-shaped elevation.

DISTRIBUTION: The type specimen was collected in the Anjanaharibe-Sud Reserve, west of Andapa, which is located in the southern part of Antsiranana province (Map 2.2). The only known male specimen was collected in Maroantsetra near the Toamasina-Antsiranana border, which is in close proximity to the type locality.

DISCUSSION: Chlond (2010a) recently described this species from a single female representative. An undetermined male specimen was observed to have morphological similarities to *G. elongata* and *G. mirabilis*: elongate, cylindrical head shape; dorsally protruding anterior pronotal lobe; dorsally oriented scutellar processes; and general dark brown to blackish and pale coloration. Although extreme sexual dimorphism exists in *Gibbosella*, we assign the male to *G. elongata* based on the small basal dark brown band on the hind tibia and broader pale bands on the dorsal laterotergites; in *G. mirabilis*, the hind tibia is dark brown on the apical one-third to one-half and the dorsal laterotergites have more narrow pale bands. Furthermore, the male was collected near the border of Toamasina and Antsiranana provinces, in close proximity to the type

locality of *G. elongata*; *G. mirabilis* has been collected from the southwestern region of Toamasina, near the Antananarivo-Toamasina border.

HOLOTYPE: Female: **Madagascar: Antsiranana:** Andapa District, Anjanaharibe-Sud Reserve, 14.69889°S 49.45483°E, 1600 m, 1960, P. Soga (MNHN).

OTHER MATERIAL EXAMINED: **Madagascar: Toamasina:** Maroantsetra/Ambodivoangy, 15.43231°S 49.74013°E, 9 m, no date provided, 1♂ (MNHN).

*Gibbosella fulva*, new species

Figs. 2.1, 2.5; Map 2.2

DIAGNOSIS: Males are recognized among other species in this genus by the shiny brown color of the body, the head about as long as wide in dorsal view, synthlipsis two times the width of an eye, medium sized ocelli, eyes almost reaching dorsal and ventral head margin, labial segment II longer than III, anterior pronotal lobe without anterolateral projections, mesosternum with large medial depression, and fossula spongiosa on the mid tibia. This species is similar to *G. brunalvus*, but can be distinguished by the dark brown pronotum, scutellum, and corium, larger body size, sparse vestiture, head about as long as wide, depressed postclypeus, and pronotal transverse furrow continuous, among several other features.

DESCRIPTION: **MALE:** Body length: 7.40 mm (holotype), 7.37–7.40 mm. COLORATION: Dark brown with labial segments III and IV, antennal segments V–VIII, and anterior part of laterotergites pale. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); about as long as wide in dorsal view; anteocular region as long as postocular (Fig. 2.13E); gula flat, conforming to rounded shape of head (Fig. 2.13E); postclypeus with deep, narrow medial longitudinal depression to middle of interocular area; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B);

ocelli medium sized, located on shallow median tubercle (Fig. 2.13I); eye about one-third of head length, almost reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel as long as scape; antenna 8-segmented; labium stout; labial segment III shorter than II, ventrally convex (Fig. 2.13E). THORAX: Anterior pronotal lobe without anterolateral projections; pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections slightly dorsally directed (Fig. 2.15C); ventrally obscured by card mount; forefemur incrassate; fossula spongiosa on mid tibia; tarsomeres I and II combined shorter than III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex medially notched, sternites medially obscured by card mount, but intersegmental sutures carinulate at least laterally between II–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE as long as basal plate; area of endosomal struts–DPS fusion basally narrow but distally wider and ovate.

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named after the shiny brown color of the body.

DISTRIBUTION: Antanambe and Seranambe in the Mananara District of Toamasina (Map 2.2).

HOLOTYPE: Male: **Madagascar: Toamasina:** Mananara-Nord District, Seranambe, 16.24683°S 49.83916°E, 16 m, Jul 1965, J. Vadon and A. Peyrieras (MNHN).

PARATYPE: **Madagascar: Toamasina:** Mananara-Nord District, Antanambe, 16.43333°S 49.85°E, 8 m, no date provided, J. Vadon and A. Peyrieras, 1♂ (MNHN).

*Gibbosella mantella*, new species

Figs. 2.1, 2.5, 2.8, 2.13C; Map 2.3

DIAGNOSIS: Males recognized among other species in *Gibbosella* by the circular head shape

in lateral view that is as long as it is wide in dorsal view, synthlipsis less than the width of an eye, large ocelli, eyes reaching dorsal and ventral head margin, labial segment II longer than III, mesosternum with large medial depression, fossula spongiosa on the mid tibia, and BPE as long as the basal plate. This species is similar to *G. betampona*, but the slightly darker pronotum, pale meso- and metasterna and corium, smaller synthlipsis width, meso- and metasterna separated by a distinct suture, presence of the fossula spongiosa on the mid tibia, and longer BPE relative to the basal plate distinguish this species from *G. betampona*.

DESCRIPTION: **MALE:** Body length: 5.42 mm (holotype). COLORATION: Dark brown with pale labial segment IV, apical half of antennal segment V, antennal segments VI–VIII, posterior margin of pronotum, stridulatory groove, meso- and metasterna, corium and basal wing veins, coxae, trochanters, femora basally, mid and hind tibiae medially, tarsi, laterotergites except posterior margin, and sternites medially and laterally. Remainder of antenna and labium, dorsal surface of pronotum, and legs brown. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD (Fig. 2.13C): Ovoid in lateral view; as long as wide in dorsal view; anteocular region as long as postocular; gula flat, conforming to rounded shape of head; postclypeus with deep, narrow medial longitudinal depression to middle of interocular area; synthlipsis less than width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli large, located on distinct median tubercle; eye about half of head length, reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel as long as scape; antenna 8-segmented; labium stout; labial segment III shorter than II, ventrally convex. THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); mesosternum with large medial depression; metasternum slightly medially longitudinally

depressed; forefemur incrassate; fossula spongiosa on mid tibia; tarsomeres I and II combined shorter than III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex medially notched; sternites medially convex, intersegmental sutures carinulate between II–IV and laterally between IV–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G); endosomal struts–DPS fusion basally narrow, distally wider and ovate (Fig. 2.21A).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Camp Mantella, Marojejy National Park, Madagascar.

DISTRIBUTION: Marojejy Nature Reserve in Antsiranana province (Map 2.3).

HOLOTYPE: Male: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 11 Feb 2005–18 Feb 2005, M. Irwin, R. Harin’Hala (00006458) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 28 Apr 2005–07 May 2005, M. Irwin, R. Harin’Hala, 1♂ (00006460) (CAS); 14 Oct 2005–22 Oct 2005, M. Irwin, R. Harin’Hala, 1♂ (00007038) (UCR). R.N.I. de Marojejy, 10.0 km NW Manantenina, 14.43333°S 49.76167°E, 750 m, 15 Oct 1996–22 Oct 1996, E. Quinter and T. Nguyen, 1♂ (00078367) (AMNH).

*Gibbosella megafrons*, new species

Figs. 2.1, 2.5, 2.8; Map 2.1

DIAGNOSIS: Males recognized among other species in *Gibbosella* by the very small body size, coloration, antecular region shorter than the postocular, postclypeus depressed, synthlipsis width about 3.5 times the width of an eye, and distal part of R absent. This species is similar to *G.*

*andasibe*, from which it differs by the entirely dark brown pronotum and scutellum, depressed postclypeus, larger synthlipsis width, presence of a fossula spongiosa on the mid tibia, and intersegmental sutures carinulate between II–VI.

DESCRIPTION: **MALE:** Body length: 3.78 mm (holotype), 3.78–5.05 mm. COLORATION: Dark brown with pale antennal segments VI and VII, posterodorsal spot on mesopleuron, spots on pleura dorsad of coxae, corium basally, coxae, trochanters, femora basally, anterior half of laterotergites, sternites III–VI medially, and anterolateral margins of sternites pale. Antennal segments IV and V, labium, scutellar processes, remainder of corium, basal wing veins, and remainder of legs brown. Wing membrane light brown. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region shorter than postocular; gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about 3.5 times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel as long as scape; antenna 8-segmented; labium stout; labial segment III subequal to II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); separation of meso- and metasterna obscured; mesosternum with medial and paramedial longitudinal depressions; metasternum slightly medially longitudinally depressed; forefemur slightly incrassate; presence of fossula spongiosa on mid tibia cannot be determined due to tibial apex obscured; tarsomeres I and II combined shorter than III; hemelytron surpassing abdominal apex; distal part of R absent. ABDOMEN: Apex medially notched; sternites medially convex,

intersegmental sutures carinulate between II–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; basal plate and BPE damaged; endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate.

ETYMOLOGY: The species epithet is a noun in the nominative case and is named for the very large synthlipsis width.

DISTRIBUTION: Ranomafana National Park and Midongy du Sud National Park in the Fianarantsoa province (Map 2.1).

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1110 m, 28 Nov 2001–06 Dec 2001, M. Irwin, R. Harin’Hala (00007168) (CAS).

PARATYPE: **Madagascar: Fianarantsoa:** Sud-Est Midongy, Sud Mont Papango (sud-ouest Befotaka), 23.85000°S 46.98333°E, 1200 m, Mar 1959, Andria Robinson, 1♂ (MNHN).

*Gibbosella mirabilis* Chłond, 2010

Fig. 2.12; Map 2.1

*Gibbosella mirabilis* Chłond, 2010: 63.

DISTRIBUTION: The only known specimen was collected from “Périnet” (currently known as Andasibe-Mantadia National Park) in the Toamasina province (Map 2.1).

DISCUSSION: This species was recently described by Chłond (2010a) from a single female representative. No other female or male specimens have been examined, and, thus, this species is not redescribed. This species is very similar to *G. elongata*, but is distinguished by the more strongly dorsally elevated anterior pronotal lobe, convergent scutellar processes, and hind tibiae with dark coloration on the apical half.

HOLOTYPE: Female: **Madagascar: Toamasina: Périnet (Andasibe)**, 18.82666°S  
48.44778°E, 1119 m, Oct 1972, A. Peyrieras (MNHN).

*Gibbosella nitida*, new species

Figs. 2.1, 2.5; Map 2.2

DIAGNOSIS: Males are recognized among other species in this genus by the small body size, synthlipsis two times the width of an eye, eye one-fourth of head length, fossula spongiosa absent on the mid tibia, and corium with relatively large shiny granulations not covered by setae, as well as small granulations with very short, erected, flattened setae. This species is similar to *G. megafrons*, but is distinguished by the smaller body size, more subpentagonal head that is as long as it is wide, anteocular length subequal to postocular, smaller synthlipsis width, longer labial segment II relative to III, and appearance of the corium.

DESCRIPTION: **MALE:** Body length: 4.75 mm (holotype). COLORATION: Dark brown with labial segment III and IV brown. Labial segment II apically and III basally, basal two-thirds of scape, coxae, trochanters, femora ventrally, and sternites II–VI medially and laterally pale. Corium with shiny areas due to relatively large shiny granulations not covered by setae, as well as dull areas due to very small granulations with very short, erect, flattened setae. VESTITURE: Dense; corium with shorter, flattened setae; other features as in generic description. Hemelytra covered by very dense, small granulations with very short and flattened setae, as well as sparse relatively large granulations without setae. STRUCTURE: HEAD: Ovate, almost pentagonal in lateral view; about as long as wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula slightly swollen ventrolaterally, not distinctly produced beyond ventral head margin (Fig. 2.13F–K); postclypeus with deep, narrow medial longitudinal depression to middle of interocular area; synthlipsis two times width of eye; interocular sulcus near hind margin of eye



(Fig. 2.13B); ocelli large, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel slightly longer than scape; antenna 8-segmented; labium stout; labial segment III shorter than II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe without anterolateral projections; pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); ventrally obscured by card mount; forefemur incrassate; fossula spongiosa absent on mid tibia; tarsomeres I and II combined shorter than III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; sternites medially obscured by card mount, but intersegmental sutures carinulate at least laterally between II–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE as long as basal plate; area of endosomal struts–DPS fusion basally narrow but distally wider and ovate subquadrate.

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the shiny hemelytra.

DISTRIBUTION: The only known specimen was collected in the Marojejy Nature Reserve in that Antsiranana province (Map 2.2).

HOLOTYPE: Male: **Madagascar: Antsiranana:** Sambava District, Marojejy, Ambinanitelo, 14.44133°S 49.73908°E, 500 m, Dec 1958, Raharizonina (MNHN).

*Gibbosella notoconica*, new species

Figs. 2.1, 2.5, 2.8, 2.20A, 2.21A; Map 2.2

DIAGNOSIS: Males are recognized among other species in this genus by a combination of the following characters: the larger body size, synthlipsis as wide as an eye, large ocelli, eyes reaching the dorsal and ventral head margins in lateral view, labial segment III ventrally straight,

conically elevated anterior pronotal paramedian lobes in lateral view, anterior pronotal lobe without anterolateral protuberances, continuous pronotal transverse suture, and mesosternum with large medial depression. This species is similar to *G. conisimilis*, but the slightly larger body size, pale antennal segment V, dark brown pronotum with pale lateral spots extending from posterior half of anterior lobe to anterior half of posterior lobe, pale meso- and metasterna, smaller synthlipsis width, interocular sulcus near the hind margin of the eye, larger ocelli, eye size relative to the head length and height, the more distinctive conical anterior pronotal paramedian lobes with a dorsal protuberance, and intersegmental sutures carinulate between II–IV and laterally between IV–VI distinguish this species from *G. conisimilis*.

DESCRIPTION: **MALE:** Body length: 9.48 mm (holotype), 8.53–9.48 mm. COLORATION: Dark brown with pale synthlipsis laterally, postocular dorsally except ocellar tubercle, neck dorsally, scape basally, antennal segments V–VIII, anterior and posterolateral margins of pronotum, lateral spots extending from posterior half of anterior pronotal lobe to anterior half of posterior lobe, stridulatory groove, meso- and metasterna, coxae, trochanters, forefemur basally and ventrally, mid and hind femora basally and subapically, tibiae medially, anterior area of laterotergites, lateral margins of sternites, and sternites medially. Labium and remainder of antenna brown. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with deep, narrow medial longitudinal depression to middle of interocular area; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli large, located on distinct median tubercle (Fig. 2.13C); eye about half of head length, reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about one-fourth longer than scape; antenna 8-segmented; labium slender; labial segment III subequal to II,

ventrally straight (Fig. 2.13G–I, K). THORAX: Anterior pronotal lobe conical in lateral view (Fig. 2.15C), without distinct anterolateral projections; pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe transversely striated; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); mesosternum with large medial depression; metasternum slightly medially longitudinally depressed; legs slender; fossula spongiosa on mid tibia; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex medially notched or sinuate; sternites medially convex, intersegmental sutures carinulate between II–IV and laterally between IV–VI; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G); area of endosomal struts–DPS fusion ovate (Fig. 2.21A).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the conical anterior pronotal paramedian lobes.

DISTRIBUTION: Marojejy Nature Reserve in Antsiranana province (Map 2.2).

DISCUSSION: The coloration is slightly variable with the postclypeus entirely dark brown, the pronotum less pale, and the brown ventrolateral longitudinal band on the sternites varying in size in some specimens.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 30 May 2005–11 Jun 2005, M. Irwin, R. Harin'Hala (00045367) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 25 Dec 2004–30 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00007240) (CAS); 18 Feb 2005–25 Feb 2005, M. Irwin, R. Harin'Hala, 1♂

(00045655) (CAS); 11 Mar 2005–18 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00044813) (SU); 28 Apr 2005–07 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00007174) (UCR); 14 Oct 2005–22 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00045558) (UCR). **Unknown:** 1♂ (00006144) (CAS).

*Gibbosella pallidacorium*, new species

Figs. 2.1, 2.5, 2.8; Map 2.1

**DIAGNOSIS:** Males are recognized among other species of *Gibbosella* by a combination of the following characters: the coloration, circular head shape in lateral view that is as long as it is wide in dorsal view, large ocelli, anterior pronotal lobe laterally carinate, meso- and metasterna not completely separated by a distinct suture, and intersegmental sutures carinate between II–IV and laterally between IV and V. This species is similar to *G. betampona*, but differs by the pale postocular and neck, pale markings on the pronotum, pale corium, pale ventral abdominal surface with dark brown longitudinal stripe laterally, circular head shape, relatively deeper postclypeal depression, eye not reaching ventral head margin, labial segment II and III subequal, anterior pronotal lobe laterally carinate, mesosternum with three longitudinal depressions, fossula spongiosa on the mid tibia, and intersegmental sutures carinate between II–IV and laterally between IV and V.

**DESCRIPTION: MALE:** Body length: 5.78 mm (holotype). **COLORATION:** Dark brown with pale postocular and neck dorsally, scape basally, apical half of antennal segment V and segments VI and VII, labial segment IV, anterior pronotal lobe anteromedially, pronotal disc, posterior pronotal lobe anterolaterally and posterolateral margin, stridulatory groove, meso- and metasterna, corium except apical margin, basal wing veins, coxae, trochanters, mid and hind femora basally and medially (forelegs missing), mid tibia medially and apically, hind tibia except basally, dorsal laterotergites except posterior margin, ventral laterotergites, and sternites except

pygophore and laterally. Labial segment III and wing membrane brown. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Circular in lateral view (Fig. 2.13C); as long as wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with deep, narrow medial longitudinal depression to middle of interocular area; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli large, located on shallow median tubercle (Fig. 2.13I); eye about half of head length, nearly reaching dorsal and ventral head surface; antenna inserted dorsally on head; pedicel about one-fourth longer than scape; antenna 8-segmented; labium slender; labial segment III subequal to II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B), slightly carinate laterally; pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe transversely striated; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna not completely separated by distinct transverse suture (Fig. 2.15F); mesosternum with medial and paramedial longitudinal depressions; metasternum slightly medially longitudinally depressed; forelegs missing, mid and hind legs slender; fossula spongiosa on mid tibia; tarsi missing; hemelytron surpassing abdominal apex; distal part of R absent. ABDOMEN: Apex sinuate; sternites medially convex, intersegmental sutures carinulate between II–IV and laterally between IV and V; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS, posteriorly obscured by semiextended endosoma; area of endosomal struts–DPS fusion circular.

ETYMOLOGY: The species epithet is a noun in the nominative case and is named for the pale corium.

DISTRIBUTION: The only known specimen was collected in Marojejy Nature Reserve in Antsiranana province (Map 2.1).

HOLOTYPE: Male: **Madagascar: Antsiranana:** R.N.I. de Marojejy, 11.0 km NW Manantenina, 14.43667°S 49.74167°E, 1225 m, 25 Oct 1996–03 Nov 1996, Eric L. Quinter (00078364) (AMNH).

*Gibbosella pallidalata*, new species

Figs. 2.1, 2.5, 2.8; Map 2.1

DIAGNOSIS: Males recognized among other species in *Gibbosella* by the coloration, 6-segmented antenna, anterior pronotal lobe without distinct anterolateral protuberances, and fossula spongiosa absent on the mid tibia. This species is similar to *G. planiscutum*, but, in addition to the diagnostic features, it differs by the continuous pronotal transverse suture, separation of the meso- and metasterna, and intersegmental sutures carinulate between II–VI.

DESCRIPTION: **MALE:** Body length: 5.23 mm (holotype), 4.88–5.34 mm. COLORATION: Pale with dark brown head, propleuron medially, mesopleuron and metapleuron except dorsad of coxae, thoracic sterna, sternites ventrolaterally, and pygophore medially. Antenna, labium, posterior pronotal lobe medially, scutellum, distal half of corium, legs except femora basally and ventrally and mid and hind tibiae medially, and posterior half of laterotergites II–VI brown.

VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula moderately swollen ventrolaterally, not distinctly produced beyond ventral head margin (Fig. 2.13F–K); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area; synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about

one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about as long as scape; antenna 6-segmented; labium stout; labial segment III shorter than II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe without distinct anterolateral projections; pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); mesosternum with medial and paramedial longitudinal depressions; metasternum slightly medially longitudinally depressed; forefemur slightly incrassate; fossula spongiosa absent on mid tibia; tarsomeres I and II combined subequal to III on forelegs, longer on mid and hind legs; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex rounded or sinuate; sternites medially convex, intersegmental sutures carinulate between II–VI; pygophore process flattened in lateral view (Fig. 2.20F), subtriangular in caudal view (Fig. 2.19G, I), apex rounded; BPE shorter than basal plate; endosomal struts not reaching posterior margin of DPS, posteriorly obscured by semiextended endosoma; area of endosomal struts–DPS fusion elongate subquadrate.

ETYMOLOGY: The species epithet is a noun in the nominative case and is named for the pale wing membrane.

DISTRIBUTION: Fianarantsoa and Toliara provinces (Map 2.1).

DISCUSSION: Some specimens have small paramedian brown spots on the anterior pronotal lobe.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 14 Apr 2002–23 Apr 2002, M. Irwin, R. Harin'Hala (00044983) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 16 Oct 2001–08 Nov 2001, M. Irwin, R. Harin'Hala, 2♂ (00007237, 00007245) (CAS); 08 Nov 2001–15 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00007159) (SU); 22 Nov 2001–28 Nov 2001, M. Irwin, R. Harin'Hala, 2♂ (00045117, 00045624) (BMNH); 04 Feb 2002–12 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00007206) (CAS); 26 Feb 2002–04 Mar 2002, M. Irwin, R. Harin'Hala, 1♂ (00007130) (AMNH); 31 Mar 2002–07 Apr 2002, M. Irwin, R. Harin'Hala, 2♂ (00045535, 00045541) (USNM); 07 Apr 2002–14 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00044918) (CAS); 14 Apr 2002–23 Apr 2002, M. Irwin, R. Harin'Hala, 2♂ (00007074, 00044825) (MNHN), 2♂ (00044914, 00045473) (BMNH); 28 Apr 2002–05 May 2002, M. Irwin, R. Harin'Hala, 2♂ (00005356, 00007215) (SU); 05 May 2002–13 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00007232) (CAS); 04 Jul 2002–14 Jul 2002, M. Irwin, R. Harin'Hala, 3♂ (00007205, 00007222, 00007243) (CAS); 14 Jul 2002–24 Jul 2002, M. Irwin, R. Harin'Hala, 1♂ (00007246) (UCR). Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1110 m, 28 Nov 2001–06 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00044934) (CAS), 1♂ (00045471) (MNHN); 15 Dec 2001–21 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00099046) (CAS); 28 Jan 2002–04 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00044827) (SU); 04 Feb 2002–12 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00007235) (CAS); 06 May 2002–15 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00007194) (CAS); 04 Jun 2002–14 Jun 2002, M. Irwin, R. Harin'Hala, 2♂ (00007262, 00007267) (USNM); 15 Jul 2002–25 Jul 2002, M. Irwin, R. Harin'Hala, 1♂ (00045533) (UCR). Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 28 Nov 2001–06 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00045580) (CAS); 31 Mar 2002–09 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00007132) (CAS); 09 Apr 2002–16 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00007228) (AMNH); 15 Jul 2002–25 Jul 2002, M. Irwin, R. Harin'Hala, 1♂ (00005358) (USNM); 07 Sep



2003–18 Sep 2003, M. Irwin, R. Harin'Hala, 2♂ (00007108, 00044902) (CAS) ; 21 Mar 2004–02 Apr 2004, M. Irwin, R. Harin'Hala, 1♂ (00007036) (SU); 18 Mar 2006–30 Mar 2006, M. Irwin, R. Harin'Hala, 1♂ (00007145) (CAS); 17 Oct 2006–01 Nov 2006, M. Irwin, R. Harin'Hala, 1♂ (00045653) (UCR). Ranomafana JIRAMA water works, 21.2485°S 47.45216°E, 690 m, 22 Nov 2001–28 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00007210) (CAS), 1♂ (00007227) (UCR). Ranomafana National Park, Talatakely, 21.25°S 47.41667°E, 900 m, 09 Jan 2001–19 Jan 2001, D.H. and K.M. Kavanaugh, R.L. Brett, E. Elsom, and F. Vargas, 2♂ (00007110, 00044853) (CAS), 2♂ (00044979, 00044996) (AMNH). **Toliara:** Berenty Special Reserve, 8 km NW Amboasary, 25.021°S 46.3055°E, 35 m, 29 Aug 2004–08 Sep 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00044876) (CAS). Parc National d'Andohahela, Col du Sedro, 3.8 km 113° ESE Mahamavo, 37.6 km 341° NNW Tolagnaro, 24.76388°S 46.75166°E, 900 m, 21 Jan 2002–25 Jan 2002, Fisher et al., 1♂ (00006463) (UCR), 1♂ (00007085) (CAS), 1♂ (00007247) (MNHN). **Unknown:** 6♂ (00007076, 00007125, 00007204, 00007216, 00007249, 00007264) (CAS).

*Gibbosella planiscutum*, new species

Figs. 2.1, 2.5, 2.8, 2.17A, 2.23B; Map 2.3

DIAGNOSIS: Males are recognized among other species in this genus by a combination of the following characters: the coloration, dense vestiture, pronotal transverse suture divided by paramedian ridges, scutellar processes horizontally directed, meso- and metasterna not entirely separated by a distinct suture, and intersegmental sutures carinulate between II–VI and laterally between VI and VII. Females are recognized by the coloration and synthlipsis about three times the width of an eye. Males are similar to *G. brunalvus*, from which it differs by the pale markings on the head (e.g., mandibular and maxillary plates, antennifers, ventral and lateral antecular and postocular surface), pale pronotum with dark brown paramedian spots on the anterior lobe and

margins of posterior lobe, depressed postclypeus, orientation of the scutellar processes, lack of a distinct separation between the meso- and metasterna, and intersegmental sutures carinulate between II–VI and laterally between VI and VII. Females are very similar to *G. brunalvus*, but are distinguished by the nearly black body, pale apex of antennal segment VIII and apex of scutellum, and larger synthlipsis width.

DESCRIPTION: **MALE:** Body length: 5.08 mm (holotype), 4.91–5.90 mm. COLORATION: Dark brown with pale mandibular and maxillary plates, antennifers, ventral and lateral anteocular and postocular surface, except at posterior margin, scape basally, antennal segments VI–VIII, anterior pronotal lobe except paramedian spots and pronotal longitudinal furrow, posterior pronotal lobe except medially, anterior propleural lobe except medially, posterior propleural lobe dorsally, mesopleuron and metapleuron dorsad of coxae, apical half of scutellum, basal half of corium, coxae, trochanters, forefemur basally and ventrally, mid and hind femora basally, tibiae basally, anterior half of laterotergites, and sternites medially and anterolaterally. Remainder of antenna, labium, remainder of legs, apical half of corium, and basal wing veins brown.

VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula moderately swollen ventrolaterally, not distinctly produced beyond ventral head margin (Fig. 2.13F–K); postclypeus with shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about as long as scape; antenna 8-segmented; labium stout; labial segment III shorter than II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow

divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions on posterior pronotal lobe transversely striated; scutellar apical projections horizontally directed (Fig. 2.15D); meso- and metasterna not completely separated by distinct transverse suture (Fig. 2.15E); mesosternum with medial and paramedial longitudinal depressions; metasternum slightly medially longitudinally depressed; fore- and mid femora incrassate; fossula spongiosa on mid tibia; tarsomeres I and II combined shorter than III; hemelytron reaching or surpassing abdominal apex; distal part of R vein present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex rounded; sternites with shallow medial depression (Fig. 2.18C), intersegmental sutures carinulate between II–VI and laterally between VI and VII; pygophore process subtriangular in lateral (Fig. 2.20A) and caudal (Fig. 2.19G, I) views, apex rounded; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G), fused anterior to fusion with DPS; area of endosomal struts–DPS fusion ovate (Fig. 2.21A). **FEMALE:** Body length: 4.78–5.06 mm. Similar to males, but differ in the following characters: **COLORATION:** Dark brown with brown scape, pedicel, antennal segments III–IV, base of segment V, apex of segment VIII, labium, and legs except trochanter and tarsi. Remainder of antennal segments, scutellar processes, trochanters, and tarsi whitish to pale brown. **STRUCTURE:** In addition to characters mentioned in the generic description: **HEAD:** Synthlipsis about three times width of eye. **THORAX:** Pronotal longitudinal furrow restricted to anterior lobe; lateral depressions on posterior pronotal lobe obsolete; forefemur with small ventral tubercles (Fig. 2.16D). **ABDOMEN:** Intersegmental sutures carinulate between sternites II–VI; external genitalia as in Fig. 2.23B; bursa copulatrix membranous, with lateral lobes (Fig. 2.23F); vermiform gland elongate, much longer than bursa copulatrix; lateral spermathecae inserted on median oviduct with slender, fragile duct and large ovoid pouch.

ETYMOLOGY: The species epithet is a noun in the nominative case and is named for the horizontally-oriented scutellar apices.

DISTRIBUTION: Antananarivo and Mahajanga provinces (Map 2.3).

DISCUSSION: In some specimens, the entire postocular is dark brown, the brown spots on the paramedian anterior pronotal lobes are absent, and the scutellum is entirely pale. Females were associated to males based on morphology and the same collection event as another male.

HOLOTYPE: Male: **Madagascar: Mahajanga:** Analamanitra Forest, 14 km NE of Misinjo, 16.13333°S 45.7°E, 20 m, 20 Nov 2007–04 Dec 2007, M. Irwin, R. Harin'Hala (00044849) (CAS).

PARATYPES: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 10 Jan 2004–20 Jan 2004, M. Irwin, R. Harin'Hala, 1♂ (00006461) (CAS), 1♂ (00007266) (SU); 20 Jan 2004–01 Feb 2004, M. Irwin, R. Harin'Hala, 1♂ (00007058) (UCR). **Mahajanga:** 160 km N of Maevatanana on RN 04, Ampijoroa National Park, 16.31933°S 46.81333°E, 43 m, 07 Feb 2005–19 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00045216) (USNM). Namoroka, 53 km from Soalala, 3 km N Vilanandro Village, 16.47333°S 45.39133°E, 122 m, 30 Nov 2007–09 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00045296) (USNM). Parc National de Namoroka, 16.9 km 317° NW Vilanandro, 16.40666°S 45.31°E, 100 m, 12 Nov 2002–16 Nov 2002, Fisher et al., 1♂ (00006320), 1♀ (00045373) (CAS), 1♀ (00045711) (UCR). Parc National de Namoroka, 17.8 km 329° WNW Vilanandro, 16.37666°S 45.32666°E, 100 m, 08 Nov 2002–12 Nov 2002, Fisher et al., 1♂ (00007209) (CAS). Réserve forestière Beanka, 50.2 km E Maintirano, 18.02638°S 44.05055°E, 250 m, 19 Oct 2009–26 Oct 2009, B.L. Fisher et al., 1♂ (00006464) (CAS). Réserve Spéciale de Bemarivo, 23.8 km 223° SW Besalampy, 16.925°S 44.36833°E, 30 m, 19 Nov 2002–23 Nov 2002, Fisher et al., 1♂ (00045515) (SU). **Unknown:** 1♂ (00007149) (CAS).

*Gibbosella quadocris*, new species

Figs. 2.1, 2.5, 2.8, 2.19A, 2.20B; Map 2.2

DIAGNOSIS: Males are recognized among other species in *Gibbosella* by the small body size, ovoid head shape in lateral view that is longer than it is wide in dorsal view, synthlipsis 1.5 times the width of an eye, labial segments II and III subequal, subquadrate median pygophore process with truncated apex in caudal view, and BPE as long as the basal plate. This species is very similar to *G. vangocris*, but is distinguished by the smaller body size, slightly shorter head, more narrowly depressed postclypeus, relative lengths of the labial segments, meso- and metasterna not distinctly or weakly separated by a distinct transverse suture, and shape of the pygophore process in caudal view.

DESCRIPTION: **MALE:** Body length: 5.39 mm (holotype), 5.33–5.78 mm. COLORATION: Dark brown with pale interocular sulcus, anterior pronotal lobe except interspersed brown markings posterolateral margin of pronotum, metasternum, corium basally, coxae, trochanters, femora basally and ventrally, mid and hind tibiae basally, laterotergites except posterior margin, and sternites (except medial pygophore and lateral and posterolateral margins of sternites II–VI). Antennal segment V (except basally) and VI–VIII white. Remainder of antenna, labium, parts of anterior pronotal lobe, posterior pronotal lobe, and rest of corium and basal wing veins brown. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about as long as scape;

antenna 8-segmented; labium stout; labial segment III subequal to II, ventrally convex (Fig. 2.13C). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna not completely or weakly separated by distinct transverse suture (Fig. 2.15E); mesosternum with medial and paramedial depressions; metasternum slightly medially longitudinally depressed; forefemur incrassate; fossula spongiosa on mid tibia; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex rounded or sinuate; sternites longitudinally depressed medially (Fig. 2.18C), intersegmental sutures carinulate between II–VI; pygophore (Figs. 2.19A, 2.20B) process subtriangular in lateral view, subquadrate in caudal view, apex truncated; BPE as long as basal plate; endosomal struts not reaching posterior margin, posteriorly obscured by semiextended endosoma; fused anterior to fusion with DPS; area of endosomal struts–DPS fusion elongate ovoid.

ETYMOLOGY: The species epithet is a noun in apposition and is named for the subquadrate median pygophore process.

DISTRIBUTION: Ranomafana National Park, Fianarantsoa province (Map 2.2).

DISCUSSION: In some specimens, the posterior pronotal lobe is dark brown, the anterior pronotal lobe have reduced pale coloration and appear almost uniformly brown, and the corium and basal wing veins range from pale to brown.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 08 Nov 2001–15 Nov 2001, M. Irwin, R. Harin'Hala (00007072) (CAS).

PARATYPES: **Madagascar: Fianarantsoa: Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 08 Nov 2001–15 Nov 2001, M. Irwin, R. Harin'Hala, 2♂ (00006457, 00007128) (CAS), 1♂ (00007258) (UCR); 28 Nov 2001–06 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00007200) (SU); 02 Jan 2002–10 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00007153) (CAS); 10 Jan 2002–14 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00007172) (CAS); 12 Feb 2002–19 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00007236) (CAS); 31 Mar 2002–07 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00045552) (CAS); 07 Apr 2002–14 Apr 2002, M. Irwin, R. Harin'Hala, 2♂ (00044888, 00045425) (AMNH); 14 Apr 2002–23 Apr 2002, M. Irwin, R. Harin'Hala, 3♂ (00044909, 00044949, 00044951) (USNM), 2♂ (00044954, 00045166) (MNHN); 28 Apr 2002–05 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00007184) (CAS); 05 May 2002–13 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00007186) (CAS); 13 Jun 2002–23 Jun 2002, M. Irwin, R. Harin'Hala, 1♂ (00044802) (CAS); 04 Jul 2002–14 Jul 2002, M. Irwin, R. Harin'Hala, 1♂ (00044895) (CAS); 24 Jul 2002–04 Aug 2002, M. Irwin, R. Harin'Hala, 1♂ (00007219) (BMNH). Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1100 m, 02 Jan 2002–10 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00007188) (CAS); 19 Feb 2002–26 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00007257) (SU); 04 Jun 2002–14 Jun 2002, M. Irwin, R. Harin'Hala, 1♂ (00044860) (UCR). Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 14 Jan 2002–21 Jan 2002, M. Irwin, R. Harin'Hala, 2♂ (00044882, 00045572) (BMNH); 21 Jan 2002–28 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00045651) (SU). Parc National Ranomafana, radio tower at forest edge, 21.25083°S 47.40716°E, 1130 m, 18 Mar 2006–30 Mar 2006, M. Irwin, R. Harin'Hala, 1♂ (00007224) (CAS). Ranomafana JIRAMA water works, 21.2485°S 47.45216°E, 690 m, 06 Dec 2001–15 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00007166) (CAS), 1♂ (00007203) (AMNH); 21 Dec 2001–24 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00006465) (UCR); 21 Jan 2002–28 Jan 2002, M.**

Irwin, R. Harin'Hala, 1♂ (00045348) (CAS). **Unknown:** 4♂ (00007117, 00007180, 00007181, 00007261) (CAS).

*Gibbosella vangocris*, new species

Figs. 2.1, 2.5, 2.8, 2.19B, 2.20C, 2.21B, 2.22A; Map 2.3

**DIAGNOSIS:** Males recognized among other *Gibbosella* species by the relatively larger body size, slightly broader postclypeal depression, synthlipsis 1.5 times the width of an eye, labial segment III shorter than II, intersegmental sutures carinulate between II–VI, and spadelike median pygophore process in caudal view. This species is very similar to *G. quadocris*, but is distinguished from *G. quadocris* by the larger body size, slightly more elongate head, more broadly depressed postclypeus, labial segment II longer than III, meso- and metasterna distinctly separated by a distinct transverse suture, and shape of the pygophore process in caudal view.

**DESCRIPTION: MALE:** Body length: 7.17 mm (holotype), 6.80–7.61 mm. **COLORATION:** Dark brown with antennal segment V apically; antennal segments VI–VIII; posterior margin of pronotum; corium basally; coxae; trochanters; basal half of femora; anterior areas of laterotergites; sternites medially, except pygophore; and anterolaterally on sternites II–VI pale. Remainder of corium and basal wing veins brown. **VESTITURE:** Sparse; other features as in generic description. **STRUCTURE: HEAD:** Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; anteocular region as long as postocular (Fig. 2.13C); gula flat, conforming to rounded shape of head (Fig. 2.13C); postclypeus with shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about 1.5 times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small, located on shallow median tubercle (Fig. 2.13I); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about as long as scape; antenna 8-



segmented; labium stout; labial segment III shorter than II, ventrally convex (Fig. 2.13C).

THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe smooth; scutellar apical projections dorsally directed (Fig. 2.15C); meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); mesosternum with medial and paramedial longitudinal depressions; metasternum medially longitudinally depressed; legs slender; fossula spongiosa on mid tibia; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Apex rounded; sternites medially convex, with shallow medial depression on posterior half of segment VI, intersegmental sutures carinulate between II–VI; pygophore (Figs. 2.19B, 2.20C) process subtriangular basally and apically flattened and curved in lateral view, spadelike in caudal view, apex rounded; BPE as long as basal plate (Fig. 2.22A); endosomal struts reaching posterior margin of DPS, with posterior ventral process (Fig. 2.22G), fused anterior to fusion with DPS; area of endosomal struts–DPS fusion ovate (Fig. 2.21B).

ETYMOLOGY: The species epithet is a noun in apposition and is named for the spadelike median pygophore process.

DISTRIBUTION: Marojejy National Park in the Antsiranana province, Ranomafana National Park in the Fianarantsoa province, and Berenty Special Reserve in the Toliara province (Map 2.3).

DISCUSSION: The coloration is slightly variable with the posterior pronotal lobe, corium, and basal wing veins ranging from pale to brown.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 16 Oct 2001–08 Nov 2001, M. Irwin, R. Harin'Hala (00006995) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 25 Feb 2005–04 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00007275) (CAS). **Fianarantsoa:** Parc National Ranomafana, Bevaohazo-Ranomana, 21.2°S 47.48333°E, 970 m, Nov 2000, M. Irwin, R. Harin'Hala, 1♂ (00044919) (CAS). Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 16 Oct 2001–08 Nov 2001, M. Irwin, R. Harin'Hala, 3♂ (00006984, 00007000, 00007001) (USNM), 13♂ (00007005, 00007014, 00007022, 00007029, 00007030, 00007033, 00007055, 00007057, 00007063, 00007070, 00007071, 00007077, 00007092) (CAS), 4♂ (00007111, 00007114, 00007126, 00007254) (MNHN), 3♂ (00007268, 00007270, 00048227) (AMNH); 08 Nov 2001–15 Nov 2001, M. Irwin, R. Harin'Hala, 3♂ (00006085, 00007006, 00007017) (MNHN), 4♂ (00007018, 00007044, 00007144, 00007274) (BMNH); 15 Nov 2001–22 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00045570) (UCR); 22 Nov 2001–28 Nov 2001, M. Irwin, R. Harin'Hala, 6♂ (00048028–00048033) (CAS); 15 Dec 2001–21 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00045586) (SU); 24 Dec 2001–02 Jan 2002, M. Irwin, R. Harin'Hala, 3♂ (00007093, 00044929, 00045720) (USNM); 19 Feb 2002–26 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00007016) (UCR); 31 Mar 2002–07 Apr 2002, M. Irwin, R. Harin'Hala, 5♂ (00048052–00048056) (CAS); 07 Apr 2002–14 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00048034) (SU), 6♂ (00048036–00048041) (CAS); 14 Apr 2002–23 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00045342) (CAS); 05 May 2002–13 May 2002, M. Irwin, R. Harin'Hala, 2♂ (00006985, 00007013) (AMNH); 13 May 2002–23 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00007027) (CAS); 13 Jun 2002–23 Jun 2002, M. Irwin, R. Harin'Hala, 1♂ (00045716) (CAS); 04 Jul 2002–14 Jul 2002, M. Irwin, R. Harin'Hala, 1♂ (00007273) (BMNH); 14 Jul 2002–24 Jul 2002, M. Irwin, R. Harin'Hala, 1♂ (00045172) (BMNH); 24 Jul 2002–04 Aug 2002, M. Irwin, R. Harin'Hala, 2♂ (00007032, 00007280) (USNM) ; 15 May 2003–28 May 2003, M. Irwin, R.

Harin'Hala, 1♂ (00006993) (UCR). Parc National Ranomafana, Vohiparara, at broken bridge,  
 21.22616°S 47.36983°E, 1100 m, 28 Nov 2001–06 Dec 2001, M. Irwin, R. Harin'Hala, 4♂  
 (00044845, 00044870, 00045020, 00045230) (BMNH), 3♂ (00045406, 00045409, 00045676)  
 (CAS), 1♂ (00045688) (UCR); 02 Jan 2002–10 Jan 2002, M. Irwin, R. Harin'Hala, 2♂  
 (00006999, 00007023) (AMNH); 28 Jan 2002–04 Feb 2002, M. Irwin, R. Harin'Hala, 4♂  
 (00048042–00048045) (MNHN), 6♂ (00048046–00048051) (CAS); 19 Mar 2002–26 Mar 2002,  
 M. Irwin, R. Harin'Hala, 2♂ (00045199, 00045384) (SU); 26 Mar 2002–31 Mar 2002, M. Irwin,  
 R. Harin'Hala, 1♂ (00007040) (CAS); 08 Apr 2002–15 Apr 2002, M. Irwin, R. Harin'Hala, 2♂  
 (00007042, 00007097) (CAS); 22 Apr 2002–29 Apr 2002, M. Irwin, R. Harin'Hala, 2♂  
 (00007272, 00007278) (CAS); 06 May 2002–15 May 2002, M. Irwin, R. Harin'Hala, 1♂  
 (00007015) (UCR); 04 Jun 2002–14 Jun 2002, M. Irwin, R. Harin'Hala, 2♂ (00007116,  
 00007119) (USNM); 15 Jul 2002–25 Jul 2002, M. Irwin, R. Harin'Hala, 2♂ (00045313,  
 00045578) (AMNH). Parc National Ranomafana, radio tower at forest edge, 21.251°S  
 47.40716°E, 1130 m, 06 Dec 2001–15 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00007011)  
 (MNHN), 1♂ (00007213) (CAS); 15 Dec 2001–21 Dec 2001, M. Irwin, R. Harin'Hala, 1♂  
 (00045588) (UCR); 21 Dec 2001–24 Dec 2001, M. Irwin, R. Harin'Hala, 2♂ (00007019,  
 00045037) (SU); 14 Jan 2002–21 Jan 2002, M. Irwin, R. Harin'Hala, 2♂ (00045516, 00045554)  
 (AMNH); 19 Feb 2002–26 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00007008) (CAS); 12 Mar  
 2002–19 Mar 2002, M. Irwin, R. Harin'Hala, 1♂ (00006994) (CAS); 23 Apr 2002–30 Apr 2002,  
 M. Irwin, R. Harin'Hala, 1♂ (00007187) (SU); 14 Jun 2002–24 Jun 2002, M. Irwin, R.  
 Harin'Hala, 2♂ (00006997, 00007007) (BMNH); 24 Jun 2002–05 Jul 2002, M. Irwin, R.  
 Harin'Hala, 1♂ (00007105) (UCR); 15 Jul 2002–25 Jul 2002, M. Irwin, R. Harin'Hala, 2♂  
 (00007004, 00007025) (USNM); 18 Feb 2003–27 Feb 2003, M. Irwin, R. Harin'Hala, 1♂  
 (00045078) (AMNH); 20 Mar 2003–03 Apr 2003, M. Irwin, R. Harin'Hala, 1♂ (00007121) (SU);

06 Jul 2003–17 Jul 2003, M. Irwin, R. Harin'Hala, 1♂ (00007269) (CAS); 27 Aug 2003–07 Sep 2003, M. Irwin, R. Harin'Hala, 1♂ (00007021) (UCR); 28 Sep 2003–08 Oct 2003, M. Irwin, R. Harin'Hala, 2♂ (00007002, 00007233) (CAS); 02 Apr 2004–15 Apr 2004, M. Irwin, R. Harin'Hala, 1♂ (00007003) (USNM); 11 Jun 2004–24 Jun 2004, M. Irwin, R. Harin'Hala, 1♂ (00045507) (CAS); 24 Jun 2004–08 Jul 2004, M. Irwin, R. Harin'Hala, 1♂ (00007169) (AMNH); 23 Sep 2004–07 Oct 2004, M. Irwin, R. Harin'Hala, 1♂ (00045660) (SU); 11 Dec 2004–18 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00045625) (UCR); 02 Feb 2005–17 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00045563) (CAS); 27 Feb 2005–13 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00007069) (UCR); 25 Jul 2006–09 Aug 2006, M. Irwin, R. Harin'Hala, 1♂ (00045371) (CAS).

Ranomafana JIRAMA water works, 21.2485°S 47.45216°E, 690 m, 22 Nov 2001–28 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00006124) (UCR); 06 Dec 2001–15 Dec 2001, M. Irwin, R. Harin'Hala, 4♂ (00006982, 00006988, 00006991, 00006996) (CAS), 2♂ (00007020, 00007031) (SU); 21 Dec 2001–24 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00007276) (CAS); 24 Dec 2001–02 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00006986) (UCR); 21 Jan 2002–28 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00044847) (SU), 1♂ (00045687) (CAS). **Toliara:** Berenty Special Reserve, 8 km NW Amboasary, 25.021°S 46.3055°E, 35 m, 29 Aug 2004–08 Sep 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00007197) (UCR), 1♂ (00007248) (CAS), 1♂ (00007250) (SU). **Unknown:** 12♂ (00006983, 00006987, 00006989, 00006990, 00006992, 00007010, 00007024, 00007026, 00007221, 00007271, 00007277, 00007279) (CAS).

*Glymmatophora* Stål, 1853

Figs. 2.1, 2.2, 2.5, 2.8, 2.12, 2.13A, 2.13D, 2.15E, 2.16B, 2.16C, 2.16E, 2.17B, 2.19C, 2.20D, 2.21C, 2.21D, 2.22B, 2.23C, 2.23D, 2.23F; Map 2.4

*Glymmatophora* Stål, 1853: 261.

TYPE SPECIES: *Glymmatophora submetallica* Stål, 1853

DISTRIBUTION: Thirty-five described species of *Glymmatophora* occur throughout Africa (Maldonado, 1990). Within Madagascar, the two species of *Glymmatophora* are known from the Toliara and Fianarantsoa provinces. Specimens have been collected in spiny forest thicket, degraded grassland, and in the Antsarongaza dry forest between 18–825 m elevation.

DISCUSSION: The largely Afrotropical genus *Glymmatophora* was described by Carl Stål in 1853. In 1914, Géza Horváth described three subgenera: (*Cyclosandalus*), (*Glymmatophora*), and (*Haematorrhophus*), the last of which is currently treated as a distinct genus. The former two subgenera were primarily diagnosed based on the presence (subgenus *Glymmatophora*) or absence (subgenus *Cyclosandalus*) of ocelli in the female and the size and shape of the fossula spongiosa (large and oblong in subgenus *Glymmatophora*; small and round in subgenus *Cyclosandalus*). Prior to this monograph, one species of *Glymmatophora* was known from Madagascar, *G. (Cyclosandalus) crassipes*. A new Madagascan species of *Glymmatophora* is here described, *G. (Glymmatophora) carolae*.

*Identification key to the males and females of Madagascan species of Glymmatophora*

1. Males only known; macropterous; ocelli present ..... (*Glymmatophora*) *carolae*, new species
- Males and females apterous; ocelli absent (Fig. 2.13D) ..... (*Cyclosandalus*) *crassipes* Horváth

*Glymmatophora (Glymmatophora) carolae*, new species

Figs. 2.1, 2.5, 2.8, 2.16C, 2.17B, 2.20D, 2.21C; Map 2.4

DIAGNOSIS: Males are recognized among other *Glymmatophora* species by the very distinctive coloration (mostly red with some black markings), anteriorly concave postclypeus, forefemur with anterior subapical and medial protuberances and macropterous condition. This

species is very similar to the Afrotropical *G. lothei* Villiers, 1948, from which it differs by the slightly smaller body size, black antenna, red abdomen with black markings (i.e., anterior band on laterotergites IV and V, laterotergite VI except posterolaterally, anterior half of laterotergite VII, lateral margin of abdominal sternite VI, and anterolateral margin of sternite VII), more medially concaved anterior pronotal margin, and forefemur with only two protuberances.

DESCRIPTION: **MALE:** Macropterous, medium body size (length: 16.91 mm, holotype; 14.90–16.91 mm). COLORATION: Red with antenna, hemelytron except corium basally and anteriorly, anterior band on laterotergites IV and V, laterotergite VI except posterolaterally, anterior half of laterotergite VII, lateral margin of abdominal sternite VI, and anterolateral margin of sternite VII black. VESTITURE: Glabrous, except antenna with long, erect brown setae and tibiae with stout pale brown setae that are denser near apex. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13D); wider than long in dorsal view; ventrally flat; anteocular region as long as postocular; gula swollen ventrolaterally, slightly produced beyond ventral margin of head (Fig. 2.13E); maxillary plate not reaching dorsal clypeal surface (Fig. 2.13E, I); postclypeus concave anteriorly with shallow, very narrow medial longitudinal depression to middle of interocular area; synthlipsis (Fig. 2.13A) about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli small, located on distinct median tubercle (Fig. 2.13H), separated by more than diameter of ocellus; eye about one-third head length, not reaching dorsal and ventral head surfaces; antennal shield concealing antennal insertion in lateral view (Fig. 2.13D); antenna inserted sublaterally on head; pedicel about one-fourth longer than scape, slightly curved. THORAX: Pronotum wider than long, anterior margin moderately concave, smooth; anterior pronotal lobe shorter than posterior lobe (Fig. 2.15B), more than half as wide as posterior lobe, without distinct anterolateral projections; pronotal longitudinal furrow reaching anterior but not posterior margin, weakly foveate posteriorly (Fig. 2.15B); pronotal transverse furrow divided

by paramedian longitudinal ridges (Fig. 2.15A, B); lateral depressions of posterior pronotal lobe distinct (Fig. 2.15B), weakly transversely striated; meso- and metasterna not completely separated by transverse suture (Fig. 2.15E); mesosternum with medial and paramedian longitudinal depressions (Fig. 2.15F); metasternum medially longitudinally depressed (Fig. 2.15F); MGE with shallow meshlike cuticle that does not extend dorsally in lateral view (Fig. 2.16B, C); forefemur with anterior subapical and medial protuberances (Fig. 2.16E), base ventrally carinate; mid and hind femora with anterior and posterior subapical and medial protuberances; foretibia with ventral spine basal to fossula spongiosa; hemelytron (Fig. 2.17B) surpassing abdominal tergite VII but not reaching apex; corium well developed; distal part of R present but not forming R+M cell; proximal parts of M and Cu fused; distal part of M extending beyond apical junction of M+Cu; base of M+Cu cell shorter than or as wide as Cu+1A cell. ABDOMEN: Apex rounded; dorsal laterotergite II expanded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; sternites shallowly depressed medially (Fig. 2.18C); spiracles ovoid; pygophore process dorsally knoblike in lateral view (Fig. 2.20D), subtriangular in caudal view (Fig. 2.19C), directed dorsoposteriad, not surpassing posterior margin of pygophore, apex rounded; BPE shorter than basal plate; DPS apex concave medially (Fig. 2.21C); endosomal struts reaching posterior margin of DPS (Fig. 2.21C), ventrally obscured by semiextend endosoma; area of endosomal struts–DPS fusion elongate subquadrate; endosoma medially sclerotized posteriorly and sclerotization extending anterolaterally (Fig. 2.21C).

ETYMOLOGY: This species epithet is a noun in the genitive case and is named in memoriam of the senior author's mother, Carol Forthman.

DISTRIBUTION: Antsarongaza dry forest in the Toliara province (Map 2.4).

DISCUSSION: Female specimens are unknown. We assign this species to the subgenus *Glymmatophora* based on the features mentioned in the generic discussion.

HOLOTYPE: Male: **Madagascar: Toliara:** Beroboka village, 45 km NE Morondava, 19.9775°S 44.82483°E, 131 m, 17 Nov 2008–25 Nov 2008, M. Irwin, R. Harin'Hala (00044889) (CAS).

PARATYPES: **Madagascar: Toliara:** Bekily, 24.22802°S 45.30858°E, 388 m, Oct 1936, A. Seyrig, 2♂ (MNHN); Beroboka village, 45 km NE Morondava, 19.9775°S 44.82483°E, 131 m, 05 Feb 2009–13 Feb 2009, M. Irwin, R. Harin'Hala, 1♂ (00045479) (CAS).

*Glymmatophora (Cyclosandalus) crassipes* Horváth, 1914

Figs. 2.1, 2.2, 2.5, 2.8, 2.12, 2.13A, 2.13D, 2.15E, 2.16B, 2.16E, 2.19C, 2.21D, 2.22B, 2.23C, 2.23D, 2.23F; Map 2.4

*Glymmatophora (Cyclosandalus) crassipes* Horváth, 1914: 128.

REVISED DIAGNOSIS: Males and females are recognized among other species in this genus by the apterous condition, lack of ocelli in both sexes, and fossula spongiosa on foretibia small and round. This species is similar to apterous forms of *G. lothei*, but the presence of two protuberances on the forefemur distinguishes *G. crassipes* from *G. lothei*.

REDESCRIPTION: **MALE:** Apterous, medium body size (length: 15.52–19.60 mm).

COLORATION: Black with head, labium, scape, pedicel basally, pronotum except pronotal longitudinal furrow, propleuron except dorsal spot and stripe on anterior lobe, meso- and metapleura dorsad of coxae, meso- and metasterna laterally, legs, laterotergite II, laterotergite III and IV posteriorly, sternites II–VI transversely basally, sternite VII posteriorly, and pygophore red. VESTITURE: Glabrous, except antenna with short, semierect to erect brown setae and tibiae with stout, pale brown setae that are denser near apex. STRUCTURE: HEAD (Fig. 2.13A, D): Subquadrate in lateral view; longer than wide in dorsal view; ventrally flat; anteocular region longer than postocular; gula swollen ventrolaterally, not produced beyond ventral margin of head,



carinate; maxillary plate not reaching dorsal clypeal surface (Fig. 2.13E, I); postclypeus flat; synthlipsis about 2.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antennal shield concealing antennal insertion in lateral view; antenna inserted sublaterally on head; pedicel slightly longer than scape, slightly curved. THORAX (Fig. 2.15E): Pronotum as long as wide, anterior margin slightly concave, smooth; anterior pronotal lobe longer than and as wide as posterior lobe (Fig. 2.15A), without distinct anterolateral projections; pronotal longitudinal furrow not reaching anterior and posterior margins, reduced to deep medial depression near posterior margin of anterior lobe (Fig. 2.15A); pronotal transverse furrow weakly to strongly divided by paramedian longitudinal ridges (Fig. 2.15A); lateral depressions of posterior pronotal lobe obsolete; meso- and metasterna not completely separated by transverse suture; mesosternum with medial and paramedian longitudinal depressions; metasternum medially longitudinally depressed; MGE with shallow meshlike cuticle that does not extend dorsally in lateral view (Fig. 2.16B, C); trochanters and fore- and mid femora basally with patches of small papillae (Fig. 2.16E); forefemur with anterior subapical and medial protuberances (Fig. 2.16E); mid and hind femora with anterior and posterior subapical and medial protuberances; foretibia with ventral spine basal to fossula spongiosa. ABDOMEN: Apex rounded; dorsal laterotergite II slightly expanded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; sternites medially flat; spiracles ovoid; pygophore (Figs. 2.19C, 2.20D) process dorsally knoblike in lateral view, subtriangular in caudal view, directed dorsoposteriad, not surpassing posterior margin of pygophore, apex rounded; BPE about as long as basal plate (Fig. 2.22B); DPS apex rounded (Fig. 2.21D); endosomal struts reaching posterior margin of DPS, with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21D); endosoma medially sclerotized posteriorly and extends anterolaterally

(Fig. 2.21D). **FEMALE:** Body length: 16.50 mm (holotype; from original description), 16.50–23.90 mm. Similar to males, but differ in the following characters: **COLORATION:** Red on prothorax, mesopleuron, metapleuron dorsad of coxa, lateral metanotal spots, scutellum laterally, meso- and metasterna, laterotergite II, posterolateral margin of laterotergites III–VII, and sternite VII posteromedially. **STRUCTURE:** **THORAX:** Fossula spongiosa larger. **ABDOMEN:** External genitalia as in plate Fig. 2.23C, D; bursa copulatrix membranous, with lateral lobes (Fig. 2.23F); vermiform gland and lateral spermathecae missing.

**DISTRIBUTION:** Most specimens were collected in the Fianarantsoa and Toliara provinces and were successfully georeferenced (map 4). Some could not be georeferenced beyond province based on locality data, with most of them collected in the Toliara province. One specimen was associated with historical locality data that could not be georeferenced (i.e., “Amboasary [Imerina]”). Imerina refers to a kingdom that once existed and likely is part of modern-day Antananarivo province. A second specimen could not be georeferenced but was collected from “P. Majanga” in the Mahajanga province. It is not clear if “P. Majanga” is referring to a park within the Mahajanga province or the town Mahajanga.

**DISCUSSION:** A few specimens without or with reduced red coloration on the scutellum, laterotergites, and sternites were examined and are treated as conspecific with *G. crassipes* based on corresponding external and internal morphology. Specimens of both color morphs were collected in close proximity.

**HOLOTYPE:** Female: **Madagascar: Toliara:** Tsivory, 24.0684°S 46.07471°E, 431 m, 1906 Fauchère (HNHM).

**OTHER MATERIAL EXAMINED:** **Madagascar: Antananarivo:** Amboasary (Imerina), 1899, G. Grandidier, 1♀ (MNHN). **Fianarantsoa:** Ikongo, 21.99483°S 47.37065°E, 536 m, 1902, G. Grandidier, 1♂ (MNHN). **Mahajanga:** P. Majanga, Nov 1965, Malzy 1♂ (MNHN). **Toliara:** 18

km NW Betroka, 23.16333°S 45.96861°E, 825 m, 04 Dec 1994–09 Dec 1994, M.A. Ivie and D.A. Pollock, 1♂ (00120258) (MTEC). Ankazoabo, 21.50348°S 45.20956°E, 278 m, 1902, J. Bastard, 1♀ (MNHN); 1920, G. Le Barbier, 1♂ (MNHN); no date provided, 1♀ (MNHN). Antaloba, no date provided, 1♀ (MNHN). Bas Mangoky, Agriculture Station, no date provided, 1♂, 1♀ (MNHN). Behara, 24.95347°S 46.38593°E, 49 m, 10 Aug 1918–11 Aug 1918, A.R., 1♀ (MNHN). Bekily, 24.22802°S 45.30858°E, 388 m, Mar 1936, A. Seyrig, 1♂ (MNHN). Berenty Reserve, W of Fort Dauphin, 25.00564°S 46.30278°E, 09 Nov 2008–24 Nov 2008, J.R. Cryan and G. Svenson, 2♂ (00088087, 00088088) (UCR). Betroka, Andalamby, 23.26833°S 46.10444°E, 822 m, Jan 1958, J. Elie, 1♀ (MNHN). Country Mahafaly, 1906, J. Bastard, 1♂ (MNHN). Lambomakandro, 22.69831°S 44.70099°E, 526 m, 1936, R. Catala 1♀ (MNHN). Mahafaly Plateau, 6.2 km 74° ENE Itampolo, 24.65361°S 43.99666°E, 80 m, 21 Feb 2002–25 Feb 2002, Fisher et al., 1♀ (00007050) (CAS). Mahafaly Plateau, Ankairano, 25.45°S 45.71667°E, 18 m, 17 Jan 1974, P. Viette & A. Peyrieras, 1♀ (MNHN). On plateau, 23.00283°S 43.70366°E, 30 m, 26 Feb 2003–08 May 2003, Frontier Wilderness Project, CAS, 1♀ (00006372) (CAS). Onilahy River, 1922, G. Petit, 1♀ (MNHN). Ranobe, 23.00943°S 43.60934°E, 20 m, 1905, F. Geay, 2♀ (MNHN). Réserve Spéciale de Cap Sainte Marie, 12.3 km 262° W Marovato, 25.58166°S 45.16833°E, 200 m, 11 Feb 2002–15 Feb 2002, Fisher et al., 1♂ (00044857) (CAS). Zombitse-Vohibasia National Park, Eastern Sakaraha, Matsabory, 22.8417°S 44.68184°E, 640 m, 13 Feb 1974, P. Viette and A. Peyrieras, 1♂ (MNHN).

*Maraenaspis* Karsch, 1892

Figs. 2.2, 2.5, 2.8, 2.13E, 2.15A, 2.19D, 2.20E, 2.21E, 2.22C; Map 2.4

*Maraenaspis* Karsch, 1892: 134.

TYPE SPECIES: *Maraenaspis typhlops* Karsch, 1892

DISTRIBUTION: Thirty-eight described species of *Maraenaspis* occur throughout the Afrotropics (Maldonado, 1990). Specimens of the only known Madagascan species *Maraenaspis* have been collected from Nosy Be Island off the northwest coast, as well as the Antsiranana, Mahajanga, and Toamasina provinces. Records are from tropical dry forest between 8–135 m elevation.

DISCUSSION: *Maraenaspis* was described by Ferdinand Karsch in 1892. *Maraenaspis bidens* (Reuter), which was originally described from an apterous female, is the only species of *Maraenaspis* known from Madagascar. The male of *M. bidens* is described here for the first time.

*Maraenaspis bidens* (Reuter), 1887

Figs. 2.2, 2.5, 2.8, 2.13E, 2.15A, 2.19D, 2.20E, 2.21E, 2.22C; Map 2.4

*Cleptria bidens* Reuter, 1887: 108.

*Maraenaspis bidens*: Villiers, 1945: 199.

REVISED DIAGNOSIS: Males and females are recognized among other species of *Maraenaspis* by a combination of the following characters: the large body size, coloration, anterior pronotal lobe with very large anterolateral projections, apterous condition, ovoid abdomen that is apically notched, and sternites convex. This species is similar to *M. neglecta* Villiers, 1948, but is distinguished by the slightly larger body size, bluish-black coloration on the head and legs, and very strong anterolateral protuberances on the anterior pronotal lobe.

REDESCRIPTION: **MALE:** Apterous; large body size (length: 21.63–22.08 mm).

COLORATION: Bluish-black suffused with violet sheen on abdominal tergites. Antennal segments III, IV, base of V, and apex of VIII brown and antennal segments V apically, VI, VII, and VIII basally pale to light brown. VESTITURE: Glabrous, except antenna with short, semierect to erect pale to brown setae and tibiae with stout pale orange-brown setae that are

denser near apex. STRUCTURE: HEAD (Fig. 2.13E): Ovoid in lateral view; longer than wide in dorsal view; ventrally flat; antecular region longer than postocular; gula swollen ventrolaterally, produced beyond ventral margin of head, carinate; maxillary plate not reaching dorsal clypeal surface; postclypeus flat; antennal shield not concealing antennal insertion in lateral view; synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about one-fourth longer than scape, slightly curved. THORAX (Fig. 2.15A): Pronotum as long as wide, smooth; anterior pronotal lobe with large anterolateral projections; pronotal longitudinal furrow reaching anterior but not posterior margins, deeply depressed near posterior margin of anterior lobe; pronotal transverse furrow divided by paramedian longitudinal ridges; lateral depressions of posterior pronotal lobe obsolete; meso- and metasterna not completely separated by transverse suture (Fig. 2.15E); mesosternum with medial and paramedian longitudinal depressions; metasternum medially longitudinally depressed; MGE with shallowly depressed meshlike cuticle that does not extend dorsally in lateral view (Fig. 2.16B, C); trochanters and fore- and mid femora basally with small papillae (Fig. 2.16E). ABDOMEN: Apex medially notched; dorsal laterotergite II expanded; intersegmental sutures carinulate between sternites II and III and laterally between III–VII; sternites medially flat; spiracles elliptical; pygophore (Figs. 2.19D, 2.20E) process hooklike in lateral view, subtriangular in caudal view, directed dorsoposteriad, not surpassing posterior margin of pygophore, apex rounded; BPE shorter than basal plate; DPS apex rounded (Fig. 2.21E); endosomal struts reaching posterior margin of DPS, ventrally obscured by semiextended endosoma; area of endosomal struts–DPS fusion circular (Fig. 2.21E); endosoma with platelike sclerite on posterior margin (Fig. 2.21E). **FEMALE:** Body length: 21.5 mm (holotype; original description), 21.30–23.00 mm; coloration and structure same as the male.

DISTRIBUTION: Natural Reserve of Lokobe on Nosy Be Island, and the Antsiranana, Mahajanga, and Toamasina provinces (Map 2.4).

DISCUSSION: This species was described from a single female specimen collected (coll. Stumpff) in “Loucoubé,” which is currently known as the Natural Reserve of Lokobe on Nosy Be Island. Putshkov & Putschkov’s (1986) Ectrichodiinae catalog indicates that the holotype for *M. bidens* is deposited in the National Museum of Natural History, Paris, France, but the type could not be located (Eric Guilbert, personal commun.). Correspondence with other major European collections did not offer any further leads, and we conclude that the type may be lost or placed within undetermined material. Given the potential that the holotype may be located among undetermined material that we have not examined, we refrain from designating a neotype.

The male and female specimens examined match Reuter’s original description of the species, except the antennal coloration. The original description states the last four antennal segments in the female as “straminous” with the “base of the fifth yellowish.” In the females examined, distal antennal segments were missing; in the males, antennal segments III, IV, base of V, and apex of VIII are brown, while segments V apically, VI, VII, and VIII basally are pale to light brown. This slight difference is considered to be variation within the species or between the sexes.

HOLOTYPE: Female: **Madagascar: Toliara:** Nosy Be, Lokobe Natural Reserve, 13.39917°S 48.31833°E, 204 m (see Discussion regarding depository).

OTHER MATERIAL EXAMINED: **Madagascar: Antsiranana:** Forêt d’Anabohazo, 21.6 km 247° WSW Maromandia, 14.30888°S 47.91444°E, 120 m, 11 Mar 2001–16 Mar 2001, Fisher et al., (00044913) (CAS). **Mahajanga:** Parc National d’Ankarafantsika, Forêt de Tsimaloto, 18.3 km 46° NE de Tsaramandroso, 16.22805°S 46.14361°E, 135 m, 02 Apr 2001–08 Apr 2001, Fisher et al., 1♂ (00045017) (CAS). **Toamasina:** Alahakato Forest, 1888, Edouard Perrot, 1♀ (MNHN). Antanambe, 16.43333°S 49.85°E, 8 m, 1898, A. Mocquerys, 1♀ (MNHN).

*Marojejycoris*, new genus

Figs. 2.2, 2.5, 2.6, 2.9, 2.13F, 2.19E, 2.20F, 2.21F, 2.22D; Map 2.5

TYPE SPECIES: *Marojejycoris notadichroa*, new species

DIAGNOSIS: Males are recognized by the small body size, dull coloration, dense vestiture, broadly depressed postclypeus, ocelli located on a shallow medial ocellar tubercle, large antennal shield that does not conceal the antennal insertion in lateral view, continuous transverse suture on the pronotum, MGE with deeply depressed meshlike cuticle that extends dorsally in lateral view, distal part of M not extending beyond apical junction of M+Cu, dorsal laterotergite II not expanded, and circular spiracles. *Marojejycoris* species are similar to the Afrotropical genus *Synavecoris*, but are distinguished by the denser vestiture, head longer than wide in dorsal view, depressed postclypeus, presence of the antennal shield, more medially excavated anterior margin of the pronotum, anterior pronotal lobe lacking carinae laterally, continuous transverse suture, meso- and metasterna separated by a distinct transverse suture, MGE extending dorsally in lateral view, unarmed legs, proximal part of M and Cu veins not fused, distal parts of M and Cu not fused, and dorsal laterotergites uniform in color and not expanded.

DESCRIPTION: **MALE:** Macropterous, small body size. **COLORATION:** Orange-brownish or brown with pale markings. **VESTITURE:** Dense, semierect to erect, long pale to brown setae on head, thorax, legs, corium of hemelytra, and abdomen; setae on tibiae stouter and denser near apex. **STRUCTURE: HEAD (Fig. 2.13F):** Ovoid or subtriangular, longer than wide in dorsal view, shorter than pronotum; anteocular region as long as postocular; ventrally with shallow anteromedial depression; clypeal apex not elevated relative to labrum; maxillary plate not reaching dorsal surface of clypeus (Fig. 2.13E, I); postclypeus broadly depressed (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli present, located on shallow medial tubercle, separated by less than diameter of ocellus; distinct constriction between postocular and neck; gula

moderately swollen ventrolaterally, not distinctly produced beyond ventral head margin; antennal shield not concealing antennal insertion in lateral view; antenna inserted dorsally on head; scape surpassing clypeal apex; pedicel slightly curved; flagellum subdivided into BFLA and DFLA; BFLA divided into two pseudosegments; DFLA divided into two or three pseudosegments; antenna thus appearing 6- or 7-segmented; labium relatively slender; labial segment III shorter than II, ventrally convex (appears dorsal when reposed). THORAX: Pronotum wider than long, anterior margin distinctly concave, collar distinct (Fig. 2.15B); anterior pronotal lobe shorter than posterior lobe (Fig. 2.15B), more than half as wide as posterior lobe, without distinct anterolateral projections, relatively smooth; pronotal longitudinal furrow reaching anterior but not posterior margin of pronotum, weakly to strongly foveate posteriorly (Fig. 2.15B); pronotal transverse furrow continuous; lateral depressions on posterior pronotal lobe distinct (Fig. 2.15B), transversely striated; scutellum with two moderately separated apical processes, disc medially depressed; meso- and metasterna separated by distinct transverse suture (Fig. 2.15F); mesosternum with medial and paramedial longitudinal depressions; metasternum medially longitudinally depressed; MGE with deeply depressed meshlike cuticle that extends dorsally in lateral view (Fig. 2.16A); fossula spongiosa on foretibia, absent on mid tibia in some species; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex; corium restricted to areas adjacent to basal wing veins, with pterostigmalike appearance on anterodistal margin (pl. 17A, C); proximal parts of M and Cu separate (Fig. 2.17A, C); proximal parts of M and Cu separate (Fig. 2.17A, C); distal part of M not extending beyond apical junction of M+Cu; base of M+Cu cell shorter than or as wide as Cu+1A cell (Fig. 2.17B, C). ABDOMEN: Apex rounded; dorsal laterotergites not expanded; sternites longitudinally depressed medially (Fig. 2.18C), intersegmental sutures carinulate; spiracles circular; pygophore (Figs. 2.19E, 2.20F) process flattened in lateral view, broadly rodlike in caudal view, directed dorsoposteriad, not



surpassing posterior margin of pygophore, apex rounded; DPS apex rounded (Fig. 2.21F); endosomal struts reaching posterior margin of DPS, with anterior and posterior ventral processes (Fig. 2.22G); endosoma sclerotized medially (Fig. 2.21F).

ETYMOLOGY: The genus is named after the type locality of the type species, Marojejy National Park, Madagascar. The gender is feminine.

DISTRIBUTION: Species are known from the Antsiranana, Fianarantsoa, and Toamasina provinces and in habitats between 10–1,130 m elevation. Macrohabitats are described as mixed tropical forests, low altitude rainforests and dense humid forests, and forested limestone ridge.

DISCUSSION: Female specimens are unknown for all species in this genus.

*Identification key to the males of species of Marojejycoris*

- 1. DFLA divided into two pseudosegments; antenna thus appearing 6-segmented .....2
- DFLA divided into three pseudosegments; antenna thus appearing 7-segmented .....3
- 2. Postclypeal depression extending to middle of interocular area (Fig. 2.13B); eye about one-fourth head length in lateral view; scape, pedicel, and antennal segments III–IV and base of V dark brown, nearly black; corium orange ..... *auranticorium*, new species
- Postclypeal depression extending to interocular sulcus; eye about one-third head length in lateral view; scape apically, antennal segments II–IV, and base of V brown; corium brown.....  
..... *notadichroa*, new species
- 3. Fossula spongiosa absent or reduced on the mid tibia .....4
- Fossula spongiosa distinctly present on the mid tibia ..... *ranomafana*, new species
- 4. Synthlipsis about two times width of eye; eye not reaching dorsal and ventral head margins.....  
..... *francais*, new species

- Synthlipsis about width of eye; eye reaching dorsal and ventral head margins (Fig. 2.13F).....  
.....*brevifrons*, new species

*Marojejycoris auranticorium*, new species

Figs. 2.2, 2.5, 2.9, 2.19E, 2.20F; Map 2.5

DIAGNOSIS: Males are recognized among other *Marojejycoris* species by the larger body size, coloration of antenna and corium, subtriangular head shape in lateral view, relatively small ocelli, and 6-segmented antenna. This species is similar to *M. notadichroa*, but differs by the uniform dull orange pronotum, orange corium, slightly more shallow postclypeal depression that extends to middle of interocular area, slightly smaller ocelli, and eye about one-fourth of the head length in lateral view.

DESCRIPTION: Body length: 9.10 mm (holotype), 8.54–9.18 mm. COLORATION: Dull orange with thorax, coxae, trochanters, and hemelytron basally light orange-brown. Scape, pedicel, antennal segments III–IV and base of V, posterior margin of pronotum, and wing membrane dark brown. Antennal segment V apically and VI white. VESTITURE: As in generic description. STRUCTURE: HEAD: Subtriangular in lateral view (Fig. 2.13G–I); postclypeus with very shallow medial longitudinal depression to middle of interocular area; synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna 6-segmented, pedicel about one-fourth longer than scape. THORAX: Posterior pronotal lobe smooth; forefemur slightly incrassate; fossula spongiosa present on mid tibia; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; BPE shorter than basal plate (Fig. 2.22D); area of endosomal struts–DPS fusion ovate (Fig. 2.21F).

ETYMOLOGY: The species epithet is a noun in the nominative case and is named for the orange corium.

DISTRIBUTION: Antsiranana and Toamasina provinces (Map 2.5).

DISCUSSION: The coloration varies throughout the body of some specimens from light orange-brown to dull orange.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Montagne Française, 12.325°S 49.33333°E, 150 m, 30 Jan 2001–15 Feb 2001, M. Irwin, R. Harin'Hala (00045023) (CAS).

PARATYPES: **Madagascar: Antsiranana:** 7 km N of Joffreville, 12.33333°S 49.25°E, 360 m, 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00045331) (CAS). Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 15 Jan 2005–26 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00044906) (CAS); 18 May 2005–30 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00045555) (UCR). Montagne Française, 12.325°S 49.33333°E, 150 m, 30 Jan 2001–15 Feb 2001, M. Irwin, R. Harin'Hala, 1♂ (00007120) (USNM); 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala, 2♂ (00007163, 00044966) (SU). Parc National Montagne d'Ambre, 12.51444°S 49.18138°E, 960 m, 19 Mar 2001–05 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00044916) (CAS). Parc National Montagne d'Ambre, 12.51444°S 49.18138°E, 960 m, 04 Mar 2001–19 Mar 2011, M. Irwin, R. Harin'Hala, 1♂ (00045016) (USNM); 19 Mar 2001–05 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006474) (UCR). **Toamasina:** Botanic Garden near entrance to Andasibe National Park, 18.92633°S 48.40783°E, 1025 m, 21 May 2001–04 Jun 2001, M. Irwin, R. Harin'Hala, 1♂ (00006125) (CAS). **Unknown:** 2♂ (00007103, 00044988) (CAS).

*Marojejycoris brevifrons*, new species

Figs. 2.2, 2.5, 2.9, 2.13F, 2.21F; Map 2.5

DIAGNOSIS: Males are recognized among other species in this genus by the small body size, coloration, ovoid head shape, 7-segmented antenna, synthlipsis width about the width of an eye, eye reaching dorsal and ventral head margins, and fossula spongiosa absent on mid tibia. This species is similar to *M. ranomafana*, but is differentiated by the slightly larger body size, dark brown head and corium, lateral areas of anterior pronotal lobe dark brown with medial area pale to light brown, abdomen uniformly light brown to dull orange-brown, smaller synthlipsis width, interocular sulcus near the hind margin of the eye, smooth posterior pronotal lobe, distal part of R present, intersegmental sutures carinulate between sternites II–V (sometimes VI) and laterally between V–VII, and BPE shorter than the basal plate.

DESCRIPTION: Body length: 7.13 mm (holotype), 5.63–7.13 mm. COLORATION: Pale brown with dark brown head (except ventral anteocular surface), scape except basally, pedicel except basally, anterior pronotal lobe laterally, dorsal half of pro- and mesopleura, scutellum except apical processes, wing membrane, femora distally, and tibiae. Postocular ventrolaterally, labium, and antennal segments III and IV orange-brown. Disc of pronotum, posterior pronotal lobe, corium (except basally), and basal wing veins brown. Antennal segments V and VI white. VESTITURE: As in generic description. STRUCTURE: HEAD (Fig. 2.13F): Ovoid in lateral view; postclypeus with very shallow medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli large; eye about one-third of head length, reaching dorsal and ventral head surfaces; antenna 7-segmented, pedicel about as long as scape. THORAX: Posterior pronotal lobe smooth; forefemur slightly incrassate; fossula spongiosa absent on mid tibia; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Intersegmental sutures carinulate

between sternites II–V (sometimes VI) and laterally between V–VII; BPE shorter than basal plate (Fig. 2.22D); area of endosomal struts–DPS fusion ovate (Fig. 2.21F).

ETYMOLOGY: The species epithet is a noun in the nominative case and is named for the short length of the anteocular.

DISTRIBUTION: Antsiranana, Fianarantsoa, and Toamasina provinces (Map 2.5).

DISCUSSION: In some specimens, the intersegmental suture between sternites V and VI is entirely carinulate rather than being restricted to the lateral area.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 09 Apr 2002–16 Apr 2002, M. Irwin, R. Harin'Hala (00006479) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 10 Jan 2005–15 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00044981) (CAS); 26 Jan 2005–04 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00007167) (SU); 04 Apr 2005–16 Apr 2005, M. Irwin, R. Harin'Hala, 1♂ (00007146) (UCR). R.N.I. de Marojejy, 8.0 km NW Manantenina, 14.43667°S 49.775°E, 450 m, 05 Oct 1996–13 Oct 1996, E. Quinter and T. Nguyen, 1♂ (00078365) (AMNH). R.N.I. de Marojejy, 10.0 km NW Manantenina, 14.43333°S 49.76167°E, 750 m, 15 Oct 1996–22 Oct 1996, E. Quinter and T. Nguyen, 1♂ (00078366) (AMNH). **Fianarantsoa:** Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 10 Jan 2002–14 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00006142) (CAS); 12 Feb 2002–19 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00007049) (UCR). Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1100 m, 04 Jun 2002–14 Jun 2002, M. Irwin, R. Harin'Hala, 1♂ (00007051) (CAS). Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 28 Nov 2001–06 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00007122) (CAS); 09 Mar 2003–20 Mar 2003, M. Irwin, R.

Harin'Hala, 1♂ (00007198) (CAS); 17 Dec 2003–30 Dec 2003, M. Irwin, R. Harin'Hala, 1♂ (00007241) (BMNH); 08 Jul 2004–22 Jul 2004, M. Irwin, R. Harin'Hala, 1♂ (00045538) (BMNH); 09 Aug 2005–25 Aug 2005, M. Irwin, R. Harin'Hala, 2♂ (00044830, 00045518) (USNM); 27 Nov 2006–02 Dec 2006, M. Irwin, R. Harin'Hala, 1♂ (00006123) (SU).

**Toamasina:** Forêt d'Analava Mandrisy, 5.9 km 195° Antanambe, 16.48555°S 49.84694°E, 10 m, 12 Nov 2005–13 Nov 2005, Brian L. Fisher et al., 1♂ (00045579) (CAS). Mobot Site, Analalava 7 km SW of Foulpointe, 17.69333°S 49.46027°E, 18 m, 30 Nov 2007–07 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00006480) (CAS). Parc National Mananara-Nord, 7.1 km 261° Antanambe, 16.455°S 49.7875°E, 225 m, 14 Nov 2005, Brian L. Fisher et al., 1♂ (00006471) (CAS).

**Unknown:** 3♂ (00007157, 00007234, 00007239) (CAS).

*Marojejycoris francais*, new species

Figs. 2.2, 2.5, 2.9, 2.22D; Map 2.5

DIAGNOSIS: Males are recognized among other species in *Marojejycoris* by the small body size, ovoid head shape, 7-segmented antenna, fossula spongiosa absent on the mid tibia, and intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI. This species is similar to *M. notadichroa*, from which it differs by the small body size, less elongated and more ovoid head, slightly more shallow postclypeal depression, interocular sulcus near hind margin of the eye, 7-segmented antenna, fossula spongiosa absent on mid tibia, and intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI.

DESCRIPTION: Body length: 6.18 mm (holotype), 6.18–6.82 mm. COLORATION: Dull orange with labial segments III and IV, thoracic sterna, scutellum, corium basally, coxae, trochanters, femora basally, and abdomen pale. Scape apically, pedicel, and antennal segments III–VI dark brown. Wing membrane brown and antennal segment VII white. VESTITURE: As in

generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13D); postclypeus with very shallow medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli large; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna 7-segmented; pedicel about one-third longer than scape. THORAX: Posterior pronotal lobe smooth; forefemur incrassate; fossula spongiosa absent on mid tibia; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI; BPE shorter than basal plate (Fig. 2.22D); area of endosomal struts–DPS fusion ovate (Fig. 2.21F).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Montagne Français, Madagascar.

DISTRIBUTION: Montagne des Français Reserve and Amber Mountain National Park in the Antsiranana province (Map 2.5).

HOLOTYPE: Male: **Madagascar: Antsiranana:** Montagne Français, 12.325°S 49.33333°E, 150 m, 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala (00006470) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Parc National Montagne d'Ambre, 12.51666°S 49.18333°E, 975 m, 25 Jan 2001–11 Feb 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 1♂ (00044900) (UCR). Parc National Montagne d'Ambre, 12.52027°S 49.17916°E, 1125 m, 04 Mar 2001–19 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006129) (CAS).

*Marojejycoris notadichroa*, new species

Figs. 2.2, 2.6, 2.9; Map 2.5

DIAGNOSIS: Males recognized among other species in this genus by a combination of the following characters: the larger body size, more deeply depressed postclypeus, postclypeal

depression extending to the interocular sulcus, and 6-segmented antenna. This species is similar to *M. auranticorium*, from which it differs by the general brown color of the antenna, bicolored pronotum (anterior lobe orange-brown and posterior lobe dark brown), dark brown corium (except basally pale), deeper postclypeal depression that extends to the interocular sulcus, and eye about one-third of the head length in lateral view.

DESCRIPTION: Body length: 8.18 mm (holotype), 7.40–8.73 mm. VESTITURE: As in generic description. COLORATION: Orange with thoracic pleura and sterna, corium basally, and abdomen pale. Scape apically, pedicel, antennal segments III–IV and basal half of V, posterior pronotal lobe, and remainder of hemelytron brown. Apical half of antennal segment V and segment VI entirely white. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13D); postclypeus with relatively deep medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; ocelli large; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna 6-segmented, pedicel about as long as scape. THORAX: Posterior pronotal lobe smooth; forefemur incrassate; fossula spongiosa present on mid tibia; distal part of R present but not forming R+M cell (Fig. 2.17B, C). ABDOMEN: Intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; BPE shorter than basal plate (Fig. 2.22D); area of endosomal struts–DPS fusion ovate (Fig. 2.21F).

ETYMOLOGY: The species epithet is a noun in the nominative case and is named for the bicolored pronotum.

DISTRIBUTION: Antsiranana and Toamasina provinces (Map 2.5).

HOLOTYPE: Male: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 10 Jan 2005–15 Jan 2005, M. Irwin, R. Harin'Hala (00045019) (CAS).



PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 09 Dec 2004–15 Dec 2004, M. Irwin, R. Harin'Hala, 2♂ (00045521, 00045672) (CAS); 15 Dec 2004–20 Dec 2004, M. Irwin, R. Harin'Hala, 2♂ (00044980, 00045013) (USNM); 25 Dec 2004–30 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00044940) (CAS); 26 Jan 2005–04 Feb 2005, M. Irwin, R. Harin'Hala, 2♂ (00007124, 00045532) (UCR); 16 Apr 2005–28 Apr 2005, M. Irwin, R. Harin'Hala, 1♂ (00044910) (SU); 18 May 2005–30 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00007115) (SU); 13 Jul 2005–28 Jul 2005, M. Irwin, R. Harin'Hala, 1♂ (00044969) (UCR). Montaigne Français, 12.325°S 49.33333°E, 150 m, 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006475) (CAS). R.N.I. de Marojejy, 10.0 km NW Manantenina, 14.43333°S 49.76167°E, 750 m, 15 Oct 1996–22 Oct 1996, E. Quinter and T. Nguyen, 1♂ (00078361) (AMNH). **Toamasina:** Forêt d'Analava Mandrisy, 5.9 km 195° Antanambe, 16.48555°S 49.84694°E, 10 m, 12 Nov 2005–13 Nov 2005, Brian L. Fisher et al., 1♂ (00007179) (CAS). **Unknown:** 3♂ (00044991, 00044993, 00045015) (CAS).

*Marojejycoris ranomafana*, new species

Figs. 2.2, 2.6, 2.9; Map 2.5

DIAGNOSIS: Males are recognized among other *Marojejycoris* species by the small body size, bicolored head and thorax in lateral view, ovoid head shape, 7-segmented antenna, striated posterior pronotal lobe, distal part of R absent, intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI, and BPE as long as the basal plate. This species is similar to *M. brevifrons*, but differs by the slightly smaller body size, pale to light brown dorsal and dark brown ventral surfaces of the head and pronotum, pale corium and membrane, wider synthlipsis, interocular sulcus posterior to hind margin of the eye, striated posterior pronotal lobe, fossula

spongiosa on mid tibiae, distal part of R absent, intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI, and BPE as long as the basal plate.

DESCRIPTION: Body length: 5.41 mm (holotype), 5.41–5.50 mm. COLORATION: Brown with head laterally and ventrally, pedicel apically, antennal segments III and IV, dorsal half of thoracic pleura, thoracic sterna, coxae, abdominal sternites II–VI laterally, and small lateral spot on sternite VII dark brown. Antennal segment VII, ventral half of thoracic pleura, scutellum except apical processes, corium, trochanters, femora basally, and remainder of abdomen pale. VESTITURE: As in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13D); postclypeus with very shallow medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; ocelli small; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna 7-segmented; pedicel about as long as scape. THORAX: Posterior pronotal lobe transversely striated; legs slender; fossula spongiosa present on mid tibia; distal part of R absent. ABDOMEN: Intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI; BPE as long as basal plate; area of endosomal struts–DPS fusion elongate subquadrate; endosoma damaged.

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Ranomafana National Park, Madagascar.

DISTRIBUTION: Ranomafana National Park in the Fianarantsoa province (Map 2.5).

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 09 Mar 2003–20 Mar 2003, M. Irwin, R. Harin'Hala (00045464) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1100 m, 04 Feb 2002–12 Feb 2002, M. Irwin, R.

Harin'Hala, 1♂ (00007155) (CAS). Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 21 Dec 2001–24 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00007137) (CAS); 26 Nov 2003–06 Dec 2003, M. Irwin, R. Harin'Hala, 1♂ (00048057) (UCR); 07 Oct 2005–16 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00048058) (UCR).

*Tanindrazanus*, new genus

Figs. 2.2, 2.3, 2.6, 2.7, 2.9, 2.10, 2.13G, 2.13H, 2.15D, 2.15F, 2.16A, 2.17C, 2.18B, 2.19F, 2.20G–I, 2.21G, 2.21H, 2.22E; Maps 2.6–2.9

TYPE SPECIES: *Tanindrazanus irwini*, new species

DIAGNOSIS: Male recognized by the subtriangular head in lateral view, maxillary plate reaching or nearly reaching dorsal clypeal surface, 7-segmented antenna (6-segmented in some specimens of *T. harinhali*), and slender labium with labial segment III ventrally (appears dorsal when folded under head) straight. *Tanindrazanus* is most similar to *Toliarus*, but can be differentiated by the the medium to large body size, maxillary plate reaching or nearly reaching the dorsal clypeal surface, 7-segmented antenna (except some *T. harinhali*), smooth or striated posterior pronotal lobe, and distinct transverse suture separating the meso- and metasterna.

DESCRIPTION: **MALE:** Macropterous, body size medium to large. COLORATION: Uniformly orange-brownish, brown, or black; red and black; orange and black; red and brown; or dark orange, brown, and black. VESTITURE: Dense, semierect to erect, long pale, orange, brown, and dark brown setae on head, thorax, legs, corium of hemelytra, and abdomen; setae on tibiae stouter and denser near apex. STRUCTURE: HEAD (Fig. 2.13G, H): Subtriangular in lateral view, longer than wide in dorsal view, shorter than pronotum; anteocular region longer than postocular; distinct constriction between postocular and neck; gula moderately swollen ventrolaterally, not distinctly produced beyond ventral head margin; maxillary plate reaching or

nearly reaching dorsal surface of clypeus; postclypeus depressed (Fig. 2.13B); ocelli present; antennal shield not concealing antennal insertion in lateral view; antenna inserted dorsally on head; scape surpassing clypeal apex; pedicel slightly curved; flagellum subdivided into BFLA and DFLA; BFLA divided into two pseudosegments; DFLA divided into three pseudosegments (may be two in some specimens of *T. harinhali*); antenna thus appearing 7-segmented (6-segmented in some specimens of *T. harinhali*); labium slender; labial segment III ventrally straight. THORAX (Fig. 2.15D, F): Pronotum wider than long, anterior margin distinctly concave, collar distinct (Fig. 2.15B); anterior pronotal lobe shorter than posterior lobe, more than half as wide as posterior lobe, relatively smooth; pronotal longitudinal furrow reaching anterior but not posterior margin of pronotum, foveate posteriorly (Fig. 2.15B); pronotal transverse furrow distinct (Fig. 2.15B); lateral depressions on posterior pronotal lobe distinct (Fig. 2.15B), transversely striated; scutellum with two broadly separated apical processes, disc medially depressed; meso- and metasterna separated by distinct transverse suture; mesosternum with medial and paramedial longitudinal depressions; metasternum medially longitudinally depressed; MGE with deeply depressed meshlike cuticle that extends dorsally in lateral view (Fig. 2.16A); fossula spongiosa on fore- and mid tibiae; hemelytron (Fig. 2.17C) with corium restricted to areas adjacent to basal wing veins, with pterostigmalike appearance on anterodistal margin; distal part of R present but not forming cell with M; proximal parts of M and Cu separate; distal part of M extending beyond apical junction of M+Cu; base of M+Cu cell shorter than or as wide as Cu+1A cell. ABDOMEN (Fig. 2.18B): Dorsal laterotergite II not distinctly expanded; sternites longitudinally depressed medially (Fig. 2.18C), intersegmental sutures carinulate; spiracles ovoid; pygophore process subtriangular caudal view (Fig. 2.19F, G, I), directed dorsoposteriad, not surpassing posterior margin of pygophore, apex rounded; DPS apex rounded (Fig. 2.20G, H); endosomal struts reaching posterior margin of DPS, with anterior and posterior ventral processes

(Fig. 2.22G); endosoma weakly to heavily sclerotized medially (Fig. 2.21G, H).

ETYMOLOGY: This genus is named after the Madagascan word “Tanindrazana”, which stands for “Fatherland” in the nation’s motto and represents national unity. The gender is masculine.

DISTRIBUTION: Species are known from all provinces and occur in habitats between 10–1,130 m elevation. Macrohabitats are described as gallery forests, tropical dry forests, mixed tropical forests, low- and high-altitude rainforests, sclerophyl forests, dwarf littoral forests, secondary forests, and dry deciduous forests.

DISCUSSION: Female specimens are unknown for this genus. Approximately half of the specimens belonging to *T. harinhali* have the DFLA subdivided into two pseudosegments instead of three and the antenna is therefore 6-segmented. Specimens from the same locality may have 6- or 7-segmented antenna and this polymorphism can be observed even within one individual.

*Identification key to the males of species of Tanindrazanus*

- 1. Clypeal apex not elevated relative to labral base (Fig. 2.13C–F, I–K).....2
  - Clypeal apex slightly to distinctly elevated relative to labral base (Fig. 2.13G, H).....3
- 2. Head elongate, nearly cylindrical; postocular broad in dorsal view (Fig. 2.13B); labial segment II longer than III; forefemora slender ..... *brunneus*, new species
  - Head subtriangular (Fig. 2.13G, H); postocular narrow in dorsal view (Fig. 2.13A); labial segment II and III subequal in length; forefemora incrassate..... *joffrevillus*, new species
- 3. Anterior pronotal lobe laterally carinate .....4
  - Anterior pronotal lobe not laterally carinate.....6
- 4. Body length  $\geq 20$  mm; head ventrally flat; labial segment II shorter than III; red and black or orange and black .....5

- Body length <20 mm; head ventrally with shallow anteromedial depression; labial segment II longer than III; dull orange .....	<i>bemara</i> , new species
5. Red and black; forefemora slightly incrassate; sternal intersegmental sutures carinulate between II and III and laterally between III–V.....	<i>hannajagadae</i> , new species
- Orange and black; forefemora slender; sternal intersegmental sutures carinulate between II–IV and laterally between IV–VI.....	<i>irwini</i> , new species
6. Eye reaching dorsal head surface.....	7
- Eye not reaching dorsal head surface .....	11
7. Eye reaching ventral head surface .....	8
- Eye not reaching ventral head surface.....	<i>tenebricus</i> , new species
8. Pronotal transverse suture complete, not divided by paramedian ridges.....	9
- Pronotal transverse suture incomplete, divided by paramedian ridges (Fig. 2.15B).....	<i>amboasaricus</i> , new species
9. Eye about half head length; posterior pronotal lobe smooth; dorsal laterotergite II very slightly expanded laterally; sternal intersegmental sutures carinulate between II–V.....	<i>vohiparara</i> , new species
- Eye about one-third head length; posterior pronotal lobe smooth or striated; dorsal laterotergite II not distinctly expanded laterally; sternal intersegmental suture carination patterns not as previously mentioned.....	10
10. Pale to dark brown coloration; synthlipsis about width of eye; postocular narrow in dorsal view; posterior pronotal lobe striated; sternal intersegmental sutures carinulate between II–IV and laterally between IV–VII .....	<i>marginatus</i> , new species

- Red and pale to dark brown coloration; synthlipsis less than width of eye; postocular broad in dorsal view; posterior pronotal lobe smooth; sternal intersegmental sutures carinulate between II–V and laterally between V–VI .....	<i>anjozorobeus</i> , new species
11. Labial segment II and III not subequal in length .....	12
- Labial segment II and III subequal in length.....	14
12. Labial segment II shorter than III .....	13
- Labial segment II longer than III.....	<i>andohahela</i> , new species
13. Body length <20 mm; red and black; postclypeus not depressed (Fig. 2.13A); pronotal transverse suture complete, not divided by paramedian ridges .....	<i>marojejy</i> , new species
- Body length ≥20 mm; dark orange-brown or dark orange-red; postclypeus medially depressed (Fig. 2.13B); pronotal transverse suture incomplete, divided by paramedian ridges (Fig. 2.15B).....	<i>mahafaly</i> , new species
14. Pronotal transverse suture complete, not divided by paramedian ridges.....	15
- Pronotal transverse suture incomplete, divided by paramedian ridges (Fig. 2.15B).....	18
15. Red and black, pale red and brown, or uniformly dark coloration; anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); forefemora slender .....	16
- Dull orange with hind femur distally and hind tibia distally black; anterior pronotal lobe without distinct anterolateral projections; forefemora moderately incrassate .....	<i>notatus</i> , new species
16. Synthlipsis width less than two times width of eye; pale red and brown or uniformly dark brown to blackish coloration.....	17
- Synthlipsis width two times width of eye; red and black .....	<i>kathrynae</i> , new species
17. Synthlipsis width about 1.25 times width of eye; uniformly dark brown to blackish coloration .....	<i>simulans</i> , new species

- Synthlipsis about width of eye; pale red and brown..... *antanarivo*, new species
- 18. Black and pale or tricolored; forefemora slightly to moderately incrassate ..... 19
- Bright orange and black; forefemora slender ..... *varicolor*, new species
- 19. Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); sternal intersegmental sutures carinulate between II–IV; black and pale ..... *nigripes*, new species
- Anterior pronotal lobe without distinct anterolateral projections; sternal intersegmental sutures carinulate between II–VI and laterally between VI and VII; tricolored, dark brown, dull orange, and pale ..... *harinhali*, new species

*Tanindrazanus amboasaricus*, new species

Figs. 2.2, 2.6, 2.9; Map 2.9

DIAGNOSIS: Males are recognized among other species in *Tanindrazanus* by a combination of the following characters: the distinct coloration, synthlipsis about the width of the eye, eye reaching the dorsal and ventral head margins, and the pronotal transverse furrow not continuous. This species is similar to *T. harinhali*, from which it differs by the larger body size; dark brown to blackish head, pronotum, and scutellum; the width of the synthlipsis about the width of an eye; and intersegmental sutures carinulate between sternites II–V.

DESCRIPTION: Medium body size (length: 16.05 mm, holotype; 14.30–16.05 mm).

COLORATION: Dark brown with labrum, basal half of scapus, costal margin, and tarsi light brown. Antennifers apically, basal part of hemelytra, trochanters, basal half of femora, and tibiae medially yellowish. Tarsomere III on forelegs with dark brown band in apical part. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; postocular narrow in



dorsal view (Fig. 2.13A); synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); eyes about one-third of head length, reaching dorsal and ventral head surfaces; ocelli large, separated by less than half of diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); pedicel about one-fifth longer than scape; labial segment III subequal to II.

THORAX: Anterior pronotal lobe without anterolateral projections; posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); legs slender; tarsomeres I and II combined subequal to III on fore- and mid legs, longer on hind legs; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–V; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate; area of endosomal struts–DPS fusion tonguelike, apically expanded and rounded.

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality Amboasary, Madagascar.

DISTRIBUTION: Ambovombe in the Amboasary District of the Toliara province (Map 2.9).

DISCUSSION: The coloration on the dorsal surface of the pronotum is sometimes dark brown.

HOLOTYPE: Male: **Madagascar: Toliara:** Amboasary District, Ambovombe, 25.17201°S 46.08971°E, 131 m, 19 Jun 1957, P. Griveaud (MNHN).

PARATYPES: **Madagascar: Toliara:** Amboasary District, Ambovombe, 25.17201°S 46.08971°E, 131 m, 19 Jun 1957, P. Griveaud, 3♂ (MNHN).

*Tanindrazanus andohahela*, new species

Figs. 2.2, 2.6, 2.9; Map 2.7

DIAGNOSIS: Males recognized among other species in this genus by the longer labial segment II relative to III and slightly incrassate forefemur. This species is similar to *T. bemaraha*,

but the shorter labial segment III and lack of anterolateral protuberances on the anterior pronotal lobe distinguish *T. andohahela* from *T. bemaraha*. This species is also similar to *T. notatus*, but differs from it by the slightly incrassate forefemur and lack of black markings on the hind legs.

DESCRIPTION: Medium body size (length: 11.10 mm, holotype; 10.34–11.10 mm).

COLORATION: Orange with antennal segments III–IV and basal half of V dark brown, hemelytron basally (except corium) brown, and antennal segment V apically and VI white.

VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-third longer than scape; labial segment III shorter than II. THORAX: Anterior pronotal lobe without distinct anterolateral projections; posterior pronotal lobe transversely striated; pronotal transverse furrow continuous; forefemur slightly incrassate; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate; area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21G).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Parc National d' Andohahela, Madagascar.

DISTRIBUTION: Antananarivo, Mahajanga, and Toliara provinces (Map 2.7).

HOLOTYPE: Male: **Madagascar: Toliara:** Parc National d'Andohahela, Ihazofotsy Parcelle III, 24.83083°S 46.53616°E, 80 m, 16 Dec 2002–26 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala (00044865) (CAS).

PARATYPES: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 17 Oct 2003–24 Oct 2003, M. Irwin, R. Harin'Hala, 1♂ (00006126) (BMNH), 1♂ (00045559) (USNM); 28 Dec 2003–10 Jan 2004, M. Irwin, R. Harin'Hala, 1♂ (00006093) (UCR); 10 Jan 2004–20 Jan 2004, M. Irwin, R. Harin'Hala, 1♂ (00007177) (CAS), 1♂ (00045466) (SU). **Mahajanga:** 160 km N of Maevatanana on RN 04, Ampijoroa National Park, 16.31933°S 46.81333°E, 43 m, 07 Feb 2005–19 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00045044) (CAS). **Unknown:** 2♂ (00044823, 00044877) (CAS).

*Tanindrazanus anjzorobeus*, new species

Figs. 2.2, 2.6; Map 2.9

DIAGNOSIS: Males are recognized among other *Tanindrazanus* species by the tricolor body (red, orange, and brown coloration), elevated clypeal apex relative to the labrum, synthlipsis smaller than the width of the eye, and broad postocular in dorsal view. This species is very similar to *T. kathrynae*, but it can be easily distinguished by the larger body size, small width of the synthlipsis, larger eyes that surpass the dorsal and ventral margins of the head in lateral view, very large ocelli separated by less than one-third of the diameter of an ocellus, and lack of anterolateral projections on the pronotum.

DESCRIPTION: Large body size (length: 17.32 mm, holotype; 16.20–17.32 mm).

COLORATION: Red with pale orange-brown mandibular and maxillary plates, clypeus, dorsal postocular near ocellar tubercle, labium, anterior pronotal lobe except lateral margins, tibiae except apically, and corium except distally. Remainder of head, lateral margins of anterior

pronotal lobe, anterior part of propleura, meso- and metapleura, scutellum, corium distally, forewing membrane, femora and apical part of tibiae, ventrolateral spot on abdominal sternites II–V, and pygophore dark brown to black. VESTITURE: As in generic description.

STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis smaller than the width of eye; interocular sulcus posterior to hind margin of eye; postocular broad in dorsal view (Fig. 2.13A); ocelli very large, separated by less than one-third diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-third of head length, distinctly surpassing dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III subequal to II.

THORAX: Anterior pronotal lobe without anterolateral projections; posterior pronotal lobe smooth; pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–V and laterally between V and VI; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE longer than basal plate; area of endosomal struts–DPS fusion ovate.

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality Anjozorobe, Madagascar.

DISTRIBUTION: Anjozorobe in the Antananarivo province (Map 2.9).

DISCUSSION: Coloration varies with the pronotal longitudinal furrow either light brown or entirely black and the abdominal sternites dark brown.

HOLOTYPE: Male: **Madagascar: Antananarivo:** Anjozorobe District, Anjozorobe, 18.2619°S 47.6849°E, 1091 m, Dec 1938 (MNHN).

PARATYPES: **Madagascar: Antananarivo:** Anjozorobe District, Anjozorobe, 18.2619°S 47.6849°E, 1091 m, Dec 1938, 3♂ (MNHN).

*Tanindrazanus antananarivo*, new species

Figs. 2.2, 2.6, 2.9; Map 2.6

DIAGNOSIS: Males recognized among other *Tanindrazanus* species by a combination of the following characters: the dark brown color with pale red abdomen, synthlipsis about the width of the eye, eye not reaching the dorsal and ventral head margins, and continuous pronotal transverse furrow. This species is similar to *T. marginatus* and *T. tenebricus*, but is easily distinguished by the dark brown and pale red color pattern and intersegmental sutures carinulate between sternites II–V and laterally between V and VI.

DESCRIPTION: Medium body size (length: 12.52 mm, holotype; 12.41–14.00 mm).

COLORATION: Dark brown with light brown clypeus, antennifer except apically, postocular and neck dorsomedially, head posteroventrally, anteocular laterally and ventrally, labium, basal half of antennal segment V, anterior margin of pronotum, prosternum, corium basally, coxae, trochanters, femora basally, tibiae apically, and tarsi. Meso- and metasterna medially and abdomen, except pygophore, pale red. Distal half of antennal segment V and segments VI and VII white. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; postocular narrow in dorsal view (Fig. 2.13A); synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); eyes about one-third of head length, not reaching dorsal and ventral head surfaces; ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); pedicel about one-fourth longer than scape; labial

segment III subequal to II. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III on fore- and mid legs, longer on hind legs; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–V and laterally between V and VI; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality province Antananarivo, Madagascar.

DISTRIBUTION: Northern region of Antananarivo province (Map 2.6).

DISCUSSION: The dorsal surface of the antennifers and postclypeus are light brown in some specimens.

HOLOTYPE: Male: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 16 Jan 2005–27 Jan 2005, M. Irwin, R. Harin'Hala (00006330) (CAS).

PARATYPES: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 24 Oct 2003–31 Oct 2003, M. Irwin, R. Harin'Hala, 1♂ (00005367) (CAS); 07 Nov 2003–15 Nov 2003, M. Irwin, R. Harin'Hala, 1♂ (00006334) (CAS); 22 Nov 2003–29 Nov 2003, M. Irwin, R. Harin'Hala, 1♂ (00006333) (CAS); 01 Nov 2004–14 Nov 2004, M. Irwin, R. Harin'Hala, 1♂ (00006336) (CAS); 07 Dec 2004–22 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00006335) (CAS); 22 Dec 2004–06 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00006083) (BMNH); 16 Jan 2005–27 Jan 2005, M. Irwin, R. Harin'Hala, 2♂ (00006331, 00006332) (USNM); 23 Mar 2005–03 Apr 2005, M. Irwin, R. Harin'Hala, 2♂ (00006958, 00006959) (UCR); 03 Apr 2005–17 Apr 2005, M. Irwin, R. Harin'Hala, 1♂ (00006127) (SU).

*Tanindrazanus bemaraha*, new species

Figs. 2.2, 2.6, 2.9; Map 2.7

DIAGNOSIS: Males are recognized among other species in *Tanindrazanus* by the longer labial segment II relative to III, anterior pronotal lobe with very small anterolateral protuberances, and anterior pronotal lobe slightly carinate laterally. This species is similar to *T. andohahela* and *T. notatus*, but differs from both species by the presence of the anterolateral protuberances and laterally carinate anterior pronotal lobe. *Tanindrazanus bemaraha* is further distinguished from *T. notatus* by the lack of black markings on the hind femur and tibia and longer labial segment II.

DESCRIPTION: Medium body size (length: 13.93 mm, holotype; 13.07–13.93 mm).

COLORATION: Orange-brown with dark brown scape except basally, pedicel, and antennal segments III and IV, as well as V basally. Antennal segment V apically and segments VI and VII pale. Wing membrane and distal wing veins brown. VESTITURE: As in generic description.

STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-third longer than scape; labial segment III shorter than II. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B), slightly carinate laterally; posterior pronotal lobe transversely striated; pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III on mid and hind tarsi, foretarsi missing; wings spread out in specimens. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII;

pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate; area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21G).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Tsingy de Bemaraha National Park, Madagascar.

DISTRIBUTION: Tsingy de Bemaraha National Park in the Mahajanga province (Map 2.7).

HOLOTYPE: Male: **Madagascar: Mahajanga:** Parc National Tsingy de Bemaraha, 10.6 km ESE 123° Antsalova, 19.70944°S 44.71806°E, 150 m, 16 Nov 2001–20 Nov 2001, Fisher et al. (00006473) (CAS).

PARATYPE: **Madagascar: Mahajanga:** Parc National Tsingy de Bemaraha, 10.6 km ESE 123° Antsalova, 19.70944°S 44.71806°E, 150 m, 16 Nov 2001–20 Nov 2001, Fisher et al., 1♂ (00005362) (CAS).

*Tanindrazanus brunneus*, new species

Figs. 2.2, 2.6, 2.9; Map 2.8

DIAGNOSIS: Males recognized among other species in this genus by the very elongate subtriangular (almost cylindrical) head in lateral view, postclypeal depression extending to interocular sulcus, broad postocular in dorsal view, synthlipsis about 2.5 width of an eye, eyes about one-fifth of the head length, and small ocelli on a shallower ocellar tubercle. This species is similar to *T. joffrevillus*, but is distinguished by the preceding characters, labial segment II longer than III, striated posterior pronotal lobe, and slender legs.

DESCRIPTION: Medium body size (length: 10.31 mm, holotype). COLORATION: Orange with pale thoracic pleura and sterna and brown pronotum and wing membrane. Scape, pedicel, antennal segments III and IV, as well as segment V basally and segment VII apically, dark brown. Antennal segment V apically, segment VI, and base of segment VII white. VESTITURE:



As in generic description. STRUCTURE: HEAD: Elongate; ventrally flat; clypeal apex not elevated relative to labrum (Fig. 2.13C–F, I–K); postclypeus with very shallow, narrow medial longitudinal depression to interocular sulcus; synthlipsis about 2.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli relatively small, separated by less than diameter of ocellus, located on shallow medial tubercle (Fig. 2.13G); eyes about one-fifth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III shorter than II. THORAX: Anterior pronotal lobe without distinct anterolateral projections; posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); legs slender; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process very flattened in lateral view (Fig. 2.20F); BPE as long as basal plate; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the color of the body.

DISTRIBUTION: The only known specimen was collected from Marojejy Nature Reserve in the Antsiranana province (Map 2.8).

HOLOTYPE: Male: **Madagascar: Antsiranana:** R.N.I. de Marojejy, 10.0 km NW Manantenina, 14.43333°S 49.76167°E, 750 m, 15 Oct 1996–22 Oct 1996, E. Quinter and T. Nguyen (00078368) (AMNH).

*Tanindrazanus hannajagoda*, new species

Figs. 2.2, 2.6, 2.9; Map 2.6

DIAGNOSIS: Males are recognized among other *Tanindrazanus* species by the large body size, red and black coloration, flat ventral head surface, labial segment II shorter than III, laterally carinate anterior pronotal lobe, slightly incrassate forefemur, and intersegmental sutures carinate between sternites II and III and laterally between III–V. This species is similar to *T. marojejy*, *T. mahafaly*, and *T. irwini*, but can be separated from the features mentioned, in particular the coloration and sternal intersegmental carination pattern.

DESCRIPTION: Large body size (length: 23.39 mm, holotype; 22.47–23.69 mm).

COLORATION: Red with black antenna (except pedicel basally, apical half of segment V, and segment VI), posterior pronotal margin, scutellar processes apically, hemelytron except base of corium, trochanters, femora basally, and posterolateral stripes on abdominal sternites III–VI.

Hind tibia dark red and distal half of antennal segment V and segment VI white. VESTITURE:

As in generic description. STRUCTURE: HEAD: Ventrally flat; clypeal apex dorsally elevated

relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular

sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli

large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H);

eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about

one-fourth longer than scape; labial segment III longer than II. THORAX: Anterior pronotal lobe

without distinct anterolateral projections, laterally carinate; posterior pronotal lobe transversely

striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B);

forefemur slightly incrassate; tarsomeres I and II combined shorter than III on fore- and mid legs,

subequal on hind legs; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex

rounded or notched medially; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II and III and laterally between III–V; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate; area of endosomal struts–DPS fusion circular.

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after the second author's daughter, Hanna Jagoda Chłond.

DISTRIBUTION: Northern region of the Antsiranana province (Map 2.6).

DISCUSSION: The coloration is variable with the hind femur dark red, the ventral intersegmental sutures black, and the hemelytron light brown in some specimens.

HOLOTYPE: Male: **Madagascar: Antsiranana:** 7 km N of Joffreville, 12.45°S 49.23333°E, 426 m, 11 Feb 2007–18 Feb 2007, M. Irwin, R. Harin'Hala (00006261) (CAS).

PARATYPES: **Madagascar: Antsiranana:** 7 km N of Joffreville, 12.45°S 49.23333°E, 426 m, 20 Jan 2007–27 Jan 2007, M. Irwin, R. Harin'Hala, 1♂ (00006263) (SU); 11 Feb 2007–18 Feb 2007, M. Irwin, R. Harin'Hala, 1♂ (00006262) (UCR). Réserve Spéciale d'Ambre, 3.5 km 235° SW Sakaramy, 12.46888°S 49.24222°E, 325 m, 26 Jan 2001–31 Jan 2001, Fisher et al., 1♂ (00006117) (CAS).

*Tanindrazanus harinhali*, new species

Figs. 2.2, 2.6, 2.9; Map 2.6

DIAGNOSIS: Males are recognized among other species in this genus by a combination of the following characters: the distinct tricolor pattern (dark brown, orange, and pale yellow), synthlipsis about 1.5 times the width of the eye, eye not reaching the dorsal and ventral head margins, ocelli separated by diameter of ocellus, and the pronotal transverse furrow not continuous. This species is similar to *T. amboasarius*, but differs by the smaller body size;

orangish head, pronotum, and scutellum; greater synthlipsis width; and intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII.

DESCRIPTION: Medium body size (length: 10.99 mm, holotype; 10.39–11.89 mm).

COLORATION: Dark brown with pale yellow mandibular and maxillary plates, postclypeus, antennifer dorsally, scape basally, scutellum medially, corium basally, coxae, trochanters, fore- and mid femora except dorsally, hind femur except apically, foretibia, mid tibia except apically, hind tibia medially, tarsi, laterotergites except posterior half of VII, ventrolateral stripes on abdominal sternites III and IV, lateral margins of sternites II–VI, and anterolateral area of VII, labial segment II basally, labial segment IV, prothorax, and dorsal half of mesopleuron orange.

VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-third of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe without distinct anterolateral projections; posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); forefemur slightly incrassate; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE shorter than basal plate; area of endosomal struts–DPS fusion elongate ovoid.

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after Rasolondalao Harin'Hala Hasinjaka, who helped collect a majority of the specimens examined for this species.

DISTRIBUTION: Fianarantsoa and Toliara provinces (Map 2.6).

DISCUSSION: The yellow coloration on the head, legs, and stripes on the abdominal sternites may be reduced in size or absent, and the clypeus, posteroventral head surface, and dorsomedial neck surface may be yellow. Approximately half of the specimens examined have the DFLA subdivided into two pseudosegments instead of three among and within individuals.

HOLOTYPE: Male: **Madagascar: Toliara:** Beza Mahafaly Reserve, Parcelle I near research station, 23.6865°S 44.591°E, 165 m, 28 Nov 2001–04 Dec 2001, R. Harin'Hala (00006418) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 03 Mar 2007–10 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00044892) (CAS). Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 16 Oct 2001–08 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00006425) (CAS); 09 Aug 2005–25 Aug 2005, M. Irwin, R. Harin'Hala, 1♂ (00006537) (UCR). Radio tower 22 km SW of Ilakaka, near Fianarantsoa/Toliara border, 22.77917°S 45.025°E, 1100 m, 06 Mar 2002–11 Mar 2002, M. Irwin, R. Harin'Hala, 1♂ (00006340) (CAS). **Toliara:** Andohahela Natl Park, Tsimelahy, Parcelle II, 24.93683°S 46.62666°E, 180 m, 27 Dec 2002–06 Jan 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006391) (CAS); 06 Jan 2003–16 Jan 2003, R. Harin'Hala, 1♂ (00006376) (CAS), 1♂ (00006377) (USNM); 08 Mar 2003–18 Mar 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006337, 00006344) (SU); 28 Mar 2003–08 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006552) (CAS); 22 Jun 2003–29 Jun 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006375) (AMNH); 17 Aug 2003–24 Aug 2003, M. Irwin, F.

Parker, R. Harin'Hala, 1♂ (00006554) (UCR); 09 Nov 2003–20 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006551) (CAS). Berenty Special Reserve, 8 km NW Amboasary, 25.00666°S 46.30333°E, 85 m, 25 Oct 2002–26 Oct 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006340) (CAS); 26 Oct 2002–02 Nov 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006349) (CAS), 1♂ (00006357) (BMNH); 30 Nov 2002–07 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00005359) (SU), 3♂ (00006353–00006355) (CAS); 26 Jan 2003–05 Feb 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006529) (UCR); 24 Mar 2003–03 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006532, 00045392) (AMNH); 03 May 2003–14 May 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006374) (CAS); 22 Nov 2003–30 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006534) (BMNH); 07 Dec 2003–14 Dec 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006541, 00006542) (USNM); 02 Jan 2004–13 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006547) (CAS); 23 Jan 2004–04 Feb 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006543, 00006546) (MNHN); 04 Feb 2004–15 Feb 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006528, 00006536) (AMNH); 15 Feb 2004–02 Mar 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006358) (SU); 27 May 2004–06 Jun 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006544, 00006545) (CAS); 06 Jun 2004–20 Jun 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006533, 00006553) (USNM); 22 Aug 2004–01 Sep 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006530) (UCR). Berenty Special Reserve, 8 km NW Amboasary, 25.021°S 46.3055°E, 35 m, 21 Dec 2003–01 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006346) (CAS); 02 Jan 2004–13 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006359) (UCR); 24 Mar 2004–04 Apr 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006549, 00006550) (CAS); 04 Apr 2004–15 Apr 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006531) (SU); 23 Jun 2004–04 Jul 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006350, 00006351) (AMNH); 04 Jul 2004–18 Jul 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006356) (CAS). Beza Mahafaly

Reserve, Parcelle II near Bellevue, 23.68983°S 44.5755°E, 180 m, 10 Nov 2001–21 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00006424) (MNHN); 18 Dec 2001–25 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00006426) (SU); 16 Jan 2002–18 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00006385) (CAS); 08 Feb 2002–15 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00006540) (UCR).

Beza Mahafaly Reserve, Parcelle I near research station, 23.6865°S 44.591°E, 165 m, 15 Oct 2001–10 Nov 2001, M.E. Irwin, F.D. Parker, R. Harin'Hala, 1♂ (00006387) (CAS); 10 Nov 2001–21 Nov 2001, R. Harin'Hala, 1♂ (00006557) (UCR); 21 Nov 2001–28 Nov 2001, R. Harin'Hala, 2♂ (00006341, 00006392) (CAS); 28 Nov 2001–04 Dec 2001, R. Harin'Hala, 1♂ (00006417) (CAS); 04 Dec 2001–11 Dec 2001, R. Harin'Hala, 2♂ (00006345, 00006415) (SU); 11 Dec 2001–18 Dec 2001, R. Harin'Hala, 2♂ (00006132, 00006379) (AMNH); 18 Dec 2001–25 Dec 2001, R. Harin'Hala, 1♂ (00006388) (CAS); 25 Dec 2001–02 Jan 2002, R. Harin'Hala, 1♂ (00006548) (BMNH); 18 Jan 2002–25 Jan 2002, M.E. Irwin, F.D. Parker, R. Harin'Hala, 1♂ (00045278) (BMNH); 25 Jan 2002–01 Feb 2002, R. Harin'Hala, 2♂ (00006389, 00006390) (USNM); 08 Feb 2002–15 Feb 2002, R. Harin'Hala, 2♂ (00006342, 00006402) (CAS); 14 Mar 2002–22 Mar 2002, R. Harin'Hala, 1♂ (00006420) (CAS); 29 Mar 2002–10 Apr 2002, R. Harin'Hala, 1♂ (00006555) (CAS); 29 Apr 2002–19 May 2002, R. Harin'Hala, 1♂ (00006386) (CAS); 19 May 2002–08 Jun 2002, R. Harin'Hala, 1♂ (00006401) (CAS); 08 Jun 2002–18 Jun 2002, R. Harin'Hala, 2♂ (00006421, 00006423) (BMNH); 18 Jun 2002–28 Jun 2002, R. Harin'Hala, 1♂ (00006393) (CAS); 07 Jul 2002–18 Jul 2002, R. Harin'Hala, 1♂ (00006412) (CAS); 28 Jul 2002–09 Aug 2002, R. Harin'Hala, 1♂ (00006338) (CAS); 09 Aug 2002–16 Aug 2002, R. Harin'Hala, 1♂ (00006427) (CAS); 16 Aug 2002–28 Aug 2002, R. Harin'Hala, 2♂ (00006403, 00006404) (USNM); 09 Sep 2002–20 Sep 2002, R. Harin'Hala, 1♂ (00006414) (CAS); 20 Sep 2002–05 Oct 2002, R. Harin'Hala, 1♂ (00006422) (UCR); 20 Oct 2002–28 Oct 2002, R. Harin'Hala, 1♂ (00006380) (CAS); 28 Oct 2002–10 Nov 2002, M.E. Irwin, F.D. Parker,

R. Harin'Hala, 2♂ (00006409, 00006410) (BMNH); 02 Dec 2002–12 Dec 2002, R. Harin'Hala, 2♂ (00006347, 00006348) (CAS); 12 Dec 2002–17 Dec 2002, R. Harin'Hala, 1♂ (00006378) (CAS); 17 Dec 2002–20 Dec 2002, R. Harin'Hala, 1♂ (00006419) (CAS); 20 Dec 2002–24 Dec 2002, R. Harin'Hala, 1♂ (00006095) (CAS); 24 Dec 2002–02 Jan 2003, R. Harin'Hala, 1♂ (00006413) (AMNH); 09 Jan 2003–23 Jan 2003, R. Harin'Hala, 4♂ (00006405–00006408) (CAS); 23 Jan 2003–04 Feb 2003, R. Harin'Hala, 5♂ (00006395–00006399) (MNHN); 04 Feb 2003–16 Feb 2003, R. Harin'Hala, 1♂ (00006400) (SU); 16 Feb 2003–02 Mar 2003, R. Harin'Hala, 1♂ (00006556) (BMNH); 13 Mar 2003–23 Mar 2003, R. Harin'Hala, 2♂ (00006381, 00006382) (USNM); 23 Mar 2003–02 Apr 2003, R. Harin'Hala, 1♂ (00006339) (CAS); 02 Apr 2003–10 Apr 2003, R. Harin'Hala, 1♂ (00006352) (CAS); 29 Apr 2003–07 May 2003, R. Harin'Hala, 1♂ (00006416) (UCR). Parc National d'Andohahela, Ihazofotsy Parcelle III, 24.83083°S 46.53616°E, 80 m, 06 May 2003–13 May 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006343) (SU); 29 Jun 2003–06 Jul 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00044843) (UCR); 03 Aug 2003–13 Aug 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00044809, 00044837) (CAS). **Unknown:** 5♂ (00006383, 00006384, 00006535, 00006538, 00006539) (CAS).

*Tanindrazanus irwini*, new species

Figs. 2.2, 2.6, 2.9, 2.15F; Map 2.7

DIAGNOSIS: Males are recognized among other species in *Tanindrazanus* by the large body size, black coloration with orange markings, flat ventral head surface, labial segment II shorter than III, laterally carinate anterior pronotal lobe, slender legs, and intersegmental sutures carinulate between sternite II–IV and laterally between IV–VI. This species is similar to *T. varicolor*, but differs by the larger body size, flat ventral head surface, labial segment II shorter



than III, laterally carinate anterior pronotal lobe, and intersegmental sutures carinulate between sternite II–VI and laterally between VI and VII.

DESCRIPTION: Large body size (length: 22.76 mm, holotype; 18.95–22.76 mm).

COLORATION: Dark brown-black with bright orange head dorsally, scape, labium except segment II basally, scutellum medially, hemelytron except posteromedial spot, femora apically, tibiae, tarsi, laterotergites II–V, anterior half of laterotergite VI, sternites III–V, and anterior half of sternite VI except posterolateral stripes. Distal half of antennal segment V and segments VI and VII white. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally flat; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III longer than II. THORAX (Fig. 2.15F): Anterior pronotal lobe without distinct anterolateral projections, laterally carinate; posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); legs slender; tarsomeres I and II combined subequal to III; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternite II–IV and laterally between IV–VI; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after Mike Irwin, who helped collect a majority of the specimens examined for this species.

DISTRIBUTION: Marojejy Nature Reserve in the Antsiranana province and near the Andasibe-Mantadia National Park in the Toamasina province (Map 2.7).

DISCUSSION: Coloration is variable with the orange areas reduced or absent on the dorsal head surface, labium, scutellum, femora, hind tibia, and abdomen in some specimens.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 30 May 2005–11 Jun 2005, M. Irwin, R. Harin'Hala (00045088) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 09 Dec 2004–15 Dec 2004, M. Irwin, R. Harin'Hala, 2♂ (00045220, 00045375) (CAS); 25 Dec 2004–30 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00006224) (SU), 1♂ (00006225) (UCR); 05 Jan 2005–10 Jan 2005, M. Irwin, R. Harin'Hala, 2♂ (00006096, 00006097) (CAS); 10 Jan 2005–15 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00006235) (UCR); 15 Jan 2005–26 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00006135) (AMNH); 04 Feb 2005–11 Feb 2005, M. Irwin, R. Harin'Hala, 3♂ (00006233, 00006234, 00006239) (CAS); 11 Feb 2005–18 Feb 2005, M. Irwin, R. Harin'Hala, 3♂ (00006217, 00006241, 00006242) (AMNH); 18 Feb 2005–25 Feb 2005, M. Irwin, R. Harin'Hala, 3♂ (00045603, 00045612, 00045712) (BMNH); 25 Feb 2005–04 Mar 2005, M. Irwin, R. Harin'Hala, 5♂ (00005366, 00006226–00006229) (CAS); 11 Mar 2005–18 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00006232) (UCR); 04 Apr 2005–16 Apr 2005, M. Irwin, R. Harin'Hala, 2♂ (00006499, 00045339) (SU); 16 Apr 2005–28 Apr 2005, M. Irwin, R. Harin'Hala, 6♂ (00007087, 00045004, 00045048, 00045072, 00045405, 00045475) (CAS); 28 Apr 2005–07 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00006221) (UCR); 07 May 2005–18 May 2005, M. Irwin, R. Harin'Hala, 3♂ (00006134, 00006237, 00006238) (USNM); 18 May 2005–30 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00006216) (BMNH), 1♂ (00006243) (USNM); 30 May 2005–11 Jun

2005, M. Irwin, R. Harin'Hala, 4♂ (00007211, 00044957, 00044982, 00045462) (CAS); 28 Jun 2005–13 Jul 2005, M. Irwin, R. Harin'Hala, 1♂ (00006218) (SU), 1♂ (00006222) (UCR), 4♂ (00006223, 00006230, 00006231, 00006236) (MNHN), 4♂ (00006240, 00006452–00006454) (CAS). **Toamasina:** Lakato env. Near Andasibe-Mantadia National Park, 19.18651°S 48.43856°E, 635 m, 09 Jan 2007–10 Jan 2007, Z. Mráček, 1♂ (MMBC).

*Tanindrazanus joffrevillus*, new species

Figs. 2.2, 2.6, 2.9; Map 2.6

DIAGNOSIS: Males recognized among other species in this genus by the dull orange coloration, ventrally flat head surface, clypeal apex not elevated relative to the labrum, smooth posterior pronotal lobe, and slightly incrassate forefemur. This species is similar to *T. andohahela*, but is distinguished by dark brown corium, ventrally flat head surface, clypeal apex not elevated relative to the labrum, larger synthlipsis width, subequal lengths of labial segments II and III, smooth posterior pronotal lobe, divided pronotal transverse suture, and intersegmental sutures carinate between sternites II–VI.

DESCRIPTION: Medium body size (length: 11.19 mm, holotype). COLORATION: Dull orange with thoracic sterna pale and hemelytron, except corium basally, brown. Antennal segments V and VI white. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally flat; clypeal apex not elevated relative to labrum (Fig. 2.13C–F, I–K); postclypeus with relatively shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eyes about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial

segment III subequal II. THORAX: Anterior pronotal lobe without distinct anterolateral projections; posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); forefemur slightly incrassate; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–VI; pygophore process very flattened in lateral view (Fig. 2.20F); BPE as long as basal plate; area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21G).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Joffreville, Madagascar.

DISTRIBUTION: Only one of the two known specimens has locality data. It is known from a locality north of Joffreville in the Antsiranana province (Map 2.6).

HOLOTYPE: Male: **Madagascar: Antsiranana:** 7 km N of Joffreville, 12.33333°S 49.25°E, 360 m, 20 Mar 2001–07 Apr 2001, M. Irwin, R. Harin’Hala (00007256) (CAS). **Unknown**, 1♂ (00044998) (CAS).

PARATYPE: **Madagascar: Unknown**, 1♂ (00044998) (CAS).

*Tanindrazanus kathrynae*, new species

Figs. 2.2, 2.6, 2.9, 2.13G, 2.15D, 2.19F, 2.20G; Map 2.7

DIAGNOSIS: Males recognized among other *Tanindrazanus* species by the red and black coloration, elevated clypeal apex relative to the labrum, broad postocular in dorsal view, and synthlipsis about two times the width of the eye. This species is very similar to *T. anjozorobeus*, but is distinguished by the smaller body size, smaller eyes that do not surpass the dorsal and ventral margins of the head in lateral view, larger width of the synthlipsis, and the very small anterolateral projections on the pronotum.

DESCRIPTION: Medium body size (length: 12.67 mm, holotype; 11.62–12.67 mm).

COLORATION: Red with pale orange-brown mandibular and maxillary plates, clypeus, anterolateral and dorsal margin of eye, labium, anterior pronotal lobe except lateral margins, and corium except distally. Remainder of head except two paramedian dorsal stripes posterior to ocelli, lateral margins of anterior pronotal lobe, anterior part of propleura, meso- and metapleura, scutellum, corium distally, forewing membrane, femora and apical one-third to one-half of fore- and mid tibiae, hind tibiae except small apical part, ventrolateral spot on abdominal sternites II–VI, and pygophore dark brown to black. VESTITURE: As in generic description. STRUCTURE: HEAD (Fig. 2.13G): Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum; postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; postocular broad in dorsal view (Fig. 2.13B); ocelli large, separated by less than diameter of ocellus, located on slightly elevated median tubercle; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about two-fifths longer than scape; labial segment III subequal to II. THORAX (Fig. 2.15D): Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–V and laterally between V and VI; pygophore process subtriangular in lateral view (Fig. 2.20G); BPE longer than basal plate; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is a noun in the genitive case and is named in memoriam of the senior author's partner's mother, Kathryn Carroll.

DISTRIBUTION: Miandritsara Forest in the Fianarantsoa province (Map 2.7).

DISCUSSION: In some specimens, the head and anterior pronotal lobe have reduced red coloration or are entirely black and the median longitudinal suture on the pronotum and the medial area of abdominal sternite VII are black.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Miandritsara Forest, 40 km S Ambositra, 20.79266°S 47.17566°E, 825 m, 13 Nov 2006–22 Nov 2006, M. Irwin, R. Harin'Hala (00006103) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** Andringitra Est, Ambalamarovandana, 22.22595°S 46.93418°E, 1500–1600m, 15 Jan 1971–25 Jan 1971, 1♂ (MNHN). Miandritsara Forest, 40 km S Ambositra, 20.79266°S 47.17566°E, 825 m, 23 Oct 2004–03 Nov 2004, M. Irwin, R. Harin'Hala, 1♂ (00045053) (UCR); 05 Jan 2005–18 Jan 2005, M. Irwin, R. Harin'Hala, 2♂ (00044858, 00045506) (CAS), 1♂ (00045608) (SU); 09 Feb 2005–22 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00045383) (AMNH); 22 Feb 2005–05 Mar 2005, M. Irwin, R. Harin'Hala, 2♂ (00044863, 00045360) (CAS), 1♂ (00045626) (USNM); 13 Apr 2005–27 Apr 2005, M. Irwin, R. Harin'Hala, 2♂ (00007062, 00044984) (CAS), 1♂ (00045549) (MNHN); 05 Jan 2006–12 Jan 2006, M. Irwin, R. Harin'Hala, 2♂ (00044818, 00045566) (BMNH); 16 Feb 2006–27 Feb 2006, M. Irwin, R. Harin'Hala, 1♂ (00044832) (BMNH); 27 Feb 2006–08 Mar 2006, M. Irwin, R. Harin'Hala, 1♂ (00045632) (SU); 14 Apr 2006–02 May 2006, M. Irwin, R. Harin'Hala, 2♂ (00006094, 00006298) (CAS), 1♂ (00006312) (MNHN); 13 Nov 2006–22 Nov 2006, M. Irwin, R. Harin'Hala, 2♂ (00006293, 00006299) (AMNH); 01 Dec 2006–10 Dec 2006, M. Irwin, R. Harin'Hala, 1♂ (00006302) (UCR); 10 Dec 2006–19 Dec 2006, M. Irwin, R. Harin'Hala, 2♂ (00006313, 00006314) (CAS); 19 Dec 2006–28 Dec 2006, M. Irwin, R. Harin'Hala, 4♂ (00007053, 00007202, 00045201, 00045365) (CAS); 15 Jan 2007–24 Jan 2007, M. Irwin, R. Harin'Hala, 2♂ (00045083, 00045482) (CAS); 24 Jan 2007–31 Jan 2007, M. Irwin, R. Harin'Hala, 1♂ (00006294) (BMNH); 31 Jan 2007–11 Feb 2007, M. Irwin, R. Harin'Hala, 2♂

(00006315, 00006316) (SU); 11 Feb 2007–20 Feb 2007, M. Irwin, R. Harin'Hala, 1♂  
(00006301) (MNHN), 3♂ (00006309–00006311) (CAS); 20 Feb 2007–27 Feb 2007, M. Irwin, R.  
Harin'Hala, 1♂ (00006297) (USNM); 08 Mar 2007–17 Mar 2007, M. Irwin, R. Harin'Hala, 4♂  
(00006303–00006306) (CAS); 17 Mar 2007–26 Mar 2007, M. Irwin, R. Harin'Hala, 1♂  
(00006300) (AMNH); 26 Mar 2007–05 Apr 2007, M. Irwin, R. Harin'Hala, 2♂ (00006307,  
00006308) (USNM); 05 Apr 2007–14 Apr 2007, M. Irwin, R. Harin'Hala, 2♂ (00006295,  
00006296) (UCR).

*Tanindrazanus mahafaly*, new species

Figs. 2.3, 2.6, 2.9; Map 2.7

DIAGNOSIS: Males are recognized among other species in this genus by a combination of the following characters: the large body size, distinct dark orange-brown to red-orange coloration, labial segment II shorter than III, slightly incrassate forefemur, and only the intersegmental suture between sternites II and III carinulate. This species is similar to *T. irwini*, from which it differs by the dark orange-red color pattern, anteromedially depressed ventral head surface, and carination pattern on the sternal intersegmental sutures.

DESCRIPTION: Large body size (length: 19.57 mm, holotype; 18.03–19.92 mm).

COLORATION: Dark orange with apical half of pedicel and basiflagellomere, anterior pronotal lobe, metapleuron, and abdomen, except laterotergites, dark reddish brown. Hemelytron bright orange and antennal segments VI and VII pale orange to white. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal

view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-third of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III longer than II.

THORAX: Anterior pronotal lobe without distinct anterolateral projections; posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); forefemur slightly incrassate; tarsomeres I and II combined subequal to III on fore- and mid legs, longer on hind legs; wings spread in specimens. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II and III; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate; area of endosomal struts–DPS fusion circular.

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Beza Mahafaly Reserve, Madagascar.

DISTRIBUTION: Beza Mahafaly Reserve in the Toliara province (Map 2.7).

DISCUSSION: In some specimens, the head, scutellum, fore- and mid legs, and abdomen are bright red-orange, while the remaining structures are dull orange.

HOLOTYPE: Male: **Madagascar: Toliara:** Beza Mahafaly Reserve, Parcelle I near research station, 23.6865°S 44.591°E, 165 m, 15 Oct 2001–10 Nov 2001, M. Irwin, F. Parker, R. Harin'Hala (00045290) (CAS).

PARATYPES: **Madagascar: Toliara:** Beza Mahafaly Reserve, Parcelle II near Bellevue, 23.68983°S 44.5755°E, 180 m, 21 Nov 2001–28 Nov 2001, R. Harin'Hala, 1♂ (00006244) (CAS). Beza Mahafaly Reserve, Parcelle I near research station, 23.6865°S 44.591°E, 165 m, 04 Dec 2001–11 Dec 2001, R. Harin'Hala, 1♂ (00006245) (CAS); 08 Feb 2002–15 Feb 2002, R. Harin'Hala, 1♂ (00006133) (CAS); 22 Feb 2002–01 Mar 2002, R. Harin'Hala, 1♂ (00006220) (SU); 18 Jun 2002–28 Jun 2002, R. Harin'Hala, 1♂ (00006219) (UCR).



*Tanindrazanus marginatus*, new species

Figs. 2.3, 2.6, 2.9; Map 2.8

DIAGNOSIS: Males are recognized among other species in this genus by the brownish coloration, synthlipsis about the width of an eye, and large eyes reaching the dorsal and ventral head surfaces. This species is similar to *T. tenebricus* and *T. nigripes*, but differs from both species by the lighter coloration and lateral carination on the intersegmental suture between sternites VI and VII. *Tanindrazanus marginatus* is further distinguished from *T. nigripes* by the smaller synthlipsis width.

DESCRIPTION: Medium body size (length: 16.43 mm, holotype; 13.62–17.43 mm).

COLORATION: Dark brown with light brown clypeus, postclypeus medially, dorsal margin of ocellar tubercle, labium, basal wing veins, trochanters, femora except apically, tibiae basally, tarsi, and lateral margin of sternites. Distal half of antennal segment V and segments VI and VII white. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-third of head length, reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III on forelegs, slightly longer on mid and hind legs; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinate between sternite

II–IV and laterally between IV–VII; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE longer than basal plate; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the pale laterotergites.

DISTRIBUTION: Antsiranana, Fianarantsoa, Toamasina, and Toliara provinces (Map 2.8).

DISCUSSION: In some specimens, the dorsal neck surface, thorax, corium, coxae, and laterotergites are light brown and the postclypeus medially dark brown.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1110 m, 08 Apr 2002–15 Apr 2002, M. Irwin, R. Harin'Hala (00006821) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 04 Feb 2005–11 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00006892) (CAS); 18 Mar 2005–25 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00006888) (CAS); 28 Apr 2005–07 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00006908) (USNM); 13 Jul 2005–28 Jul 2005, M. Irwin, R. Harin'Hala, 1♂ (00006087) (AMNH); 14 Oct 2005–22 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00006850) (SU). **Montaigne Francais**, 12.325°S 49.33333°E, 150 m, 30 Jan 2001–15 Feb 2001, M. Irwin, R. Harin'Hala, 2♂ (00006833, 00006847) (CAS), 1♂ (00006921) (AMNH); 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006920) (UCR). **Parc National Montagne d'Ambre**, 12.51444°S 49.18138°E, 960 m, 12 Feb 2001–04 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006932) (CAS). **Parc National Montagne d'Ambre**, 12.52027°S 49.17916°E, 1125 m, 30 May 2001–06 Jun 2001, M. Irwin, R. Harin'Hala, 1♂ (00006834) (UCR). **Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 13 Jan 2007–20 Jan 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006931) (MNHN); 04 Feb 2007–10 Feb 2007, M. Irwin,

F. Parker, R. Harin'Hala, 4♂ (00006805, 00006806, 00006855, 00006860) (CAS); 10 Feb 2007–17 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006803, 00006804, 00006861) (AMNH); 03 Mar 2007–10 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006941) (UCR); 17 Mar 2007–24 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006936) (SU); 24 Mar 2007–31 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006810) (CAS); 07 Apr 2007–14 Apr 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006937) (CAS); 14 Apr 2007–21 Apr 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006942) (CAS); 06 May 2007–14 May 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006801) (BMNH); 14 May 2007–21 May 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006943) (MNHN); 02 Jun 2007–10 Jun 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006930) (CAS); 07 Jul 2007–14 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006934) (CAS); 14 Jul 2007–19 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006926) (SU); 19 Jul 2007–23 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006927) (SU); 23 Jul 2007–28 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006928) (AMNH); 03 Aug 2007–11 Aug 2007, M. Irwin, F. Parker, R. Harin'Hala, 4♂ (00006811, 00006857, 00006858, 00006938) (USNM); 27 Aug 2007–06 Sep 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006929) (CAS); 06 Sep 2007–13 Sep 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006856, 00006944) (MNHN); 27 Sep 2007–04 Oct 2007, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006807–00006809) (CAS); 25 Oct 2007–08 Nov 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006922) (CAS); 08 Nov 2007–22 Nov 2007, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006812, 00006859, 00006945) (BMNH); 28 Nov 2007–13 Dec 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006935) (USNM); 03 Jan 2008–09 Jan 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006907) (SU); 09 Jan 2008–17 Jan 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006836) (UCR); 01 Apr 2008–09 Apr 2008, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006887, 00006890) (CAS). Manombo Special Reserve camp site, 32 km SSE

of Farafangana, 23.02183°S 47.72°E, 36 m, 10 Oct 2004–21 Oct 2004, M. Irwin, R. Harin'Hala, 1♂ (00045347) (CAS); 23 Nov 2004–05 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00045266) (UCR); 15 Dec 2004–26 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00045207) (AMNH); 26 Dec 2004–05 Jan 2005, M. Irwin, R. Harin'Hala, 3♂ (00045006, 00045195, 00045522) (CAS); 16 Jan 2005–20 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00045467) (SU); 15 Feb 2005–27 Feb 2005, M. Irwin, R. Harin'Hala, 2♂ (00044964, 00045376) (CAS); 24 Apr 2005–10 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00044810) (BMNH); 10 May 2005–22 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00045152) (BMNH); 06 Jun 2005–24 Jun 2005, M. Irwin, R. Harin'Hala, 2♂ (00007190, 00045382) (USNM); 24 Jun 2005–10 Jul 2005, M. Irwin, R. Harin'Hala, 1♂ (00044842) (MNHN); 10 Jul 2005–27 Jul 2005, M. Irwin, R. Harin'Hala, 1♂ (00045119) (CAS); 09 Oct 2005–16 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00044946) (UCR). Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 15 Nov 2001–22 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00045415) (CAS); 28 Jan 2002–04 Feb 2002, M. Irwin, R. Harin'Hala, 2♂ (00006879, 00006880) (CAS); 12 Feb 2002–19 Feb 2002, M. Irwin, R. Harin'Hala, 2♂ (00006835, 00006940) (SU); 14 Apr 2002–23 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00045707) (CAS); 03 Jun 2002–13 Jun 2002, M. Irwin, R. Harin'Hala, 1♂ (00006904) (MNHN); 13 Jun 2002–23 Jun 2002, M. Irwin, R. Harin'Hala, 1♂ (00006905) (CAS); 16 Feb 2003–26 Feb 2003, M. Irwin, R. Harin'Hala, 2♂ (00006883, 00006884) (USNM); 21 Mar 2003–12 Apr 2003, M. Irwin, R. Harin'Hala, 1♂ (00006891) (BMNH); 04 May 2003–16 May 2003, M. Irwin, R. Harin'Hala, 2♂ (00006885, 00006886) (AMNH); 15 May 2003–28 May 2003, M. Irwin, R. Harin'Hala, 1♂ (00006960) (CAS); 28 May 2003–06 Jun 2003, M. Irwin, R. Harin'Hala, 1♂ (00045034) (UCR). Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1110 m, 19 Feb 2002–26 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00006902) (CAS); 04 Mar 2002–12 Mar 2002, M. Irwin, R. Harin'Hala, 1♂ (00006837) (SU);

26 Mar 2002–31 Mar 2002, M. Irwin, R. Harin'Hala, 3♂ (00006844–00006846) (CAS), 1♂ (00006924) (USNM); 31 Mar 2002–08 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00006903) (USNM); 08 Apr 2002–15 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00006799) (AMNH), 17♂ (00006815–00006820, 00006822–00006830) (CAS), 3♂ (00006843, 00006925, 00006933) (UCR). Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 06 Dec 2001–15 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00006917) (CAS); 21 Dec 2001–24 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00044908) (UCR); 14 Jan 2002–21 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00045718) (CAS); 28 Jan 2002–04 Feb 2002, M. Irwin, R. Harin'Hala, 1♂ (00099047) (CAS); 09 Apr 2002–16 Apr 2002, M. Irwin, R. Harin'Hala, 1♂ (00006896) (USNM); 30 Apr 2002–07 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00006839) (AMNH); 07 May 2002–14 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00006802) (CAS); 18 Feb 2003–27 Feb 2003, M. Irwin, R. Harin'Hala, 1♂ (00007112) (MNHN); 09 Mar 2003–20 Mar 2003, M. Irwin, R. Harin'Hala, 1♂ (00006895) (BMNH); 07 May 2003–17 May 2003, M. Irwin, R. Harin'Hala, 2♂ (00006874, 00006875) (SU); 06 Aug 2003–16 Aug 2003, M. Irwin, R. Harin'Hala, 3♂ (00006911–00006913) (CAS), 1♂ (00006919) (BMNH); 28 Sep 2003–08 Oct 2003, M. Irwin, R. Harin'Hala, 1♂ (00006898) (CAS); 18 Oct 2003–26 Oct 2003, M. Irwin, R. Harin'Hala, 1♂ (00006899) (CAS); 17 Dec 2003–30 Dec 2003, M. Irwin, R. Harin'Hala, 2♂ (00006831, 00006832) (CAS); 30 Dec 2003–11 Jan 2004, M. Irwin, R. Harin'Hala, 1♂ (00006914) (CAS); 11 Jan 2004–21 Jan 2004, M. Irwin, R. Harin'Hala, 2♂ (00006881, 00006882) (MNHN); 31 Jan 2004–12 Feb 2004, M. Irwin, R. Harin'Hala, 1♂ (00006894) (CAS); 21 Mar 2004–02 Apr 2004, M. Irwin, R. Harin'Hala, 2♂ (00006796, 00006797) (AMNH); 02 Apr 2004–15 Apr 2004, M. Irwin, R. Harin'Hala, 1♂ (00006840) (USNM); 24 Jun 2004–08 Jul 2004, M. Irwin, R. Harin'Hala, 2♂ (00006838, 00006897) (BMNH); 07 Oct 2004–20 Oct 2004, M. Irwin, R. Harin'Hala, 1♂ (00006918) (UCR); 14 Jan 2005–02 Feb 2005, M. Irwin, R. Harin'Hala, 1♂

(00006868) (SU); 27 Feb 2005–13 Mar 2005, M. Irwin, R. Harin'Hala, 3♂ (00006841, 00006842, 00006916) (CAS); 28 Mar 2005–10 Apr 2005, M. Irwin, R. Harin'Hala, 1♂ (00006900) (BMNH); 25 Sep 2005–07 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00006869) (CAS); 06 Nov 2005–13 Nov 2005, M. Irwin, R. Harin'Hala, 1♂ (00006946) (CAS); 13 Nov 2005–24 Nov 2005, M. Irwin, R. Harin'Hala, 1♂ (00006865) (MNHN); 21 Dec 2005–30 Dec 2005, M. Irwin, R. Harin'Hala, 1♂ (00006866) (BMNH); 20 Jan 2006–29 Jan 2006, M. Irwin, R. Harin'Hala, 2♂ (00006872, 00006873) (SU); 29 Jan 2006–07 Feb 2006, M. Irwin, R. Harin'Hala, 1♂ (00006915) (CAS); 15 Feb 2006–26 Feb 2006, M. Irwin, R. Harin'Hala, 9♂ (00006947–00006950, 00006953–00006957) (CAS); 05 Mar 2006–18 Mar 2006, M. Irwin, R. Harin'Hala, 2♂ (00006854, 00006867) (UCR); 18 Mar 2006–30 Mar 2006, M. Irwin, R. Harin'Hala, 3♂ (00006862–00006864) (MNHN); 13 Apr 2006–24 Apr 2006, M. Irwin, R. Harin'Hala, 1♂ (00006870) (AMNH); 02 May 2006–13 May 2006, M. Irwin, R. Harin'Hala, 2♂ (00006871, 00006901) (MNHN); 20 May 2006–29 May 2006, M. Irwin, R. Harin'Hala, 1♂ (00006893) (CAS); 09 Aug 2006–23 Aug 2006, M. Irwin, R. Harin'Hala, 1♂ (00006800) (USNM); 23 Aug 2006–07 Sep 2006, M. Irwin, R. Harin'Hala, 1♂ (00006853) (CAS); 07 Sep 2006–20 Sep 2006, M. Irwin, R. Harin'Hala, 3♂ (00006851, 00006852, 00006951) (BMNH); 20 Sep 2006–05 Oct 2006, M. Irwin, R. Harin'Hala, 1♂ (00006952) (SU); 01 Nov 2006–11 Nov 2006, M. Irwin, R. Harin'Hala, 1♂ (00044923) (UCR). Parc National d'Isalo, 9.1 km 354° N Ranohira, 22.48166°S 45.46166°E, 725 m, 27 Jan 2003–31 Jan 2003, Fisher et al., 1♂ (00005363) (AMNH), 3♂ (00006876–00006878) (CAS). Ranomafana JIRAMA water works, 21.2485°S 47.45216°E, 690 m, 16 Oct 2001–08 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00005361) (CAS). **Toamasina:** Botanic Garden near entrance to Andasibe National Park, 18.92633°S 48.40783°E, 1025 m, 01 Sep 2001–05 Sep 2001, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006889) (CAS). Fampanambo, 15.37550°S 49.62175°E, 108 m, 1962, J. Vadon, 1♂ (MRAC). Mobot Site,

Analalava 7 km SW of Foulpointe, 17.69333°S 49.46027°E, 18 m, 21 Dec 2007–28 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00044874) (SU); 18 Jan 2008–25 Jan 2008, M. Irwin, R. Harin'Hala, 2♂ (00006848, 00006909) (CAS), 1♂ (00044801) (USNM); 25 Jan 2008–01 Feb 2008, M. Irwin, R. Harin'Hala, 1♂ (00006923) (CAS), 1♂ (00006939) (AMNH); 21 Mar 2008–28 Mar 2008, M. Irwin, R. Harin'Hala, 1♂ (00006798) (UCR). Mobot Site, Analalava 7 km SW of Foulpointe, 17.70889°S 49.45806°E, 24 m, 15 Sep 2007–21 Sep 2007, M. Irwin, R. Harin'Hala, 1♂ (00006906) (CAS). **Toliara:** Kirindy forest, 60 km NE of Morondava, 20.0665°S 44.65767°E, 45 m, 18 Oct 2003–30 Oct 2003, M. Irwin, R. Harin'Hala, 1♂ (00045067) (CAS). **Unknown:** 3♂ (00006813, 00006814, 00006849) (CAS).

*Tanindrazanus marojejy*, new species

Figs. 2.3, 2.6, 2.9, 2.20H; Map 2.6

DIAGNOSIS: Males recognized among other *Tanindrazanus* species by the black body with red abdomen and pale orange hemelytra, ventrally flat head surface, broad postocular in dorsal view, labial segment II shorter than segment III, and continuous pronotal transverse furrow. This species is similar to *T. anjozorobeus* and *T. kathrynae*, but is distinguished by the black head, thorax, and legs; pale orange hemelytra; flat ventral head surface; synthlipsis about 1.5 times width of eye; labial segment II shorter than III; and intersegmental sutures carinulate between II and III and laterally between III–VI.

DESCRIPTION: Medium body size (length: 17.53 mm, holotype; 17.05–19.72 mm).

COLORATION: Black with antennal segment VI and basal half of segment VII, hemelytron, tarsi, and pretarsi orange and distal half of antennal segment VII pale orange. Abdomen red with anterior margin of sternite II, sternite VII (except small anterolateral spot), laterotergite VII (except anterior margin), and pygophore black. VESTITURE: As in generic description.

STRUCTURE: HEAD: Ventrally flat; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); synthlipsis about 1.5 times width of eye; interocular sulcus posterior to hind margin of eye; ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); postclypeus with shallow, broad depression to middle of interocular area (Fig. 2.13B); postocular moderately broad in dorsal view; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III longer than II. THORAX: Anterior pronotal lobe without distinct anterolateral projections; posterior pronotal lobe transversely striated; pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III; hemelytron almost reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between II and III and laterally between III–VI; pygophore process subtriangular in lateral view (Fig. 2.20H); BPE shorter than basal plate; area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21G).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Marojejy Nature Reserve, Madagascar.

DISTRIBUTION: Marojejy Nature Reserve in the Antsiranana province (Map 2.6).

HOLOTYPE: Male: **Madagascar: Antsiranana:** R.N.I. de Marojejy, 11.0 km NW Manantenina, 14.43667°S 49.74167°E, 1225 m, 25 Oct 1996–03 Nov 1996, Eric L. Quinter (00078359) (AMNH).

PARATYPES: **Madagascar: Antsiranana:** Sambava District, R.N. XII Marojejy, Ambatosoratra, 14.38735°S 49.75717°E, 1700m, Nov 1960, P. Soga, 3♂ (MNHN).



*Tanindrazanus nigripes*, new species

Figs. 2.3, 2.6, 2.9, 2.22E; Map 2.8

DIAGNOSIS: Males recognized among other *Tanindrazanus* species by the dark brown to black coloration with pale yellow markings, synthlipsis about 1.5 times the width of an eye, and eyes not reaching the dorsal and ventral head margins. This species is similar to *T. tenebricus*, but the larger width of the synthlipsis, eyes not reaching the dorsal and ventral head margins, and intersegmental sutures carinulate between sternites II–IV distinguishes *T. marginatus* from *T. tenebricus*.

DESCRIPTION: Medium body size (length: 12.63 mm, holotype; 11.78–12.63 mm).

COLORATION: Dark brown to black with mandibular plates dorsally, postocular dorsally except ocellar tubercle, and fore- and mid femora ventrally and laterally pale yellow. Labium light brown except segment II basally and distal two-thirds of antennal segment V and segments VI and VII white. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with shallow, broad medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about 1.5 times width of eye; interocular sulcus at (Fig. 2.13B) or posterior to hind margin of eye; postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-third of head length, not reaching dorsal and ventral head surfaces; pedicel about two-fifths longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); forefemur moderately incrassate; tarsomeres I and II combined subequal to III on fore- and mid legs, longer on hind legs; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal

laterotergite II not expanded; intersegmental sutures carinulate between sternites II–IV; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE as long as basal plate (Fig. 2.22E); area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the blackish coloration of the body.

DISTRIBUTION: Antsiranana and Toamasina provinces (Map 2.8).

DISCUSSION: Coloration is variable with the clypeus, postclypeus, coxae, intersternal sutures, median longitudinal depression on sternites, and medial sternites pale yellow in some specimens.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Sakalava Beach, dwarf littoral forest, 12.26277°S 49.3975°E, 10 m, 31 May 2001–07 Jun 2001, M. Irwin, R. Harin'Hala (00006712) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Forêt Ambato, 26.6 km 33° NE Ambanja, 13.46444°S 48.55166°E, 150 m, 08 Dec 2004, B.L. Fisher, 1♂ (00044939) (CAS). Montagne Français, 12.325°S 49.33333°E, 150 m, 15 Feb 2001–06 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006705) (SU), 1♂ (00006717) (USNM); 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00045437) (UCR). Parc National Montagne d'Ambre, 12.51444°S 49.18138°E, 960 m, 26 Jan 2001–29 Jan 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 3♂ (00006715, 00007165, 00007195) (CAS), 1♂ (00007231) (AMNH); 11 Feb 2001–12 Feb 2001, M. Irwin, R. Harin'Hala, 2♂ (00006727, 00045465) (CAS); 04 Mar 2001–19 Mar 2001, M. Irwin, R. Harin'Hala, 4♂ (00006704, 00006710, 00006726, 00045231) (CAS), 1♂ (000045328) (BMNH), 1♂ (000045436) (MNHN), 1♂ (00045500) (SU). Parc National Montagne d'Ambre, 12.51666°S 49.18333°E, 975 m, 25 Jan 2001–11 Feb 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 1♂ (00005368) (UCR); 25 Jan 2001–29 Jan 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 2♂

(00007047, 00007134) (CAS); 11 Feb 2001–04 Mar 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 1♂ (00006145) (BMNH), 1♂ (00006718) (MNHN). Parc National Montagne d'Ambre, 12.52027°S 49.17916°E, 1125 m, 19 Mar 2001–15 Apr 2001, M. Irwin, R. Harin'Hala, 2♂ (00006714, 00006716) (CAS), 1♂ (00006721) (BMNH), 1♂ (00006722) (AMNH); 21 Apr 2001–26 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006703) (UCR); 14 May 2001–30 May 2001, M. Irwin, R. Harin'Hala, 1♂ (00006711) (USNM); 30 May 2001–06 Jun 2001, M. Irwin, R. Harin'Hala, 1♂ (00006706) (CAS), 1♂ (00006713) (MNHN). Sakalava Beach, dwarf littoral forest, 12.26277°S 49.3975°E, 10 m, 07 Apr 2001–22 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006708) (SU); 13 May 2001–16 May 2001, M. Irwin, R. Harin'Hala, 1♂ (00006708) (USNM); 07 Jun 2001–25 Jun 2001, M. Irwin, R. Harin'Hala, 1♂ (00006709) (AMNH); 13 Aug 2001–20 Aug 2001, M. Irwin, R. Harin'Hala, 1♂ (00045396) (CAS); 20 Aug 2001–28 Aug 2001, M. Irwin, R. Harin'Hala, 1♂ (00007143) (UCR). **Toamasina:** Botanic Garden near entrance to Andasibe National Park, 18.92633°S 48.40783°E, 1025 m, 08 Oct 2001–16 Oct 2001, M. Irwin, R. Harin'Hala, 1♂ (00006725) (CAS). **Unknown:** 2♂ (00006707, 00045240) (CAS).

*Tanindrazanus notatus*, new species

Figs. 2.3, 2.6, 2.10, 2.21G; Map 2.6

DIAGNOSIS: Males are recognized among other species in this genus by the distinct dull orange-brown coloration with black markings on the hind legs. *Tanindrazanus notatus* is similar to *T. andohahela* and *T. bemaraha*, but the black apical half of the hind femur and tibia and the subequal lengths of labial segments II and III differentiate *T. notatus* from these species.

DESCRIPTION: Medium body size (length: 10.17 mm, holotype; 10.17–12.24).

COLORATION: Orange with apical half of scape, pedicel, antennal segments III–VI, apical half of hind femur, and hind tibia, except basally, dark brown. Apex of corium, membrane, and distal

wing veins brown. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe without distinct anterolateral projections; posterior pronotal lobe transversely striated; pronotal transverse furrow continuous; forefemur moderately incrassate; tarsomeres I and II combined subequal to III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE shorter than basal plate; area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21G).

ETYMOLOGY: The species epithet is a past participle in the nominative case and is named for the black markings on the hind legs.

DISTRIBUTION: Beroboka village from the Toliara province (Map 2.6).

DISCUSSION: The two specimens examined differ in size (10.17 and 12.24 mm) but are morphologically similar and were collected from the same locality.

HOLOTYPE: Male: **Madagascar: Toliara:** Beroboka village, 45 km NE Morondava, 19.9775°S 44.82483°E, 131 m, 05 Feb 2009–13 Feb 2009, M. Irwin, R. Harin'Hala (00045368) (CAS).

PARATYPE: **Madagascar: Toliara:** Beroboka village, 45 km NE Morondava, 19.9775°S 44.82483°E, 131 m, 19 Dec 2008–27 Dec 2008, M. Irwin, R. Harin'Hala, 1♂ (00044868) (CAS).

*Tanindrazanus simulans*, new species

Figs. 2.3, 2.6, 2.10

DIAGNOSIS: Males are recognized among other *Tanindrazanus* species by the nearly uniform dark brown to blackish coloration, synthlipsis about 1.25 times wider than the width of an eye, eyes reaching ventral and almost reaching the dorsal head margins, and a continuous pronotal transverse furrow. This species is similar to *T. tenebricus* and *T. nigripes*, but is distinguished by the width of synthlipsis, eyes reaching the ventral head margin but not the dorsal margin, and the pedicel about one-fifth longer than scape.

DESCRIPTION: Medium body size (length: 15.40 mm, holotype; 15.10–15.40 mm).

COLORATION: Blackish except postocular part of the head around ocellar tubercle and tarsi

pale brown. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with

shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G,

H); postclypeus with relatively deep, broad medial longitudinal depression to middle of

interocular area (Fig. 2.13B); synthlipsis about 1.25 times width of eye; interocular sulcus near

hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large,

separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye

about one-third of head length, reaching ventral but not dorsal head surface; pedicel about one-

fifth longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe with

very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated;

pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III

on forelegs, longer on mid and hind legs; hemelytra surpassing abdominal apex. ABDOMEN: Apex

rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternite

II–IV and laterally between IV–VI; pygophore process subtriangular in lateral view (Fig. 2.20H,

I); BPE longer than basal plate; area of endosomal struts–DPS fusion ovate.

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named after the similarity to *T. tenebricus*.

DISTRIBUTION: The two known specimens were labeled as originating from “Ivondro.” There are at least four localities in the Fianarantsoa, Toamasina, and Toliara provinces that include the name “Ivondro”; we were unable to determine the exact locality.

HOLOTYPE: Male: **Madagascar: Unknown:** Reg. S.E. Ivondro, 1939, A. Seyrig (MNHN).

PARATYPE: **Madagascar: Unknown:** Reg. S.E. Ivondro, Feb 1940, A. Seyrig, 1♂ (MNHN).

*Tanindrazanus tenebricus*, new species

Figs. 2.3, 2.6, 2.10, 2.18B; Map 2.8

DIAGNOSIS: Males are recognized among other *Tanindrazanus* species by the blackish coloration with pale markings on the head and pronotum, synthlipsis about the width of an eye, and large eyes reaching only dorsal head. This species is similar to *T. nigripes* and *T. simulans*, from which it differs by the smaller synthlipsis width and eyes reaching the dorsal but not the ventral head surface. *Tanindrazanus tenebricus* is further differentiated from *T. nigripes* by the intersegmental sutures carinulate between sternite II–IV and laterally between IV–VI.

DESCRIPTION: Medium body size (length: 16.37 mm, holotype; 15.40–17.02 mm).

COLORATION: Blackish with mandibular plates dorsally, postocular dorsally except ocellar tubercle, posterolateral margin of pronotum, and tarsi pale yellow. Labium light brown, except segment II basally, and antennal segment VI light brown and segment VII white. VESTITURE:

As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum (Fig. 2.13G, H); postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13H); postocular narrow in dorsal view

(Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about one-third of head length, reaching dorsal but not ventral head surface; pedicel about one-fourth longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); legs slender; tarsomeres I and II combined subequal to III on forelegs, longer on mid and hind legs; wings spread out in specimens. ABDOMEN (Fig. 2.18B): Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternite II–IV and laterally between IV–VI; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE longer than basal plate; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the dark coloration of the body.

DISTRIBUTION: Specimens are known from the southeastern region of the Fianarantsoa province (Map 2.8). Two specimens were labeled as originating from “Ivondro”. There are at least four localities in the Fianarantsoa, Toamasina, and Toliara provinces that include the name “Ivondro”; we were unable to determine the exact locality.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Manombo Special Reserve camp site, 32 km SSE of Farafangana, 23.02183°S 47.72°E, 36 m, 16 Jan 2005–20 Jan 2005, M. Irwin, R. Harin’Hala (00045483) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 10 Feb 2007–17 Feb 2007, M. Irwin, F. Parker, R. Harin’Hala, 1♂ (00006723) (CAS); 03 Mar 2007–10 Mar 2007, M. Irwin, F. Parker, R. Harin’Hala, 1♂ (00006720) (SU); 17 Mar 2007–24 Mar 2007, M. Irwin, F. Parker, R. Harin’Hala, 1♂ (00006724) (UCR). Manombo Special Reserve camp site, 32 km SSE of

Farafangana, 23.02183°S 47.72°E, 36 m, 10 Oct 2004–21 Oct 2004, M. Irwin, R. Harin'Hala, 2♂ (00007141, 00044862) (CAS); 23 Nov 2004–05 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00044995) (AMNH); 05 Jan 2005–16 Jan 2005, M. Irwin, R. Harin'Hala, 2♂ (00007046, 00045656) (CAS); 16 Jan 2005–20 Jan 2005, M. Irwin, R. Harin'Hala, 2♂ (00045488, 00045723) (BMNH); 13 Mar 2005–26 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00045301) (UCR); 10 Apr 2005–24 Apr 2005, M. Irwin, R. Harin'Hala, 2♂ (00044873, 00045022) (CAS), 1♂ (00045267) (AMNH); 22 May 2005–06 Jun 2005, M. Irwin, R. Harin'Hala, 7♂ (00007118, 00045270, 00045399, 00045403, 00045496, 00045606, 00045721) (CAS); 10 Aug 2005–25 Aug 2005, M. Irwin, R. Harin'Hala, 2♂ (00044828, 00045684) (MNHN); 11 Sep 2005–25 Sep 2005, M. Irwin, R. Harin'Hala, 1♂ (00045104) (CAS), 1♂ (00045196) (USNM); 25 Sep 2005–02 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00007101) (SU), 1♂ (00045477) (UCR). **Unknown:** Reg. S.E. Ivondro, Dec 1938, A. Seyrig, 1♂ (MNHN); 1939, A. Seyrig, 1♂ (MNHN).

*Tanindrazanus varicolor*, new species

Figs. 2.3, 2.6, 2.7, 2.10, 2.13H, 2.16A, 2.17C, 2.20I, 2.21H; Map 2.7

**DIAGNOSIS:** Males are recognized among other species in this genus by the distinct black and orange coloration, labial segments II and III subequal in length, and the anterior pronotal lobe with very small anterolateral protuberances. This species is similar to *T. irwini*, but differs by the smaller body size, anteromedial depression on the ventral head surface, and intersegmental sutures carinulate between sternite II–VI and laterally between VI and VII, as well as the diagnostic features mentioned.

**DESCRIPTION:** Medium body size (length: 15.97 mm, holotype; 15.89–16.12 mm).

**COLORATION:** Dark brown-black with bright orange head, scape, labium, pronotum, scutellum medially, stridulatory groove, meso- and metasternal longitudinal depressions, hemelytron except



posteromedial spot, femora except medially on mid leg, tibiae, tarsi, anterior half of laterotergites, sternite III medially, and sternites II–VII anterolaterally. Distal half of antennal segment VII white. VESTITURE: As in generic description. STRUCTURE: HEAD (Fig. 2.13H): Ventrally with shallow anteromedial depression; clypeal apex dorsally elevated relative to labrum; postclypeus with relatively deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); legs slender; tarsomeres I and II combined subequal to III on fore- and mid legs, longer on hind legs; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; dorsal laterotergite II not expanded; intersegmental sutures carinulate between sternite II–VI and laterally between VI and VII; pygophore process subtriangular in lateral view (Fig. 2.20I); BPE as long as basal plate; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the variable color forms of this species.

DISTRIBUTION: Toamasina and the northern region of the Antsiranana provinces (Map 2.7).

DISCUSSION: Some specimens are predominately orange in coloration while others have reduced to absent orange markings on the head, thorax, scutellum, femora, tibiae, and abdomen.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Parc National Montagne d'Ambre, 12.51444°S 49.18138°E, 960 m, 26 Jan 2001–29 Jan 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala (00006271) (CAS).

PARATYPES: **Madagascar: Antsiranana:** 7 km N of Joffreville, 12.33333°S 49.25°E, 360 m, 20 Mar 2001–07 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006287) (CAS); 27 Apr 2001–13 May 2001, M. Irwin, R. Harin'Hala, 1♂ (00006288) (UCR); 13 May 2001–16 May 2001, M. Irwin, R. Harin'Hala, 1♂ (00006484) (USNM). Montagne Française, 12.325°S 49.33333°E, 150 m, 30 Jan 2001–15 Feb 2001, M. Irwin, R. Harin'Hala, 1♂ (00006279) (SU); 15 Feb 2001–06 Mar 2001, M. Irwin, R. Harin'Hala, 2♂ (00006274, 00006281) (CAS); 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006259) (UCR). Parc National Montagne d'Ambre, 12.52027°S 49.17916°E, 1125 m, 11 Feb 2001–04 Mar 2001, M. Irwin, R. Harin'Hala, 2♂ (00006254, 00006264) (CAS); 19 Mar 2001–15 Apr 2001, M. Irwin, R. Harin'Hala, 3♂ (00006247, 00006256, 00006257) (CAS), 1♂ (00006282) (MNHN); 05 Apr 2001–21 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006136) (BMNH), 1♂ (00006267) (AMNH); 21 Apr 2001–26 Apr 2001, M. Irwin, R. Harin'Hala, 2♂ (00006482, 00006483) (UCR). Parc National Montagne d'Ambre, 12.61666°S 49.15°E, 1116 m, Dec 1968, J. Vadon and A. Peyrieras, 1♂ (MNHN). Parc National Montagne d'Ambre, 12.51666°S 49.18333°E, 975 m, 25 Jan 2001–11 Feb 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 2♂ (00006255, 00006486) (CAS); 04 Mar 2001–19 Mar 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 1♂ (00006266) (UCR). Parc National Montagne d'Ambre, 12.51444°S 49.18138°E, 960 m, 23 Jan 2001, M.E. Irwin, 2♂ (00006116, 00006258) (USNM); 26 Jan 2001–29 Jan 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 2♂ (00006253, 00006272) (CAS), 1♂ (00007065) (SU); 11 Feb 2001–12 Feb 2001, M. Irwin, R. Harin'Hala, 2♂ (00006277, 00006278) (UCR); 04 Mar 2001–19 Mar 2001, M. Irwin, R. Harin'Hala, 3♂ (00006252, 00006275, 00006276) (CAS), 1♂ (00006292) (AMNH); 19 Mar

2001–05 Apr 2001, M. Irwin, R. Harin'Hala, 3♂ (00006289, 00006291, 00006702) (CAS). Sakalava Beach, dwarf littoral forest, 12.26277°S 49.3975°E, 10 m, 15 Feb 2001–06 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006283) (CAS); 06 Mar 2001–20 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006260) (CAS); 20 Mar 2001–07 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006273) (BMNH); 07 Apr 2001–22 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006265) (MNHN); 13 May 2001–16 May 2001, M. Irwin, R. Harin'Hala, 1♂ (00006498) (AMNH); 31 May 2001–07 Jun 2001, M. Irwin, R. Harin'Hala, 1♂ (00006251) (SU); 13 Aug 2001–20 Aug 2001, M. Irwin, R. Harin'Hala, 1♂ (00045641) (USNM); 20 Aug 2001–28 Aug 2001, M. Irwin, R. Harin'Hala, 1♂ (00006485) (UCR). **Toamasina:** 7 km SE of Andasibe National Park headquarters, 18.96266°S 48.45266°E, 1050 m, 23 Mar 2001–07 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006269) (CAS); 09 Apr 2001–23 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00006115) (UCR). Botanic Garden near entrance to Andasibe National Park, 18.92633°S 48.40783°E, 1025 m, 31 Jul 2001–15 Aug 2001, M. Irwin, R. Harin'Hala, 1♂ (00006250) (UCR); 08 Oct 2001–16 Oct 2001, M. Irwin, R. Harin'Hala, 1♂ (00006246) (BMNH), 1♂ (00006248) (MNHN), 5♂ (00006249, 00006280, 00006284–00006286) (CAS), 1♂ (00006290) (SU); 07 Nov 2001–16 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00005360) (CAS), 1♂ (00006268) (AMNH). **Unknown:** 1♂ (00006270) (CAS).

*Tanindrazanus vohiparara*, new species

Figs. 2.3, 2.7, 2.10; Map 2.8

DIAGNOSIS: Males are recognized among other species *Tanindrazanus* by the blackish coloration with pale yellow markings, large eyes reaching dorsal and ventral head margins, smooth posterior pronotal lobe, a very slightly expanded dorsal laterotergite II, and the intersegmental sutures carinulate between sternites II–V. This species is similar to *T. nigripes*,

particularly a variant of this species, but is distinguished by the eye about half of head length, smooth posterior pronotal lobe, and expanded dorsal laterotergite II.

DESCRIPTION: Medium body size (length: 11.20 mm, holotype). COLORATION: Black with pale yellow scape basally, antennal segment VII, labial segment III and VI, thoracic sterna except medial longitudinal depression on meso- and metasterna, coxae, trochanters, forefemur ventrally and basally, mid and hind femora except medially, foretibia ventrally, anterior margins of laterotergites, and abdominal sternites except large lateral spots on II–VI and apex of VII. VESTITURE: As in generic description. STRUCTURE: HEAD: Ventrally with shallow anteromedial depression; clypeal apex slightly elevated relative to labrum (Fig. 2.13G, H); postclypeus with deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, separated by less than diameter of ocellus, located on distinct median tubercle (Fig. 2.13H); eye about half of head length, reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape; labial segment III subequal to II. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow continuous; legs slender; tarsomeres I and II combined subequal to III; wings spread out in specimen. ABDOMEN: Apex rounded; dorsal laterotergite II very slightly expanded; intersegmental sutures carinulate between sternites II–V; pygophore process subtriangular in lateral view (Fig. 2.20H, I); BPE shorter than basal plate; endosomal struts ventrally obscured by semiextended endosoma; area of endosomal struts–DPS fusion ovate (Fig. 2.21H).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality that is part of Ranomafana National Park, Madagascar.

DISTRIBUTION: Ranomafana National Park in the Fianarantsoa province (Map 2.8).

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1100 m, 04 Mar 2002–12 Mar 2002, M. Irwin, R. Harin'Hala (00006139) (CAS).

PARATYPE: **Madagascar: Fianarantsoa:** Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 23 Oct 2005–30 Oct 2005, M. Irwin, R. Harin'Hala, 1♂ (00045227) (CAS).

*Toliarus*, new genus

Figs. 2.3, 2.7, 2.10, 2.13I, 2.19G, 2.20J, 2.21I, 2.22F; Map 2.9

TYPE SPECIES: *Toliarus trichrous*, new species

DIAGNOSIS: Males recognized by a combination of the following characters: the small body size, ocelli on a shallow median tubercle, large antennal shield that does not conceal antennal insertion in lateral view, 6-segmented antenna, and punctate posterior pronotal lobe. This genus is very similar to *Tanindrazanus*, from which it differs by the smaller body size, 6-segmented antenna, and punctate posterior pronotal lobe.

DESCRIPTION: **MALE:** Macropterous, small body size. COLORATION: Red and black or orange and black. VESTITURE: Dense, semierect to erect, long pale to dark brown setae on head, thorax, legs, corium, and abdomen; setae on tibiae stouter and denser near apex.

STRUCTURE: HEAD (Fig. 2.13I): Subtriangular in lateral view, longer than wide in dorsal view, shorter than pronotum; anteocular region longer than postocular; clypeal apex not elevated relative to labrum; maxillary plate not reaching dorsal surface of clypeus (except nearly reaching in a some specimens of *T. trichrous*); postclypeus depressed (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli present, located on shallow median tubercle; distinct constriction between postocular and neck; gula moderately swollen ventrolaterally, not distinctly produced

beyond ventral head margin; antennal shield not concealing antennal insertion in lateral view; antenna inserted dorsally on head; scape surpassing clypeal apex, pedicel slightly curved; flagellum subdivided into BFLA and DFLA; BFLA divided into two pseudosegments; DFLA divided into two pseudosegments; antenna thus appearing 6-segmented; labium slender; labial segment III subequal to II, ventrally straight. THORAX: Pronotum wider than long, anterior margin distinctly concave, collar distinct (Fig. 2.15B); anterior pronotal lobe shorter than posterior lobe (Fig. 2.15B), more than half as wide as posterior lobe, relatively smooth; posterior pronotal lobe weakly punctate; pronotal longitudinal furrow reaching anterior but not posterior margin, foveate posteriorly (Fig. 2.15B); pronotal transverse furrow distinct (Fig. 2.15B); lateral depressions on pronotum distinct (Fig. 2.15B), transversely striated; scutellum with two broadly separated apical processes, disc medially depressed; meso- and metasterna shallowly separated by transverse suture (Fig. 2.15F); mesosternum with medial and paramedial longitudinal depressions; metasternum medially longitudinally depressed; MGE with deeply depressed meshlike cuticle that extends dorsally in lateral view (Fig. 2.16A); fossula spongiosa on fore- and mid tibiae; corium restricted to areas adjacent to basal wing veins, with pterostigmalike appearance on anterodistal margin (Fig. 2.17A, C); distal part of R present but not forming cell with M (Fig. 2.17B, C); proximal parts of M and Cu veins separate (Fig. 2.17A, C); distal part of M extending beyond apical junction of M+Cu (Fig. 2.17A, C); base of M+Cu cell shorter than or as wide as Cu+1A cell (Fig. 2.17B, C). ABDOMEN: Dorsal laterotergite II not expanded; sternites shallowly depressed medially (Fig. 2.18C), intersegmental sutures carinulate; spiracles ovoid; pygophore (Figs. 2.19G, 2.20J) process subtriangular in lateral view, directed dorsoposteriad, not surpassing posterior margin of pygophore; DPS apex rounded (Fig. 2.21I); endosomal struts reaching posterior margin of DPS, with anterior and posterior ventral processes (Fig. 2.22G); endosoma medially sclerotized (Fig. 2.21I).

ETYMOLOGY: This genus is named after the locality province of the type species, Toliara, Madagascar. The gender is masculine.

DISTRIBUTION: Species are known from the southern region of the Toliara province and from the Antsiranana province and in habitats between 35–180 m elevation. Macrohabitats are described as gallery forests, spiny forests, dry deciduous forests, and transitional forests.

DISCUSSION: Female specimens are unknown for species currently in this genus. The size of the maxillary plates is polymorphic in *T. trichrous*; in approximately 40% of the specimens, the maxillary plates nearly reach the dorsal clypeal surface when viewed laterally (Fig. 2.13H).

*Identification key to the males of species of Toliarus*

1. Orange with blackish markings, ocelli separated by a distance about the diameter of an ocellus, medium body size (>10 mm) ..... *karinae*, new species  
- Tricolored (red and black with dark brown fore wing membrane), ocelli separated by more than diameter of an ocellus; small body size (<10 mm) ..... *trichrous*, new species

*Toliarus karinae*, new species

Figs. 2.3, 2.7; Map 2.9

DIAGNOSIS: Males are similar to *T. trichrous*, but is easily distinguished by the larger body size, distinct orange body with blackish markings on the hemelytra and tibiae, medium sized ocelli, and weak punctuation on the posterior pronotal lobe.

DESCRIPTION: Medium body size (length: 12.10 mm, holotype). COLORATION: Orange with antenna except scape ventrally, clavus, forewing membrane, and apices of tibiae blackish.

Antennifer apically white. VESTITURE: As in generic description. STRUCTURE: HEAD:

Ventrally with shallow anteromedial depression; postclypeus with relatively deep, narrow medial

longitudinal depression to middle of interocular area; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli medium sized, separated by a distance about the diameter of an ocellus; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape. THORAX: Anterior pronotal lobe without distinct anterolateral projections; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); forefemur incrassate; tarsomeres I and II subequal to III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process subquadrate in caudal view, with truncate apex; BPE as long as basal plate (Fig. 2.22F); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21I).

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after the second author's wife, Karina Wieczorek.

DISTRIBUTION: The only known specimen was collected from the Ankarana Reserve in the Antsiranana province (Map 2.9).

HOLOTYPE: Male: **Madagascar: Antsiranana:** Ankarana Reserve, 12.91688°S 49.14355°E, 155m, no date provided, J. Vadon and A. Peyrieras (MNHN).

*Toliarus trichrous*, new species

Figs. 2.3, 2.7, 2.10, 2.13I, 2.19G, 2.20J, 2.21I, 2.22F; Map 2.9

DIAGNOSIS: Males are recognized from *T. karinae* by the smaller body size, red and black coloration, and more punctate posterior pronotal lobe. Males are most similar to *Tanindrazanus anjozorobeus* and *Tanindrazanus kathrynae*, but the smaller body size, red anteocular region and abdominal sternites (except black pygophore), black postocular region, 6-segmented antenna, and punctate posterior pronotal lobe distinguishes *T. trichrous* from these species.



DESCRIPTION: Small body size (length: 7.22 mm, holotype; 7.22–8.97 mm).

COLORATION: Red with black head ventrolaterally below eye, postocular except ventrolateral spot, neck except dorsomedially, anterior pronotal and propleural lobes, prosternum, medial pronotal longitudinal furrow on posterior pronotal lobe, meso- and metapleura, scutellum, apical half of hind femur, hind tibia medially, abdominal sternite VII apically, and pygophore.

Hemelytron dark brown, except red corium. VESTITURE: As in generic description.

STRUCTURE: HEAD (Fig. 2.13I): Ventrally with shallow anteromedial depression; postclypeus with relatively deep, narrow medial longitudinal depression to middle of interocular area; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli small, separated by more than diameter of ocellus; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; pedicel about one-fourth longer than scape. THORAX: Anterior pronotal lobe without distinct anterolateral projections; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); forefemur incrassate; tarsomeres I and II subequal to III; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process subtriangular in lateral view (Fig. 2.20J), apex rounded; BPE as long as basal plate (Fig. 2.22F); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21I).

ETYMOLOGY: The species epithet is an adjective and is named for the body coloration.

DISTRIBUTION: Specimens were collected from the southern region of Toliara (Map 2.9).

DISCUSSION: Coloration varies with the posterior pronotal lobe and posterior propleural lobe either uniformly black, black with red lateral margins, or uniformly red. The hind femur may be more extensively black, and the black medial spot on abdominal sternite VII varies in size.

HOLOTYPE: Male: **Madagascar: Toliara:** Beza Mahafaly Reserve, Parcelle II near Bellevue, 23.68983°S 44.5755°E, 180 m, 21 Nov 2001–28 Nov 2001, R. Harin'Hala (00006163) (CAS).

PARATYPES: **Madagascar: Toliara:** Andohahela Nat'yl Park, Tsimelahy, Parcelle II, 24.93683°S 46.62666°E, 180 m, 09 Dec 2002–16 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006494) (CAS); 16 Dec 2002–17 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006185, 00006186) (CAS); 15 Feb 2003–26 Feb 2003, M. Irwin, F. Parker, R. Harin'Hala, 5♂ (00006513–00006517) (CAS); 18 Mar 2003–28 Mar 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006184, 00006519) (USNM); 28 Mar 2003–08 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006155, 00006497) (BMNH); 19 Apr 2003–26 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006168, 00006169) (MNHN); 29 Jun 2003–10 Jul 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006183, 00006518) (AMNH); 17 Aug 2003–24 Aug 2003, M. Irwin, F. Parker, R. Harin'Hala, 5♂ (00006176–00006180) (CAS); 11 Oct 2003–19 Oct 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006162) (SU); 15 Jan 2004–28 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006493) (UCR). Berenty Special Reserve, 8 km NW Amboasary, 25.00666°S 46.30333°E, 85 m, 25 Oct 2002–26 Oct 2002, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006153, 00006156) (CAS); 09 Nov 2002–16 Nov 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006167) (UCR); 30 Nov 2002–07 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006171) (CAS); 14 Dec 2002–16 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006181, 00006182, 00006501) (AMNH); 26 Jan 2003–05 Feb 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006507) (SU); 24 Apr 2003–03 May 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006506) (CAS); 25 May 2003–04 Jun 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006202) (USNM); 22 Nov 2003–30 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006490) (BMNH); 30 Nov 2003–07 Dec 2003, M. Irwin, F. Parker, R.

Harin'Hala, 1♂ (00006520) (BMNH); 07 Dec 2003–14 Dec 2003, M. Irwin, F. Parker, R.  
 Harin'Hala, 1♂ (00006492) (UCR); 21 Dec 2003–02 Jan 2004, M. Irwin, F. Parker, R.  
 Harin'Hala, 2♂ (00006504, 00006522) (CAS); 13 Jan 2004–23 Jan 2004, M. Irwin, F. Parker, R.  
 Harin'Hala, 2♂ (00006505, 00006510) (MNHN); 04 Feb 2004–15 Feb 2004, M. Irwin, F. Parker,  
 R. Harin'Hala, 1♂ (00006500) (CAS); 02 Mar 2004–11 Mar 2004, M. Irwin, F. Parker, R.  
 Harin'Hala, 1♂ (00006509) (CAS); 05 Apr 2004–15 Apr 2004, M. Irwin, F. Parker, R.  
 Harin'Hala, 1♂ (00006503) (AMNH); 06 Jun 2004–20 Jun 2004, M. Irwin, F. Parker, R.  
 Harin'Hala, 1♂ (00006489) (CAS); 11 Jul 2004–25 Jul 2004, M. Irwin, F. Parker, R. Harin'Hala,  
 3♂ (00006511, 00006512, 00006521) (CAS); 07 Aug 2004–22 Aug 2004, M. Irwin, F. Parker, R.  
 Harin'Hala, 2♂ (00006487, 00006488) (USNM); 22 Aug 2004–01 Sep 2004, M. Irwin, F. Parker,  
 R. Harin'Hala, 2♂ (00006495, 00006496) (SU); 01 Sep 2004–08 Sep 2004, M. Irwin, F. Parker,  
 R. Harin'Hala, 1♂ (00006508) (UCR). Berenty Special Reserve, 8 km NW Amboasary, 25.021°S  
 46.3055°E, 35 m, 10 Jun 2003–19 Jun 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006502)  
 (CAS); 08 Nov 2003–15 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00005364) (CAS);  
 21 Dec 2003–01 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006175, 00006189,  
 00006190) (MNHN); 24 Mar 2004–04 Apr 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
 (00006491) (SU); 23 Jun 2004–04 Jul 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006157)  
 (BMNH), 1♂ (00006158) (CAS); 04 Jul 2004–18 Jul 2004, M. Irwin, F. Parker, R. Harin'Hala,  
 1♂ (00006170) (UCR). Beza Mahafaly Reserve, Parcelle II near Bellevue, 23.68983°S  
 44.5755°E, 180 m, 10 Nov 2001–21 Nov 2001, M. Irwin, R. Harin'Hala, 1♂ (00006208) (CAS);  
 04 Dec 2001–11 Dec 2001, M. Irwin, R. Harin'Hala, 1♂ (00006204) (UCR). Beza Mahafaly  
 Reserve, Parcelle I near research station, 23.6865°S 44.591°E, 165 m, 21 Nov 2001–28 Nov  
 2001, R. Harin'Hala, 3♂ (00006154, 00006161, 00006188) (CAS); 04 Dec 2001–11 Dec 2001,  
 R. Harin'Hala, 1♂ (00006213) (CAS); 18 Dec 2001–25 Dec 2001, R. Harin'Hala, 2♂ (00006193,

00006203) (SU); 22 Feb 2002–01 Mar 2002, R. Harin'Hala, 1♂ (00045647) (CAS); 29 Apr 2002–19 May 2002, R. Harin'Hala, 1♂ (00006205) (BMNH); 28 Jul 2002–09 Aug 2002, R. Harin'Hala, 1♂ (00006160) (CAS); 09 Sep 2002–20 Sep 2002, R. Harin'Hala, 1♂ (00006209) (CAS); 28 Oct 2002–10 Nov 2002, M.E. Irwin, F.D. Parker, R. Harin'Hala, 1♂ (00006191) (CAS); 10 Nov 2002–22 Nov 2002, R. Harin'Hala, 2♂ (00006199, 00006215) (USNM); 02 Dec 2002–12 Dec 2002, R. Harin'Hala, 3♂ (00006150, 00006151, 00006212) (AMNH); 12 Dec 2002–17 Dec 2002, R. Harin'Hala, 1♂ (00006187) (CAS); 17 Dec 2002–20 Dec 2002, R. Harin'Hala, 1♂ (00006214) (CAS); 20 Dec 2002–24 Dec 2002, R. Harin'Hala, 1♂ (00006164) (UCR); 24 Dec 2002–02 Jan 2003, R. Harin'Hala, 2♂ (00006210, 00006211) (CAS); 09 Jan 2003–23 Jan 2003, R. Harin'Hala, 1♂ (00006192) (CAS); 04 Feb 2003–16 Feb 2003, R. Harin'Hala, 1♂ (00006194) (BMNH); 02 Mar 2003–13 Mar 2003, R. Harin'Hala, 2♂ (00006159, 00006165) (CAS); 13 Mar 2003–23 Mar 2003, R. Harin'Hala, 2♂ (00006197, 00006198) (BMNH); 23 Mar 2003–02 Apr 2003, R. Harin'Hala, 2♂ (00006086, 00006152) (CAS); 02 Apr 2003–10 Apr 2003, R. Harin'Hala, 3♂ (00006166) (SU), 3♂ (00006172–00006174) (CAS); 10 Apr 2003–21 Apr 2003, R. Harin'Hala, 2♂ (00006206, 00006207) (CAS); 21 Apr 2003–29 Apr 2003, R. Harin'Hala, 1♂ (00006196) (UCR). Cap Ste Marie Special Reserve, 74 km S of Tsihombe, 25.58766°S 45.163°E, 37 m, 18 May 2003–26 May 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006200) (CAS), 1♂ (00006201) (UCR). Parc National d'Andohahela, Ihazofotsy Parcelle III, 24.83083°S 46.53616°E, 80 m, 15 Jan 2003–26 Jan 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00007162) (BMNH), 3♂ (00045213, 00045297, 00045312) (MNHN); 24 Aug 2003–03 Sep 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045633) (CAS); 12 Nov 2003–23 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045242) (UCR); 17 Dec 2003–19 Dec 2003, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00007208, 00044928, 00045630) (USNM); 14 Jan 2004–28 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00007185) (AMNH), 6♂

(00044953, 00045131, 00045203, 00045298, 00045597, 00045634) (CAS), 2♂ (00045700, 00045726) (SU). **Unknown:** 1♂ (00006195) (CAS).

*Toxopus* Bergroth, 1905

Figs. 2.3, 2.4, 2.7, 2.10, 2.11, 2.12, 2.13B, 2.13J, 2.13K, 2.14, 2.15B, 2.19H, 2.19I, 2.20K, 2.20L, 2.21J–L, 2.22G, 2.22H, 2.23E; Maps 2.10–2.13

*Toxopus* Bergroth, 1905: 374.

TYPE SPECIES: *Toxopus politus* Bergroth, 1905

REVISED DIAGNOSIS: Male recognized by the shiny dark brown to blackish coloration with pale yellow markings, meso- and metasterna not completely separated by distinct transverse suture, dorsal laterotergite II expanded laterally, and laterotergites transversely bicolored. Females recognized by the apterous condition, shiny dark brown to blackish coloration sometimes with pale yellow markings, head about as wide as anterior margin of pronotum, 7-segmented antenna with short vestiture, scape as long as distance between anterior margin of eye and apex of head, eye about half head height, anterior pronotal lobe much longer than posterior lobe, fore- and mid trochanters with small papillae, and abdominal tergites I–III declinate toward posterior with remaining tergites oriented horizontally. This genus is most similar to *Rochonia* Distant, 1913, from which it differs by the relatively smooth head and pronotum, hemelytron reaching or surpassing abdominal apex, and dorsal laterotergite II expanded.

REDESCRIPTION: **MALE:** Macropterous, medium body size. **COLORATION:** Shiny dark brown to black with pale yellow markings. **VESTITURE:** Sparse to dense, semierect to erect, long pale to dark brown setae on head, thorax, legs, corium, and abdomen; setae on tibiae stouter and denser near apex. **STRUCTURE:** HEAD (Fig. 2.13B, J, K): Ovoid or subquadrate; shorter than pronotum; clypeal apex not elevated relative to labrum; maxillary plate not reaching dorsal

surface of clypeus; postclypeus depressed; distinct constriction between postocular and neck; gula moderately swollen ventrolaterally, not distinctly produced beyond ventral head margin; ocelli present; antennal shield not concealing antennal insertion in lateral view; scape surpassing clypeal apex; flagellum subdivided into BFLA and DFLA; BFLA divided into two pseudosegments; DFLA divided into three pseudosegments; antenna thus appearing 7-segmented; labial segment III subequal to II. THORAX (Fig. 2.15B): Pronotum wider than long, anterior margin distinctly concave, collar distinct; anterior pronotal lobe shorter than posterior lobe, more than half as wide as posterior lobe, relatively smooth; pronotal longitudinal furrow reaching anterior but not posterior margin of pronotum, foveate posteriorly; pronotal transverse furrow distinct; lateral depressions on posterior pronotal lobe distinct; scutellum with two moderately separated apical processes, disc medially depressed; meso- and metasterna not completely separated by distinct transverse suture (Fig. 2.15E); mesosternum with medial and paramedial longitudinal depressions; metasternum medially longitudinally depressed; MGE with deeply depressed meshlike cuticle that extends dorsally in lateral view (Fig. 2.16A); forefemur slightly to strongly incrassate; fossula spongiosa on fore- and mid tibiae; tarsomeres I and II combined subequal to III; corium restricted to areas adjacent to basal wing veins, with pterostigmalike appearance on anterodistal margin (Fig. 2.17A, C); distal part of R present but not forming cell with M (Fig. 2.17B, C); proximal parts of M and Cu separate (Fig. 2.17A, C); distal part of M extending beyond apical junction of M+Cu (Fig. 2.17A, C); base of M+Cu cell shorter than or as wide as Cu+1A cell (Fig. 2.17B, C). ABDOMEN: Dorsal laterotergite II expanded; sternites shallowly depressed medially (Fig. 2.18C), intersegmental sutures carinulate; spiracles ovoid; pygophore process subtriangular in lateral view (Fig. 2.20K, L), directed dorsoposteriad, apex rounded; DPS apex rounded (Fig. 2.21J–L). **FEMALE:** Differs from males in the following characteristics: apterous; reduced yellow markings; head larger, about as wide as anterior pronotal margin,

shorter or longer than pronotum; postclypeus not depressed (Fig. 2.13A); antennal shield not expanded; antennal vestiture much shorter; scape as long as distance between anterior eye margin and apex of head; ocelli absent; eyes small, about half head height; labial segments II shorter than or subequal to segment III; collar barely distinct; anterior pronotal lobe longer than and as wide as posterior lobe (Fig. 2.15A), without distinct anterolateral projections; pronotal longitudinal furrow reduced to deep medial depression near posterior margin of anterior pronotal lobe (Fig. 2.15A); lateral depressions on posterior pronotal lobe obsolete; scutellum weakly developed; meso- and metasterna platelike with shallow, wide medial longitudinal depression; MGE reduced, shallow, not extending dorsally in lateral view (Fig. 2.16B, C); fore- and mid trochanters with patches of small papillae (Fig. 2.16D, E); hind femur slightly curved in dorsal view; fossula spongiosa larger; dorsal laterotergite II not expanded; abdominal tergites I–III declinate toward posterior, intersegmental sutures carinulate, remaining tergites oriented horizontally, intersegments not carinulate; sternites flat; external genitalia short, platelike (Fig. 2.23E).

DISTRIBUTION: Species are known from all provinces and in habitats between 7–1,360 m elevation. Macrohabitats include tropical dry forests, dwarf littoral forests, low-altitude dense humid forests, low-altitude littoral rainforests, high-altitude rainforests, spiny forests, gallery forests, sclerophyl forests, montane rainforest, dry deciduous forests, and transitional forests.

DISCUSSION: Bergroth (1905) described this genus from a single female specimen. He emphasized the overall head structure and strongly curved hind tibia as diagnostic for this genus. However, head shape and structure is variable among species in some ectrichodiine genera (e.g., *Microstemmatoides* Putshkov, 1985, *Daraxa* Stål, 1859, *Gibbosella*) and the leg armature can vary among and within species. As examples, the presence and location of patches of papillae on the femora and trochanters may differ among species of *Rhiginia* Stål, 1859, males and females of the same species in *Racelda* Signoret, 1863, and apterous and macropterous forms within

species of *Glymmatophora*. Similarly, femoral and tibial spines may vary in presence, size, or number in species of *Glymmatophora*. Finally, the curvature of the hind tibia can vary between species in a genus (straight or slightly curved in *Pothea* Amyot and Serville, 1843, *Katanga* Schouteden, 1903, *Centraspis* Schaum, 1862) and within species (straight or slightly curved in females of *Ectrichodia crux* [Thunberg] 1783). Curvature and robustness of the hind tibia in females of *Toxopus politus* Bergroth, 1905, appears to be fairly extreme, and we found this character variable in two female specimens that we assign to *Toxopus*: one specimen has a more robust and strongly curved hind tibia than the other. We are unable to definitively assign either specimen to *T. politus* based on the original description; the holotype of *T. politus* is missing and could not be examined for confirmation (see discussion for *T. politus*). Comparison of these two specimens with a female with cylindrical head and slender, slightly curved hind tibia revealed many morphological similarities: apterous, shiny blackish coloration, head about as wide as anterior margin of pronotum, 7-segmented antenna with short vestiture, scape as long as distance between anterior eye margin and apex of head, eyes small, and abdominal tergites I–III declinate toward posterior, among several other features. Given the abundance of morphological similarities between these females and known variability in head shape and leg structure in other Ectrichodiinae genera, the genus is here redescribed to accommodate females with various head shapes and slender, less curved, and unarmed hind tibia. Based on morphology, geographic proximity, and molecules, we also associated the female with the cylindrical head and slender hind legs with males of a new species, *T. griswoldi*, enabling us to describe *Toxopus* males for the first time. Attempts to associate the other two females with conspecific males based on morphology, geography, and molecules failed. These two females remain undescribed for the time being since, based on head shape, presence of a ventral medial depression on the head, and geographic proximity, they could represent females of any of the following species: *T.*



*ampitavananima*, *T. basalis*, *T. brucei*, *T. namoroka*, *T. parkeri*, *T. signoretii*, *T. simulans*, *T. steineri*, *T. tibialis*, and *T. vazimba*.

The monotypic genus *Rochonia* was described by Distant from one macropterous male specimen from the Seychelles in 1913. The holotype of *R. galeatus* Distant, 1913, which is fairly damaged and has the ventral surface not visible, was examined at the BMNH. It is clear that *Rochonia* shares many morphological features with *Toxopus*, while few characters distinguish the two genera (see differential diagnosis for *Toxopus*). Based on these morphological similarities, *Toxopus* may in fact belong to *Rochonia*. We refrain from synonymizing *Toxopus* with *Rochonia* until additional specimens of *Rochonia* become available that will allow examination of ventral characters, antennal segmentation, and intraspecific variation. Future phylogenetic analyses including *Rochonia* and *Toxopus* may also provide support for synonymy.

*Identification key to the males of species of Toxopus*

1. Head cylindrical (Fig. 2.13J) .....	2
- Head subquadrate (Fig. 2.13A) or ovoid (Fig. 2.13F).....	9
2. Head ventrally flat.....	3
- Head with medial depression either restricted to anteocular region or extending to postocular .....	4
3. Pedicel not distinctly curved; sternal intersegmental sutures carinulate between II–IV and laterally between IV–VI.....	<i>insignis</i> , new species
- Pedicel slightly curved; sternal intersegmental sutures carinulate between II and III and laterally between III–VI.....	<i>fisheri</i> , new species

4. Head with ventral anteromedial depression; postocular broad in dorsal view (Fig. 2.13B); sublateral antennal insertion; pronotal transverse suture incomplete, divided by paramedian ridges (Fig. 2.15B).....	5
- Head with ventral medial depression extending to postocular; postocular narrow in dorsal view (Fig. 2.13A); dorsal antennal insertion; pronotal transverse suture complete, not divided by paramedian ridges .....	<i>melobrunneus</i> , new species
5. Posterior pronotal lobe smooth; sternal intersegmental sutures carinulate between II–IV and laterally between IV–VI.....	<i>toliara</i> , new species
- Posterior pronotal lobe striated; sternal intersegmental suture carination pattern not as previously mentioned.....	6
6. Synthlipsis width 1.5 times width of eye; sternal intersegmental sutures carinulate between II and III and laterally between III and IV.....	<i>antsiranana</i> , new species
- Synthlipsis width two times width of eye; sternal intersegmental suture carination pattern not as previously mentioned .....	7
7. Postclypeal depression extending to interocular sulcus; sternal intersegmental sutures carinulate between II and III and laterally between III–V .....	<i>ambohitantely</i> , new species
- Postclypeal depression extending to middle of interocular area (Fig. 2.13B); sternal intersegmental suture carination pattern not as previously mentioned.....	8
8. Postclypeal depression relatively broad; ventral margin of labial segment III convex (Fig. 2.13J, D, E, F); sternal intersegmental sutures carinulate between II and III and laterally between III–VII.....	<i>italaviana</i> , new species
- Postclypeal depression narrow; ventral margin of labial segment III straight (Fig. 2.13G–I, K); sternal intersegmental sutures carinulate between II–IV and laterally between IV–VII.....	<i>griswoldi</i> , new species

9. Antecular region as long as postocular (Fig. 2.13J).....	10
- Antecular region longer than postocular (Fig. 2.13K).....	11
10. Postclypeal depression extending to interocular sulcus; ventral margin of labial segment III straight (Fig. 2.13G–I, K); posterior pronotal lobe striated .....	<i>pallidus</i> , new species
- Postclypeal depression extending to middle of interocular area; ventral margin of labial segment III convex (Fig. 2.13C–F, J); posterior pronotal lobe smooth .....	<i>farafangana</i> , new species
11. Head about as long as wide in dorsal view .....	12
- Head longer than wide in dorsal view .....	16
12. Synthlipsis about width of eye; eye reaching dorsal head surface in lateral view; sternal intersegmental sutures carinulate between II and III and laterally between III–V .....	<i>toamasina</i> , new species
- Synthlipsis at least 1.5 times width of eye; eye not reaching dorsal head surface in lateral view; carination patterns on the sternal intersegmental sutures not as previously mentioned .....	13
13. Synthlipsis about 1.5 times width of eye .....	14
- Synthlipsis about two times width of eye .....	15
14. Head ventrally with anteromedial depression; eye not distinctly reaching ventral head surface in lateral view; sternal intersegmental sutures carinulate between II–VI .....	<i>tibialis</i> , new species
- Head with medial depression extending along entire ventral head surface; eye reaching or nearly reaching ventral head surface in lateral view; sternal intersegmental sutures carinulate between II–IV and laterally between IV–VI .....	<i>basalis</i> , new species
15. Posterior pronotal lobe weakly transversely striated; sternal intersegmental sutures carinulate between II–VI .....	<i>parkeri</i> , new species
- Posterior pronotal lobe smooth; sternal intersegmental sutures carinulate between II–IV and laterally between IV–VI.....	<i>brucei</i> , new species

16. Only sternal intersegmental suture between II and III carinulate .....	17
- Multiple sternal intersegmental sutures carinulate .....	18
17. Synthlipsis about two times width of eye; postocular broad in dorsal view (Fig. 2.13B); posterior pronotal lobe smooth .....	<i>miandritsara</i> , new species
- Synthlipsis about 1.5 times width of eye; postocular narrow in dorsal view (Fig. 2.13A); posterior pronotal lobe weakly transversely striated .....	<i>simulans</i> , new species
18. Synthlipsis at least 1.5 times width of eye; eye not distinctly reaching dorsal head surface... 19	
- Synthlipsis about width of eye; eye reaching or nearly reaching dorsal head surface .....	<i>signoretii</i> (Reuter), new combination
19. Synthlipsis about two times width of eye; anterior pronotal lobe with distinct, small anterolateral protuberances (Fig. 2.15B); posterior pronotal lobe smooth .....	20
- Synthlipsis about 1.5 times width of eye; anterior pronotal lobe without distinct anterolateral protuberances; posterior pronotal lobe weakly transversely striated .....	<i>ampitavananima</i> , new species
20. Ventral surface of labial segment III straight (Fig. 2.13G–I, K); sternal intersegmental sutures carinulate between II–VI and laterally between VI and VII.....	21
- Ventral surface of labial segment III convex (Fig. 2.13C–F, J); sternal intersegmental sutures carinulate between II–IV and laterally between IV–VI.....	<i>steineri</i> , new species
21. Interocular sulcus near hind margin of eye (Fig. 2.13B); head, including postclypeal depression, anterior pronotal lobe, and legs reddish brown.....	<i>namoroka</i> , new species
- Interocular sulcus anterior to hind margin of eye; head dark brown with postclypeal depression and labium pale; anterior pronotal lobe generally dark brown; legs dark brown and pale.....	<i>vazimba</i> , new species

*Identification key to the females of species of Toxopus*

1. Hind tibia distinctly curved, armed with small papillae (Fig. 2.16D, E); shiny dark brown to black; basal part of scape and antennal segments V–VII entirely, except basal half of V, pale; dorsal surface of femora basally, foretibia ventrally except basally and apically, rings on mid and hind tibiae, and tarsi pale to brown (based on original description)..... *politus* Bergroth
- Hind tibia not distinctly curved, unarmed; shiny blackish with extensive pale to brown markings  
..... *griswoldi*, new species

*Toxopus ambohitantely*, new species

Figs. 2.3, 2.7, 2.10, 2.13J; Map 2.11

DIAGNOSIS: Males are recognized among other *Toxopus* species by the moderately dense vestiture, coloration, elongate cylindrical head in lateral view, relatively deep postclypeal depression that reaches the interocular sulcus, sublateral antennal insertion, labial segment III ventrally convex, and intersegmental sutures carinulate between sternites II and III and laterally between III–V. This species is similar to *T. italaviana*, *T. griswoldi*, *T. insignis*, *T. fisheri*, *T. antsiranana*, and *T. toliara*, but is distinguished from them by the intersegmental sutures carinulate between sternites II and III and laterally between III–V, subtriangular pygophore process, and endosoma medially sclerotized.

DESCRIPTION: **MALE:** Body length: 9.93 mm (holotype), 9.83–10.18 mm. COLORATION: Blackish with pale yellow corium basally, trochanters, femora except dorsoapical half, fore- and mid tibiae medially, hind tibia subbasally and medially, anterior half of laterotergites, abdominal sternites medially (except anterior, posterior, and lateral margins), sternites II–VI anterolaterally, and lateral margin of sternite VII. Antennal segments VI and VII, hemelytron, and tarsi brown. VESTITURE: Moderately dense; other features as in generic description. STRUCTURE: HEAD

(Fig. 2.13J): Cylindrical in lateral view; longer than wide in dorsal view; ventrally with shallow anteromedial depression; antocular region longer than postocular; postclypeus with deep, narrow medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus posterior to hind margin of eye; postocular broad in dorsal view (Fig. 2.13B); ocelli small, located on shallow median tubercle, separated by a distance less than or equal to diameter of ocellus; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about one-fourth longer than scape, slightly curved; labium slender; labial segment III ventrally convex. THORAX: Anterior pronotal lobe with relatively small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded, sinuate, or medially notched; intersegmental sutures carinate between sternites II and III and laterally between III–V; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts not reaching posterior margin of DPS (Fig. 2.21L), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion ovate; endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Ambohitantely, Madagascar.

DISTRIBUTION: Antananarivo, Antsiranana, Mahajanga, and Toamasina provinces (Map 2.11).

DISCUSSION: In some specimens, the base of the scape, antennal segments VI and VII, scutellar processes, femora, tibiae, and corium are more pale yellow than dark brown. The labium may be light brown.

HOLOTYPE: Male: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 15 Oct 2004–01 Nov 2004, M. Irwin, F. Parker, R. Harin'Hala (00006782) (CAS).

PARATYPES: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 31 Oct 2003–07 Nov 2003, M. Irwin, R. Harin'Hala, 3♂ (00006788–00006790) (CAS); 05 Mar 2004–19 Mar 2004, M. Irwin, R. Harin'Hala, 1♂ (00006791) (CAS); 19 Mar 2004–28 Mar 2004, M. Irwin, R. Harin'Hala, 1♂ (00006778) (SU); 15 Oct 2004–01 Nov 2004, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006779–00006781) (USNM), 2♂ (00006783, 00006784) (CAS), 3♂ (00006785–00006787) (MNHN); 01 Nov 2004–14 Nov 2004, M. Irwin, R. Harin'Hala, 1♂ (00006792) (UCR); 17 Apr 2005–26 Apr 2005, M. Irwin, R. Harin'Hala, 2♂ (00044905, 00045071) (AMNH), 2♂ (00045364, 00045369) (BMNH). **Antsiranana:** Parc National Montagne d'Ambre, 12.51666°S 49.18333°E, 975 m, 04 Mar 2001–19 Mar 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 1♂ (00006120) (CAS). Sakalava Beach, dwarf littoral forest, 12.26277°S 49.3975°E, 10 m, 15 Feb 2001–06 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006794) (USNM); 22 Apr 2001–27 Apr 2001, M. Irwin, R. Harin'Hala, 1♂ (00045583) (SU). **Mahajanga:** Reserve Speciale de Bemarivo, 23.8 km 223° SW Besalampy, 16.925°S 44.36833°E, 30 m, 19 Nov 2002–23 Nov 2002, Fisher et al., 1♂ (00006793) (CAS). **Toamasina:** 7 km SE of Andasibe National Park headquarters, 18.96266°S 48.45266°E, 1050 m, 07 Jan 2001–22 Jan 2001, M. Irwin, R. Harin'Hala, 1♂ (00007142) (CAS). Botanic Garden near entrance to Andasibe National Park, 18.92633°S 48.40783°E, 1025 m, 01 Sep 2001–05 Sep 2001, M. Irwin, R. Harin'Hala, 1♂ (00006795) (MNHN); 19 Sep 2001–26 Sep 2001, M. Irwin, R. Harin'Hala, 1♂ (00045254) (UCR).

*Toxopus ampitavananima*, new species

Figs. 2.3, 2.7, 2.10, 2.22G; Map 2.12

DIAGNOSIS: Males are recognized among other *Toxopus* species by the subquadrate head shape in lateral view that is longer than wide in dorsal view, anterior pronotal lobe without anterolateral protuberances, and intersegmental sutures carinulate between sternites II and III and laterally between III–VI. This species is similar to *T. vazimba*, but the smaller synthlipsis width, interocular sulcus located near the hind margin of the eye, eye one-half of the head length in lateral view, anterior pronotal lobe lacking distinct anterolateral protuberances, striated posterior pronotal lobe, intersegmental sutures carinulate between sternites II and III and laterally between III–VI, and BPE as long as the basal plate distinguish *T. ampitavananima* from *T. vazimba*.

DESCRIPTION: **MALE:** Body length: 10.67 mm (holotype), 10.50–11.91 mm.

COLORATION: Blackish with pale yellow mandibular and maxillary plates dorsally, postclypeal medial longitudinal depression, anterolateral area anterior of interocular sulcus, scape basally, pedicel basally, antennal segment III basally, antennal segment IV apically, antennal segments VI and VII, anterior pronotal lobe anterolaterally, posterior margin of posterior pronotal lobe, stridulatory process, meso- and metasterna, scutellar processes apically, corium basally, coxae, trochanters, forefemur basally and ventrally, mid femur basally and subapically, hind femur except medially, foretibia except basally, mid tibia except subbasally, hind tibia apically, tarsi, anterior half of laterotergites, abdominal sternites II–V except medially and laterally, ventrolateral anterior margin of sternite VI, and lateral margin of abdomen except apex of sternite VII. Labium and remainder of pronotum brown. VESTITURE: Moderately dense; other features as in generic description. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13K); longer than wide in dorsal view; ventrally with deep medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to



interocular sulcus; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on shallow median tubercle (Fig. 2.13K), separated by less than diameter of ocellus; eye about half of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe without anterolateral projections; posterior pronotal lobe weakly to strongly transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe weakly to strongly transversely striated; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II and III and laterally between III–VI; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21K); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Ampitavananima Forest, Madagascar.

DISTRIBUTION: Antsiranana, Fianarantsoa, Toamasina, and Toliara provinces (Map 2.12).

DISCUSSION: The coloration in this species is variable: the pale yellow coloration on the head and scutellum may be reduced, the anterior pronotal lobe may have more pale yellow coloration throughout, the posterior pronotal lobe may be dark brown, and the entire ventromedial abdominal surface may be pale yellow or nearly so.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 14 Apr 2007–21 Apr 2007, M. Irwin, F. Parker, R. Harin'Hala (00006768) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Parc National Montagne d'Ambre, 12.51444°S 49.18138°E, 960 m, 04 Mar 2001–19 Mar 2001, M. Irwin, R. Harin'Hala, 1♂ (00006719) (CAS). **Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 13 Jan 2007–20 Jan 2007, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006771–00006773) (CAS); 04 Feb 2007–10 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006762–00006764) (CAS); 10 Feb 2007–17 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006766) (UCR); 17 Feb 2007–24 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006740) (CAS); 10 Mar 2007–17 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 4♂ (00006743–00006746) (BMNH), 2♂ (00006755, 00006756) (UCR); 17 Mar 2007–24 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006741) (SU); 24 Mar 2007–31 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006769) (CAS); 07 Apr 2007–14 Apr 2007, M. Irwin, F. Parker, R. Harin'Hala, 4♂ (00006737–00006739, 00006753) (MNHN), 1♂ (00006754) (CAS); 21 Apr 2007–29 Apr 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006761) (CAS); 06 May 2007–14 May 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006770) (USNM); 14 May 2007–21 May 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006775) (USNM); 21 May 2007–25 May 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006752) (USNM); 25 May 2007–02 Jun 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006767) (USNM); 02 Jun 2007–10 Jun 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006760) (CAS); 01 Jul 2007–07 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006765) (CAS); 14 Jul 2007–19 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006742) (CAS); 19 Jul 2007–23 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006747, 00006748) (SU); 23 Jul 2007–28 Jul 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂

(00006121, 00006735) (CAS); 13 Sep 2007–20 Sep 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
(00006776) (CAS); 27 Sep 2007–04 Oct 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
(00006774) (CAS); 04 Oct 2007–11 Oct 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
(00006751) (CAS); 11 Oct 2007–18 Oct 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂  
(00006758, 00044833) (AMNH); 25 Oct 2007–08 Nov 2007, M. Irwin, F. Parker, R. Harin'Hala,  
1♂ (00006736) (AMNH); 03 Jan 2008–09 Jan 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
(00006731) (AMNH); 09 Jan 2008–17 Jan 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
(00006750) (CAS); 17 Jan 2008–24 Jan 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
(00006749) (CAS); 02 Mar 2008–08 Mar 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂  
(00006759) (SU); 16 Mar 2008–22 Mar 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006732)  
(UCR). Manombo Special Reserve camp site, 32 km SSE of Farafangana, 23.02183°S 47.72°E,  
36 m, 26 Dec 2004–05 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00045458) (CAS); 10 May 2005–  
22 May 2005, M. Irwin, R. Harin'Hala, 2♂ (00045398, 00045685) (CAS); 22 May 2005–06 Jun  
2005, M. Irwin, R. Harin'Hala, 1♂ (00045378) (SU); 10 Jul 2005–27 Jul 2005, M. Irwin, R.  
Harin'Hala, 1♂ (00007067) (UCR). **Toamasina:** Mobot Site, Analalava 7 km SW of Foulpointe,  
17.69333°S 49.46027°E, 18 m, 29 Feb 2008–07 Mar 2008, M. Irwin, R. Harin'Hala, 1♂  
(00044815) (CAS). **Toliara:** Forêt Ivohibe 55.0 km N Tolagnaro, 24.56888°S 47.20388°E, 200  
m, 02 Dec 2006–04 Dec 2006, B.L. Fisher et al., 1♂ (00006733) (CAS). **Unknown:** 1♂  
(00006757) (CAS).

*Toxopus antsiranana*, new species

Figs. 2.3, 2.7, 2.10; Map 2.10

DIAGNOSIS: Males are recognized among other species in this genus by a combination of the following characters: the coloration, elongate cylindrical head in lateral view, synthlipsis 1.5

times width of an eye, antenna inserted sublaterally on the head, labial segment III ventrally straight, and intersegmental sutures carinulate between sternites II and III and laterally between III and IV. Males are similar to males of *T. italaviana*, *T. griswoldi*, *T. insignis*, *T. fisheri*, and *T. toliara*, but are distinguished by the smaller synthlipsis width and intersegmental sutures carinulate between sternites II and III and laterally between III and IV.

DESCRIPTION: **MALE:** Body length: 12.21 mm (holotype), 11.10–12.21 mm.

COLORATION: Blackish with pale yellow mandibular plates dorsally, postclypeus between antennifers, antennifer, labial segment IV except basally, scape basally, pedicel basally, base of antennal segment III, antennal segment VII, metasternum laterally, corium basally, mid and hind coxae posteriorly, trochanters, forefemur basally and subapically, mid and hind femora except medially, foretibia medially and subbasally, mid tibia subbasally and subapically, hind tibia basally and subapically, tarsi, anterior half of laterotergites, and abdominal sternites medially and anterolaterally. VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD: Cylindrical in lateral view (Fig. 2.13J); longer than wide in dorsal view; ventrally with shallow anteromedial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli small, located on shallow median tubercle (Fig. 2.13J), separated by a distance less than or equal to diameter of ocellus; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about one-fourth longer than scape, slightly curved; labium slender; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe weakly transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral

depressions of posterior pronotal lobe weakly transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded or sinuate; intersegmental sutures carinulate between sternites II and III and laterally between III and IV; pygophore process curved in lateral view (Fig. 2.20K), elongate in caudal view (Fig. 2.19H), surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21K); endosoma with sclerotized, hairlike, denticles on posterior margin (Fig. 2.21J, K).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the Antsiranana province in Madagascar, where the type specimen was collected.

DISTRIBUTION: Toamasina and southern region of Antsiranana provinces (Map 2.10).

DISCUSSION: Coloration varies with the dorsal area between antennifers sometimes brown.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Forêt Ambanitaza, 26.1 km 347° Antalaha, 14.67944°S 50.18361°E, 240 m, 26 Nov 2004, B.L. Fisher (00044804) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Forêt Ambanitaza, 26.1 km 347° Antalaha, 14.67944°S 50.18361°E, 240 m, 26 Nov 2004, B.L. Fisher, 1♂ (00045129) (CAS). Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 25 Dec 2004–30 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00044835) (CAS); 05 Jan 2005–10 Jan 2005, M. Irwin, R. Harin'Hala, 3♂ (00006128, 00006970, 00007099) (UCR); 25 Feb 2005–04 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00045652) (CAS); 18 Mar 2005–25 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00006098) (CAS); 25 Mar 2005–04 Apr 2005, M. Irwin, R. Harin'Hala, 1♂ (00006100) (SU); 18 May 2005–30 May 2005, M. Irwin, R. Harin'Hala, 1♂ (00007189) (SU). **Toamasina:** Fampanambo, 15.37550°S 49.62175°E, 108 m, Feb 1959, J. Vadon, 2♂, (MRAC); Feb 1961, J. Vadon, 2♂, (MRAC). Ivoloina Zoological Park, 12 km NW of

Toamasina, 18.05933°S 49.35790°E, 29 m, 16 Nov 2000 – 20 Nov 2000, Dolin and Andreeva, 1♂ (TLMF). Nosy Mangabe, 15.49324°S 49.76776°E, 266 m, 27 Dec 2003, R. Dolin, 1♂ (TLMF). Périnet (Andasibe), 18.82666°S 48.44778°E, 1119 m, 19 Dec 2001 – 31 Dec 2001, V. Dolin, 1♂ (TLMF). **Unknown:** 1♂ (00007191) (CAS).

*Toxopus basalis*, new species

Figs. 2.3, 2.7, 2.10, 2.22H; Map 2.11

DIAGNOSIS: Males recognized among other species in *Toxopus* by the coloration, sparse vestiture, head as long as wide, synthlipsis 1.5 times width of an eye, interocular sulcus anterior to the hind margin of the eye, eye reaching the ventral head margin, striated posterior pronotal lobe, and intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI. Males are similar to males of *T. steineri*, *T. namoroka*, and *T. vazimba*, but differs from these species by the characters mentioned. This species is most similar to *T. brucei*, but is distinguished by the smaller width of the synthlipsis relative to the eye, interocular sulcus anterior to the hind margin of the eye, larger eye in lateral view, eye not reaching ventral head margin, and striated posterior pronotal lobe.

DESCRIPTION: **MALE:** Body length: 13.01 mm (holotype), 11.49–13.01 mm.

COLORATION: Blackish with pale yellow postclypeal medial longitudinal depression, labial segment IV, scape basally, meso- and metasterna, corium basally, mid and hind coxae, trochanters, forefemur ventrally and basally, mid and hind femora basally, tibiae medially, tarsi, anterior half of laterotergites, abdominal sternites II–VI medially (except at margins of II and III and posteromedial spots on III–VI), and anterolateral spots on sternites. VESTITURE: Very sparse; other features as in generic description. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13K); as long as wide in dorsal view; ventrally with deep medial depression; anteocular

region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about 1.5 times width of eye; interocular sulcus anterior to hind margin of eye; postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on shallow median tubercle (Fig. 2.13K), separated by less than diameter of ocellus; eye about half of head length, not reaching dorsal head surface, reaching ventral head surface; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe weakly transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe weakly transversely striated; hemelytron surpassing tergite VII and not reaching, reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts not reaching posterior margin of DPS (Fig. 2.21L), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate ovoid (Fig. 2.21L); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the pale basal part of the femora.

DISTRIBUTION: Mahajanga and northern region of Toliara provinces (Map 2.11).

DISCUSSION: The coloration of the medial longitudinal depression on the postclypeus is narrowly or broadly pale yellow in some specimens. The variability of antennal color patterns could not be fully assessed because parts of the antennae are missing in most specimens. In one specimen, the basal antennal segments are light brown with segments V–VII pale yellow. This may be a variant as all other specimens have a dark brown to black scape and pedicel. It is likely

that segments V–VII are always pale yellow.

HOLOTYPE: Male: **Madagascar: Toliara:** Beroboka village, 45 km NE Morondava, 19.9775°S 44.82483°E, 131 m, 03 Dec 2008–11 Dec 2008, M. Irwin, R. Harin'Hala (00045571) (CAS).

PARATYPES: **Madagascar: Mahajanga:** 160 km N of Maevatanana on RN 04, Ampijoroa National Park, 16.31933°S 46.81333°E, 43 m, 07 Feb 2005–19 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00045511) (CAS). **Toliara:** Beroboka village, 45 km NE Morondava, 19.9775°S 44.66533°E, 128 m, 16 Oct 2008–24 Oct 2008, M. Irwin, R. Harin'Hala, 1♂ (00045574) (CAS); 09 Nov 2008–17 Nov 2008, M. Irwin, R. Harin'Hala, 1♂ (00044958) (CAS); 17 Nov 2008–25 Nov 2008, M. Irwin, R. Harin'Hala, 2♂ (00007161, 00045159) (USNM); 19 Dec 2008–27 Dec 2008, M. Irwin, R. Harin'Hala, 1♂ (00044937) (UCR). Beroboka village, 45 km NE Morondava, 19.9775°S 44.82483°E, 131 m, 30 Oct 2007–06 Nov 2007, M. Irwin, R. Harin'Hala, 2♂ (00006972, 00006975) (BMNH), 1♂ (00006977) (CAS); 22 Nov 2007–30 Nov 2007, M. Irwin, R. Harin'Hala, 2♂ (00007073, 00044898) (CAS), 2♂ (00045223, 00045525) (SU); 17 Nov 2008–25 Nov 2008, M. Irwin, R. Harin'Hala, 1♂ (00045070) (CAS); 25 Nov 2008–03 Dec 2008, M. Irwin, R. Harin'Hala, 1♂ (00045705) (UCR). **Unknown:** 1♂ (00045556) (CAS).

*Toxopus brucei*, new species

Figs. 2.3, 2.7, 2.10; Map 2.12

DIAGNOSIS: Males recognized among other species in this genus by the coloration, sparse vestiture, head as long as wide, smooth posterior pronotal lobe, and intersegmental sutures carinate between sternites II–IV and laterally between IV–VI. This species is similar to *T. steineri*, *T. namoroka*, and *T. vazimba*, but differs from these species by the features mentioned.



This species is most similar to *T. basalis*, from which it differs by the larger synthlipsis width, interocular sulcus near the hind margin of the eye, smaller eye width in lateral view, eye not reaching ventral head margin, and smooth posterior pronotal lobe.

DESCRIPTION: **MALE:** Body length: 12.61 mm (holotype), 11.00–12.70 mm.

COLORATION: Blackish with pale yellow scape basally, antennal segments VI and VII, corium basally, trochanters, forefemur ventrally, mid and hind femora basally, tibiae medially, tarsi, anterior half of laterotergites, abdominal sternites III–VI medially except at margins, and anterolateral spots on sternites. VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13K); as long as wide in dorsal view; ventrally with deep medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on shallow median tubercle (Fig. 2.13K), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe smooth; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded or sinuate; intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts not reaching posterior margin of DPS (Fig. 2.21L), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate ovoid (Fig. 2.21L);

endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: This species epithet is a noun in the genitive case and is named after the senior author's uncle, Thomas Allen Bruce.

DISTRIBUTION: Mahajanga and south-central region of Toamasina provinces (Map 2.12).

DISCUSSION: The coloration is slightly variable with antennal segment V pale yellow, the corium, except basally, brown, the meso- and metasterna and sternite II with a medial yellow spot, and the medial spots on the sternites are not bordered by dark brown margins.

HOLOTYPE: Male: **Madagascar: Mahajanga:** Besalampy District, Analangidro dry forest, 7 km NE of Besalampy, 16.6915°S 44.5235°E, 09 Oct 2007–16 Oct 2007, M. Irwin, R. Harin'Hala (00007133) (CAS).

PARATYPES: **Madagascar: Mahajanga:** Besalampy District, Analangidro dry forest, 7 km NE of Besalampy, 16.6915°S 44.5235°E, 61 m, 02 Oct 2007–09 Oct 2007, M. Irwin, R. Harin'Hala, 1♂ (00007039) (CAS); 09 Oct 2007–16 Oct 2007, M. Irwin, R. Harin'Hala, 1♂ (00007217) (SU); 14 Dec 2007–21 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00045338) (UCR). Besalampy District, Marofototra dry forest, 17 km W of Besalampy, 16.72166°S 44.42366°E, 52 m, 26 Nov 2007–03 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00044822) (CAS), 1♂ (00044972) (BMNH); 10 Dec 2007–17 Dec 2007, M. Irwin, R. Harin'Hala, 2♂ (00007043, 00007156) (AMNH), 2♂ (00007160, 00007170) (USNM), 2♂ (00007182, 00007223) (CAS), 1♂ (00045011) (BMNH); 17 Dec 2007–24 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00045155) (SU). Besalampy District, Marofototra palm forest, 17 km W of Besalampy, 16.71666°S 44.41666°E, 11 m, 10 Dec 2007–17 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00045286) (CAS). Maintirano District Asondrodava dry forest, 15 km N of Maintirano, 17.96533°S 44.0355°E, 61 m, 03 Dec 2007–10 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00006140) (CAS); 10 Dec 2007–17 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00007012) (UCR). Maintirano District, 50 km E of Maintirano,

18.004°S 44.452°E, 274 m, 25 Oct 2007–01 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00007193) (CAS). Namoroka village, Befatika Andranovory, 7 km NW Vilanandro village, 16.47333°S 45.39133°E, 122 m, 09 Nov 2007–16 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00007131) (CAS). Reserve Speciale de Bemarivo, 23.8 km 223° SW Besalampy, 16.925°S 44.36833°E, 30 m, 19 Nov 2002–23 Nov 2002, Fisher et al., 1♂ (00006149) (CAS). **Toamasina:** Botanic Garden near entrance to Andasibe National Park, 18.92633°S 48.40783°E, 1025 m, 05 Sep 2001–19 Sep 2001, M. Irwin, R. Harin'Hala, 1♂ (00044971) (CAS).

*Toxopus farafangana*, new species

Figs. 2.3, 2.7, 2.10; Map 2.11

DIAGNOSIS: Males are recognized among other *Toxopus* species by the coloration, ovoid head shape in lateral view, antecular region as long as postocular, broad postocular, labial segment III ventrally convex, smooth posterior pronotal lobe, and BPE shorter than the basal plate. Males are similar to *T. pallidus*, but the dark brown to blackish posterior propleural lobe and thoracic sterna, predominately dark brown to blackish ventral abdominal surface, interocular sulcus near the hind margin of the eye, labial segment III ventrally convex, smooth posterior pronotal lobe, and BPE shorter than the basal plate distinguish *T. farafangana* from *T. pallidus*.

DESCRIPTION: **MALE:** Body length: 10.19 mm (holotype), 9.05–10.66 mm.

COLORATION: Blackish with pale yellow anterior margin of antennifer, antennal segment V (except basally) and VI and VII, corium basally, mid and hind trochanters, forefemur ventrally, mid and hind femora basally and subapically, foretibia subbasally and apically, mid and hind tibiae apically and medially, tarsi, anterior half of laterotergites, small posterior paramedian stripes on abdominal sternites II–VII, lateral margin of sternites II and III, and anterolateral margin of sternites IV–VII. Labium except basal two-thirds of segment II, posterior pronotal lobe,

and remainder of corium brown. VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; ventrally with shallow anteromedial depression; anteocular region as long as postocular (Fig. 2.13J); postclypeus with deep, broad medial longitudinal depression to middle of interocular area; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular moderately broad in dorsal view (Fig. 2.13B); ocelli large, located on shallow median tubercle (Fig. 2.13K), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium slender; labial segment III shorter than II, ventrally convex (Fig. 2.13J). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe weakly transversely striated; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinate between sternites II and III; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore, apex rounded; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion ovate; endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in apposition and is named after a locality near which known specimens were collected.

DISTRIBUTION: Southeastern region of the Fianarantsoa province (Map 2.11).

DISCUSSION: In some specimens, the coloration is slightly variable with the posterior pronotal lobe dark brown and the abdominal ventrolateral pale yellow spots sometimes larger on some sternites or absent.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 17 Feb 2007–24 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala (00007034) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 13 Jan 2007–20 Jan 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00044948, 00044974) (CAS); 20 Jan 2007–26 Jan 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006147, 00007265) (CAS); 26 Jan 2007–04 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00044798, 00045114) (CAS), 2♂ (00045438, 00045524) (BMNH); 26 Jan 2007–04 Feb 2007, M. Irwin, R. Harin'Hala, 1♂ (00045049) (UCR); 04 Feb 2007–10 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00007080, 00007207) (CAS); 17 Feb 2007–24 Feb 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00007230) (SU); 24 Feb 2007–03 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045346) (CAS); 03 Mar 2007–10 Mar 2007, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00007173, 00007175, 00007252) (USNM); 06 Sep 2007–13 Sep 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00007052) (CAS); 08 Nov 2007–22 Nov 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045221) (SU); 20 Dec 2007–27 Dec 2007, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00045003, 00045332) (AMNH); 03 Jan 2008–09 Jan 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00044965) (CAS); 09 Jan 2008–17 Jan 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00044938) (UCR); 16 Mar 2008–22 Mar 2008, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006729, 00044968) (CAS). Manombo Special Reserve camp site, 32 km SSE of Farafangana, 23.02183°S 47.72°E, 36 m, 12 Nov 2004–23 Nov 2004, M. Irwin, R. Harin'Hala, 1♂ (00045531) (CAS); 15 Dec 2004–26 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00044890) (SU); 16 Jan 2005–20 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00045258) (UCR).

*Toxopus fisheri*, new species

Figs. 2.3, 2.7, 2.10, 2.19H, 2.20K, 2.21J; Map 2.12

DIAGNOSIS: Males are recognized among other species in this genus by the sparse vestiture, coloration, elongate cylindrical head in lateral view, ventrally flat head surface, sublateral antennal insertion, labial segment III ventrally straight, and intersegmental sutures carinulate between sternites II and III and laterally between III–VI. This species is similar to *T. italaviana*, *T. griswoldi*, *T. antsiranana*, and *T. toliara*, but differs from these species by the ventrally flat head surface. *Toxopus fisheri* are also similar to *T. insignis*, from which it differs by the dark brown to blackish postclypeal depression, pale yellow maxillary plates and antennifers, and intersegmental sutures carinulate between sternites II and III and laterally between III–VI.

DESCRIPTION: **MALE:** Body length: 10.19 mm (holotype), 9.84–11.09 mm.

COLORATION: Blackish with pale yellow maxillary plates dorsally, antennifer, scape basally, pedicel basally, base of antennal segment III, antennal segment VII, corium basally, coxae, trochanters, forefemur basally and subapically, mid and hind femora except medially, foretibia medially except lateral spots, mid and hind tibiae subbasally and medially, tarsi, laterotergites II and VII, anterior half of laterotergites III–VI, and abdominal sternites laterally and medially except intersternal sutures. Antennal segments V, VI, and apex of VII and labium brown.

VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Cylindrical in lateral view (Fig. 2.13J); longer than wide in dorsal view; ventrally flat; anteocular region longer than postocular (Fig. 2.13K); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli small, located on shallow median tubercle (Fig. 2.13J), separated by a distance less than or equal to diameter of ocellus; eye about one-fourth of head length, not reaching dorsal and

ventral head surfaces; antenna inserted sublaterally on head; pedicel about two-fifths longer than scape, slightly curved; labium slender; labial segment III ventrally straight (Fig. 2.13K).

THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe weakly transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II and III and laterally between III–VI; pygophore process curved in lateral view (Fig. 2.20K), elongate in caudal view (Fig. 2.19H), surpassing dorsal margin of pygophore; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion ovate (Fig. 2.21J); endosoma with sclerotized, hairlike, denticles on posterior margin (Fig. 2.21J).

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after Brian Fisher, who collected the majority of specimens examined for this species.

DISTRIBUTION: Antsiranana province (Map 2.12).

DISCUSSION: The coloration varies with the small medial longitudinal stripe on the postclypeus, meso- and metasterna, and mid tibia medially pale yellow in some specimens.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Forêt de Binara, 7.5 km 230° SW Daraina, 13.255°S 49.61666°E, 375 m, 01 Dec 2003, B.L. Fisher (00045439) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Forêt Bekaraoka, 6.8 km 60° ENE Daraina, 13.00277°S 49.01166°E, 150 m, 07 Dec 2003, B.L. Fisher, 1♂ (00045028) (UCR), 1♂ (00045100) (SU), 2♂ (00045143, 00045592) (CAS). Forêt d'Ampondrabe, 26.3 km 10° NNE Daraina, 12.97°S 49.7°E, 175 m, 10 Dec 2003, B.L. Fisher, 1♂ (00045087) (CAS), 2♂ (00045141, 00045148) (SU), 1♂ (00045636) (UCR). Forêt de Binara, 7.5 km 230° SW Daraina,

13.255°S 49.61666°E, 375 m, 01 Dec 2003, B.L. Fisher, 1♂ (00045469) (CAS). Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 15 Dec 2004–20 Dec 2004, M. Irwin, R. Harin'Hala, 1♂ (00006969) (CAS); 05 Jan 2005–10 Jan 2005, M. Irwin, R. Harin'Hala, 1♂ (00006968) (UCR). Rés. Analamerana, 28.4 km 99° Anivorano-Nord, 12.74666°S 49.49472°E, 60 m, 05 Dec 2004, B.L. Fisher, 1♂ (00045431) (CAS).

*Toxopus griswoldi*, new species

Figs. 2.3, 2.4, 2.7, 2.10, 2.23E; Map 2.10

DIAGNOSIS: Males recognized among other *Toxopus* species by the small body size, sparse vestiture, coloration, elongate cylindrical head in lateral view, sublateral antennal insertion, labial segment III ventrally straight, and intersegmental sutures carinulate between sternites II–IV and laterally between IV–VII. In addition to the generic diagnostic characters, the slender, slightly curved hind tibia distinguishes females of this species from other known *Toxopus* females. Males are similar to *T. italaviana*, *T. insignis*, *T. fisheri*, *T. antsiranana*, and *T. toliara*, but differs from these species by the small body size. The sparse vestiture further distinguishes this species from *T. italaviana*, *T. antsiranana*, and *T. toliara*, while the intersegmental sutures carinulate between sternites II–IV and laterally between IV–VII differentiate *T. griswoldi* from *T. insignis* and *T. fisheri*.

DESCRIPTION: **MALE:** Body length: 9.28 mm (holotype), 8.74–9.76 mm. **COLORATION:** Blackish with pale yellow clypeus; antennifer; scape except apically, antennal segment VII except apically, collar, posterolateral pronotal margins, meso- and metasterna medially, scutellar processes, corium basally, coxae, trochanters, forefemur basally and subapically, mid and hind femora except medially, tibiae medially, tarsi, laterotergite II, anterior area of laterotergites III–



VII, and abdominal sternites medially and laterally. Antennal segment VI and apex of VII and labium brown. VESTITURE: Very sparse; other features as in generic description.

STRUCTURE: HEAD: Cylindrical in lateral view (Fig. 2.13J); longer than wide in dorsal view; ventrally with shallow anteromedial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area (Fig. 2.13B); synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli small, located on shallow median tubercle (Fig. 2.13J), separated by diameter of ocellus; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about one-fourth longer than scape, slightly curved; labium slender; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–IV and laterally between IV–VII; pygophore process curved in lateral view (Fig. 2.20K), elongate in caudal view (Fig. 2.19H), not surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21K); endosoma with sclerotized, hairlike, denticles on posterior margin (Fig. 2.21J, K). **FEMALE:** Body length: 11.75 mm; in addition to characters mentioned in the generic description: STRUCTURE: HEAD: Synthlipsis about 3.5 times width of eye. THORAX: Hind tibia slender, slightly curved. ABDOMEN: Intersegmental sutures carinulate laterally between sternites II–VI; external genitalia as in plate 23E; bursa copulatrix membranous, damaged; vermiform gland and lateral spermatheca missing.

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after Charles Griswold, who collected the only known female specimen known for this species.

DISTRIBUTION: Mahajanga, Toliara, and the northern area of the Antananarivo provinces (Map 2.10).

DISCUSSION: Some specimens have pale yellow mandibular plates. The female specimen was associated with males based on morphology, geographic proximity, and molecular data (COI pairwise genetic distance 0.76%).

HOLOTYPE: Male: **Madagascar: Toliara:** Cap Ste Marie Special Reserve, 74 km S of Tsihombe, 25.58766°S 45.163°E, 37 m, 23 Nov 2003–30 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala (00045173) (CAS).

PARATYPES: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 28 Dec 2003–10 Jan 2004, M. Irwin, R. Harin'Hala, 1♂ (00045139) (CAS); 14 Nov 2004–29 Nov 2004, M. Irwin, R. Harin'Hala, 1♂ (00045132) (UCR).

**Mahajanga:** Ambovomamy Belambo, 20 km NW of Port Berger, 15.45116°S 47.61333°E, 33 m, 28 Oct 2007–06 Nov 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045234) (CAS); 23 Nov 2007–08 Dec 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00044962) (SU); 08 Dec 2007–11 Dec 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045233) (AMNH); 26 Oct 2008–01 Nov 2008, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00044994, 00045075) (CAS), 2♂ (00045386, 00045645) (USNM); 16 Nov 2008–23 Nov 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045056) (CAS); 23 Nov 2008–30 Nov 2008, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045692) (UCR). Analamanitra Forest, 14 km NE of Misinjo, 16.13333°S 45.7°E, 20 m, 23 Oct 2007–30 Oct 2007, M. Irwin, R. Harin'Hala, 1♂ (00045008) (CAS); 30 Oct 2007–16 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00007148) (AMNH); 16 Nov 2007–20 Nov 2007, M. Irwin, R. Harin'Hala, 3♂ (00007096, 00007176, 00045025) (MNHN); 20 Nov 2007–04 Dec 2007, M.

Irwin, R. Harin'Hala, 2♂ (00044799, 00044896) (USNM). Anjiaabo, 3 km N Baly village, 16.059°S 45.27416°E, 7 m, 07 Nov 2007–24 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00007214) (CAS). Besalampy District, Analangidro dry forest, 7 km NE of Besalampy, 16.6915°S 44.5235°E, 61 m, 24 Sep 2007–02 Oct 2007, M. Irwin, R. Harin'Hala, 1♂ (00044806) (CAS); 18 Nov 2007–25 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00044886) (SU). Besalampy District, Marofototra dry forest, 17 km W of Besalampy, 16.72166°S 44.42366°E, 52 m, 19 Nov 2007–26 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00044880) (CAS); 26 Nov 2007–03 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00044885) (SU). Namoroka 53 km from Soalala 3 km N Vilanandro Village, 16.47333°S 45.39133°E, 122 m, 16 Nov 2007–23 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00007171) (BMNH), 1♂ (00007212) (CAS). Namoroka village, Befatika Andranovory, 7 km NW Vilanandro village, 16.47333°S 45.39133°E, 122 m, 02 Nov 2007–09 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00045002) (CAS). Parc National de Namoroka, 16.9 km 317° NW Vilanandro, 16.40666°S 45.31°E, 100 m, 12 Nov 2002–16 Nov 2002, Fisher et al., 1♂ (00045095) (CAS), 1♂ (00045133) (SU); 12 Nov 2002–16 Nov 2002, Fisher et al., 1♀ (00045042) (CAS). Parc National de Namoroka, 17.8 km 329° WNW Vilanandro, 16.37666°S 45.32666°E, 100 m, 08 Nov 2002–12 Nov 2002, Fisher et al., 1♂ (00006435) (CAS). **Toliara:** Beroboka village, 45 km NE Morondava, 19.9775°S 44.66533°E, 128 m, 30 Sep 2008–08 Oct 2008, M. Irwin, R. Harin'Hala, 1♂ (00044841) (CAS); 08 Oct 2008–16 Oct 2008, M. Irwin, R. Harin'Hala, 1♂ (00045649) (AMNH); 09 Nov 2008–17 Nov 2008, M. Irwin, R. Harin'Hala, 2♂ (00044915, 00045565) (CAS); 11 Dec 2008–19 Dec 2008, M. Irwin, R. Harin'Hala, 1♂ (00045695) (UCR). Beroboka village, 45 km NE Morondava, 19.9775°S 44.82483°E, 131 m, 14 Oct 2007–18 Oct 2007, M. Irwin, R. Harin'Hala, 1♂ (00045110) (CAS); 18 Oct 2007–26 Oct 2007, M. Irwin, R. Harin'Hala, 2♂ (00044859, 00045096) (BMNH); 17 Nov 2008–25 Nov 2008, M. Irwin, R. Harin'Hala, 1♂ (00044814) (AMNH), 1♂ (00045041) (CAS); 05 Feb 2009–13 Feb

2009, M. Irwin, R. Harin'Hala, 1♂ (00045208) (CAS). Cap Ste Marie Special Reserve, 74 km S of Tsihombe, 25.58766°S 45.163°E, 37 m, 23 Nov 2003–30 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045379) (CAS). Parc National d'Andohahela, Ihazofotsy Parcelle III, 24.83083°S 46.53616°E, 80 m, 16 Dec 2002–26 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045333) (CAS); 21 Apr 2003–28 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00045635) (UCR).

*Toxopus insignis*, new species

Figs. 2.4, 2.7, 2.10, 2.14; Map 2.13

DIAGNOSIS: Males are recognized among other species in *Toxopus* by the sparse vestiture, coloration, elongate cylindrical head in lateral view, sublateral antennal insertion, straight pedicel, and labial segment III ventrally straight. This species is similar to *T. italaviana*, *T. griswoldi*, *T. fisheri*, *T. antsiranana*, and *T. toliara*, from which it differs by the straight pedicel. The ventrally flat head surface further differentiates *T. insignis* males from the mentioned species, except for *T. fisheri*, which can be discerned from by the pale yellow postclypeal depression, dark brown maxillary plates and antennifers, and intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI.

DESCRIPTION: **MALE:** Body length: 10.43 mm (holotype), 9.73–11.41 mm.

COLORATION: Blackish with pale yellow postclypeal medial longitudinal depression, interocular sulcus, scape basally, posterior margin of mesosternum, corium basally, mid and hind coxae anteriorly, trochanters, forefemur ventrally and subapically, mid and hind femora basally and subapically, fore- and mid tibiae medially, small subbasal and subapical band on hind tibia, anterior half of laterotergites, and sternites laterally and medially except intersternal suture.

Antennal segments VI and VII, labium, and rest of hind tibia brown. VESTITURE: Very sparse;

other features as in generic description. STRUCTURE: HEAD: Cylindrical in lateral view (Fig. 2.13J); longer than wide in dorsal view; ventrally flat; antecular region longer than postocular (Fig. 2.13K); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli small, located on shallow median tubercle (Fig. 2.13J), separated by less than diameter of ocellus; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about one-third longer than scape, not distinctly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI; pygophore process curved in lateral view (Fig. 2.20K), elongate in caudal view (Fig. 2.19H), not surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior ventral processes (Fig. 2.22G), posteriorly obscured by endosoma; area of endosomal struts–DPS fusion short, subquadrate; endosoma with sclerotized, hairlike, denticles on posterior margin (Fig. 2.21J, K).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the pale interocular sulcus.

DISTRIBUTION: Mahajanga and northern Antananarivo provinces (Map 2.13).

DISCUSSION: The pale yellow subapical band on the femora is sometimes absent.

HOLOTYPE: Male: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 14 Nov 2004–29 Nov 2004, M. Irwin, R. Harin'Hala (00045091) (CAS).

PARATYPES: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 31 Oct 2003–07 Nov 2003, M. Irwin, R. Harin'Hala, 1♂ (00045115) (CAS); 07 Nov 2003–15 Nov 2003, M. Irwin, R. Harin'Hala, 1♂ (00045662) (UCR); 15 Oct 2004–01 Nov 2004, M. Irwin, R. Harin'Hala, 2♂ (00007098, 00045080) (CAS), 2♂ (00045147, 00045265) (SU), 3♂ (00045426, 00045594, 00045670) (AMNH); 01 Nov 2004–14 Nov 2004, M. Irwin, R. Harin'Hala, 1♂ (00045051) (UCR), 5♂ (00045057, 00045125, 00045130, 00045140, 00045671) (CAS), 1♂ (00045702) (SU); 14 Nov 2004–29 Nov 2004, M. Irwin, R. Harin'Hala, 3♂ (00044920, 00045009, 00045065) (AMNH), 6♂ (00044920, 00045009, 00045065, 00045073, 00045081, 00045082) (BMNH), 14♂ (00045090, 00045098, 00045108, 00045113, 00045116, 00045121, 00045122, 00045135, 00045138, 00045149, 00045160, 00045162, 00045168, 00045169, 00045664, 00045665, 00045668) (CAS), 2♂ (00045679, 00045690) (SU), 2♂ (00045696, 00045697) (UCR); 29 Nov 2004–07 Dec 2004, M. Irwin, R. Harin'Hala, 2♂ (00044856, 00045030) (UCR), 3♂ (00045031, 00045040, 00045043) (USNM), 13♂ (00045046, 00045055, 00045058, 00045060, 00045066, 00045084, 00045103, 00045126, 00045146, 00045150, 00045151, 00045158, 00045164) (CAS), 6♂ (00045165, 00045659, 00045661, 00045666, 00045673, 00045703) (MNHN), 2♂ (00045724, 00045725) (SU); 22 Dec 2004–06 Jan 2005, M. Irwin, R. Harin'Hala, 3♂ (00007068, 00045145, 00045587) (USNM).

**Mahajanga:** Analamanitra Forest, 14 km NE of Misinjo, 16.13333°S 45.7°E, 20 m, 16 Nov 2007–20 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00007229) (CAS). Parc National Tsingy de Bemaraha, 3.4 km 93°E Bekopaka, Tombeau Vazimba, 19.14194°S 44.82805°E, 50 m, 06 Nov 2001–10 Nov 2001, Fisher et al., 1♂ (00045054) (CAS). Réserve forestière Beanka, 50.7 km E

Maintirano, 17.88027°S 44.46888°E, 140 m, 28 Oct 2009–31 Oct 2009, B.L. Fisher et al., 1♂ (00006910) (CAS); 1♂ (00007081) (UCR).

*Toxopus italaviana*, new species

Figs. 2.4, 2.7, 2.10, 2.13B, 2.15B; Map 2.12

DIAGNOSIS: Males are recognized among other *Toxopus* species by the moderately dense vestiture, elongate cylindrical head shape in lateral view, broad postclypeal longitudinal depression, relatively large ocelli, antenna inserted sublaterally on the head, and labial segment III ventrally convex. This species is similar to *T. griswoldi*, *T. insignis*, *T. fisheri*, *T. antsiranana*, and *T. toliara*, but is distinguished by the diagnostic characters mentioned. The slightly larger ocelli further distinguish *T. italaviana* from the other species, except *T. toliara*. The intersegmental sutures carinulate between sternites II and III and laterally between III–VII also distinguish males of *T. italaviana* from these species, with the exception of *T. griswoldi*.

DESCRIPTION: **MALE:** Body length: 10.58 mm (holotype), 10.25–10.63 mm.

COLORATION: Blackish with pale yellow mandibular and maxillary plates dorsally, postclypeus except laterally above eye, antennifer, scape basally, antennal segment VII basally, labial segment II ventroapically, labial segment III ventrobasally, anterolateral projections of pronotum, mesosternum anteromedially and posterolaterally, metasternum medially, corium basally, coxae, trochanters, forefemur basally and ventrally, mid femur basally and subapically, hind femur basally and apically, foretibia except basally, mid tibia except apically, hind tibia except medially and apically, laterotergite II, anterior half of dorsal laterotergites III–VII, ventral laterotergites III–VII, and abdominal sternites medially and laterally. Antennal segment VI and apex of VII, remainder of labium, and corium brown. VESTITURE: Moderately dense; other features as in generic description. STRUCTURE: HEAD (Fig. 2.13B): Cylindrical in lateral view

(Fig. 2.13J); longer than wide in dorsal view; ventrally with shallow anteromedial depression; anteocular region longer than postocular (Fig. 2.13K); synthlipsis about two times width of eye; interocular sulcus near hind margin of eye; postclypeus with shallow, broad medial longitudinal depression to middle of interocular area; postocular moderately broad in dorsal view; ocelli large, located on shallow median tubercle (Fig. 2.13J), separated by a distance less than or equal to diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about two-fifths longer than scape, slightly curved; labium stout; labial segment III ventrally convex (Fig. 2.13J). THORAX (Fig. 2.15B): Anterior pronotal lobe with small anterolateral projections; posterior pronotal lobe weakly transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges; lateral depressions of posterior pronotal lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II and III and laterally between III–VII; pygophore process curved in lateral view (Fig. 2.20K), elongate in caudal view (Fig. 2.19H), not surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior ventral processes (Fig. 2.22G), posteriorly obscured by endosoma; area of endosomal struts–DPS fusion short, subquadrate; endosoma with sclerotized, hairlike, denticles on posterior margin (Fig. 2.21J, K).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Italaviana, Madagascar.

DISTRIBUTION: Italaviana, near the Antananarivo-Fianarantsoa border (Map 2.12).

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Italaviana, 35 km SSE of Antsirabe, 20.17333°S 47.086°E, 1360 m, 27 Feb 2005–13 Mar 2005, M. Irwin, R. Harin'Hala (00006433) (CAS).



PARATYPES: **Madagascar: Fianarantsoa:** Italaviana, 35 km SSE of Antsirabe, 20.17333°S 47.086°E, 1360 m, 09 Dec 2002–19 Dec 2002, M. Irwin, R. Harin'Hala, 1♂ (00006965) (CAS); 10 Mar 2003–20 Mar 2003, M. Irwin, R. Harin'Hala, 1♂ (00006431) (CAS), 1♂ (00006966) (UCR); 15 Jul 2003–26 Jul 2003, M. Irwin, R. Harin'Hala, 4♂ (00006438, 00006439, 00006441, 00006445) (CAS), 2♂ (00006449, 00006450) (USNM); 16 Oct 2003–26 Oct 2003, M. Irwin, R. Harin'Hala, 2♂ (00006961, 00006963) (SU); 22 Jan 2004–04 Feb 2004, M. Irwin, R. Harin'Hala, 1♂ (00006962) (CAS); 24 Mar 2004–08 Apr 2004, M. Irwin, R. Harin'Hala, 1♂ (00006428) (CAS), 2♂ (00006429, 00006430) (BMNH); 27 Feb 2005–13 Mar 2005, M. Irwin, R. Harin'Hala, 1♂ (00006432) (UCR).

*Toxopus melobrunneus*, new species

Figs. 2.4, 2.7, 2.10; Map 2.11

DIAGNOSIS: Males are recognized among other species in *Toxopus* by a combination of the following characters: the large body size, coloration, cylindrical head shape in lateral view, smooth posterior pronotal lobe and lateral depressions, continuous pronotal transverse furrow, and intersegmental sutures carinulate between sternites II and III. Males are similar to males of *T. italaviana*, *T. griswoldi*, *T. insignis*, *T. fisheri*, *T. ambohitantely*, *T. antsiranana*, and *T. toliara*, but the continuous pronotal transverse suture and intersegmental sutures carinulate between sternites II and III distinguish *T. melobrunneus* from these species. The subtriangular pygophore process and medial endosomal sclerotization pattern further differentiate *T. melobrunneus* from these species, except from *T. ambohitantely*.

DESCRIPTION: **MALE:** Body length: 14.95 mm (holotype), 13.96–14.95 mm.

COLORATION: Blackish with pale yellow area anterior to interocular sulcus, lateral and posterolateral margins of ocellar tubercle, anterolaterally from interocular sulcus, antennal

segments V (except basally) and VI and VII, margins and longitudinal depression of metasternum, corium basally, femora basally, foretibia ventrally, anterior area of laterotergites III–VII, abdominal sternites II medially, anterolateral spots on sternites II–VII, and ventrolateral spots on sternites III–VII. VESTITURE: Dense; other features as in generic description.

STRUCTURE: HEAD: Cylindrical in lateral view (Fig. 2.13J); longer than wide in dorsal view; ventrally with deep medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about 1.5 times width of eye; interocular sulcus posterior to hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on distinct median tubercle (Fig. 2.13H), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium slender; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow continuous; lateral depressions of posterior pronotal lobe smooth; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II and III; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate ovoid (Fig. 2.21L); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the mostly dark brown legs.

DISTRIBUTION: Northern Fianarantsoa and southeastern Toliara provinces (Map 2.11).

DISCUSSION: Some specimens have more pale yellow on the ventral femoral and tibial surfaces and laterotergites, and can be reduced on the abdomen.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 30 Dec 2005–08 Jan 2006, M. Irwin, R. Harin'Hala (00007059) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Belle Vue at Talatakely, 21.2665°S 47.42016°E, 1020 m, 04 May 2003–16 May 2003, M. Irwin, R. Harin'Hala, 1♂ (00007037) (CAS), 1♂ (00007129) (UCR). Parc National Ranomafana, radio tower at forest edge, 21.251°S 47.40716°E, 1130 m, 07 May 2002–14 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00044926) (CAS), 1♂ (00044926) (USNM); 25 Jul 2002–03 Aug 2002, M. Irwin, R. Harin'Hala, 1♂ (00045238) (SU); 06 Jul 2003–17 Jul 2003, M. Irwin, R. Harin'Hala, 1♂ (00045291) (CAS); 12 Feb 2004–23 Feb 2004, M. Irwin, R. Harin'Hala, 1♂ (00007009) (CAS); 24 Jun 2004–08 Jul 2004, M. Irwin, R. Harin'Hala, 1♂ (00045275) (USNM); 12 Jul 2005–25 Jul 2005, M. Irwin, R. Harin'Hala, 1♂ (00044927) (USNM); 06 Nov 2005–13 Nov 2005, M. Irwin, R. Harin'Hala, 1♂ (00007127) (CAS); 07 Sep 2006–20 Sep 2006, M. Irwin, R. Harin'Hala, 2♂ (00007028, 00007196) (SU); 11 Nov 2006–18 Nov 2006, M. Irwin, R. Harin'Hala, 2♂ (00044935, 00044941) (UCR). **Toliara:** Forêt Ivohibe 55.0 km N Tolagnaro, 24.56888°S 47.20388°E, 200 m, 02 Dec 2006–04 Dec 2006, B.L. Fisher et al., 1♂ (00006728) (CAS). **Unknown:** 3♂ (00044925, 00044932, 00045243) (CAS).

*Toxopus miandritsara*, new species

Figs. 2.4, 2.7, 2.11; Map 2.10

DIAGNOSIS: Males are recognized among other species in this genus by a combination of the following characters: the coloration, ovoid head shape in lateral view, antecular region longer

than postocular, broad postocular, antenna inserted slightly sublaterally on the head, and smooth posterior pronotal lobe and lateral depressions. This species is similar to *T. simulans* and *T. toamasina*, but differs from these species by the light brown proximal part of the veins on the hemelytra, dark brown ventral abdominal surface, larger synthlipsis width, broad postocular, slight sublateral antennal insertion on the head, and smooth posterior pronotal lobe and lateral depressions. *Toxopus miandritsara* is further differentiated from *T. toamasina* by the intersegmental sutures carinulate between sternites II and III.

DESCRIPTION: **MALE:** Body length: 10.06 mm (holotype), 9.51–10.06 mm.

COLORATION: Blackish with pale yellow corium basally, proximal wing veins, mid and hind coxae basally, mid and hind trochanters, femora basally and ventrally, foretibia subbasally and apically, mid and hind tibiae medially and apically, anterior one-third of laterotergites, and lateral margin of abdominal sternites II–VI. Mandibular and maxillary plates, labium, remainder of corium, and tarsi brown. VESTITURE: Dense; other features as in generic description.

STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; ventrally with shallow medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli large, located on shallow median tubercle (Fig. 2.13J), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted slightly sublaterally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally convex (Fig. 2.13J).

THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions on pronotum smooth; hemelytron reaching or surpassing abdominal

apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II and III; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion hexagonal; endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Miandritsara Forest, Madagascar.

DISTRIBUTION: Miandritsara Forest in the Fianarantsoa province (Map 2.10).

DISCUSSION: The transverse suture between the meso- and metasterna and the ventrolateral spots on the anterior abdominal sternites are pale yellow in some specimens.

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Miandritsara Forest, 40 km S Ambositra, 20.79266°S 47.17566°E, 825 m, 26 Mar 2007–05 Apr 2007, M. Irwin, R. Harin'Hala (00044852) (CAS).

PARATYPES: **Madagascar: Fianarantsoa:** Miandritsara Forest, 40 km S Ambositra, 20.79266°S 47.17566°E, 825 m, 13 Nov 2006–22 Nov 2006, M. Irwin, R. Harin'Hala, 1♂ (00044848) (CAS); 22 Nov 2006–01 Dec 2006, M. Irwin, R. Harin'Hala, 1♂ (00044826) (CAS); 10 Dec 2006–19 Dec 2006, M. Irwin, R. Harin'Hala, 1♂ (00006148) (UCR); 28 Dec 2006–04 Jan 2007, M. Irwin, R. Harin'Hala, 2♂ (00044834, 00044970) (CAS), 1♂ (00044973) (SU); 31 Jan 2007–11 Feb 2007, M. Irwin, R. Harin'Hala, 1♂ (00006481) (UCR).

*Toxopus namoroka*, new species

Figs. 2.4, 2.7, 2.11; Map 2.13

DIAGNOSIS: Males are recognized among other *Toxopus* species by a combination of the following characters: the distinctive dark and pale coloration with red-brownish head and thoracic

markings, sparse vestiture, narrow postclypeal depression, smooth posterior pronotal lobe, and intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII. Males are similar to males of *T. steineri*, *T. basalis*, *T. vazimba*, and *T. brucei*, but differs from these species by the very distinctive red-brown head and thorax, narrow postclypeal depression, eye about one-fourth of the head length, and intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII.

DESCRIPTION: **MALE:** Body length: 11.44 mm (holotype), 11.44–11.56 mm.

COLORATION: Blackish with reddish-brown head, labium, scape, pedicel, antennal segment V, anterior pronotal lobe except pronotal longitudinal furrow, dorsal half of anterior propleural lobe, anterior margin of mesopleuron, scutellum, and legs (except coxae posteriorly and femora basally). Antennal segments VI and VII, corium basally, anterior half of laterotergites, abdominal sternites III–VI medially except at margins, and anterolateral spots on sternites pale yellow.

VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD:

Subquadrate in lateral view (Fig. 2.13K); longer than wide in dorsal view; ventrally with deep medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with shallow, narrow medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on shallow median tubercle (Fig. 2.13K), separated by less than diameter of ocellus; eye about one-fourth of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-third longer than scape, slightly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe smooth; wings spread out in specimens. ABDOMEN:

Apex rounded or sinuate; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts not reaching posterior margin of DPS (Fig. 2.21L), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate ovoid (Fig. 2.21L); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the type locality, Tsingy de Namoroka National Park, Madagascar.

DISTRIBUTION: Tsingy de Namoroka National Park in the Mahajanga province (Map 2.13).

DISCUSSION: The medial pale yellow spots on the abdominal sternites may be divided by a brown medial longitudinal stripe, and the pale yellow spot on sternite VI may be reduced.

HOLOTYPE: Male: **Madagascar: Mahajanga:** Parc National de Namoroka, 16.9 km 317° NW Vilanandro, 16.40666°S 45.31°E, 100 m, 12 Nov 2002–16 Nov 2002, Fisher et al. (00045251) (CAS).

PARATYPE: **Madagascar: Mahajanga:** Parc National de Namoroka, 16.9 km 317° NW Vilanandro, 16.40666°S 45.31°E, 100 m, 12 Nov 2002–16 Nov 2002, Fisher et al., 1♂ (00006477) (CAS).

*Toxopus pallidus*, new species

Figs. 2.4, 2.7, 2.11; Map 2.12

DIAGNOSIS: Males are recognized among other species in *Toxopus* by the smaller body size, coloration, ovoid head shape in lateral view, antecular regions as long as postocular, labial segment III ventrally straight, and BPE as long as the basal plate. This species is similar to *T. farafangana*, from which it differs by the the pale yellow posterior propleural lobe and thoracic

sterna, predominately pale to pale yellow ventral abdominal surface, interocular sulcus posterior to the hind margin of the eye, labial segment III ventrally straight, striated posterior pronotal lobe, and BPE as long as the basal plate. This species is also similar to *T. simulans*, but is distinguished by the slightly smaller body size, ventral head depression restricted to the anteocular region, anteocular as long as postocular, broader postocular, labial segment III ventrally straight, and intersegmental sutures carinulate between sternites II and III.

DESCRIPTION: **MALE:** Body length: 9.10 mm (holotype), 8.85–9.10 mm. COLORATION: Blackish with pale yellow scape basally, antennal segment V (except basally) and VI and VII; posterior propleural lobe, ventral half of mesopleuron, metapleuron, thoracic sterna, corium basally, coxae, trochanters, fore- and mid femora ventrally and basally, hind femur basally, foretibia ventrally, mid and hind tibiae except subbasally and apically, anterior margin of dorsal laterotergites, ventral laterotergites, abdominal sternites except lateral medial spots, and margins of pygophore. Labium and posterior pronotal lobe brown. VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; ventrally with shallow anteromedial depression; anteocular region as long as postocular (Fig. 2.13J); postclypeus with deep, narrow medial longitudinal depression to interocular sulcus; synthlipsis about 1.5 times width of eye; interocular sulcus posterior to hind margin of eye (Fig. 2.13B); postocular broad in dorsal view (Fig. 2.13B); ocelli large, located on shallow median tubercle (Fig. 2.13J), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal



lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II and III; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), ventrally obscured by endosoma; area of endosomal struts–DPS fusion ovate; endosoma missing.

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the pale body coloration.

DISTRIBUTION: Ranomafana National Park in the Fianarantsoa province (Map 2.12).

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** Parc National de Ranomafana, Vatoharanana River, 4.1 km 231° SW Ranomafana, 21.29°S 47.43333°E, 1100 m, 27 Mar 2003–31 Mar 2003, Fisher et al. (00045432) (CAS).

PARATYPE: **Madagascar: Fianarantsoa:** Parc National Ranomafana, Vohiparara, at broken bridge, 21.22616°S 47.36983°E, 1110 m, 08 Apr 2002–15 Apr 2002, M. Irwin, R. Harin’Hala, 1♂ (00007192) (CAS).

*Toxopus parkeri*, new species

Figs. 2.4, 2.7, 2.11, 2.13K; Map 2.11

DIAGNOSIS: Males are recognized among other species in *Toxopus* by a combination of the following characters: the coloration, dense vestiture, subquadrate head in lateral view that is as long as wide in dorsal view, and ventrally depressed head surface, synthlipsis two times width of an eye, interocular sulcus anterior to the hind margin of the eye, intersegmental sutures carinulate between sternites II–VI; and BPE as long as basal plate. Males are very similar to *T. basalis*, *T. tibialis*, and *T. brucei*, but differ by the smaller body size, larger synthlipsis width, and interocular sulcus located anterior to the hind margin of the eye. The smaller eye size relative to the head

length, the eyes not reaching the dorsal and ventral margins, larger synthlipsis width, and intersegmental sutures carinulate between sternites II–VI separate *T. parkeri* from *T. basalis*. The striated lateral depressions on the posterior pronotal lobe and intersegmental sutures carinulate between sternites II–VI differentiate *T. parkeri* from *T. brucei*.

DESCRIPTION: **MALE:** Body length: 12.22 mm (holotype), 11.52–13.34 mm.

COLORATION: Blackish with pale yellow clypeus, postclypeal medial longitudinal depression, bases of scape and pedicel, antennal segment V (except basally) and VI and VII, meso- and metasterna, corium basally, forecoxa posteriorly, mid and hind coxae except laterally, trochanters, forefemur except dorsally, mid and hind femora basally, foretibia except basally, mid and hind tibiae medially, tarsi, anterior half of laterotergites, and abdominal sternites (except pygophore) medially and laterally. Scape and pedicel medially, antennal segment IV and base of V, and postclypeal medial longitudinal depression brown. VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD (Fig. 2.13K): Subquadrate in lateral view; as long as wide in dorsal view; ventrally with shallow medial depression; anteocular region longer than postocular; postclypeus with shallow, broad medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus anterior to hind margin of eye; postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on shallow median tubercle, separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally straight. THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded or sinuate; intersegmental sutures carinulate between

sternites II–VI; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts not reaching posterior margin of DPS (Fig. 2.21L), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate ovoid (Fig. 2.21L); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after Frank Parker, who is one of the main CAS collectors for the taxonomic inventory of Madagascar.

DISTRIBUTION: Antananarivo, Mahajanga, and Toliara provinces (Map 2.11).

DISCUSSION: The antennae and postclypeus vary with respect to the extent of the dark brown coloration.

HOLOTYPE: Male: **Madagascar: Toliara:** Parc National d'Andohahela, Ihazofotsy Parcelle III, 24.83083°S 46.53616°E, 80 m, 16 Dec 2002–26 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala (00044875) (CAS).

PARATYPES: **Madagascar: Antananarivo:** 46 km NE of Ankazobe: Ambohitantely, 18.198°S 47.2815°E, 700 m, 20 Jan 2004–01 Feb 2004, M. Irwin, R. Harin'Hala, 1♂ (00006973) (UCR), 1♂ (00006974) (CAS). **Mahajanga:** Maintirano District Asondrodava dry forest, 15 km N of Maintirano, 17.96533°S 44.0355°E, 61 m, 03 Dec 2007–10 Dec 2007, M. Irwin, R. Harin'Hala, 1♂ (00007075) (CAS). **Toliara:** Parc National d'Andohahela, Ihazofotsy Parcelle III, 24.83083°S 46.53616°E, 80 m, 16 Dec 2002–26 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00044884) (CAS); 24 Feb 2003–06 Mar 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00007091, 00044897) (SU), 2♂ (00045204, 00045249) (CAS), 1♂ (00045381) (UCR); 21 Apr 2003–28 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00045052, 00045377) (USNM). **Unknown:** 1♂ (00006976) (CAS).

*Toxopus politus* Bergroth, 1905

Map 2.13

*Toxopus politus* Bergroth, 1905: 375

DISTRIBUTION: According to the original description, the type specimen was collected from Nosy Be Island off the northwest coast of Madagascar (Map 2.13).

DISCUSSION: We do not redescribe this species, since the only known specimen could not be located, and none of the specimens examined by us correspond to this species. Many of Bergroth's type specimens are deposited in the Zoological Museum, University of Helsinki, but Jansson & Coscarón (1989: appendix I) reported the female holotype of *T. politus* to be missing (or away on loan) from that collection. The specimen is still absent (Larry Huldén, personal commun.). Correspondence with other major European collections did not offer any further leads, and we speculate that the type specimen may be lost. The three female specimens of *Toxopus* examined during our study do not appear to be conspecific with *T. politus* based on differences in color patterns, ventral abdominal structure (segment VI medially longer than V in *T. politus*; subequal in female specimens), and locality information. A potential neotype designation will therefore have to await availability of additional *Toxopus* specimens.

*Toxopus signoretii* (Reuter), 1887, new combination

Fig. 2.12; Map 2.13

*Cleptria signoretii* Reuter, 1887: 107

REVISED DIAGNOSIS: Males are recognized among other *Toxopus* species by a combination of the following characters: the coloration, large body size, subquadrate head shape in lateral view that is as long as it is wide in dorsal view, eye reaching dorsal head surface, and synthlipsis about the width of an eye. This species is very similar to *T. tibialis*, *T. basalis*, and *T. parkeri*, but

differs by the slightly larger body size, smaller synthlipsis width, and intersegmental sutures carinulate between sternites entirely between II–III and laterally between III–VI. *Toxopus signoretii* can be further differentiated from *T. parkeri* by the sparse vestiture, interocular sulcus near the hind margin of the eye, and eye about half width of the head lengths and not reaching the dorsal head surface. The interocular sulcus near the hind margin of the eye and the eye not reaching the dorsal but reaching the ventral head surface further distinguish *T. signoretii* from *T. basalis*. This species differs from *T. tibialis* by the sparse vestiture, ventral medial longitudinal depression on the head extending onto postocular surface, and eye about half width of the head lengths and not reaching the dorsal head surface.

REDESCRIPTION: **MALE:** Body length: 12.20 mm (holotype). COLORATION: Brownish with pale yellow postclypeal medial longitudinal depression, base of scape, corium basally, trochanters, forefemur basally and ventrally, mid and hind femora except dorsomedially, foretibia except basally, mid and hind tibiae medially, mid tarsi (fore- and hind missing), anterior half of dorsal laterotergites, and abdominal sternites (except pygophore) paramedially and laterally.

VESTITURE: Sparse; other features as in generic description. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13K); about as long as wide in dorsal view; ventrally with longitudinal depression; antecular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); ocelli large, located on distinct median tubercle (Fig. 2.13H), separated by less than diameter of ocellus; postocular narrow in dorsal view (Fig. 2.13A); eye about half of head length, reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about as long as scape, curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); pronotal transverse furrow divided by paramedian

longitudinal ridges (Fig. 2.15B); posterior pronotal lobe smooth; lateral depressions of posterior pronotal lobe smooth; hemelytron nearly reaching abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites entirely between II–III and laterally between III–VI.

DISTRIBUTION: The holotype and one juvenile were collected from Nosy Be Island off the northwest coast of Madagascar (Map 2.13). One specimen was collected from Vohemar, which is on the northeast coast of Madagascar.

DISCUSSION: The redescription is based on habitus images and Reuter's (1887) original description of the holotype. The original description indicates that one immature specimen was collected (coll. Stumpff) in "Loucoubé," which is currently known as the Natural Reserve of Lokobe on Nosy Be Island. Reuter described this species in the genus *Cleptria* Stål, 1859, but did not provide arguments for assigning it to that genus. This species is transferred to *Toxopus* based on the 7-segmented antenna, head being longer than or as long as the width, subequal lengths of labial segments II and III, and apterous females, which are features differentiating *Toxopus* from *Cleptria*.

HOLOTYPE: Male: **Madagascar: Antsiranana:** Nosy Be, 13.31502°S 48.25926°E, 155 m (NMW).

PARATYPE: **Madagascar: Antsiranana:** Nosy Be, 13.31502°S 48.25926°E, 155 m, 1 juv. (MNHN).

OTHER MATERIAL EXAMINED: **Madagascar: Antsiranana:** Vohémar, 13.37028°S 50.00337°E, 6 m, 1♂ (MNHN). **Unknown:** 1♂ (NMW).

*Toxopus simulans*, new species

Figs. 2.4, 2.7, 2.11; Map 2.13

DIAGNOSIS: Males recognized among other *Toxopus* species by the coloration, ovoid head shape in lateral view that is longer than wide in dorsal view, anteocular region longer than postocular, narrow postocular, and labial segment III ventrally convex. This species is most similar to *T. pallidus*, but the slightly larger body size, ventral medial depression extending along the entire head surface, longer anteocular region relative to the postocular, interocular sulcus located near the hind margin of the eye, narrower postocular in dorsal view, and ventrally convex labial segment III distinguishes this species from *T. pallidus*.

DESCRIPTION: **MALE:** Body length: 11.53 mm (holotype), 10.89–11.62 mm.

COLORATION: Blackish with pale yellow antennal segments V (except basally) and VI and VII, meso- and metasterna, coxae, trochanters, forefemur ventrally, mid and hind femora except large subapical band, mid and hind tibiae medially, tarsi, anterior one-third of dorsal laterotergites, ventral laterotergites, and abdominal sternites (except pygophore) medially and laterally.

Mandibular and maxillary plates, postclypeal medial longitudinal depression, labium, posterior pronotal lobe, corium, and lateral longitudinal band across sternites varying shades of brown.

VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); longer than wide in dorsal view; ventrally with deep medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, narrow medial longitudinal depression to interocular sulcus; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on distinct median tubercle (Fig. 2.13H), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium

stout; labial segment III ventrally convex (Fig. 2.13J). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe weakly transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe weakly transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinate between sternites II and III; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion short, subquadrate; endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a participle in the nominative case and is named for the similar body coloration with *Toxopus pallidus*.

DISTRIBUTION: Toamasina and northern Antsiranana provinces (Map 2.13).

HOLOTYPE: Male: **Madagascar: Antsiranana:** Parc National Montagne d’Ambre, 12.51444°S 49.18138°E, 960 m, 04 Mar 2001–19 Mar 2001, M. Irwin, R. Harin’Hala (00045508) (CAS).

PARATYPES: **Madagascar: Antsiranana:** Parc National Montagne d’Ambre, 12.52027°S 49.17916°E, 1125 m, 14 May 2001–30 May 2001, M. Irwin, R. Harin’Hala, 1♂ (00007226) (CAS). Sakalava Beach, dwarf littoral forest, 12.26277°S 49.3975°E, 10 m, 20 Aug 2001–28 Aug 2001, M. Irwin, R. Harin’Hala, 1♂ (00045363) (CAS). 7 km N of Joffreville, 12.33333°S 49.25°E, 360 m, 13 May 2001–16 May 2001, M. Irwin, R. Harin’Hala, 1♂ (00045123) (SU).

**Toamasina:** Botanic Garden near entrance to Andasibe National Park, 18.92633°S 48.40783°E, 1025 m, 01 Nov 2001–07 Nov 2001, M. Irwin, R. Harin’Hala, 1♂ (00007048) (UCR).

**Unknown:** 1♂ (00045124) (CAS).



*Toxopus steineri*, new species

Figs. 2.4, 2.7, 2.11; Map 2.12

DIAGNOSIS: Males recognized among other species in *Toxopus* by the coloration, dense vestiture, narrow postclypeal depression, ventrally straight labial segment III, smooth posterior pronotal lobe, and intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI. This species is similar to *T. namoroka*, *T. basalis*, *T. vazimba*, and *T. brucei*, but the dark brown to blackish ventral abdominal surface, narrow postclypeal depression, and ventrally straight labial segment III distinguishes *T. steineri* from these other species.

DESCRIPTION: **MALE:** Body length: 11.14 mm (holotype), 11.00–11.59 mm.

COLORATION: Blackish with pale yellow bases of scape and pedicel, antennal segment V distally and segments VI and VII, posterolateral margins of pronotum, lateral areas of transverse sulcus between meso- and metasterna, coxae, trochanters, base forefemur, forefemur subapically and ventrally, mid and hind femora basally and apically, foretibia medially, mid tibia except subbasally and apically, hind tibia basally, foretarsi, mid and hind tarsi basally, corium basally, anterior one-third to half of laterotergites, and abdominal sternites laterally except posterior two-thirds of segments VI and VII. Maxillary plates dorsally, postclypeal longitudinal depression, paramedially along interocular sulcus, basal half of antennal segment V, labium, mid and hind tarsi apically, and anterior margin of corium (except basally) dark yellow to brown.

VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13K); longer than wide in dorsal view; ventrally with deep longitudinal depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, narrow medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on shallow median tubercle (Fig. 2.13K), separated by less than

diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about two-fifths longer than scape, slightly curved; labium stout; labial segment III ventrally convex (Fig. 2.13J). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe weakly transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), ventrally obscured by semiextended endosoma; area of endosomal struts–DPS fusion subquadrate; endosoma sclerotized medially (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in the genitive case and is named after Ranomafana, Madagascar, which is near the type locality.

DISTRIBUTION: Known from a single locality west of Ranomafana in the Fianarantsoa province (Map 2.12).

HOLOTYPE: Male: **Madagascar: Fianarantsoa:** 7 km W Ranomafana, 21.25923°S 47.38664°E, 900 m, 23 Feb 1990–28 Feb 1990, W.E. Steiner (00026334) (USNM).

PARATYPES: **Madagascar: Fianarantsoa:** 7 km W Ranomafana, 21.25923°S 47.38664°E, 900 m, 23 Feb 1990–28 Feb 1990, W.E. Steiner, 3♂ (00026332, 00026333, 00026335) (USNM).

*Toxopus tibialis*, new species

Figs. 2.4, 2.7, 2.11; Map 2.11

DIAGNOSIS: Males are recognized among other species in *Toxopus* by a combination of the following characters: the relatively larger body size, moderately dense vestiture, head as long as it

is wide in dorsal view, ventral depression on the head restricted to the anteocular region, eye about one-third of head length, eye not reaching dorsal or ventral head margins, and intersegmental sutures carinulate between sternites II–VI. Males are most similar to *T. basalis* and *T. parkeri*, from which it differs by the larger body size, ventral anteromedial depression on the head, and interocular sulcus located near the hind margin of the eye. The denser vestiture, smaller eye size that does not reach the dorsal and ventral head margins, and intersegmental sutures carinulate between sternites II–VI further differentiates *T. tibialis* from *T. basalis*. A smaller synthlipsis width further distinguished *T. tibialis* from *T. parkeri*.

DESCRIPTION: **MALE:** Body length: 14.52 mm (holotype), 13.89–15.40 mm.

COLORATION: Blackish with pale yellow clypeus, dorsal margins of mandibular and maxillary plates, postclypeal medial longitudinal depression, antennifer, scape basally, antennal segments V–VII, mesosternum except medial longitudinal depression, metasternum medially, corium basally, mid and hind coxae, trochanters, base of forefemur, for femur ventrally and subbasally, mid and hind femora except small apical and large medial bands, tibiae medially, tarsi, anterior half of laterotergites, and abdominal sternites (except pygophore) medially, and lateral margin of sternites. Labium brown. VESTITURE: Moderately dense; other features as in generic description. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13K); as long as wide in dorsal view; ventrally with deep anteromedial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about 1.5 times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on distinct median tubercle (Fig. 2.13H), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III

ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with very small anterolateral projections (Fig. 2.15B); posterior pronotal lobe weakly transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe transversely striated; hemelytron reaching or surpassing abdominal apex.

ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–VI; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts not reaching posterior margin of DPS (Fig. 2.21L), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate ovoid (Fig. 2.21L); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is an adjective in the nominative case and is named for the pale coloration of the tibia.

DISTRIBUTION: Antsiranana and Mahajanga provinces in Madagascar (Map 2.11).

DISCUSSION: The clypeus, mandibular and maxillary plates, and postclypeus are slightly darker in some specimens.

HOLOTYPE: Male: Male: **Madagascar: Antsiranana:** Forêt Ambanitaza, 26.1 km 347° Antalaha, 14.67944°S 50.18361°E, 240 m, 26 Nov 2004, B.L. Fisher (00045457) (CAS).

PARATYPES: **Madagascar: Antsiranana:** 7 km N of Joffreville, 12.33333°S 49.25°E, 360 m, 22 Jan 2001–26 Jan 2001, M.E. Irwin, E.L. Schlinger, R. Harin'Hala, 1♂ (00044821) (CAS). Forêt d'Ampondrabe, 26.3 km 10° NNE Daraina, 12.97°S 49.7°E, 175 m, 10 Dec 2003, B.L. Fisher, 1♂ (00045701) (CAS). Marojejy National Park, 5 km W Manantenina village, Camp Mantella, 14.43816°S 49.774°E, 490 m, 11 Feb 2005–18 Feb 2005, M. Irwin, R. Harin'Hala, 1♂ (00045492) (SU). **Mahajanga:** Réserve forestière Beanka, 50.2 km E Maintirano, 18.02638°S 44.05055°E, 250 m, 19 Oct 2009–26 Oct 2009, B.L. Fisher et al., 1♂ (00044854) (UCR).

**Unknown:** 1♂ (00045637) (CAS).

*Toxopus toamasina*, new species

Figs. 2.4, 2.7, 2.11; Map 2.10

DIAGNOSIS: Males are recognized among other species in this genus by the coloration, ovoid head shape in lateral view that is as long as it is wide in dorsal view, synthlipsis about as wide as an eye, narrow postocular, eye about half the length of the head in lateral view and reaching the dorsal and ventral head margins, and intersegmental sutures carinulate between sternites II and III and laterally between III–V. This species is similar to *T. simulans*, but differs by the generally darker coloration, apex of corium light brown, head as long as it is wide, smaller synthlipsis width, eye one-half of the head length in lateral view, eye height relative to dorsal and ventral head margins, labial segment III ventrally straight, and intersegmental sutures carinulate between sternites II and III and laterally between III–V.

DESCRIPTION: **MALE:** Body length: 12.04 mm (holotype), 11.22–12.41 mm.

COLORATION: Blackish with pale yellow margins of ocellar tubercle except posteromedial area, antennal segments V (except basally) and VI and VII, mid and hind coxae, trochanters, forefemur except dorsally, mid and hind femora except subapically, mid and hind tibiae except subbasally and apically, tarsi, anterior half of dorsal laterotergites, ventral laterotergites, paramedial spots on abdominal sternites II–VI, and lateral margin of abdomen. Labium, posterolateral pronotal lobe, and corium apically brown. VESTITURE: Moderately dense; other features as in generic description. STRUCTURE: HEAD: Ovoid in lateral view (Fig. 2.13F); as long as wide in dorsal view; ventrally with deep medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on distinct median tubercle (Fig. 2.13H), separated by less than diameter of ocellus; eye about half of head length,

reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe transversely striated; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe transversely striated; hemelytron surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II and III and laterally between III–V; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore; BPE as long as basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21J, K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21K); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the Toamasina province in Madagascar, where all known specimens were collected.

DISTRIBUTION: Eastern region the Toamasina province (Map 2.10).

DISCUSSION: In some specimens, the scutellar processes and abdominal sternite II are pale yellow, the pale yellow ventrolateral abdominal spots vary in size.

HOLOTYPE: Male: **Madagascar: Toamasina:** Parcelle E3, Tampolo, 17.28333°S 49.43333°E, 10 m, 14 Apr 2004, Malagasy Ant Team (00006478) (CAS).

PARATYPES: **Madagascar: Toamasina:** Mobot Site, Analalava 7 km SW of Foulpointe, 17.69333°S 49.46027°E, 18 m, 03 Jan 2008–11 Jan 2008, M. Irwin, R. Harin'Hala, 2♂ (00007095, 00044942) (CAS); 11 Jan 2008–18 Jan 2008, M. Irwin, R. Harin'Hala, 1♂ (00006394) (CAS); 22 Feb 2008–29 Feb 2008, M. Irwin, R. Harin'Hala, 1♂ (00045546) (SU); 04 Apr 2008–11 Apr 2008, M. Irwin, R. Harin'Hala, 1♂ (00007056) (SU); 18 Apr 2008–25 Apr 2008, M. Irwin, R. Harin'Hala, 1♂ (00007078) (UCR); 25 Apr 2008–02 May 2008, M. Irwin, R.

Harin'Hala, 1♂ (00045537) (UCR). Parcelle E3, Tampolo, 17.28333°S 49.43333°E, 10 m, 14 Apr 2004, Malagasy Ant Team, 1♂ (00007218) (CAS).

*Toxopus toliara*, new species

Figs. 2.4, 2.7, 2.11, 2.21K; Map 2.10

DIAGNOSIS: Males are recognized among other *Toxopus* species by the coloration, moderately dense vestiture, elongate cylindrical head in lateral view, relatively large ocelli, antenna inserted sublaterally on the head, ventrally straight labial segment III, and smooth posterior pronotal lobe. This species is similar to *T. italaviana*, *T. griswoldi*, *T. insignis*, *T. fisheri*, and *T. antsiranana*, but is distinguished by the diagnostic features mentioned.

DESCRIPTION: **MALE:** Body length: 11.02 mm (holotype), 10.39–12.54 mm.

COLORATION: Blackish with pale yellow maxillary plates dorsally; antennifers postclypeal medial longitudinal depression, paramedian longitudinal stripes on postclypeus anterior of interocular sulcus that extend laterally along interocular sulcus, labium, scape basally, antennal segment VII, collar, pronotum anterolaterally, anterolateral margin of posterior pronotal lobe, meso- and metasterna medially except transverse sulcus, corium basally, coxae, trochanters, fore- and mid femora basally and subapically, hind femur except medially, foretibia basally and medially except lateral spots, base of mid tibia, mid tibia subbasally and medially, hind tibia basally and subapically, tarsi, anterior half of laterotergites, and abdominal sternites laterally and medially, except intersternal sutures. Antennal segments V and VI and remainder of corium brown. VESTITURE: Moderately dense; other features as in generic description. STRUCTURE: HEAD: Cylindrical in lateral view (Fig. 2.13J); longer than wide in dorsal view; ventrally with shallow anteromedial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with shallow, narrow medial longitudinal depression to middle of interocular area

(Fig. 2.13B); synthlipsis about two times width of eye; interocular sulcus near hind margin of eye (Fig. 2.13B); postocular moderately broad in dorsal view; ocelli large, located on shallow median tubercle (Fig. 2.13J), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted sublaterally on head; pedicel about two-fifths longer than scape, slightly curved; labium slender; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with small anterolateral projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe weakly to strongly transversely striated; hemelytron reaching or surpassing abdominal apex. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–IV and laterally between IV–VI; pygophore process curved in lateral view (Fig. 2.20K), elongate in caudal view (Fig. 2.19H), surpassing posterior margin of pygophore; BPE shorter than basal plate; endosomal struts reaching posterior margin of DPS (Fig. 2.21K), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate subquadrate (Fig. 2.21K); endosoma with sclerotized, hairlike, denticles on posterior margin (Fig. 2.21K).

ETYMOLOGY: The species epithet is a noun in apposition and is named after the Toliara province in Madagascar, where a majority of the specimens examined were collected.

DISTRIBUTION: Fianarantsoa, Mahajanga, and Toliara provinces (Map 2.10).

DISCUSSION: The coloration is highly variable with the pale yellow markings more extensive or reduced on the head, pronotum, scutellar processes, and legs.

HOLOTYPE: Male: **Madagascar: Toliara:** Berenty Special Reserve, 8 km NW Amboasary, 25.021°S 46.3055°E, 35 m, 28 Feb 2004–24 Mar 2004, M. Irwin, F. Parker, R. Harin'Hala (00006598) (CAS).



PARATYPES: **Madagascar: Fianarantsoa:** 50 km S of Farafangana, Mahabo Mananivo, Ampitavananima Forest, 23.12983°S 47.717°E, 34 m, 06 Sep 2007–13 Sep 2007, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006614) (CAS). **Mahajanga:** Besalampy District, Marofotra dry forest, 17 km W of Besalampy, 16.72166°S 44.42366°E, 52 m, 19 Nov 2007–26 Nov 2007, M. Irwin, R. Harin'Hala, 1♂ (00044907) (CAS). **Toliara:** Andohahela Natl Park, Tsimelahy, Parcelle II, 24.93683°S 46.62666°E, 180 m, 27 Dec 2002–06 Jan 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006639, 00006643) (CAS); 05 Feb 2003–15 Feb 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006674) (CAS); 15 Feb 2003–26 Feb 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006628) (BMNH); 18 Mar 2003–28 Mar 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006626) (CAS); 29 Jun 2003–10 Jul 2003, M. Irwin, F. Parker, R. Harin'Hala, 6♂ (00006618, 00006627, 00006629, 00006630, 00006632, 00006634) (USNM); 17 Aug 2003–24 Aug 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006676) (CAS); 01 Oct 2003–11 Oct 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006647, 00006669) (SU); 30 Oct 2003–09 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006660, 00006666) (CAS); 30 Nov 2003–11 Dec 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006673, 00006675) (UCR); 21 Dec 2003–23 Dec 2003, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006579–00006581) (AMNH); 15 Jan 2004–28 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006609–00006611) (CAS); 28 Jan 2004–12 Feb 2004, M. Irwin, F. Parker, R. Harin'Hala, 6♂ (00006625, 00006633, 00006636, 00006638, 00006640, 00006644) (BMNH); 06 Mar 2004–18 Mar 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006617) (UCR). Berenty Special Reserve, 8 km NW Amboasary, 25.00666°S 46.30333°E, 85 m, 25 Oct 2002–26 Oct 2002, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006649, 00006663) (AMNH); 02 Nov 2002–09 Nov 2002, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006667, 00006681) (CAS); 30 Nov 2002–07 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006622) (UCR); 07 Dec 2002–14 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂

(00006665) (SU); 14 Dec 2002–16 Dec 2002, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006615) (CAS); 27 Dec 2002–07 Jan 2003, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006436, 00006442, 00006446) (USNM), 11♂ (00006448, 00006451, 00006601–00006608, 00006664) (CAS); 26 Jan 2003–05 Feb 2003, M. Irwin, F. Parker, R. Harin'Hala, 7♂ (00006695–00006701) (CAS); 24 Mar 2003–03 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 6♂ (00006566–00006570, 00045618) (CAS); 14 Apr 2003–24 Apr 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00005357, 00006624) (AMNH); 24 Apr 2003–03 May 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006599, 00006600) (MNHN); 14 May 2003–25 May 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006595, 00006596) (CAS); 25 May 2003–04 Jun 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006679) (UCR); 04 Jun 2003–10 Jun 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006646, 00006678) (CAS); 08 Nov 2003–15 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006677) (CAS); 22 Nov 2003–30 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006525–00006527) (MNHN); 14 Dec 2003–21 Dec 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006577, 00006578) (CAS); 21 Dec 2003–02 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006592, 00006593) (CAS); 02 Jan 2004–13 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006558) (SU); 13 Jan 2004–23 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006561, 00006562) (CAS); 04 Feb 2004–15 Feb 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006571, 00006572) (CAS); 15 Feb 2004–02 Mar 2004, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006619, 00006621, 00006680) (BMNH); 02 Mar 2004–11 Mar 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006559, 00006560) (CAS); 05 Apr 2004–15 Apr 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006563, 00006564) (MNHN); 30 Apr 2004–01 May 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006443) (SU); 14 May 2004–27 May 2004, M. Irwin, F. Parker, R. Harin'Hala, 4♂ (00006582–00006585) (CAS); 20 Jun 2004–30 Jun 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006657) (CAS); 30 Jun 2004–11 Jul 2004, M. Irwin, F.

Parker, R. Harin'Hala, 2♂ (00006691, 00006692) (CAS); 25 Jul 2004–07 Aug 2004, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006574–00006576) (MNHN); 22 Aug 2004–01 Sep 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006565) (SU); 01 Sep 2004–08 Sep 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006591) (UCR). Berenty Special Reserve, 8 km NW Amboasary, 25.021°S 46.3055°E, 35 m, 10 Jun 2003–19 Jun 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006524) (CAS); 22 Nov 2003–30 Nov 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006694) (UCR); 07 Dec 2003–14 Dec 2003, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006689, 00006690) (AMNH); 21 Dec 2003–01 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006682, 00006683) (CAS); 02 Jan 2004–13 Jan 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006686) (SU); 15 Feb 2004–28 Feb 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006620) (CAS); 28 Feb 2004–24 Mar 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006597) (CAS); 24 Mar 2004–04 Apr 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006573, 00006685) (USNM); 04 Apr 2004–15 Apr 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006594) (CAS); 29 Apr 2004–11 Jun 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006612, 00006613) (BMNH); 23 Jun 2004–04 Jul 2004, M. Irwin, F. Parker, R. Harin'Hala, 3♂ (00006468, 00006650, 00006684) (AMNH); 04 Jul 2004–18 Jul 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006623) (MNHN); 31 Jul 2004–15 Aug 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006693) (CAS); 29 Aug 2004–08 Sep 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006589) (SU); 08 Sep 2004–21 Sep 2004, M. Irwin, F. Parker, R. Harin'Hala, 2♂ (00006687, 00006688) (CAS); 21 Sep 2004–07 Oct 2004, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006590) (UCR). Beza Mahafaly Reserve, Parcelle II near Bellevue, 23.68983°S 44.5755°E, 180 m, 16 Jan 2002–18 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00006637) (CAS); 18 Jan 2002–25 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00044812) (UCR). Beza Mahafaly Reserve, Parcelle I near research station, 23.6865°S 44.591°E, 165 m, 15 Oct 2001–10 Nov 2001, M.E.

Irwin, F.D. Parker, R. Harin'Hala, 1♂ (00045584) (CAS); 10 Nov 2001–21 Nov 2001, R. Harin'Hala, 1♂ (00006586) (UCR); 21 Nov 2001–28 Nov 2001, R. Harin'Hala, 2♂ (00006099, 00006118) (SU); 28 Nov 2001–04 Dec 2001, R. Harin'Hala, 1♂ (00006653) (CAS); 04 Dec 2001–11 Dec 2001, R. Harin'Hala, 2♂ (00006645, 00006670) (BMNH); 18 Dec 2001–25 Dec 2001, R. Harin'Hala, 2♂ (00006119, 00006437) (CAS), 2♂ (00006635, 00006642) (MNHN); 16 Jan 2002–18 Jan 2002, R. Harin'Hala, 2♂ (00006654, 00006659) (USNM); 01 Feb 2002–08 Feb 2002, R. Harin'Hala, 1♂ (00006668) (USNM); 10 Apr 2002–29 Apr 2002, R. Harin'Hala, 2♂ (00006658, 00006662) (CAS); 08 Jun 2002–18 Jun 2002, R. Harin'Hala, 2♂ (00006440, 00006444) (AMNH); 28 Jun 2002–07 Jul 2002, R. Harin'Hala, 1♂ (00006434) (UCR); 07 Jul 2002–18 Jul 2002, R. Harin'Hala, 1♂ (00006652) (CAS); 09 Sep 2002–20 Sep 2002, R. Harin'Hala, 2♂ (00006648, 00006651) (SU); 20 Sep 2002–05 Oct 2002, R. Harin'Hala, 1♂ (00006656) (CAS); 09 Jan 2003–23 Jan 2003, R. Harin'Hala, 1♂ (00006641) (CAS); 21 Apr 2003–29 Apr 2003, R. Harin'Hala, 1♂ (00006616) (UCR). Cap Ste Marie Special Reserve, 74 km S of Tsihombe, 25.58766°S 45.163°E, 37 m, 02 Jun 2003–10 Jun 2003, M. Irwin, F. Parker, R. Harin'Hala, 1♂ (00006447) (CAS). Mikea Forest, NW of Manombo, 22.90366°S 43.4755°E, 30 m, 06 Jan 2002–16 Jan 2002, M. Irwin, R. Harin'Hala, 1♂ (00006661) (CAS); 28 Mar 2002–08 Apr 2002, M. Irwin, R. Harin'Hala, 2♂ (00006587, 00006588) (SU); 29 Apr 2002–09 May 2002, M. Irwin, R. Harin'Hala, 1♂ (00006672) (UCR); 06 Mar 2003–17 Mar 2003, M. Irwin, R. Harin'Hala, 1♂ (00006671) (UCR); 27 Jul 2003–03 Aug 2003, M. Irwin, R. Harin'Hala, 2♂ (00006631, 00006655) (CAS).

*Toxopus vazimba*, new species

Figs. 2.4, 2.7, 2.11, 2.19I, 2.20L, 2.21L; Map 2.10

DIAGNOSIS: Males recognized among other species in *Toxopus* by the coloration, dense vestiture, interocular sulcus anterior to the hind margin of the eye, smooth posterior pronotal lobe, and intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII. This species is similar to *T. steineri*, *T. namoroka*, *T. basalis*, and *T. brucei*, but is distinguished from these species by the features mentioned.

DESCRIPTION: **MALE:** Body length: 11.27 mm (holotype), 11.27–12.71 mm.

COLORATION: Blackish with pale yellow clypeus, postclypeal medial longitudinal depression, antennifer, scape basally, pedicel basally, antennal segment VII, anterolateral and anteromedial margins of pronotum, meso- and metasterna, corium basally, coxae, trochanters, forefemur ventrally and basally, mid and hind femora basally, fore- and mid tibiae except lateral stripes, hind tibia except apically, tarsi, anterior areas of laterotergites, abdominal sternites medially, lateral margin of sternite II, anterolateral and posterolateral margins of sternites III–VI, and anterolateral margin of sternite VII. Labium and antennal segment VI brown. VESTITURE: Dense; other features as in generic description. STRUCTURE: HEAD: Subquadrate in lateral view (Fig. 2.13K); longer than wide in dorsal view; ventrally with deep medial depression; anteocular region longer than postocular (Fig. 2.13K); postclypeus with deep, broad medial longitudinal depression to interocular sulcus; synthlipsis about two times width of eye; interocular sulcus anterior to hind margin of eye; postocular narrow in dorsal view (Fig. 2.13A); ocelli large, located on shallow median tubercle (Fig. 2.13K), separated by less than diameter of ocellus; eye about one-third of head length, not reaching dorsal and ventral head surfaces; antenna inserted dorsally on head; pedicel about one-fourth longer than scape, slightly curved; labium stout; labial segment III ventrally straight (Fig. 2.13K). THORAX: Anterior pronotal lobe with very small anterolateral

projections (Fig. 2.15B); posterior pronotal lobe smooth; pronotal transverse furrow divided by paramedian longitudinal ridges (Fig. 2.15B); lateral depressions of posterior pronotal lobe weakly transversely striated; wings spread out in specimens. ABDOMEN: Apex rounded; intersegmental sutures carinulate between sternites II–VI and laterally between VI and VII; pygophore process subtriangular in caudal view (Fig. 2.19I), not surpassing posterior margin of pygophore, apex rounded; BPE shorter than basal plate; endosomal struts not reaching posterior margin of DPS (Fig. 2.21L), with anterior and posterior ventral processes (Fig. 2.22G); area of endosomal struts–DPS fusion elongate ovoid (Fig. 2.21L); endosoma medially sclerotized (Fig. 2.21L).

ETYMOLOGY: The species epithet is a noun in apposition and is named after Tombeau Vazimba, Madagascar.

DISTRIBUTION: Known from Tsingy de Bemaraha National Park in the Mahajanga province in Madagascar (Map 2.10).

HOLOTYPE: Male: **Madagascar: Mahajanga:** Parc National Tsingy de Bemaraha, 3.4 km 93°E Bekopaka, Tombeau Vazimba, 19.14194°S 44.82805°E, 50 m, 06 Nov 2001–10 Nov 2001, Fisher et al. (00006964) (CAS).

PARATYPES: **Madagascar: Mahajanga:** Parc National Tsingy de Bemaraha, 3.4 km 93°E Bekopaka, Tombeau Vazimba, 19.14194°S 44.82805°E, 50 m, 06 Nov 2001–10 Nov 2001, Fisher et al., 1♂ (00006472) (CAS), 1♂ (00006730) (SU), 1♂ (00006967) (UCR). Parc National Tsingy de Bemaraha, 10.6 km ESE 123° Antsalova, 19.70944°S 44.71806°E, 150 m, 16 Nov 2001–20 Nov 2001, Fisher et al., 1♂ (00007109) (CAS).

## Conclusion

The Madagascan invertebrate fauna has not been thoroughly documented, but the situation is improving thanks to large taxonomic surveys such as the one that has enabled this taxonomic

project. Prior to our monograph, only 10 described species of millipede assassin bugs were known from the island. Here, we described 63 new species and three new genera of Ectrichodiinae that are endemic to Madagascar. This drastic increase in described diversity presents an opportunity to investigate the timing and origin of Madagascan Ectrichodiinae using phylogenetic, divergence dating, and biogeographic analyses; a companion paper investigating these questions is in press (Forthman & Weirauch, in press). Phylogenetic studies of Madagascan plants, vertebrates, and some invertebrates support close relationships to either Afrotropical or Oriental lineages (see Vences, 2004; Yoder & Nowak, 2006; Warren et al., 2010). Assuming their monophyly, Afrotropical-Madagascan relationships are evident in the genera *Glymmatophora* and *Maraenaspis*; both genera have the bulk of species restricted to mainland Africa. Other Madagascan genera, e.g., *Gibbosella* and *Marojejycoris*, share morphological similarities with Afrotropical taxa, e.g., *Synavecoris*, such as the drab coloration, ovoid head shape, and circular spiracles. However, male specimens of *Gibbosella* also share some morphological features (e.g., distal part of M and Cu fused or nearly so; dorsal laterotergites with posterior tubercles) with Oriental taxa, such as *Caecina* Stål, 1863, and *Bannania* Hsiao, 1973. Only formal phylogenetic analyses including comprehensive taxon sampling in the Afrotropical and Oriental regions, as well as morphological and molecular characters, will shed light on the phylogenetic relationships and origins of Madagascan Ectrichodiinae.

## References

- Álvarez-Padilla, F., Ubick, D., Griswold, C.E. 2012. *Noideattella* and *Tolegnaro*, two new genera of goblin spiders from Madagascar, with comments on the gamasomorphoid and silhouettelloid oonopids (Araneae, Oonopidae). *American Museum Novitates* 3745: 1–76.
- Bergroth, E. 1905. Rhynchota Aethiopica. IV. *Annales de la Société Entomologique de Belgique* 49: 368–378.
- Buerki, S., Devey, D.S., Callmander, M.W., Phillipson, P.B., Forest, F. 2013. Spatio-temporal history of the endemic genera of Madagascar. *Botanical Journal of the Linnean Society* 171: 304–329.
- Carpintero, D.J., Maldonado, J. 1990. Contributions to the knowledge of American Ectrichodiinae. II. Notes about *Rhiginia* and *Pothea* (Hemiptera: Reduviidae). *The Journal of Agriculture of the University of Puerto Rico (USA)* 74: 449–456.
- Carpintero, D.J., Maldonado, J. 1991. Contributions to the knowledge of American Ectrichodiinae. III. The genus *Cricetopareis* Breddin, 1903 (Hemiptera: Reduviidae). *The Journal of Agriculture of the University of Puerto Rico (USA)* 75: 81–486.
- Chłond, D. 2010a. A new, remarkable genus and two new species of Ectrichodiinae (Hemiptera: Heteroptera: Reduviidae) from Madagascar. *Zootaxa* 2522: 61–68.
- Chłond, D. 2010b. *Planeocoris*, a new genus of Stenopodainae (Hemiptera: Heteroptera: Reduviidae) from Madagascar. *Zootaxa* 2400, 41–48.
- Chłond, D. 2010c. A new species of the Madagascan genus *Censorinus* (Heteroptera: Reduviidae: Reduviinae). *Acta Entomologica Musei Nationalis Pragae* 50: 1–6.
- Chłond, D. 2011a. Description of two new reduviids from Madagascar (Heteroptera: Reduviidae: Stenopodainae). *Oriental Insects* 45: 293–303.
- Chłond, D. 2011b. A new species of the Madagascan genus *Rodepirea* Villiers, 1962 (Hemiptera: Heteroptera: Reduviidae: Physoderinae). *Heteropterus Revista de Entomología* 11: 235–240.
- Chłond, D. 2014. A review of the taxonomic position of *Vesbius seyrigi* Villiers, 1970, with a description of two new species of *Peyrierocoris* Chłond & Junkiert, 2010 (Hemiptera: Heteroptera: Reduviidae: Harpactorinae). *Zootaxa* 3785: 288–300.
- Chłond, D., Baňář, P. 2013. *Myrmicella*, a new genus of Harpactorinae (Hemiptera: Heteroptera: Reduviidae) from Madagascar. *Zootaxa* 3718: 483–495.
- Chłond, D., Guilbert, E. 2012. *Sicardicoris*, a new Madagascan subgenus of *Peyrierocoris* Chłond & Junkiert 2010 (Hemiptera: Heteroptera: Reduviidae: Harpactorinae). *Zootaxa* 3313: 44–52.



- Chłond, D., Junkiert, Ł. 2010. *Peyrierocoris* a new genus of Harpactorinae (Hemiptera: Heteroptera: Reduviidae) from Madagascar. *Zootaxa* 2487: 33–42.
- Chłond, D., Junkiert, Ł. 2011. A new species of the Madagascan genus *Catala* Villiers, 1951 (Heteroptera: Reduviidae: Stenopodainae). *Genus* 22: 545–550.
- Damgaard, J., Andersen, N.M., Cheng, L., Sperling, F.A.H. 2000. Phylogeny of sea skaters, *Halobates* Eschscholtz (Hemiptera, Gerridae), based on mtDNA sequence and morphology. *Zoological Journal of the Linnean Society* 130: 511–526.
- DeSalle, R., Egan, M.G., Siddall, M. 2005. The unholy trinity: taxonomy, species delimitation and DNA barcoding. *Philosophical Transactions of the Royal Society B* 360: 1905–1916.
- Distant, W.L. 1913. No. IX Rhynchota. Part I: Suborder Heteroptera. *Transactions of the Linnean Society of London, 2nd Series: Zoology* 16: 139–191.
- Dougherty, V. 1995. A review of the New World Ectrichodiinae genera (Hemiptera: Reduviidae). *Transactions of the American Entomological Society* 121: 173–225.
- Forero, D., Weirauch, C. 2012. Comparative genitalic morphology in the New World resin bugs Apiomerini (Hemiptera, Heteroptera, Reduviidae, Harpactorinae). *Deutsche Entomologische Zeitschrift* 59: 5–41.
- Forthman, M., Weirauch, C. (in press). Phylogenetics and biogeography of the endemic Madagascan millipede assassin bugs (Hemiptera: Reduviidae: Ectrichodiinae). *Molecular Phylogenetics and Evolution*.
- Ganzhorn, J.U., Lowry II, P.P., Schatz, G.E., Sommer, S. 2001. The biodiversity of Madagascar: one of the world's hottest hotspots on its way out. *Oryx* 35: 346–348.
- Gil-Santana, H.R. 2014. *Pothea berengeri* sp. nov. from Brazil, with taxonomic notes on *Pothea furtadoi* Gil-Santana & Costa and *Pothea jaguaris* (Carpintero) and reinstatement of *Parapothea* Carpintero as junior synonym of *Pothea* Amyot & Serville (Hemiptera: Heteroptera: Reduviidae: Ectrichodiinae). *Zootaxa* 3826: 497–516.
- Gil-Santana, H.R. 2015. First record of the genus *Pseudopothea* from South America, with description of a new species from Brazil (Hemiptera: Heteroptera: Reduviidae: Ectrichodiinae). *Zootaxa* 3904: 541–552.
- Gil-Santana, H.R., Baena, M. 2009. Two new species of *Brontostoma* Kirkaldy (Hemiptera: Heteroptera: Reduviidae: Ectrichodiinae) from Bolivia, with description of the male genitalia of two other species of the genus, and description of the female of *B. doughertyae* Gil-Santana, Lopes, Marques & Jurberg. *Zootaxa* 1979: 41–52.
- Gil-Santana, H.R., Costa, L.A.A. 2005. *Pothea furtadoi* sp. nov. (Hemiptera, Heteroptera, Reduviidae, Ectrichodiinae) de Mato Grosso e Minas Gerais, Brasil. *Revista Brasileira de Zoologia* 22: 400–404.

- Gil-Santana, H.R., Costa, L.A.A., Marques, O.M. 2004. *Brontostoma bahiensis* sp. nov. de Ectrichodiinae (Hemiptera, Heteroptera, Reduviidae) do Estado da Bahia, Brasil. *Revista Brasileira de Zoologia* 21: 127–130.
- Gil-Santana, H.R., Lopes, C.M., Marques, O.M., Jurberg, J. 2005. Descrição de *Brontostoma doughertyae* sp. nov. e estudo morfológico comparativo com *B. rubrum* (Amyot & Serville, 1843) (Hemiptera: Heteroptera: Reduviidae: Ectrichodiinae). *Entomología y Vectores* 12: 75–94.
- Gil-Santana, H.R., Baena, M., Grillo, H. 2013. *Berengeria* Gil-Santana & Coletto-Silva, a junior synonym of *Ectrichodiella* Fracker & Bruner, with new records and taxonomic notes on Ectrichodiinae from Brazil, and with keys to Ectrichodiinae and Reduviinae genera of the New World (Hemiptera: Heteroptera: Reduviidae). *Zootaxa* 3652: 60–74.
- Goodman, S.M., Benstead, J.P. 2005. Updated estimates of biotic diversity and endemism for Madagascar. *Oryx* 39: 73–77.
- Hill, L. 2014. Revision of *Silhouettanus* with description of nine new species (Hemiptera: Heteroptera: Schizopteridae). *Zootaxa* 3815: 353–385.
- Horváth, G. 1914. Reduviidae novae Africanae. *Annales Historico-Naturales Musei Nationalis Hungarici* 12: 109–145.
- Hwang, W.S., Weirauch, C. 2010. Revision of the Malagasy *Durevius* Villiers with descriptions of two new species (Hemiptera: Reduviidae: Reduviinae). *Insect Systematics and Evolution* 41: 123–141.
- Jansson, A., Coscarón, M.D.C. 1989. Lists of the insect types in the Zoological Museum, University of Helsinki. 11. Heteroptera: Reduviidae. *Acta Entomologica Fennica* 55: 3–21.
- Karsch, F. 1892. Kurze Charakteristik neuer Wanzen aus Kamerun, gesammelt durch Herrn Dr. Paul Preuss. *Entomologische Nachrichten* 18: 129–136.
- Katoh, K., Toh, H. 2008. Recent developments in the MAFFT multiple sequence alignment program. *Briefings in Bioinformatics* 9: 286–298.
- Katoh, K., Kuma, K., Toh, H., Miyata, T. 2005. MAFFT version 5: improvement in accuracy of multiple sequence alignment. *Nucleic Acids Research* 33: 511–518.
- Kerzhner, I.M., Günther, H. 2003. Two new species of micropterous Ectrichodiinae from Sichuan, China (Heteroptera: Reduviidae). *Zoosystematica Rossica* 12: 227–230.
- Krishnankutty, S.M., Dietrich, C.H. 2011. Taxonomic revision and phylogeny of an endemic leafhopper genus *Nesocerus* (Hemiptera: Cicadellidae: Idiocerinae) from Madagascar. *Zoological Journal of the Linnean Society* 162: 499–543.
- Labina, E.S., Kerzhner, I.M. 2000. A complex case of homonymy in the Emesinae from Madagascar (Heteroptera: Reduviidae). *Zoosystematica Rossica* 9: 226.

- Maldonado, J. 1990. Systematic catalogue of the Reduviidae of the world (Insecta: Heteroptera). *Caribbean Journal of Science*, Special Edition: 1–694.
- Maldonado, J. 1995. Notes about the Old World genus *Hexamerocerus* Reuter (Heteroptera: Reduviidae: Ectrichodiinae). *Proceedings of the Entomological Society of Washington* 97: 112–116.
- Meier, R., Shiyang, K., Vaidya, G., Ng, P.K.L. 2006. DNA barcoding and taxonomy in Diptera: a tale of high intraspecific variability and low identification success. *Systematic Biology* 55: 715–728.
- Mugrabi, D.F., Azevedo, C.O. 2010. Insecta, Hymenoptera, Bethylidae: range extension and filling gaps in Madagascar. *Check List* 6: 062–063.
- Murugan, C., Livingstone, D. 1995. Description of a new genus of Ectrichodiinae and two new species of the genus *Haematorrhophus* Stål from South India (Heteroptera: Reduviidae). *Journal of the Bombay Natural History Society* 92: 386–389.
- Myers, N., Mittermeier, R.A., Mittermeier, C.G., da Fonseca, G.A.B., Kent, J. 2000. Biodiversity hotspots for conservation priorities. *Nature* 403: 853–858.
- Phillipson, P.B., Schatz, G.E., Lowry II, P.P., Labat, J. 2006. A catalogue of the vascular plants of Madagascar. In Ghazanfar, S.A. & Beentje, H. (eds.): *Taxonomy and Ecology of African Plants, their Conservation and Sustainable Use. Proceedings of the 17<sup>th</sup> AETFAT Congress*. Kew Publishing Ltd, London, pp. 613–627.
- Putshkov, V.G., Putshkov, P.V. 1986–1989. A Catalogue of the Reduviidae (Heteroptera) of the World. 6 Vols. Vinity, Lyubertsy.
- Quek, S.-P. 2009. Borneo. In Gillespie, R.G., & Clague, D.A. (eds.): *Encyclopedia of Islands*. University of California Press, Berkeley, pp. 111–116.
- Rakotomanana, H., Jenkins, R.K.B., Ratsimbazafy, J. 2013. Conservation challenges for Madagascar in the next decade. In Raven, N.S., Sodhi, N.S., & Gibson, L. (eds.): *Conservation Biology: Voices from the Tropics*. John Wiley and Sons Ltd., Oxford, pp. 33–39.
- Rédei, D., Tsai, J.-F. 2012. The assassin bug genus *Haematoloecha* in Taiwan, with notes on species occurring in the neighboring areas (Hemiptera: Heteroptera: Reduviidae: Ectrichodiinae). *Zootaxa* 3332: 1–26.
- Rédei, D., Ren, S., Bu, W. 2012. A new synonymy in the genus *Rhysostethus* (Hemiptera: Heteroptera: Reduviidae). *Acta Entomologica Musei Nationalis Pragae* 52: 341–348.
- Reuter, O.M. 1887. Ad cognitionem Heteropterorum Madagascariensium. *Entomologisk Tidskrift* 8: 77–109.

- Scales, I.R. 2014. Conservation as the crossroads: biological diversity, environmental change and natural resource use in Madagascar. In Scales, I.R. (ed): *Conservation and Environmental Management in Madagascar*. Routledge, NY, pp. 1–13.
- Schatz, G.E. 2002. Taxonomy and herbaria in service of plant conservation: lessons from Madagascar's endemic families. *Annals of the Missouri Botanical Garden* 89: 145–152.
- Simon, C., Frati, F., Beckenbach, A., Crespi, B., Liu, H., Flook, P. 1994. Evolution, weighting, and phylogenetic utility of mitochondrial gene sequences and a compilation of conserved polymerase chain reaction primers. *Annals of the Entomological Society of America* 87: 651–701.
- Smith, M.A., Fisher, B.L., Hebert, P.D.N. 2005. DNA barcoding for effective biodiversity assessment of a hyperdiverse arthropod group: the ants of Madagascar. *Philosophical Transactions of the Royal Society B* 360: 1825–1834.
- Stål, C. 1853. Nya genera bland Hemiptera. *Öfversigt af Kongliga Vetenskaps-Akademiens Förhandlingar* 10: 259–267.
- Sussman, R.W., Green, G.M., Sussman, L.K. 1996. The use of satellite imagery and anthropology to assess the causes of deforestation in Madagascar. In Sponsel, L.E., Headland, T.N., & Bailey, R.C. (eds.): *Tropical Deforestation: The Human Dimension*. Colombia University Press, NY, pp. 296–315.
- Vences, M. 2004. Origin of Madagascar's extant fauna: a perspective from amphibians, reptiles and other non-flying vertebrates. *Italian Journal of Zoology* 71(S2): 217–228.
- Villiers, A. 1948. Les Réduviides de Madagascar. I. Stenopoditae. Gen. *Pygolampis* et *Sastrapada*. *Bulletin de la Société Entomologique de France* 53: 136–139.
- Villiers, A. 1968a. Faune de Madagascar. XXVIII. Insectes Hémiptères Reduviidae. (1ère partie). *Office de la Recherche Scientifique et Technique Outre-Mer, Centre National de la Recherche Scientifique* 28: 1–198.
- Villiers, A. 1968b. Reduviidae (Hemiptera Heteroptera). *Exploration du Parc National Albert, 2ème série* 21: 17–43.
- Villiers, A. 1979. Faune de Madagascar. 49. Insectes Hémiptères Reduviidae. (2ème partie). *Office de la Recherche Scientifique et Technique Outre-Mer, Centre National de la Recherche Scientifique* 49: 1–202.
- Warren, B.H., Strasberg, D., Bruggemann, J.H., Prys-Jones, R.P., Thébaud, C. 2010. Why does the biota of the Madagascar region have such a strong Asiatic flavour? *Cladistics* 26: 526–538.
- Weirauch, C. 2008a. *Mangabea barbiger*, new species of Collartidini (Heteroptera, Reduviidae, Emesinae) from Madagascar. In Grozeva, S. & Simov, S. (eds.): *Advances in Heteroptera Research*. Pensoft Publishers, Sofia, pp. 327–337.

- Weirauch, C. 2008b. Cladistic analysis of Reduviidae (Heteroptera: Cimicomorpha) based on morphological characters. *Systematic Entomology* 33: 229–274.
- Weirauch, C., Rabitsch, W., Rédei, D. 2009. *Austrokatanga*, gen. nov., new genus of Ectrichodiinae (Hemiptera: Heteroptera: Reduviidae) from Australia. *Zootaxa* 2094: 1-15.
- Weirauch, C., Berenger, J.M., Berniker, L., Forero, D., Forthman, M., Frankenberg, S., Freedman, A., Gordon, E., Hoey-Chamberlain, R., Hwang, W.S., Michael, A., Udah, O., Watson, C., Zhang, G., Zhang, J. 2014. An illustrated identification key to assassin bug subfamilies and tribes. *Canadian Journal of Arthropod Identification* No. 26, doi:10.3752/cjai.2014.26.
- Whitten, T., Damanik, S.J., Anwar, J., Hisyam, N. 2000. *The Ecology of Indonesia Series. Vol. 1. The Ecology of Sumatra*. Tuttle Publishing, North Clarendon.
- Yoder, A.D., Nowak, M.D. 2006. Has vicariance or dispersal been the predominant biogeographic force in Madagascar? Only time will tell. *Annual Review of Ecology, Evolution, and Systematics* 37: 405–431.
- Zhang, G., Weirauch, C. 2011. Matching dimorphic sexes and immature stages with adults: resolving the systematics of the *Bekilya* group of Malagasy assassin bugs (Hemiptera: Reduviidae: Peiratinae). *Systematic Entomology* 36: 115–138.

Figure 2.1. Dorsal habitus of *Gibbosella* species, *Glymmatophora carolae*, and *Glymmatophora crassipes*.

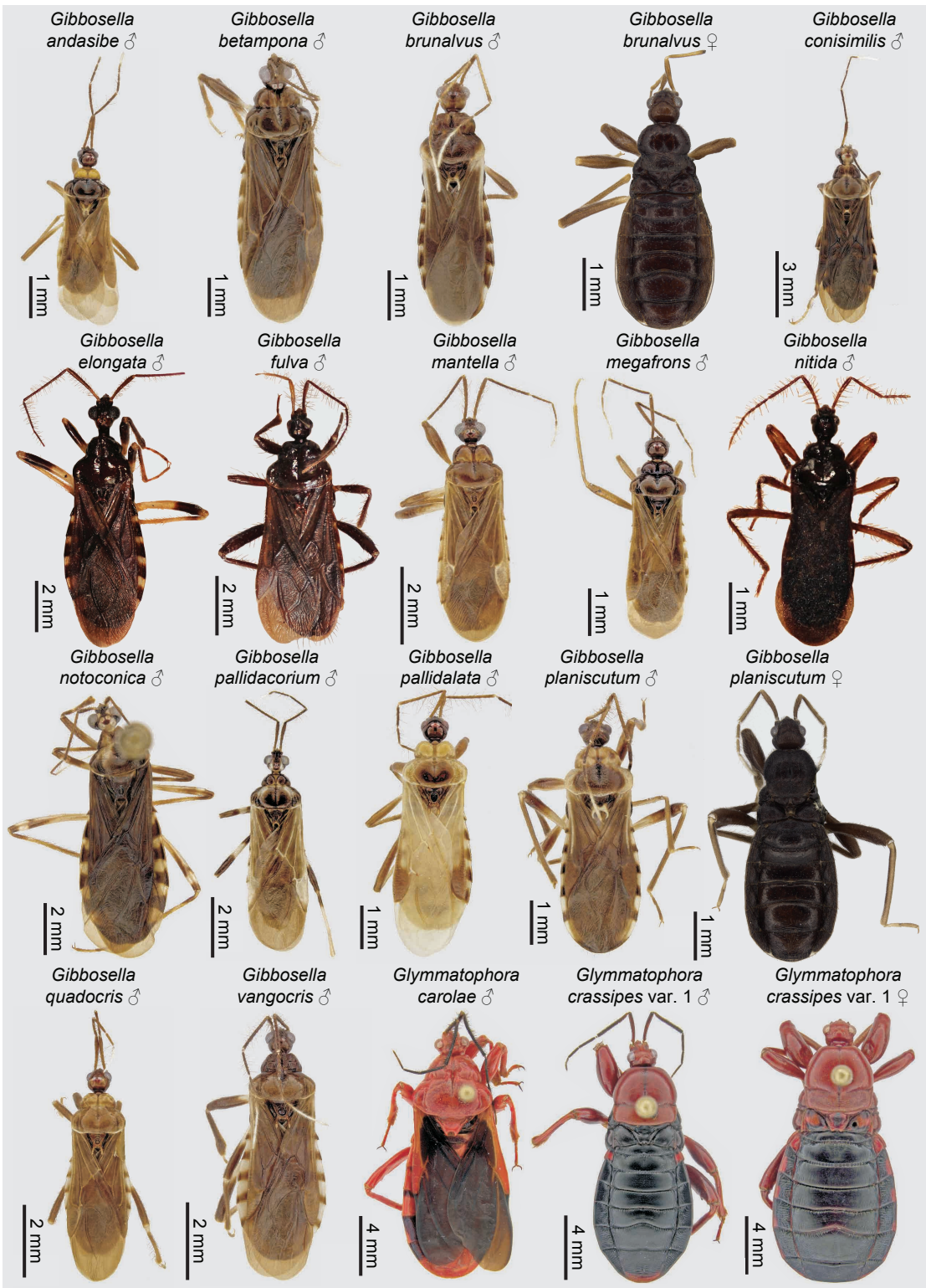


Figure 2.2. Dorsal habitus of *Glymmatophora crassipes*, *Maraenaspis bidens*, *Marojejycoris* species, and *Tanindrazanus* species (*T. amboasarius* – *T. kathrynae*).



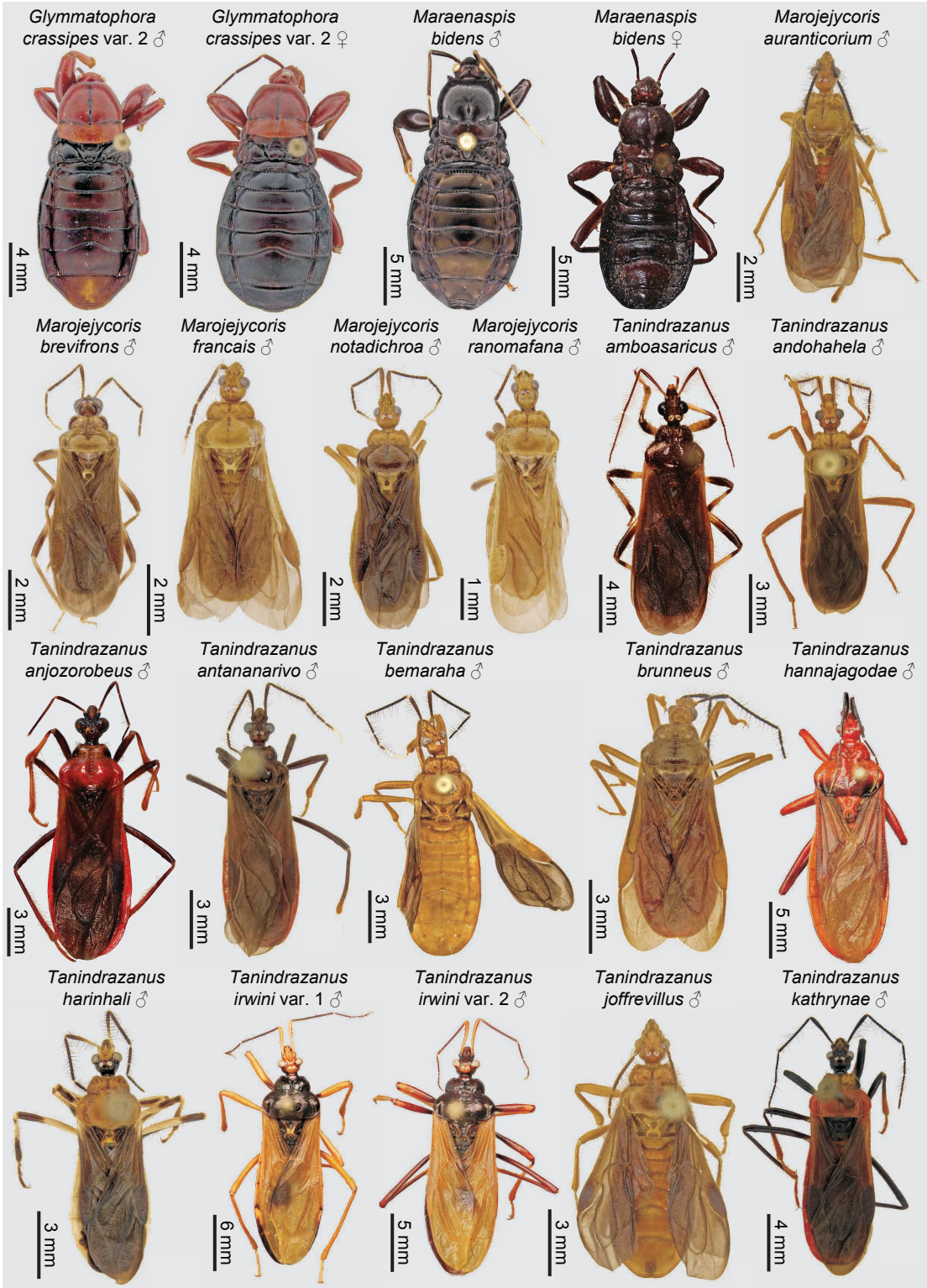


Figure 2.3. Dorsal habitus of *Tanindrazanus* species (*T. mahafaly* – *T. vohiparara*), *Toliarus* species, and *Toxopus* species (*T. ambohitantely* – *T. griswoldi* [male]).

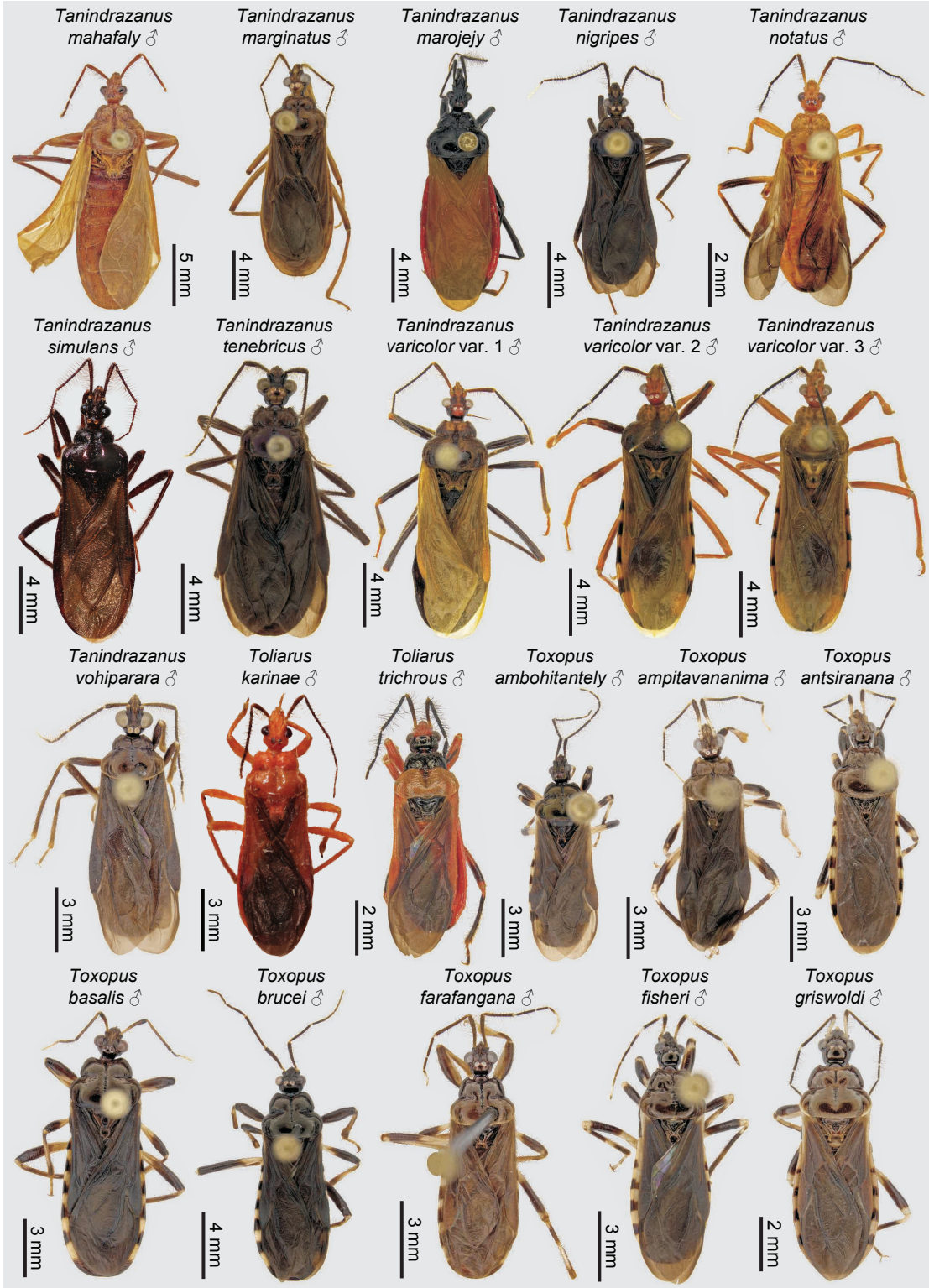


Figure 2.4. Dorsal habitus of *Toxopus* species (*T. griswoldi* [female] – *T. vazimba*).



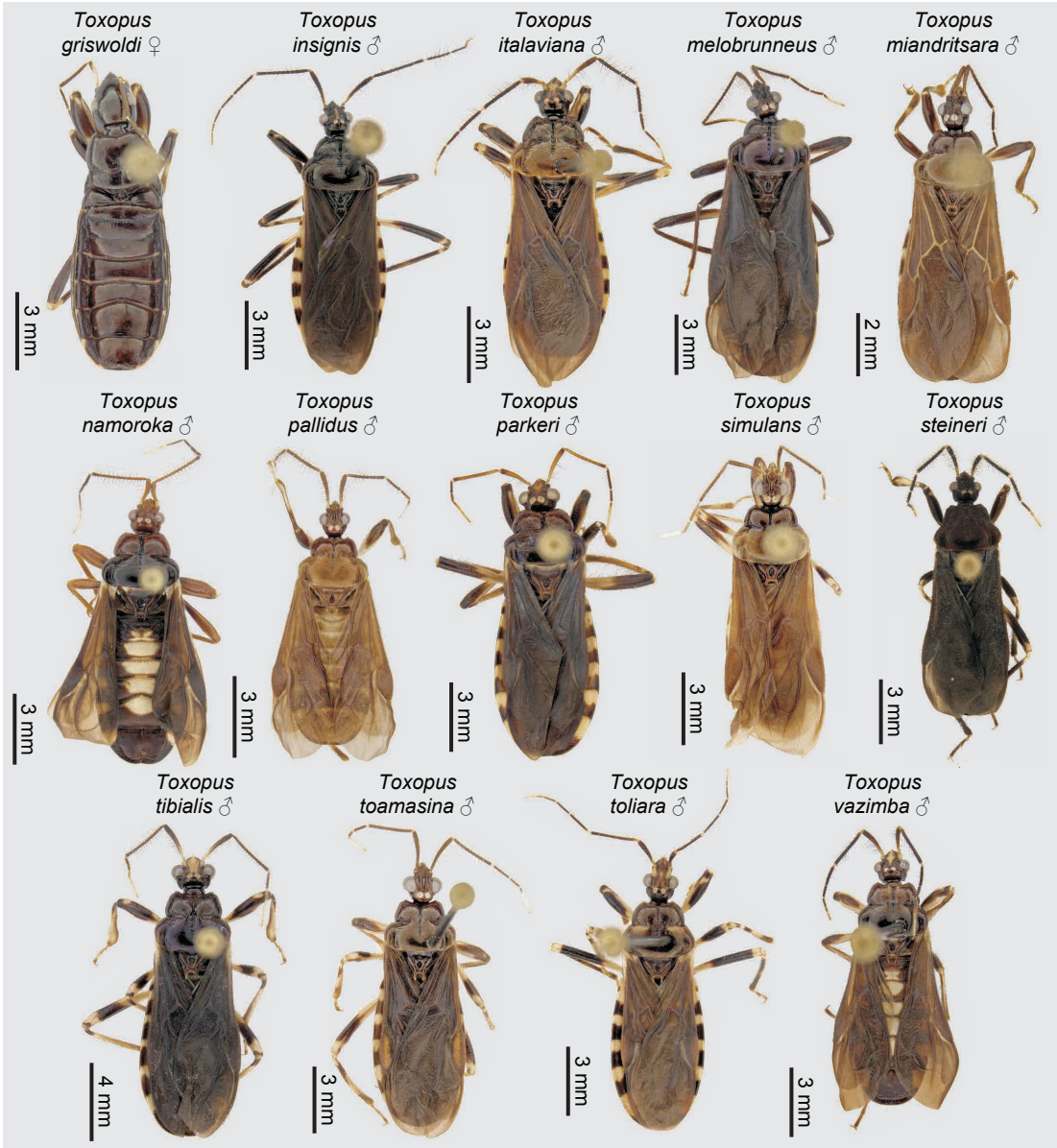


Figure 2.5. Lateral habitus of *Gibbosella* species, *Glymmatophora* species, *Maraenaspis bidens*, and *Marojejycoris* species (*M. auranticorium* – *M. francais*).

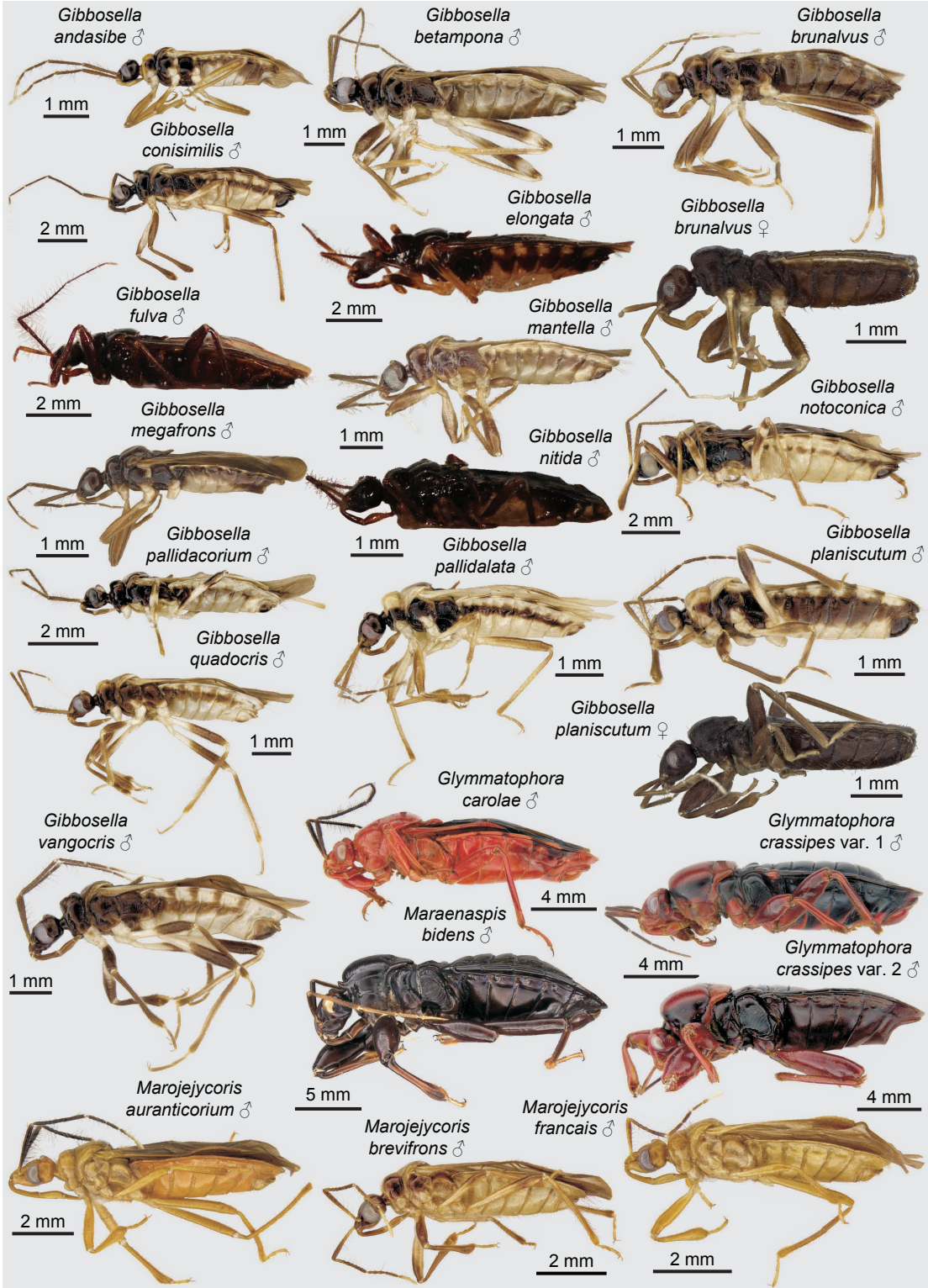


Figure 2.6. Lateral habitus of *Marojejycoris notadichroa*, *Marojejycoris francais*, and *Tanindrazanus* species (*T. amboasaricus* – *T. varicolor*).



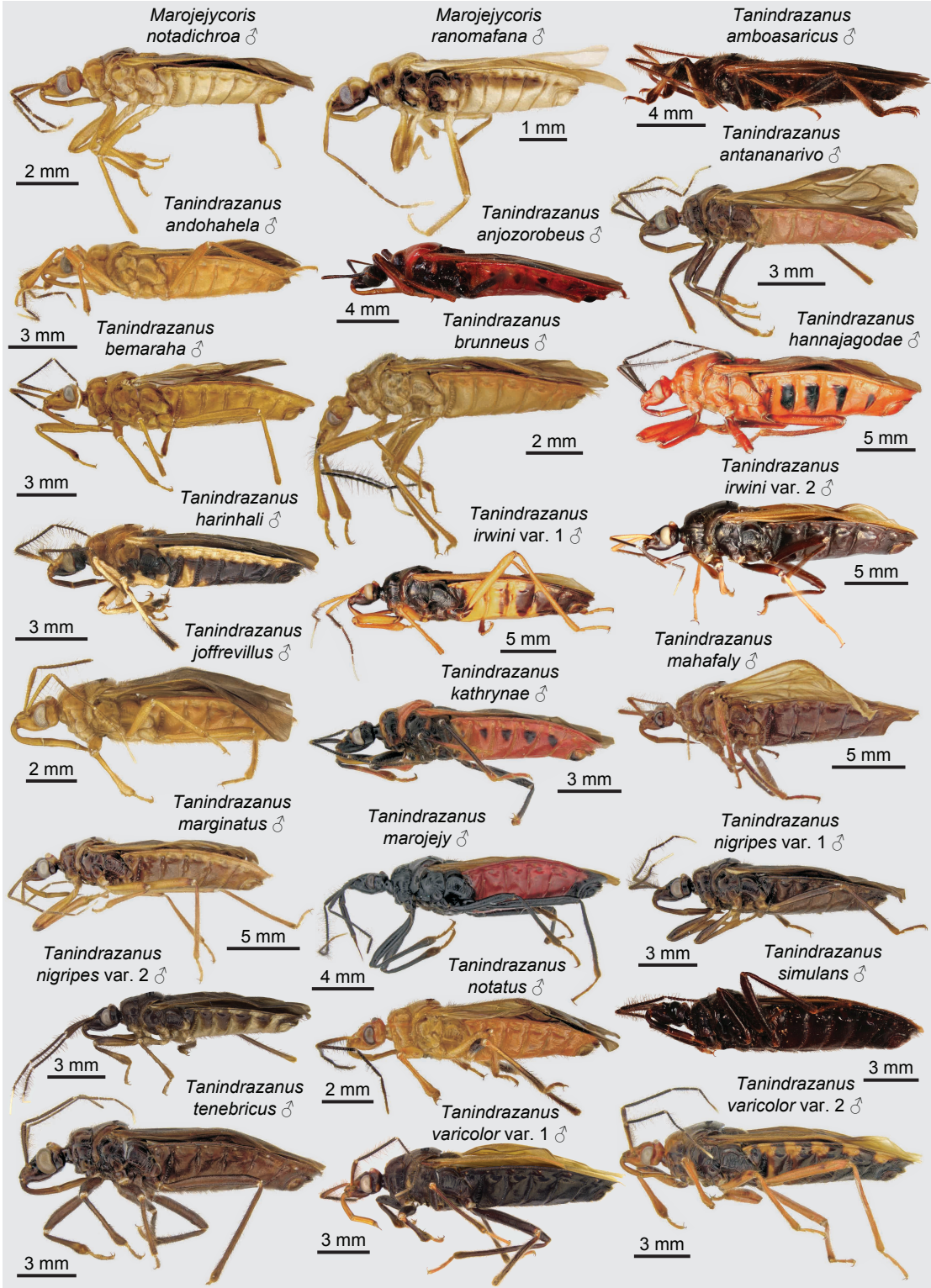


Figure 2.7. Lateral habitus of *Tanindrazanus varicolor*, *Tanindrazanus vohiparara*, *Toliarus* species, and *Toxopus* species.

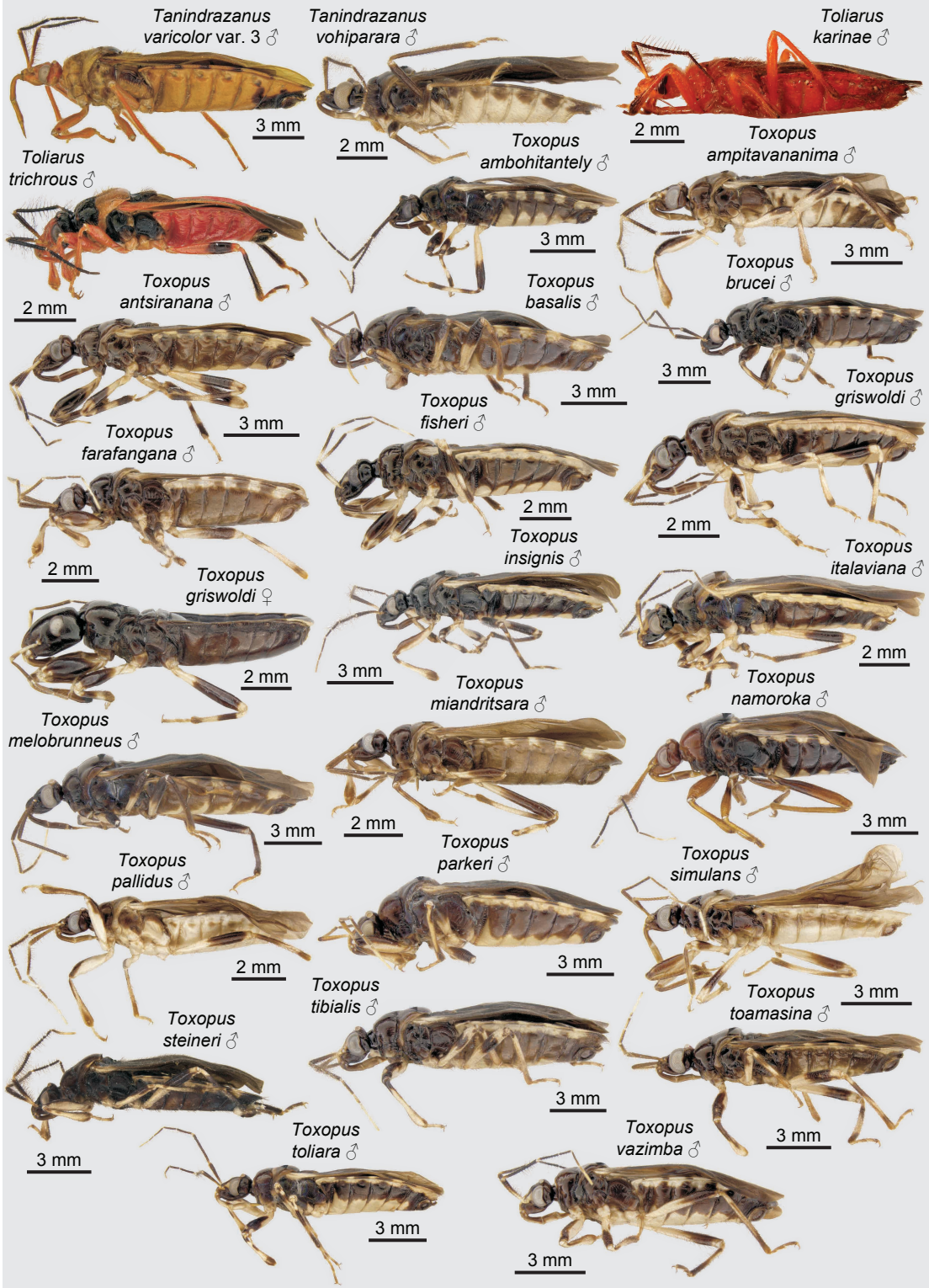


Figure 2.8. Ventral habitus of *Gibbosella* species, *Glymmatophora* species, and *Maraenaspis bidens*.



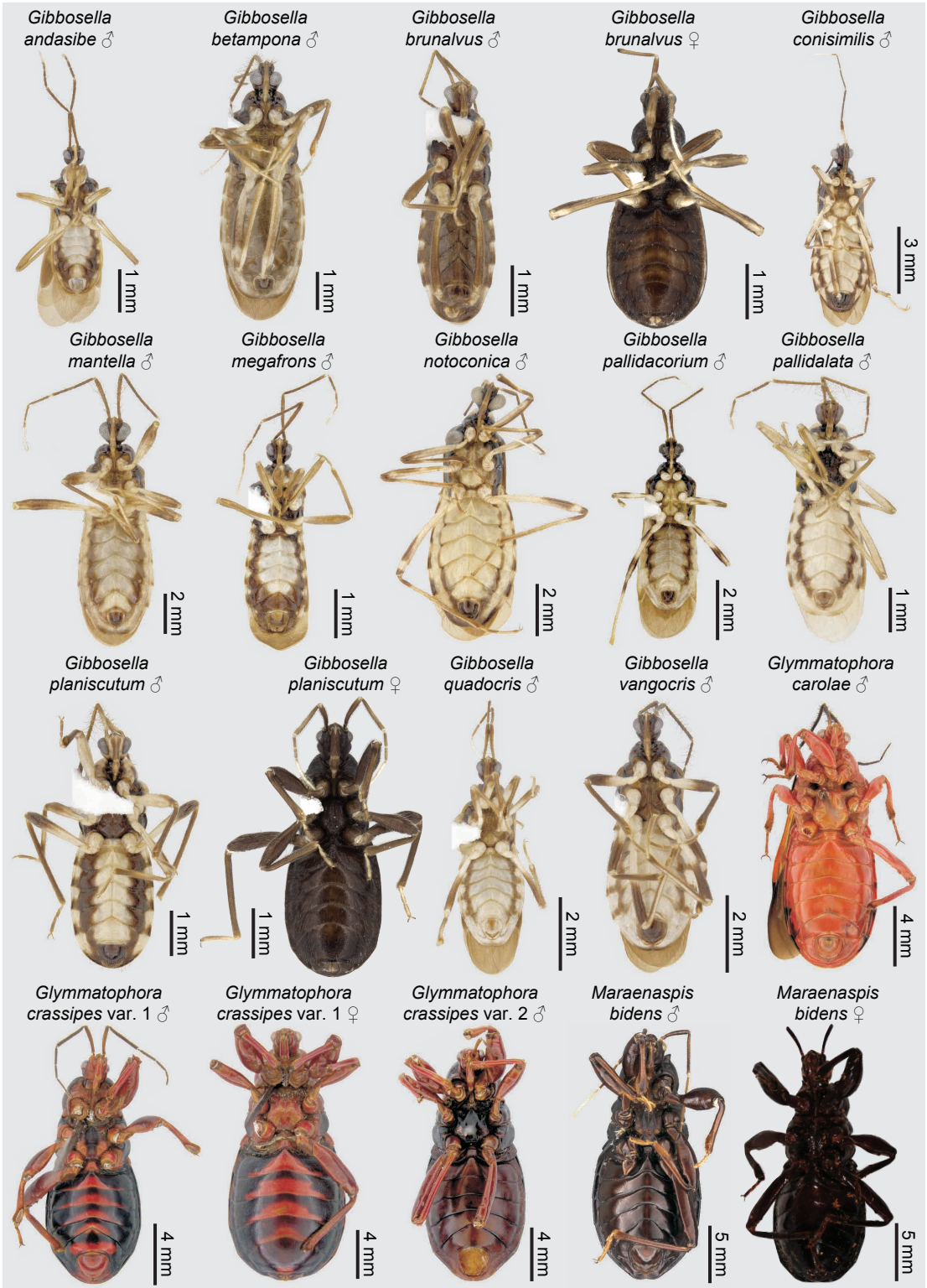


Figure 2.9. Ventral habitus of *Marojejycoris* species and *Tanindrazanus* species (*T. amboasarius* – *T. nigripes*).

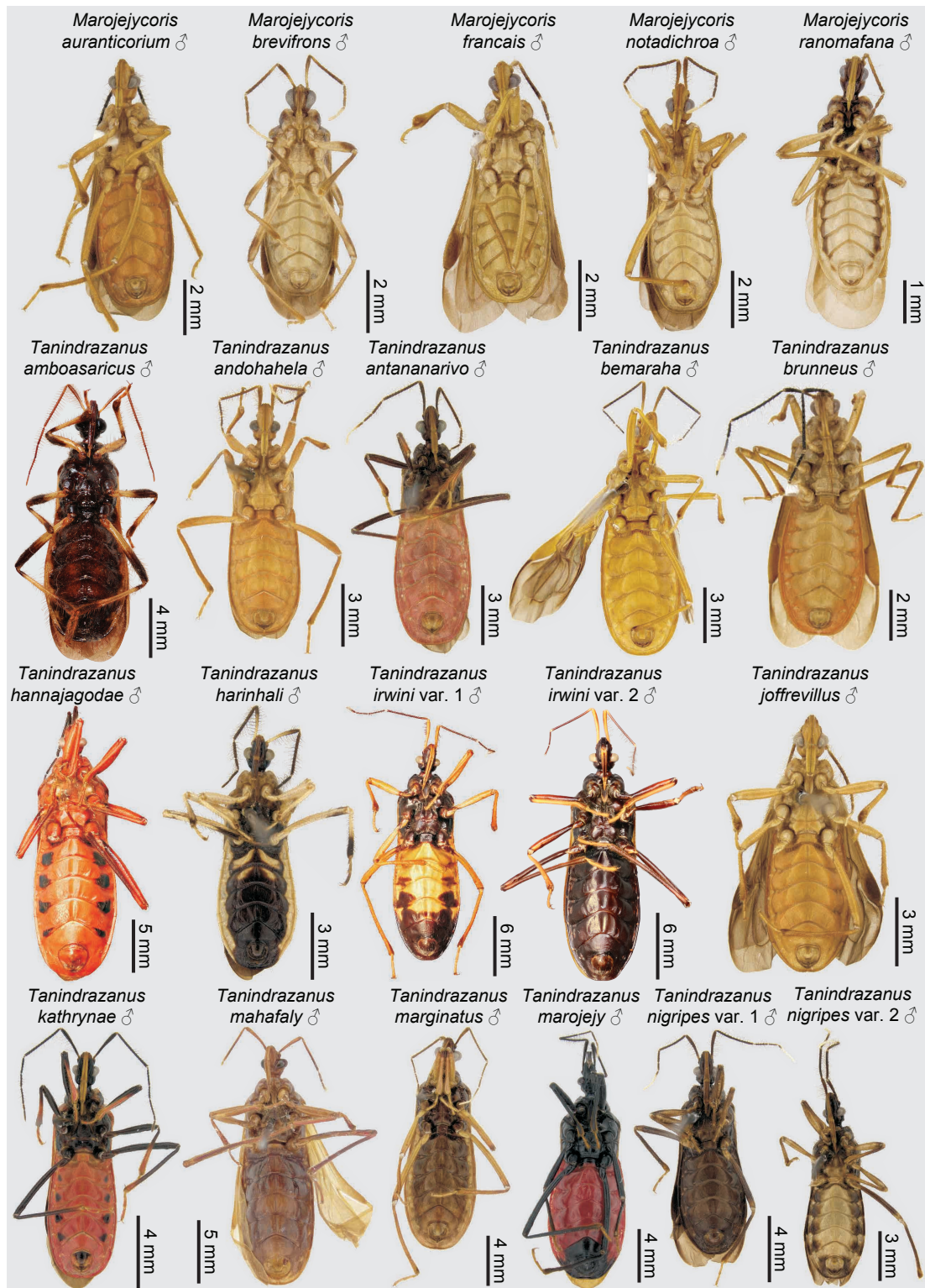


Figure 2.10. Ventral habitus of *Tanindrazanus* species (*T. notatus* – *T. vohiparara*), *Toliarus trichrous*, and *Toxopus* species (*T. ambohitantely* – *T. melobrunneus*).



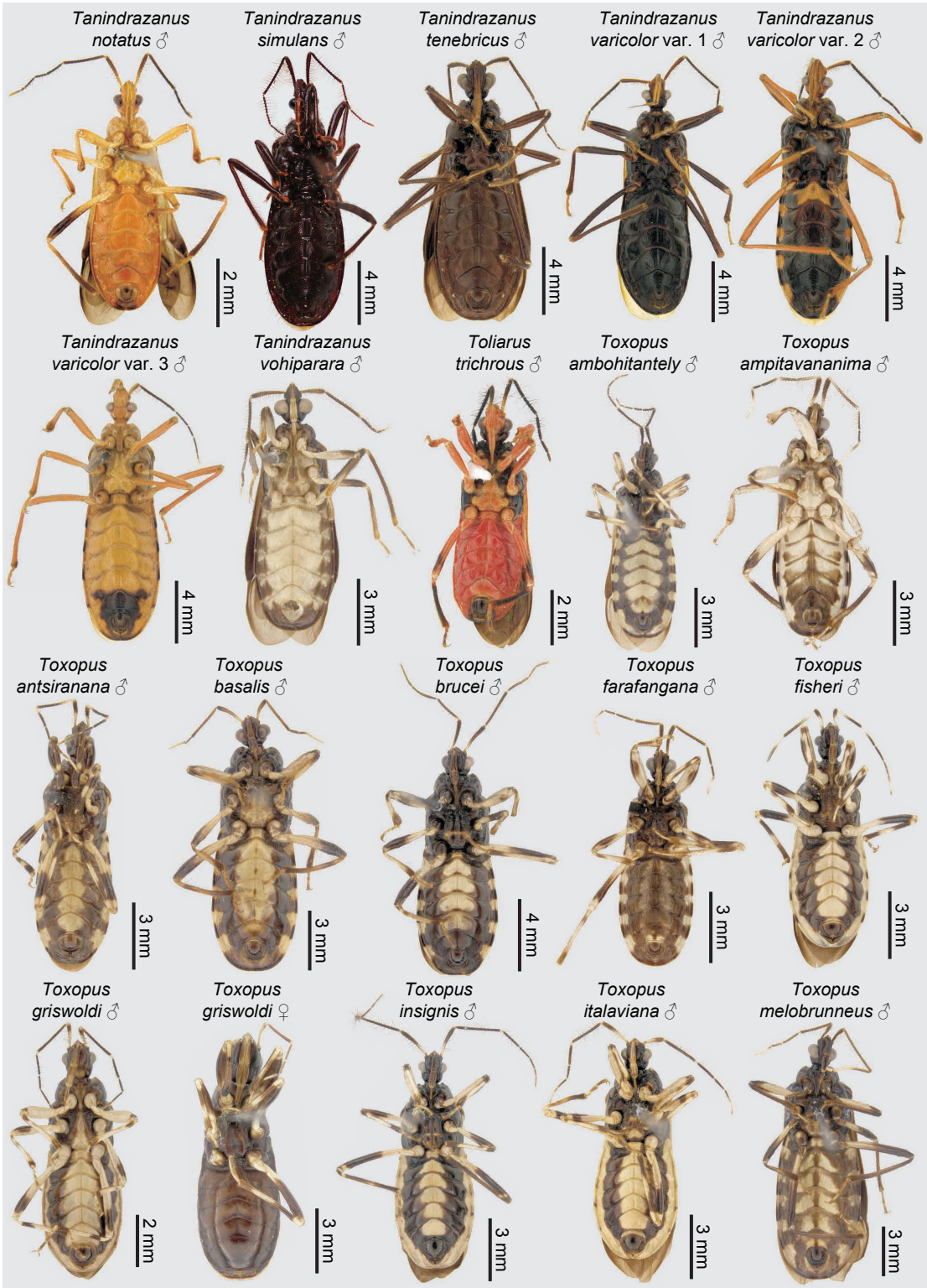


Figure 2.11. Ventral habitus of *Toxopus* species (*T. miandritsara* – *T. vazimba*).

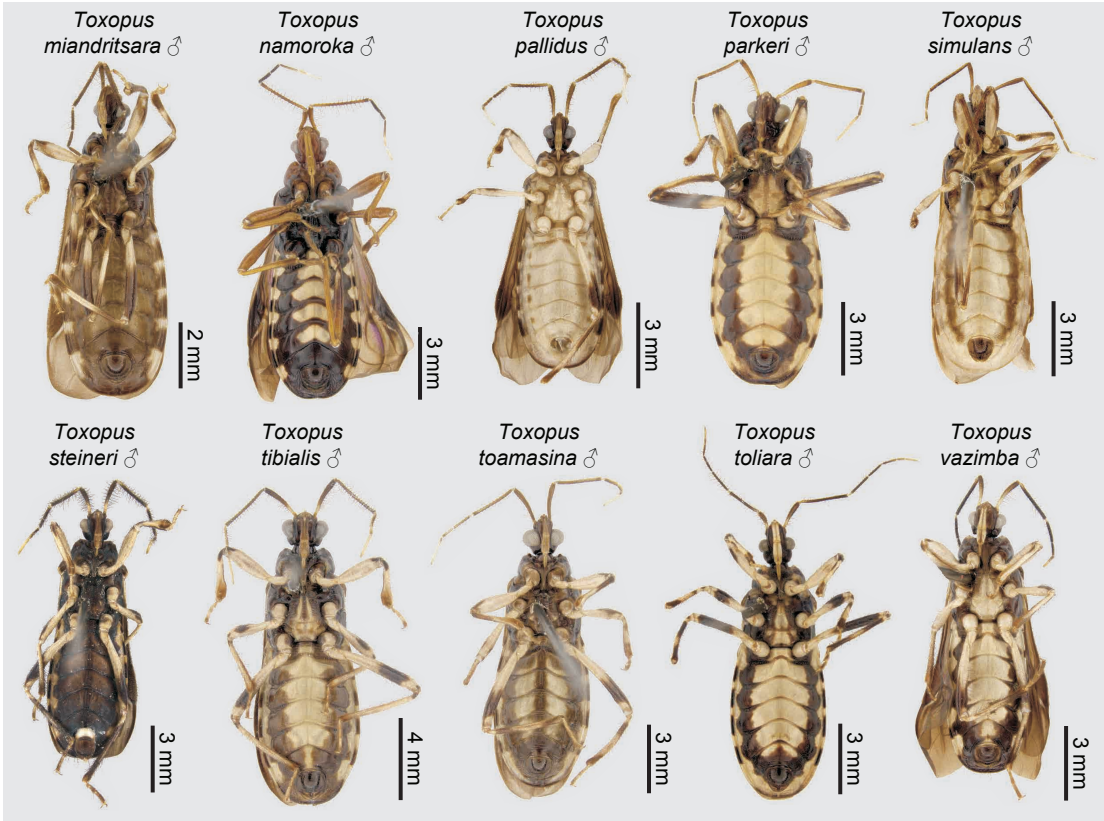


Figure 2.12. Dorsal and lateral habitus of holotypes of *Gibbosella elongata*, *Gi. mirabilis*, *Glymmatophora crassipes*, *Synavecoris dimorphus*, and *Toxopus signoretii*.



Figure 2.13. Dorsal (A., B.) and lateral (C.–K.) head structures. A. *Glymmatophora crassipes*, B. *Toxopus italaviana*, C. *Gibbosella mantella*, D. *Glymmatophora crassipes*, E. *Maraenaspis bidens*, F. *Marojejycoris brevifrons*, G. *Tanindrazanus kathrynae*, H. *Tanindrazanus varicolor*, I. *Toliarus trichrous*, J. *Toxopus ambohitantely*, K. *Toxopus parkeri*.



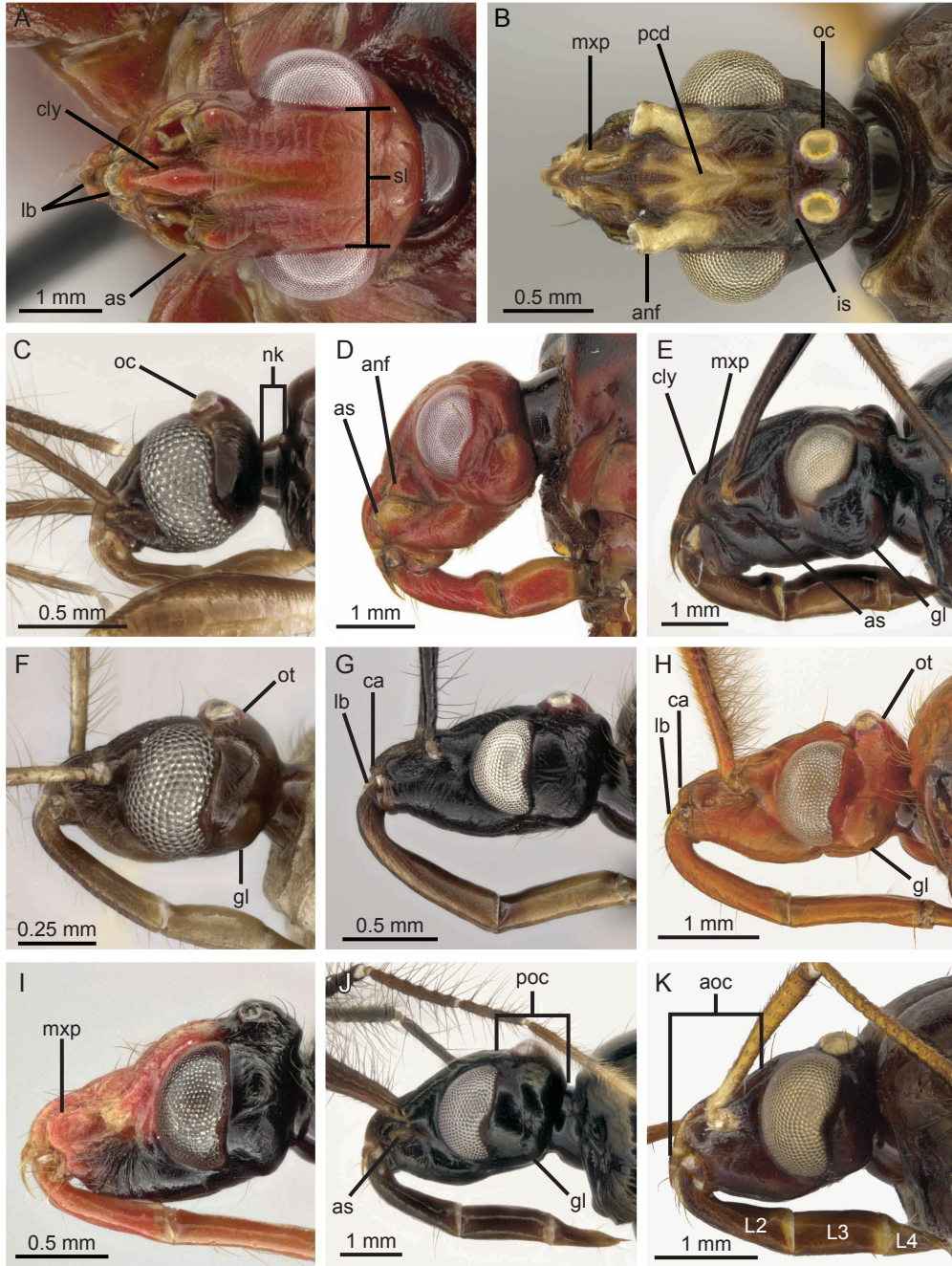


Figure 2.14. Lateral antennal morphology of *Toxopus insignis*. A. Antenna, B. basiflagellomere, C. distiflagellomere.



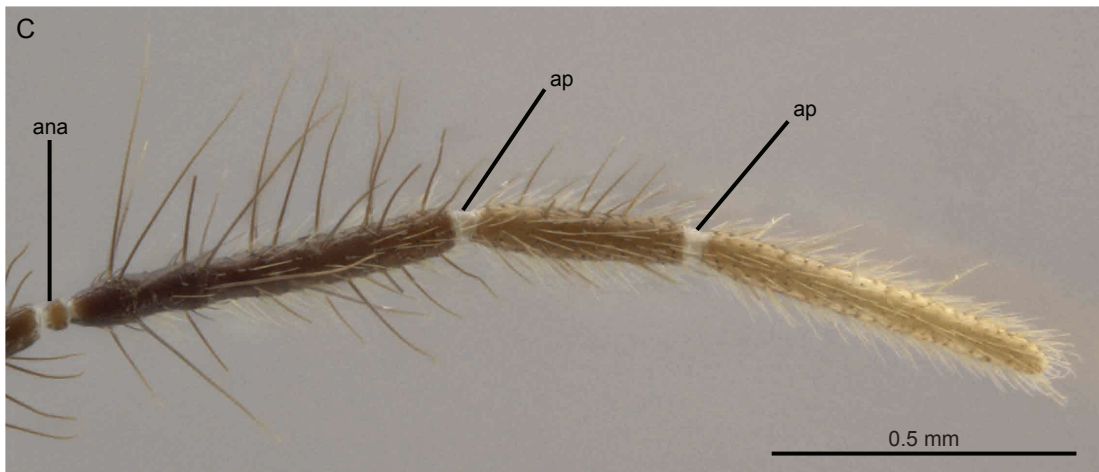
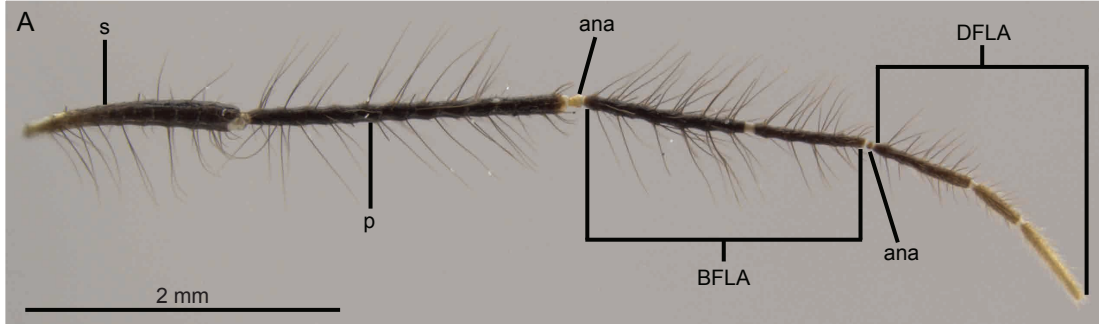


Figure 2.15. Dorsal (A., B.), lateral (C., D.), and ventral (E., F.) thoracic structures. A. *Maraenaspis bidens*, B. *Toxopus italaviana*, C. *Gibbosella conisimilis*, D. *Tanindrazanus kathrynae*, E. *Glymmatophora crassipes*, F. *Tanindrazanus irwini*.

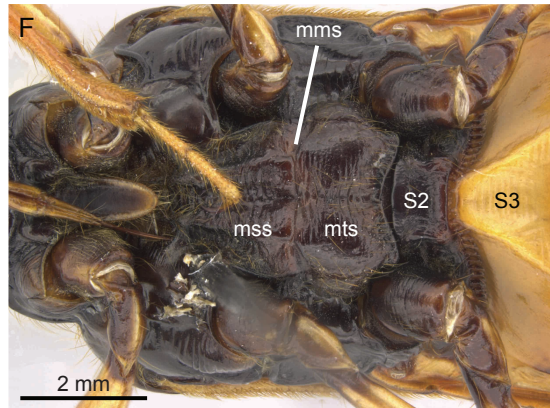
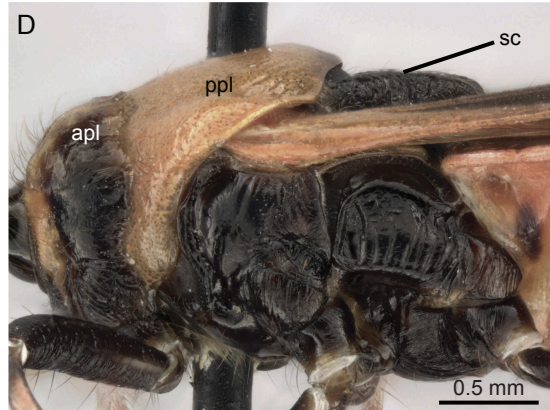
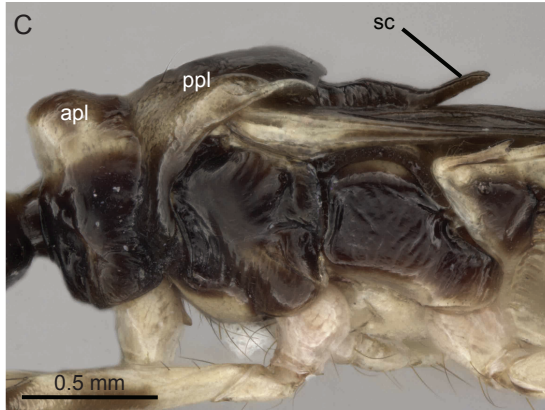
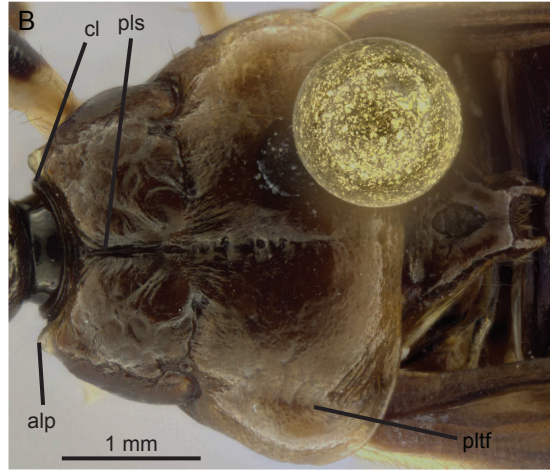
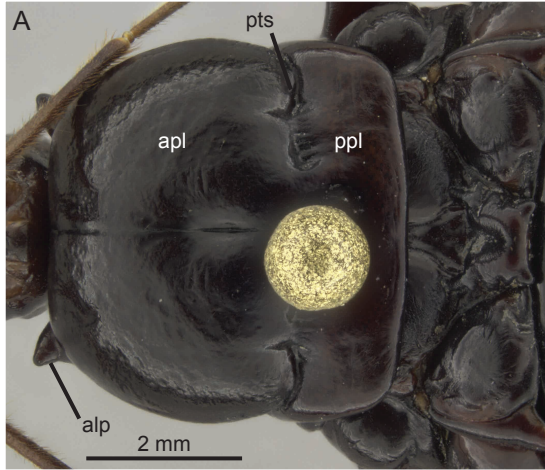


Figure 2.16. Metathoracic gland evaporatorium (A.–C.) and femoral armature (D., E.). A. *Tanindrazanus varicolor* (lateral), B. *Glymmatophora crassipes* (lateral), C. *Glymmatophora carolae* (ventral), D. *Gibbosella* sp., E. *Glymmatophora crassipes*.



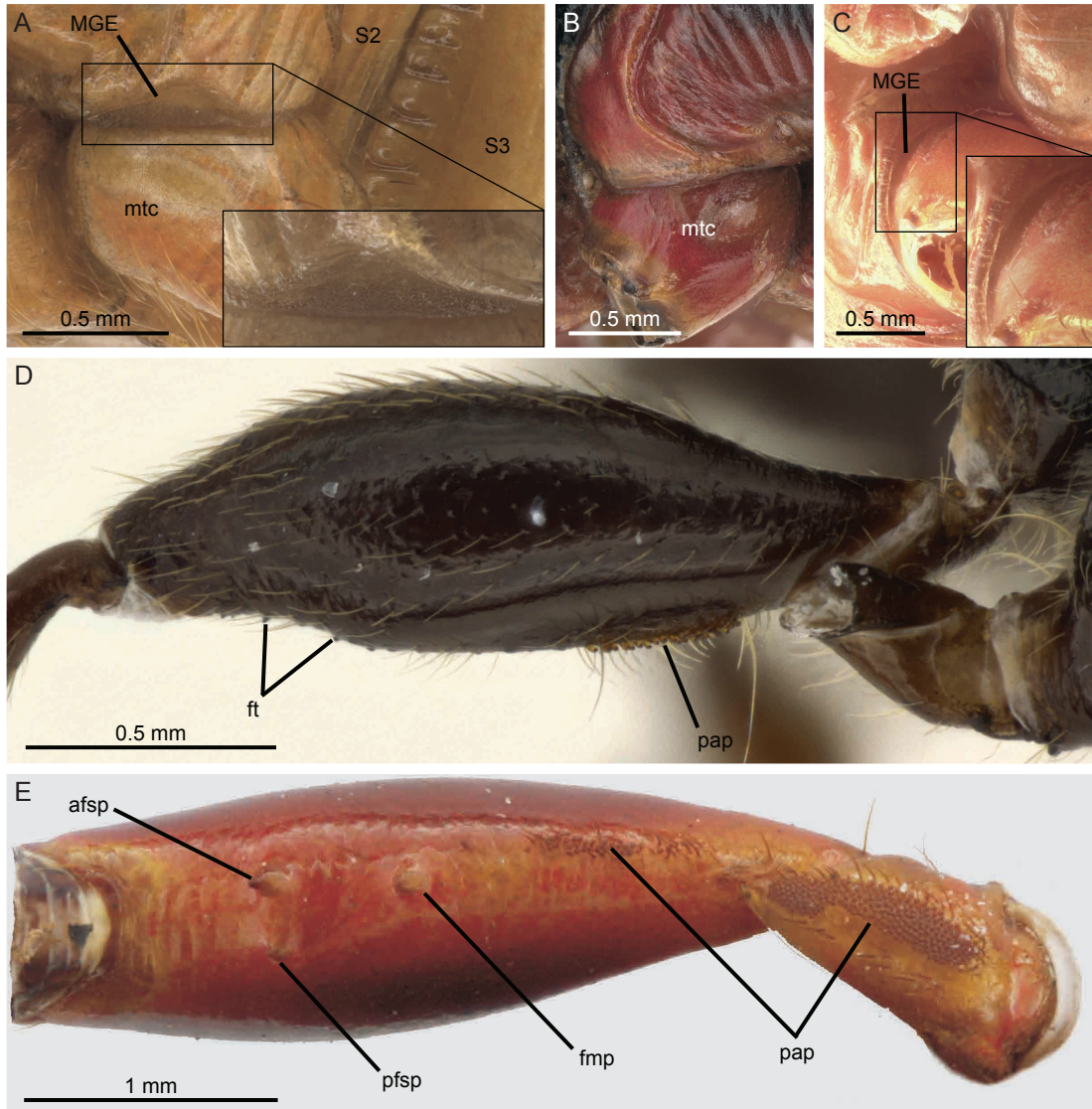


Figure 2.17. Wing morphology. A. *Gibbosella planiscutum*, B. *Glymmatophora carolae*, C. *Tanindrazanus varicolor*.

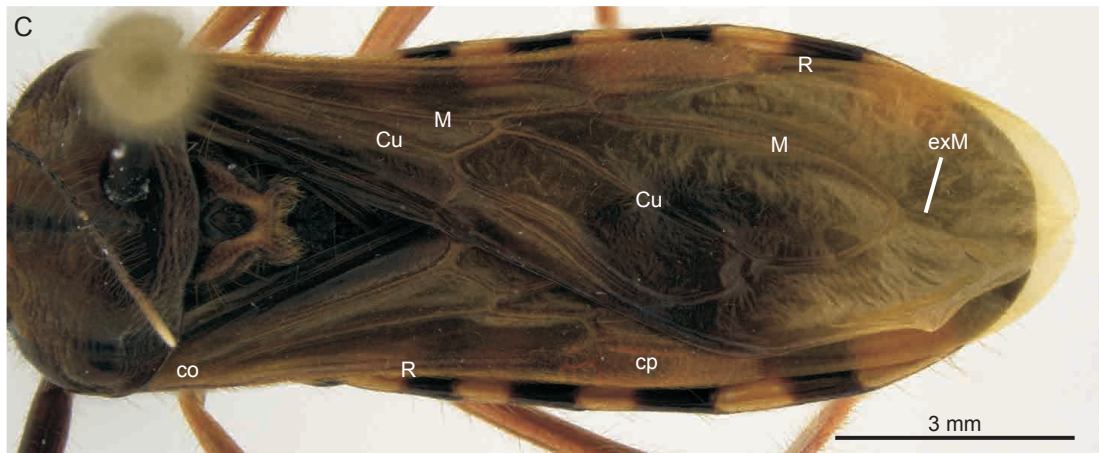
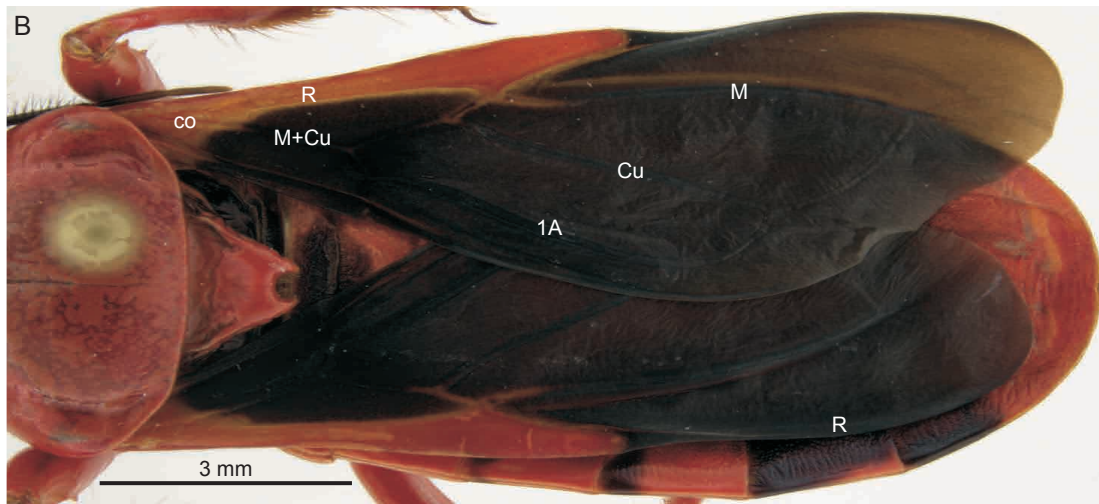
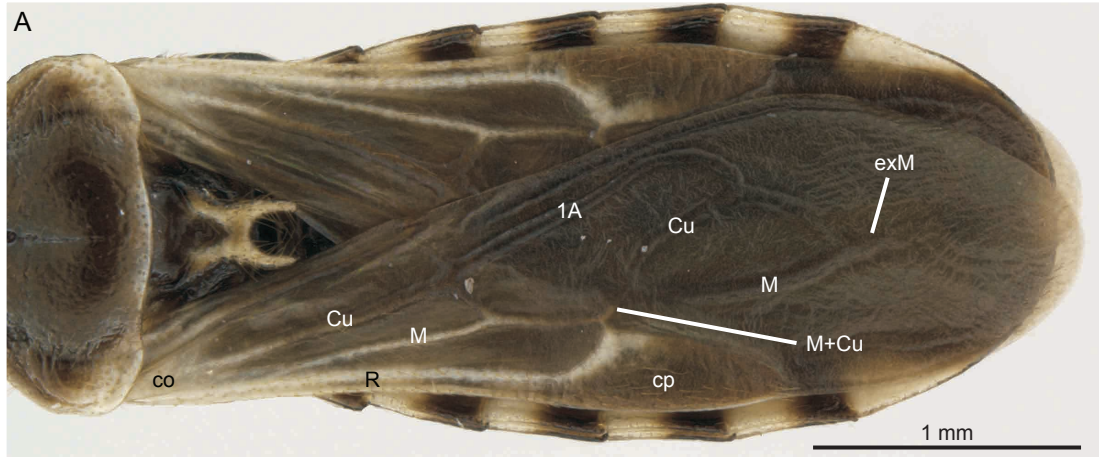


Figure 2.18. Lateral (A., B.) and ventral (C.) abdominal morphology. A. *Gibbosella conisimilis*, B. *Tanindrazanus tenebricus*, C. *Distirogaster* sp.



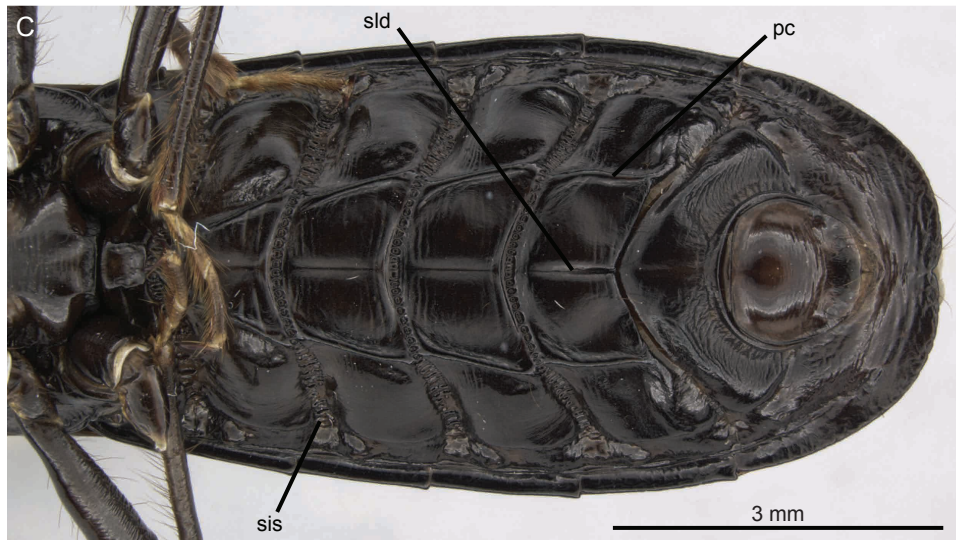
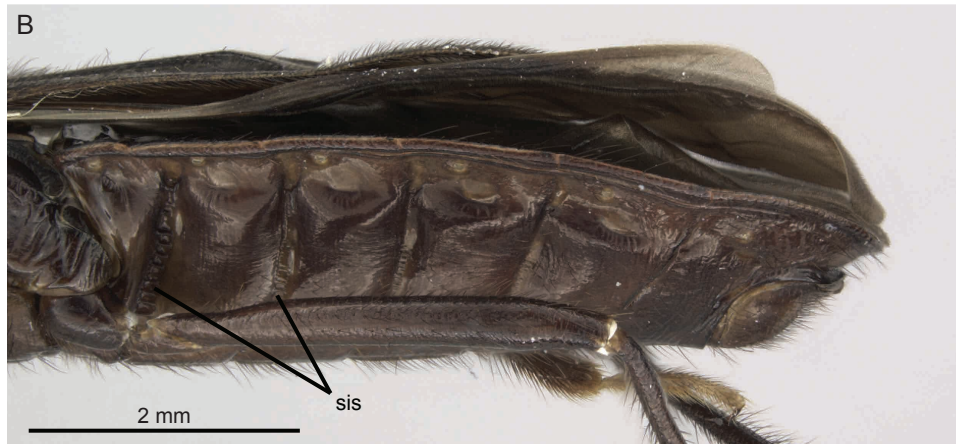
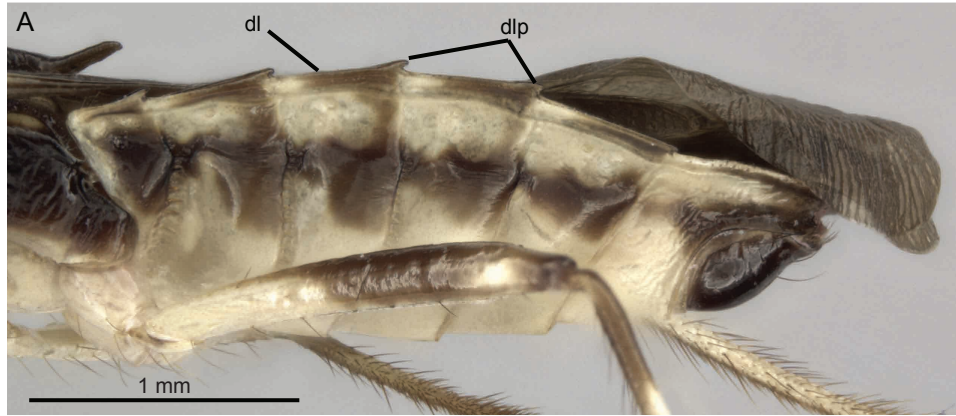


Figure 2.19. Pygophore morphology in caudal view. A. *Gibbosella quadocris*, B. *Gibbosella vangocris*, C. *Glymmatophora crassipes*, D. *Maraenaspis bidens*, E. *Marojejycoris auranticorium*, F. *Tanindrazanus kathrynae*, G. *Toliarus trichrous*, H. *Toxopus fisheri*, I. *Toxopus vazimba*.

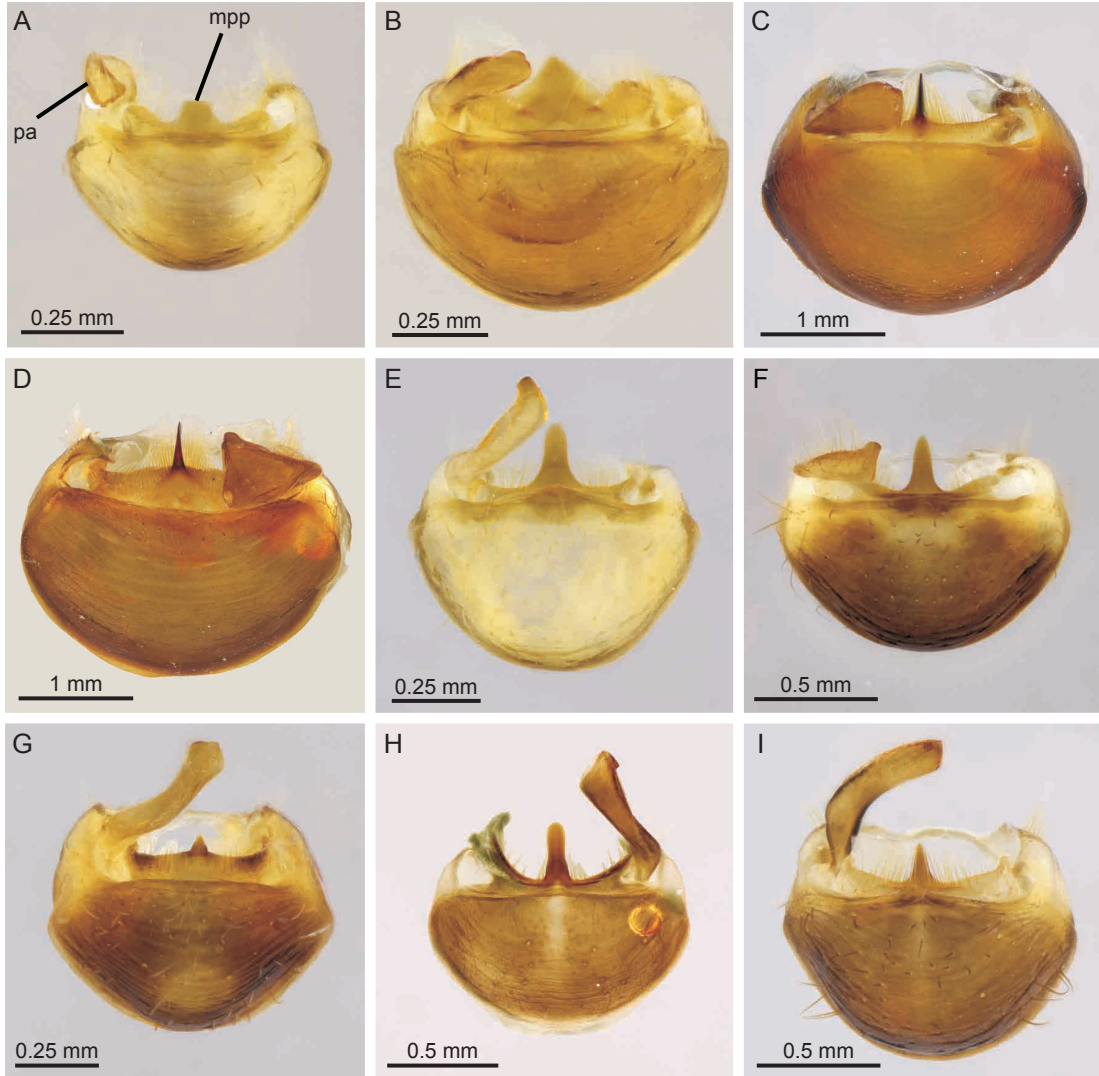


Figure 2.20. Pygophore morphology in lateral view. A. *Gibbosella notoconica*, B. *Gibbosella quadocris*, C. *Gibbosella vangocris*, D. *Glymmatophora carolae*, E. *Maraenaspis bidens*, F. *Marojejycoris auranticorium*, G. *Tanindrazanus kathrynae*, H. *Tanindrazanus marojejy*, I. *Tanindrazanus varicolor*, J. *Toliarus trichrous*, K. *Toxopus fisheri*, L. *Toxopus vazimba*.

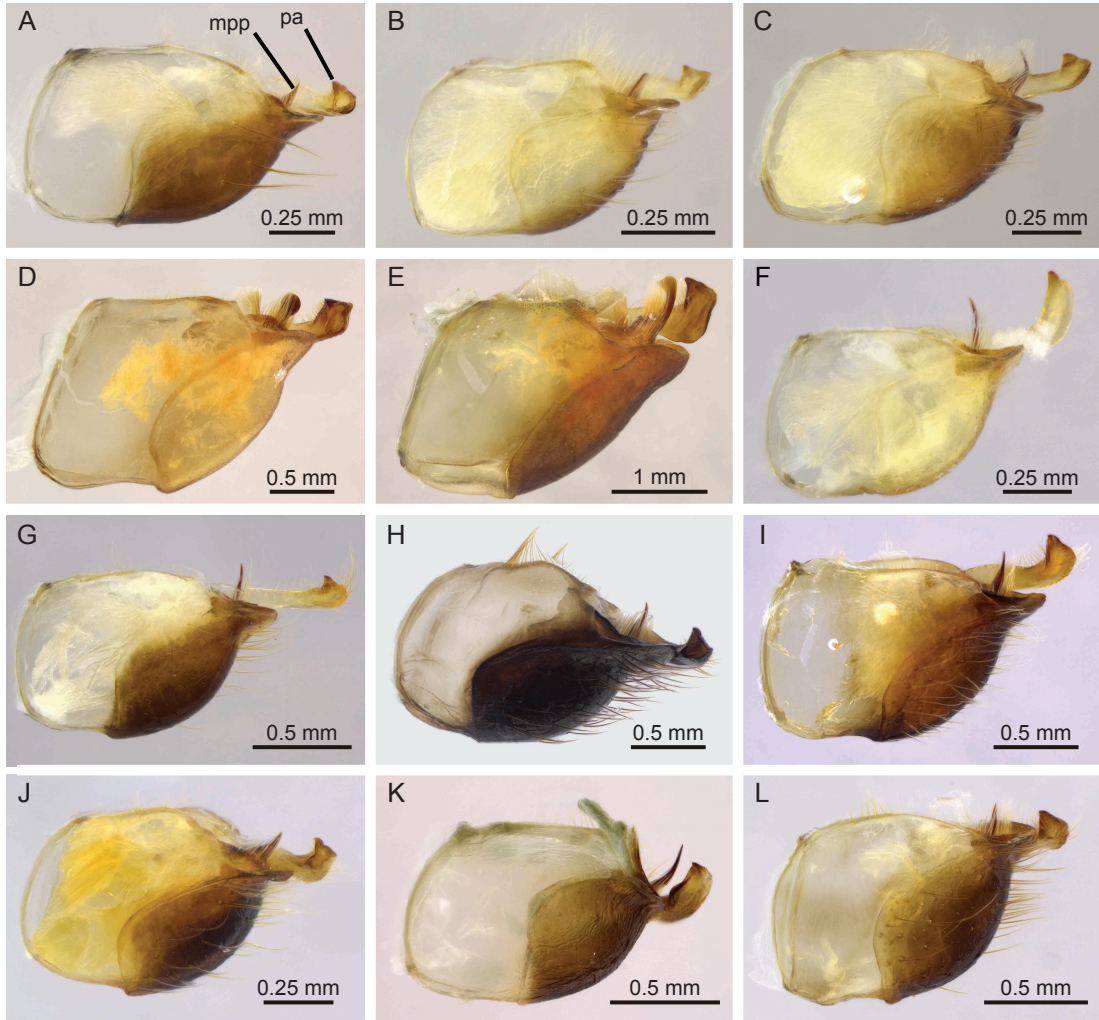


Figure 2.21. Phallus morphology in dorsal view. A. *Gibbosella notoconica*, B. *Gibbosella vangocris*, C. *Glymmatophora carolae*, D. *Glymmatophora crassipes*, E. *Maraenaspis bidens*, F. *Marojejycoris brevifrons*, G. *Tanindrazanus notatus*, H. *Tanindrazanus varicolor*, I. *Toliarus trichrous*, J. *Toxopus fisheri*, K. *Toxopus toliara*, L. *Toxopus vazimba*.



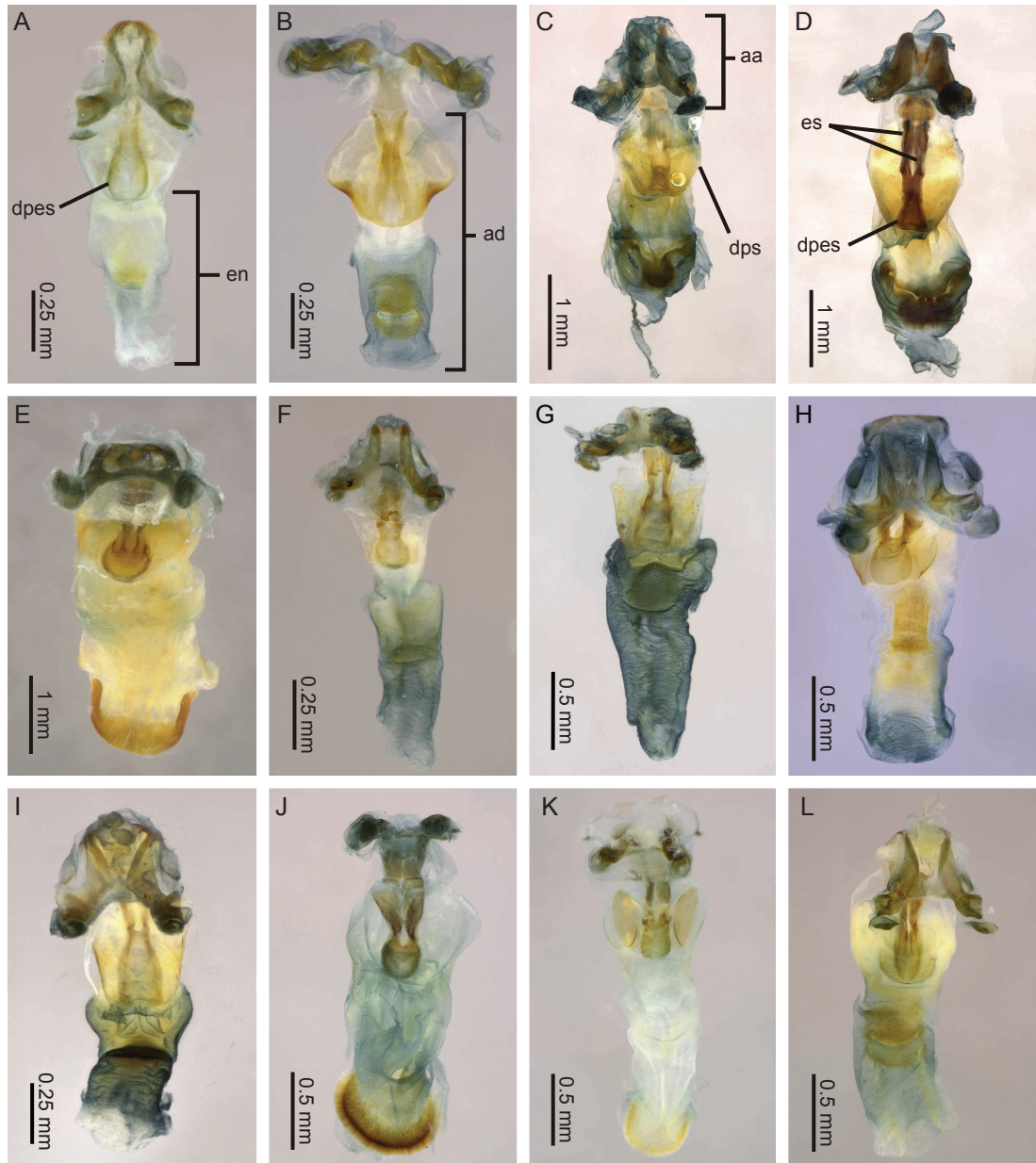


Figure 2.22. Phallus morphology in lateral view. A. *Gibbosella vangocris*, B. *Glymmatophora crassipes*, C. *Maraenaspis bidens*, D. *Marojejycoris francais*, E. *Tanindrazanus nigripes*, F. *Toliarus trichrous*, G. *Toxopus ampitavananima*, H. *Toxopus basalis*.



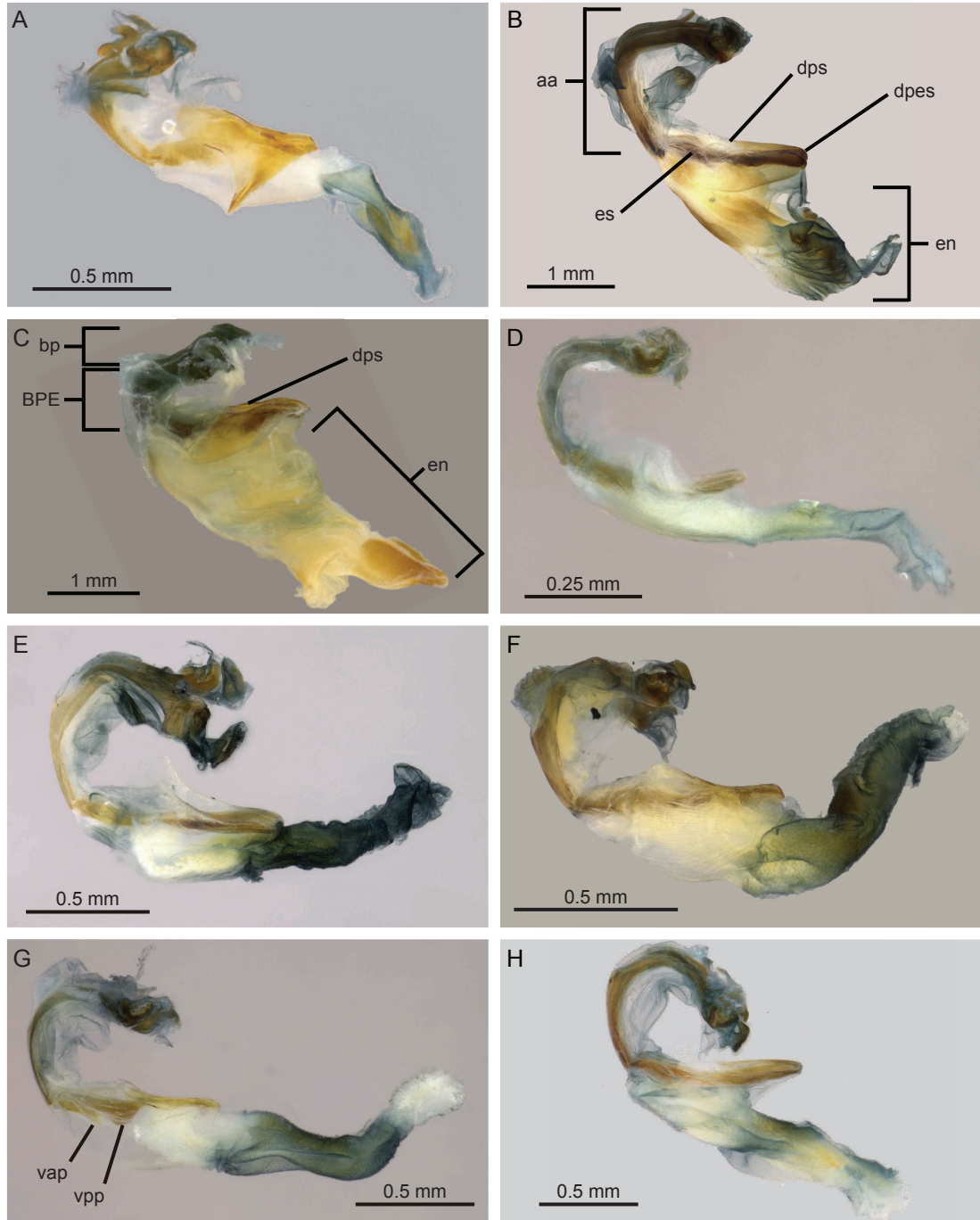


Figure 2.23. Female external (A.–E., caudal view) and internal (F., dorsal view) genitalic morphology. A. *Gibbosella brunalvus*, B. *Gibbosella planiscutum*, C. *Glymmatophora crassipes* var. 1, D. *Glymmatophora crassipes* var. 2, E. *Toxopus griswoldi*, F. *Glymmatophora crassipes* var. 2.

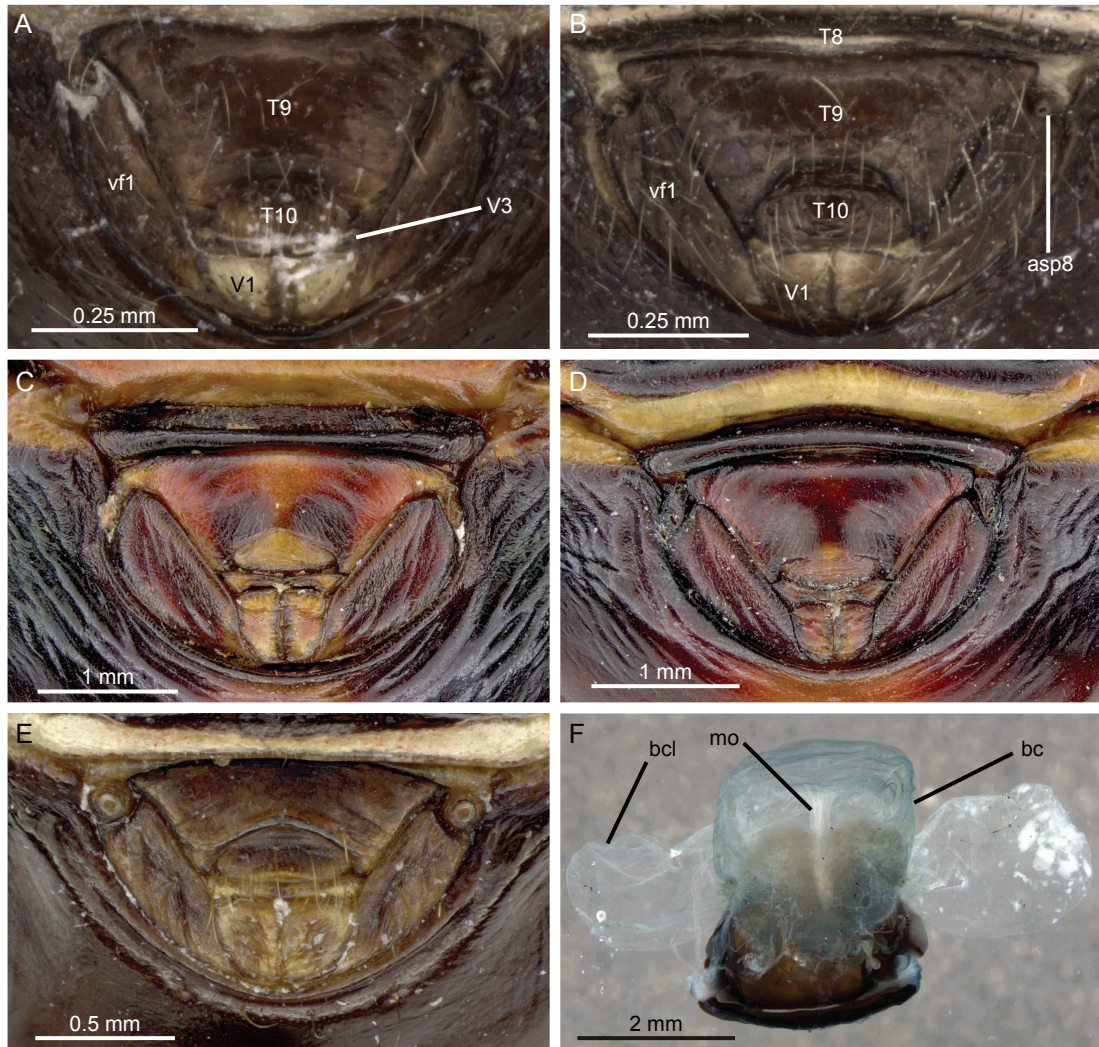
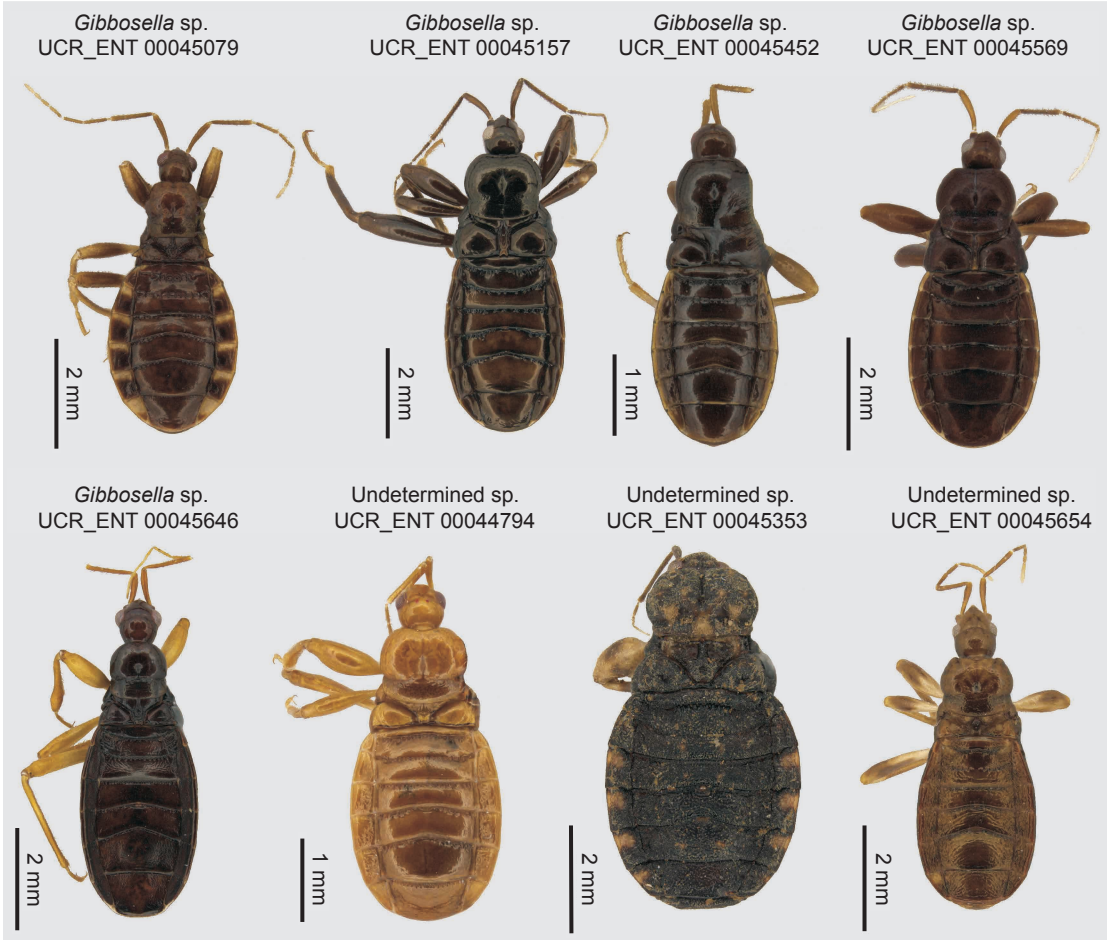
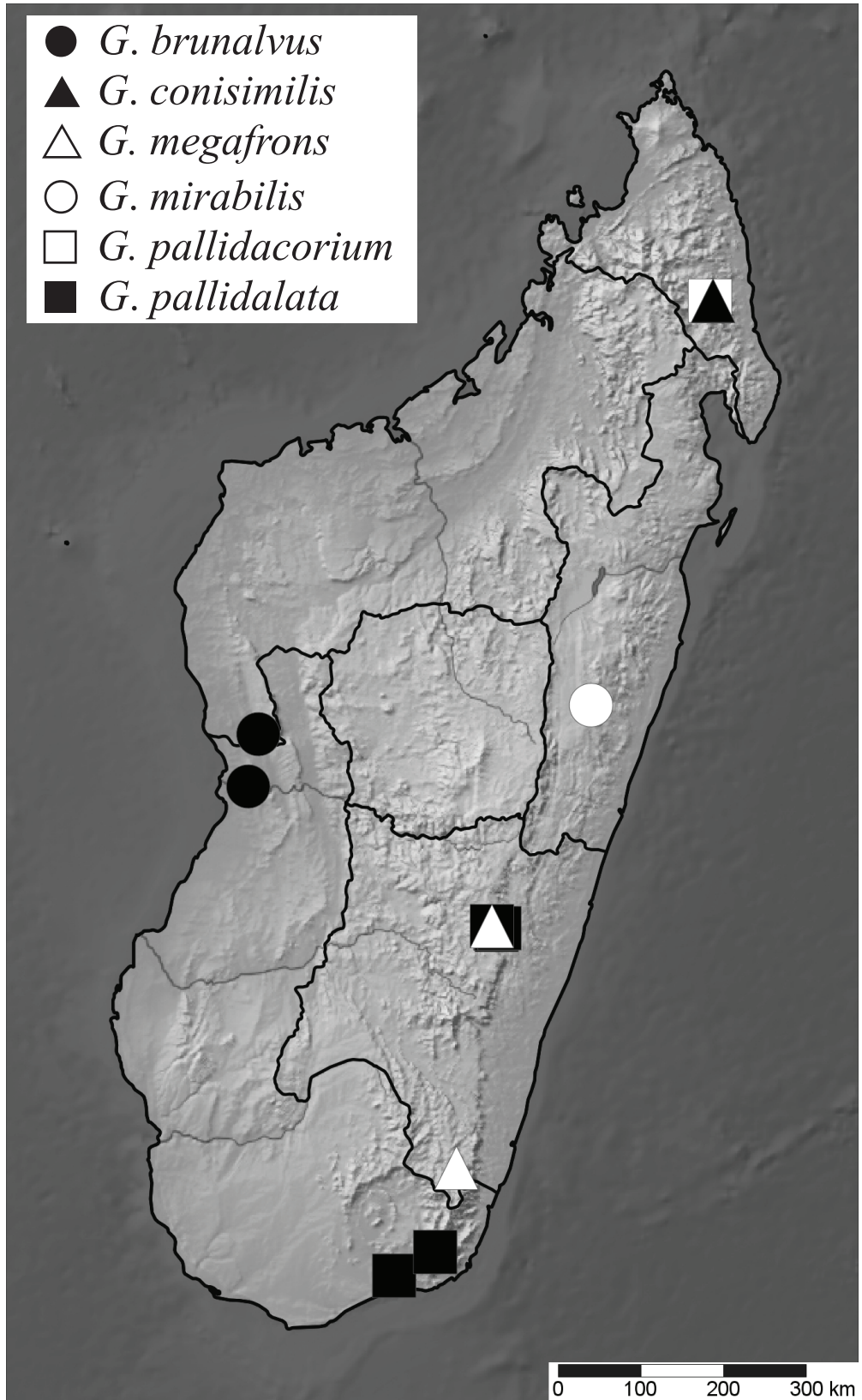


Figure 2.24. Dorsal habitus of the eight females that cannot be associated with male-based species using molecular, morphological, and geographic data. Five females are assigned to *Gibbosella* based on morphological similarities, while three females cannot be assigned to any genera.



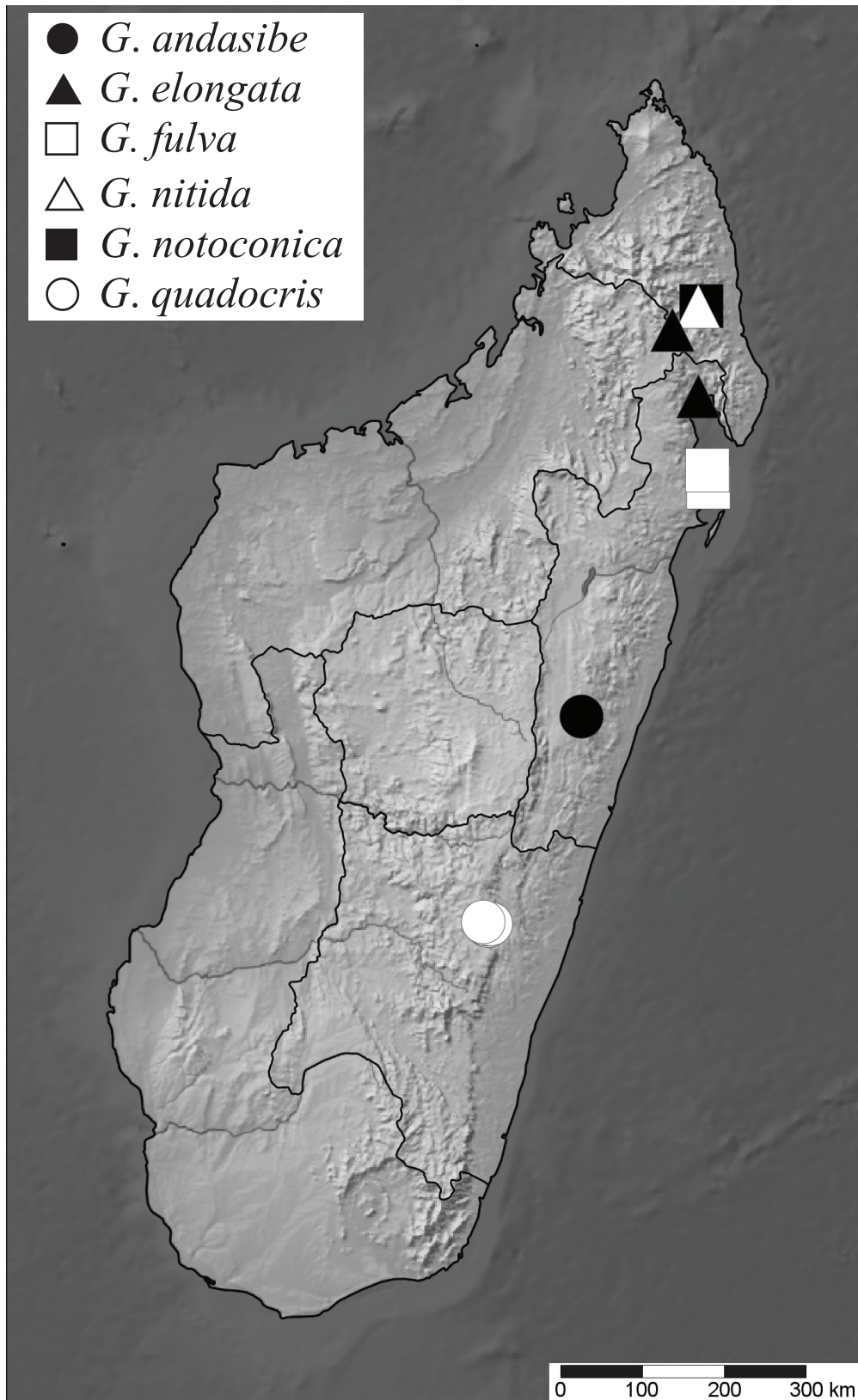
Map 2.1. Localities of *Gibbosella brunalvus*, *G. conisimilis*, *G. megafrons*, *G. mirabilis*, *G. pallidacorium*, and *G. pallidalata*.



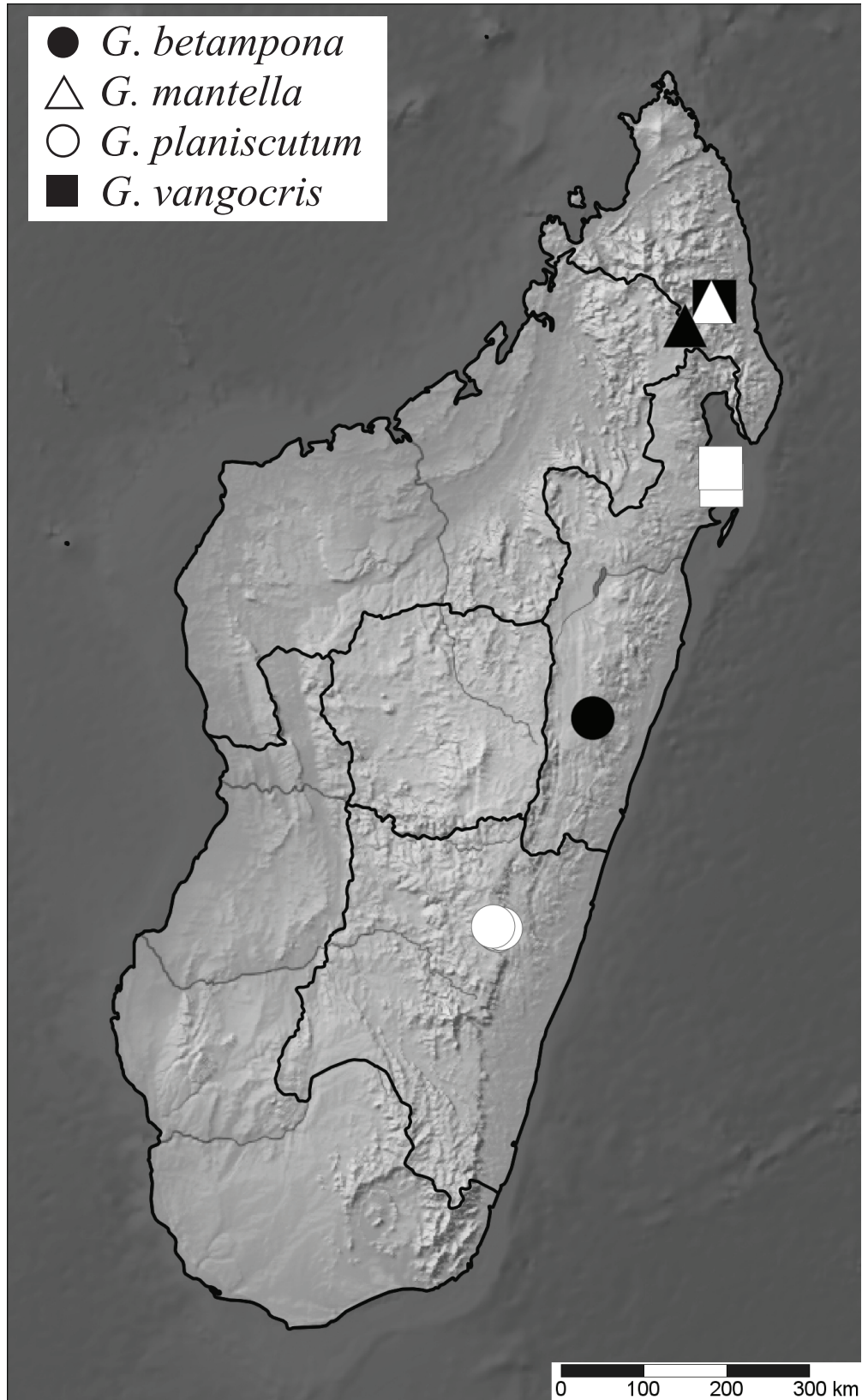


Map 2.2. Localities of *Gibbosella andasibe*, *G. elongata*, *G. fulva*, *G. nitida*, *G. notoconica* and *G. quadocris*.

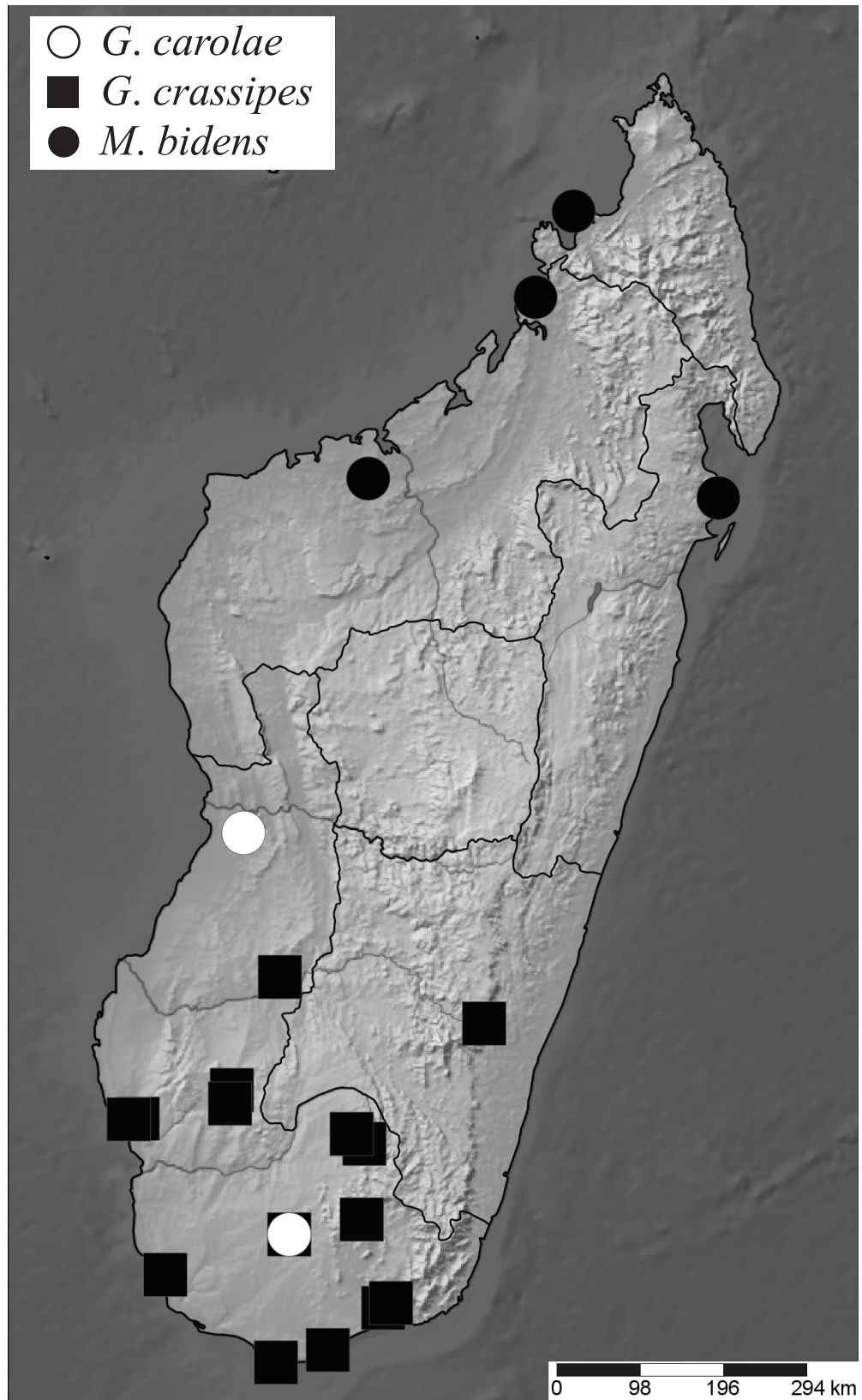




Map 2.3. Localities of *Gibbosella betampona*, *G. mantella*, *G. planiscutum*, and *G. vangocris*.

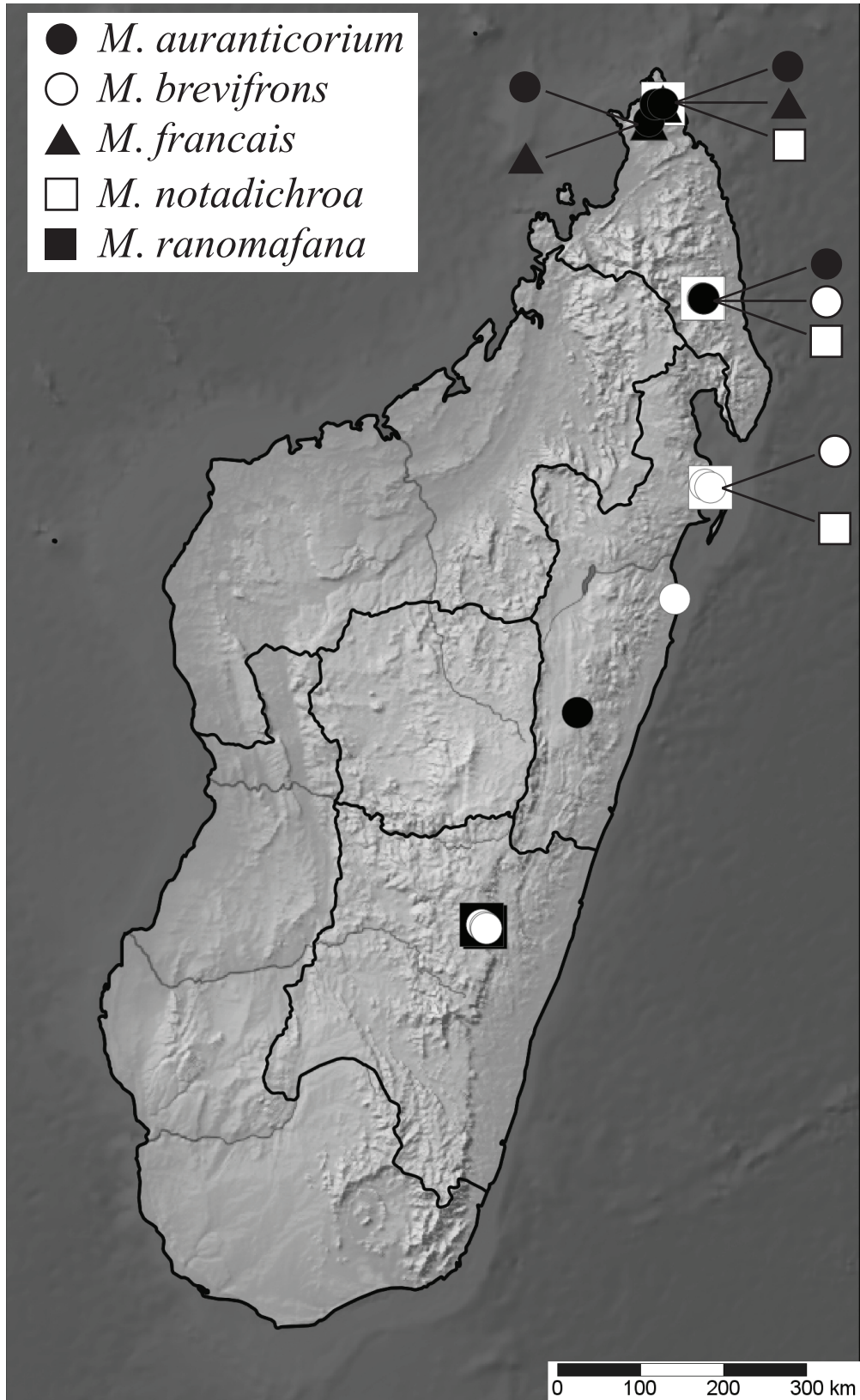


Map 2.4. Localities of *Glymmatophora* species and *Maraenaspis bidens*.



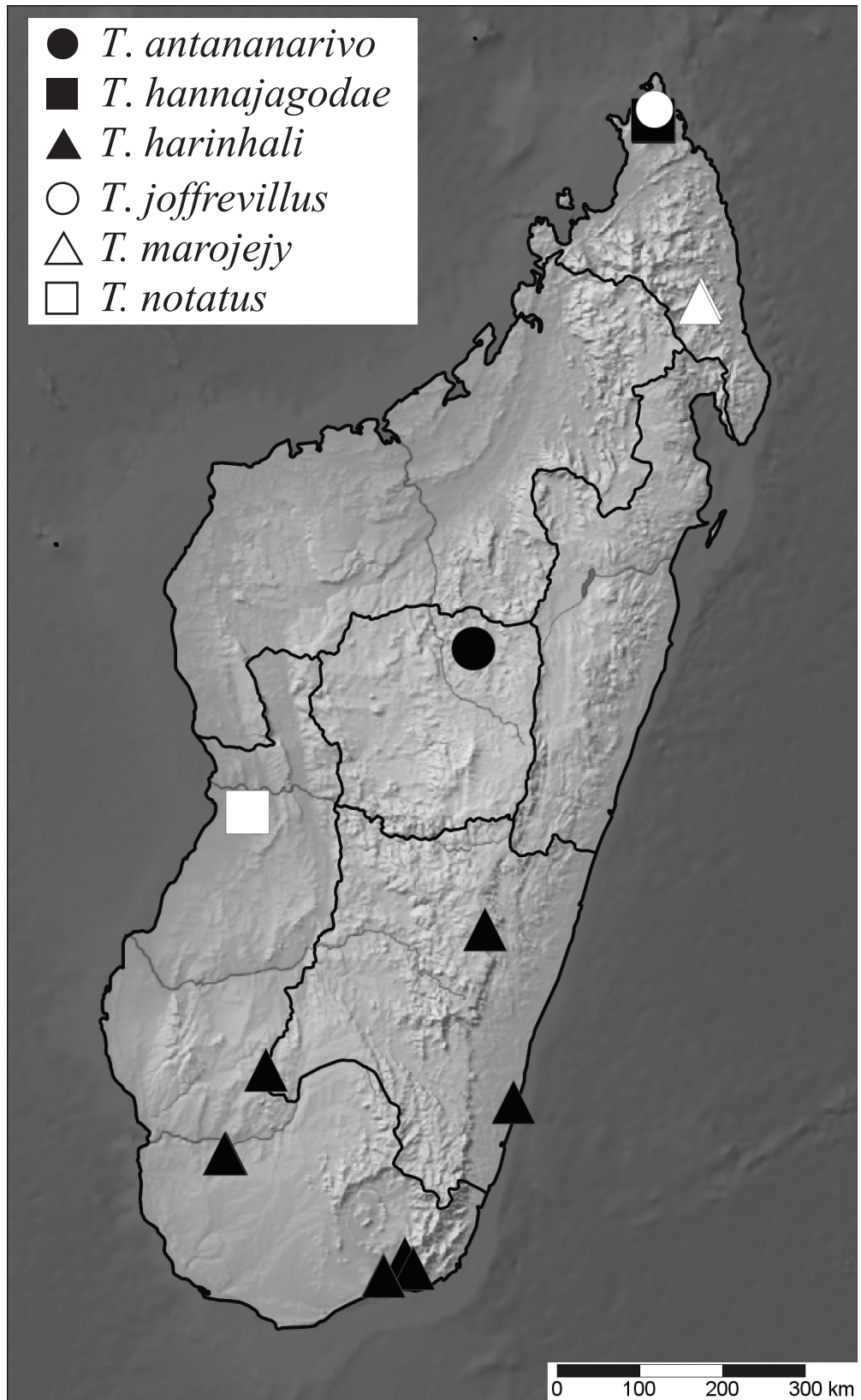
Map 2.5. Localities of *Marojejycoris* species.



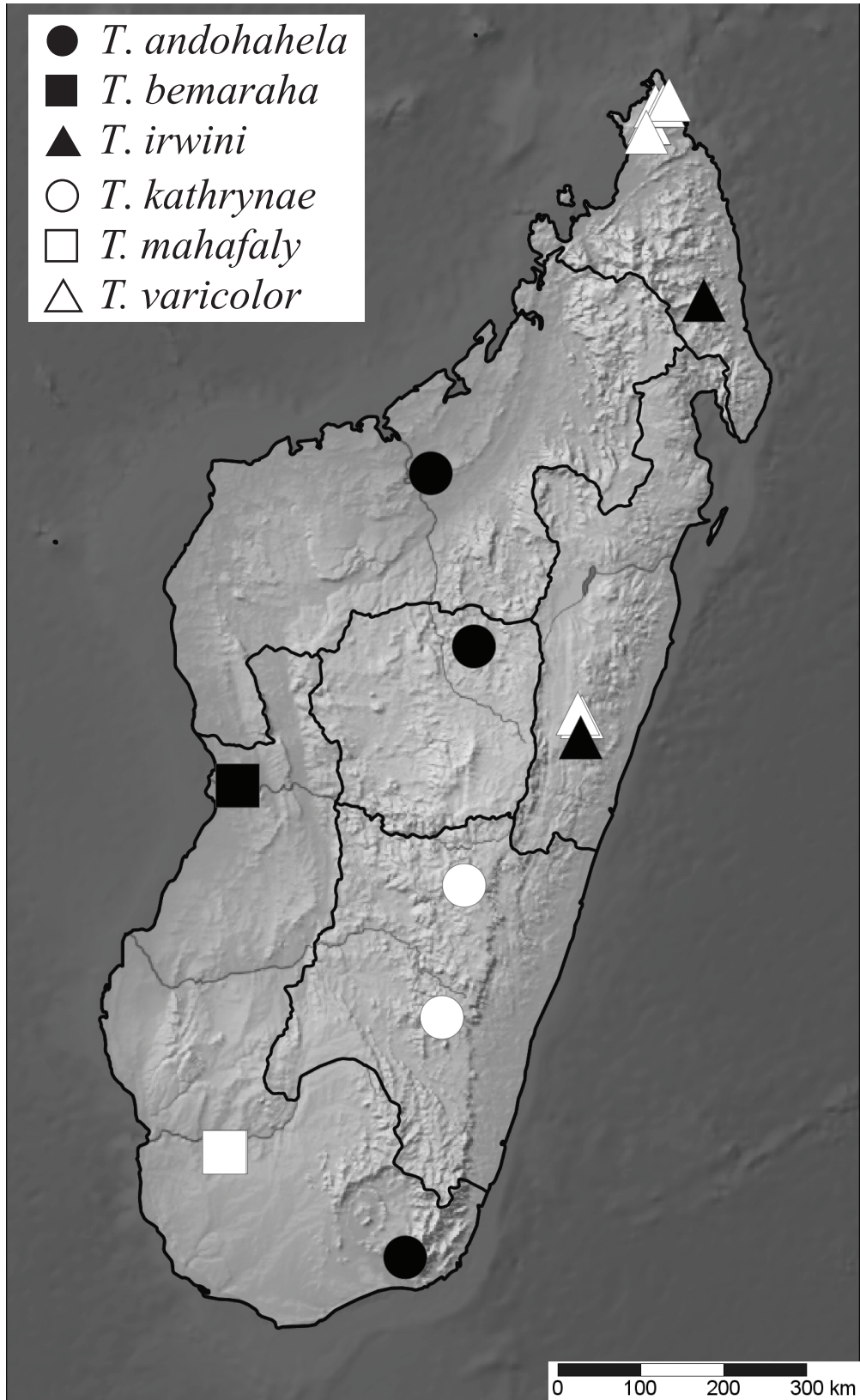


Map 2.6. Localities of *Tanindrazanus antananarivo*, *T. hannajagodae*, *T. harinhali*, *T. joffrevillus*, *T. marojejy*, and *T. notatus*.

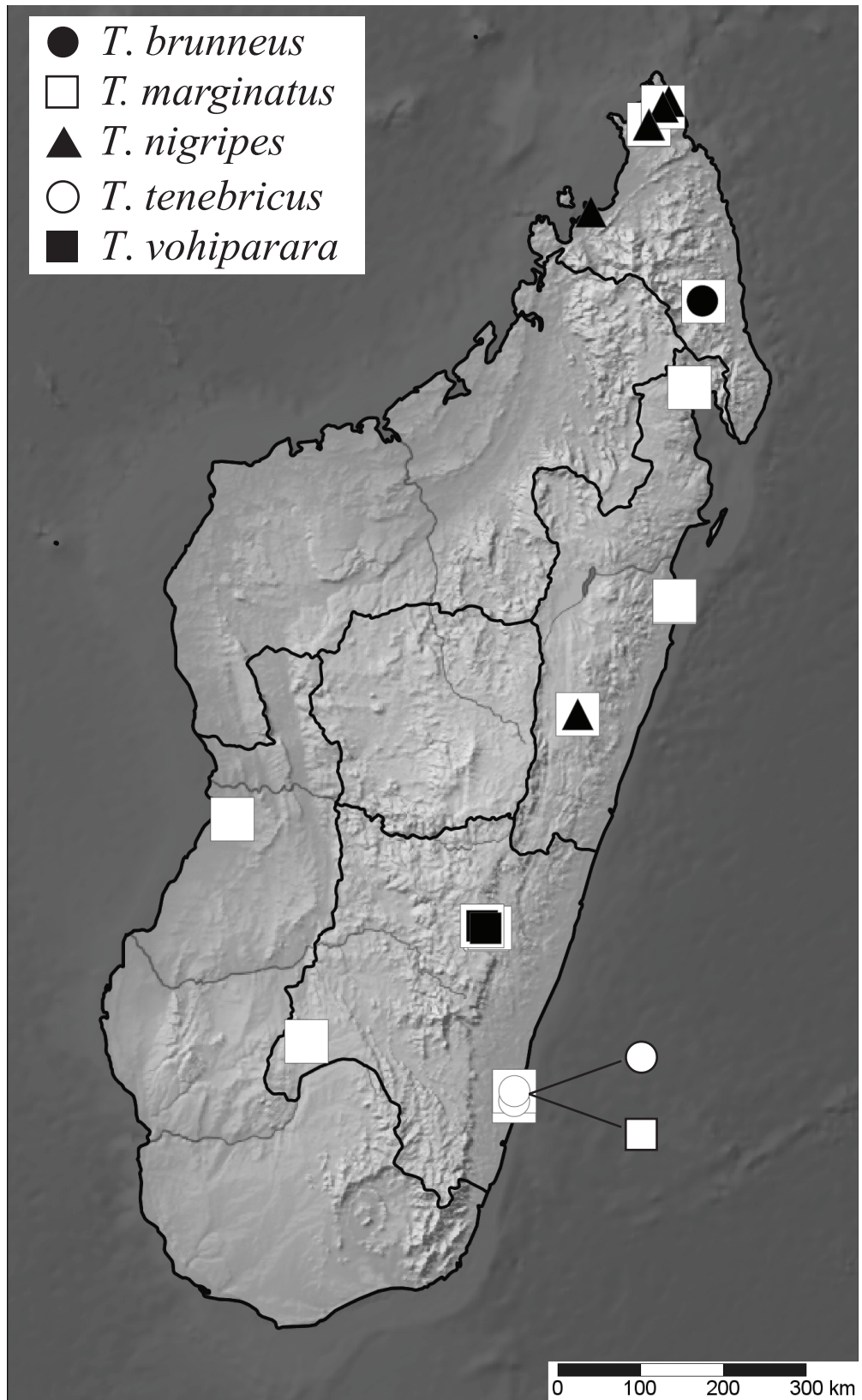




Map 2.7. Localities of *Tanindrazanus andohahela*, *T. bemaraha*, *T. irwini*, *T. kathrynae*, *T. mahafaly*, and *T. varicolor*.



Map 2.8. Localities of *Tanindrazanus brunneus*, *T. marginatus*, *T. nigripes*, *T. tenebricus*, and *T. vohiparara*.



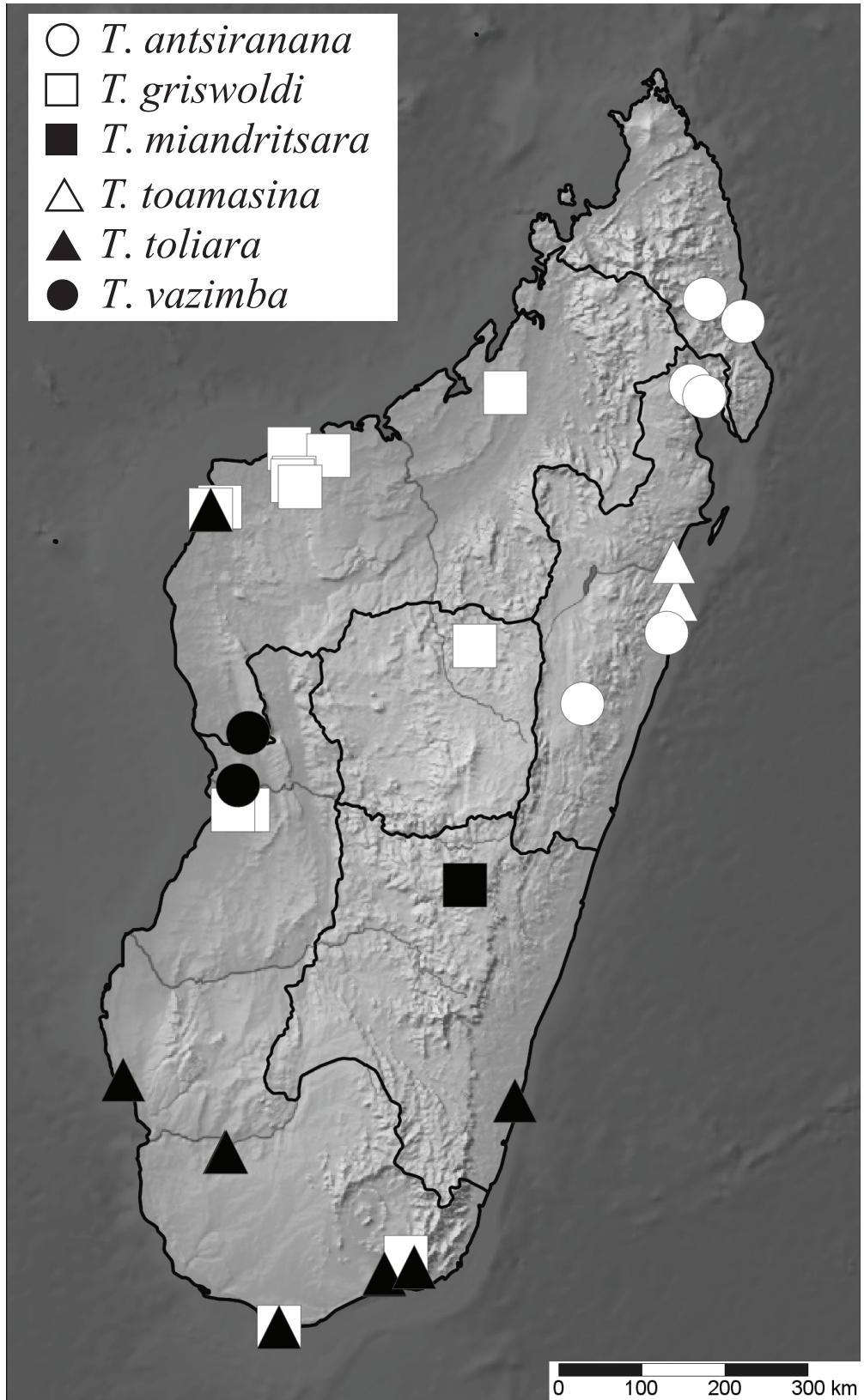
Map 2.9. Localities of *Tanindrazanus amboasaricus*, *Ta. anjzorobeus*, and *Toliarus* (*To.*) species.



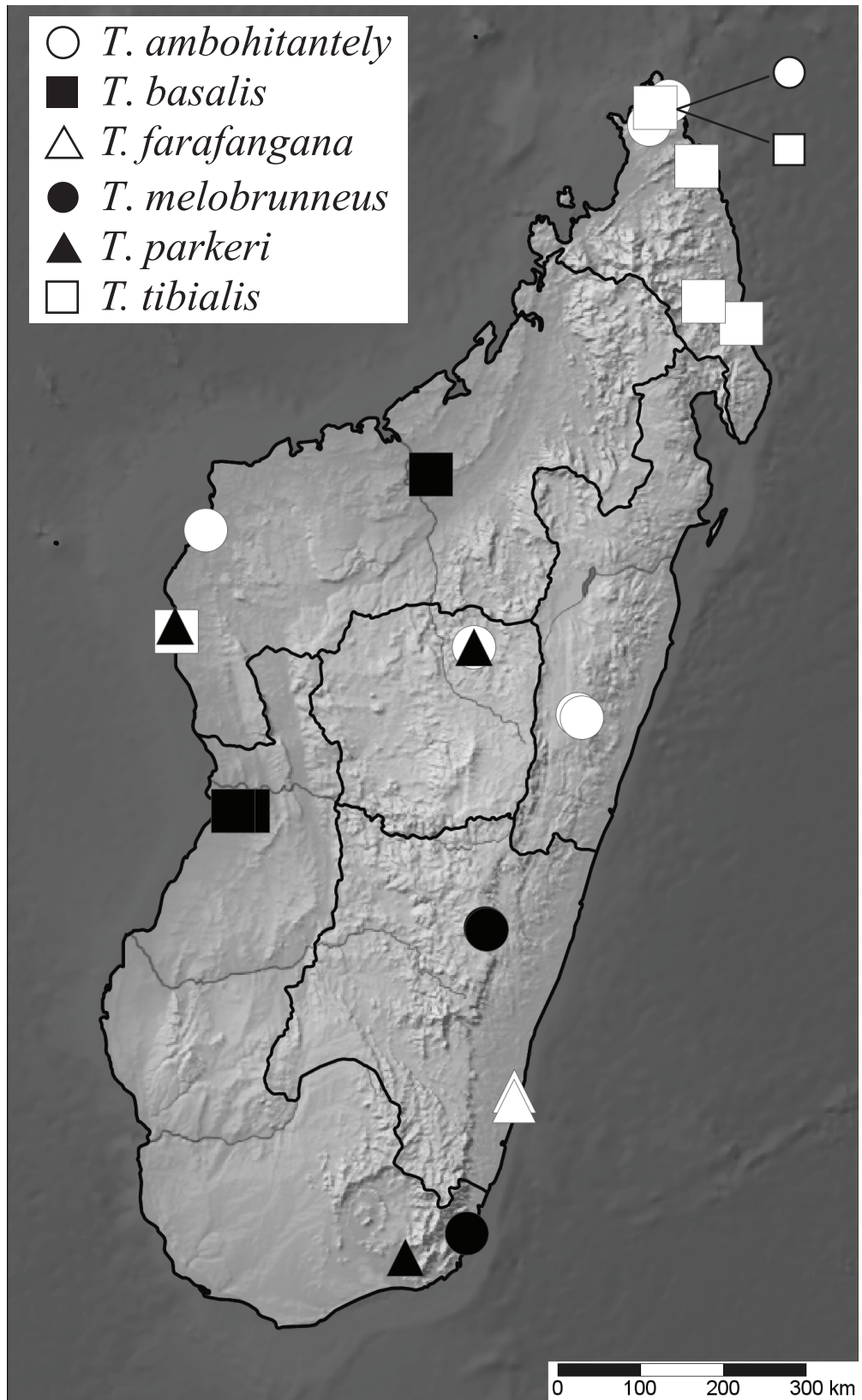


Map 2.10. Localities of *Toxopus antsiranana*, *T. griswoldi*, *T. miandritsara*, *T. toamasina*, *T. toliara*, and *T. vazimba*.

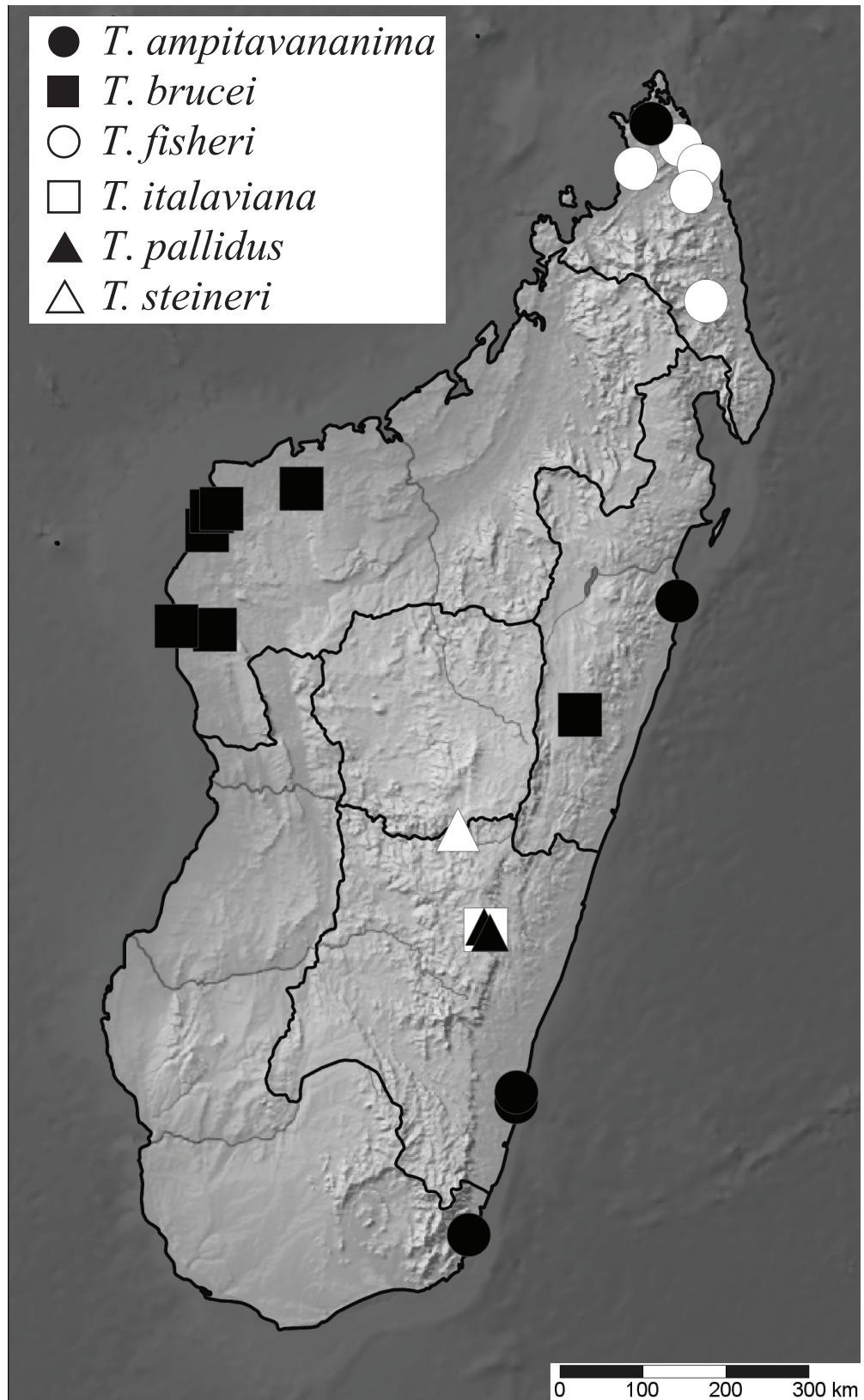




Map 2.11. Localities of *Toxopus ambohitantely*, *T. basalis*, *T. farafangana*, *T. melobrunneus*, *T. parkeri*, and *T. tibialis*.



Map 2.12. Localities of *Toxopus ampitavananima*, *T. brucei*, *T. fisheri*, *T. italaviana*, *T. pallidus*, and *T. steineri*.



Map 2.13. Localities of *Toxopus insignis*, *T. namoroka*, *T. politus*, *T. signoretii*, and *T. simulans*.



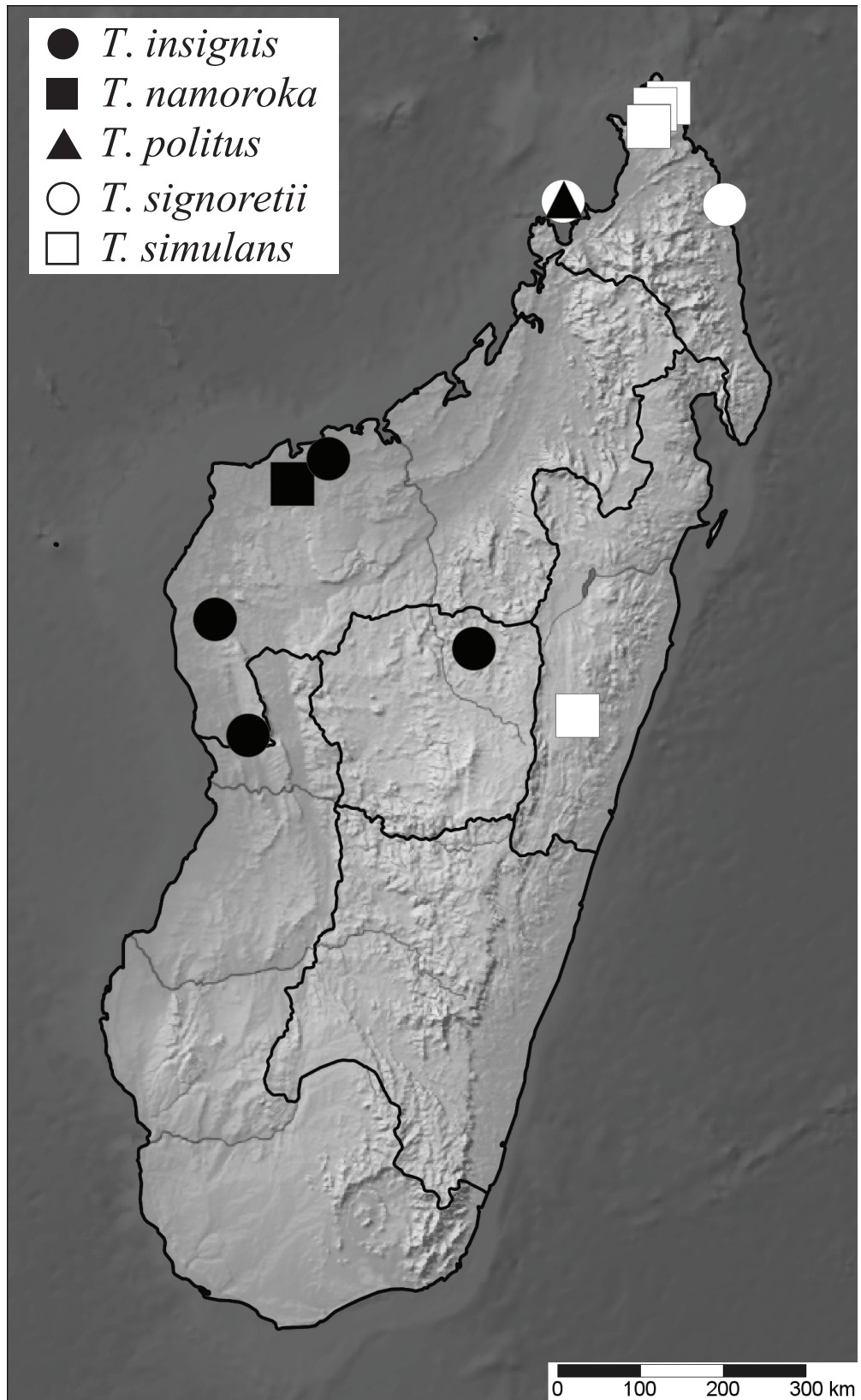


Table 2.1. Measurements (in mm). Abbreviations: Cly-Abd, clypeus to abdomen; AOc, anteocular; POc, postocular; APr, anterior pronotal lobe; PPr, posterior pronotal lobe; Scu, scutellum; Sca, scape; Ped, pedicel; Syn, synthlipsis; SD, standard deviation.

Species	Length											Width				
	Total	Clyp-Abd	Head	AOc	POc	APr	PPr	Scu	Sca	Ped	Head	Syn	APr	PPr	Abd	
<i>Gibbosella andasibe</i> n. sp.																
Male (N = 1)	Measurements	4.31	3.64	0.60	0.11	0.23	0.30	0.52	0.36	0.77	0.76	0.55	0.31	0.65	1.08	1.25
<i>Gibbosella betampona</i> n. sp.																
Male (N = 1)	Measurements	7.02	6.40	0.91	0.24	0.23	0.54	0.82	0.75	1.16	1.35	0.88	0.34	1.19	1.77	2.28
<i>Gibbosella brunahvus</i> n. sp.																
Male (N = 5)	Mean	5.05	4.93	0.85	0.21	0.27	0.44	0.67	0.55	0.95	1.04	0.69	0.36	0.93	1.36	1.60
	SD	0.10	0.15	0.03	0.02	0.02	0.02	0.04	0.05	0.03	0.02	0.03	0.01	0.04	0.07	0.06
	Range	0.24	0.34	0.06	0.04	0.05	0.04	0.09	0.13	0.06	0.05	0.07	0.02	0.09	0.15	0.13
	Min.	4.92	4.79	0.81	0.20	0.25	0.42	0.62	0.49	0.91	1.03	0.66	0.34	0.88	1.30	1.54
	Max.	5.16	5.13	0.88	0.24	0.30	0.46	0.71	0.62	0.98	1.08	0.73	0.36	0.97	1.45	1.67
Female (N = 1)	Measurements	4.54	4.54	0.92	0.25	0.30	0.69	0.22	0.31	0.64	0.69	0.64	0.39	0.97	0.90	1.79
<i>Gibbosella conisimilis</i> n. sp.																
Male (N = 3)	Mean	8.82	8.07	1.32	0.32	0.36	0.61	1.14	0.99	1.22	1.62	1.10	0.51	1.34	2.19	2.76
	SD	0.25	0.32	0.05	0.00	0.01	0.03	0.06	0.13	0.03	0.02	0.02	0.03	0.05	0.09	0.16
	Range	0.49	0.58	0.10	0.01	0.02	0.06	0.12	0.25	0.06	0.04	0.04	0.07	0.10	0.17	0.30
	Min.	8.61	7.86	1.27	0.32	0.36	0.57	1.10	0.88	1.20	1.60	1.08	0.48	1.29	2.12	2.65
	Max.	9.10	8.44	1.37	0.33	0.37	0.64	1.21	1.12	1.26	1.64	1.12	0.54	1.40	2.30	2.95
<i>Gibbosella elongata</i> Chlond, 2010																
Male (N = 1)	Measurements	10.50	9.90	1.70	0.62	0.47	0.80	1.28	0.91	1.89	2.21	1.40	0.48	1.49	2.44	3.45
<i>Gibbosella fulva</i> n. sp.																
Male (N = 2)	Mean	7.58	7.39	1.14	0.37	0.42	0.71	1.02	0.80	1.03	1.56	0.90	0.39	1.41	2.21	2.44
	SD	0.03	0.02	0.02	0.01	0.00	0.01	0.01	0.00	0.03	0.01	0.01	0.00	0.01	0.01	0.02
	Range	0.04	0.03	0.03	0.01	0.00	0.01	0.02	0.00	0.04	0.02	0.01	0.00	0.01	0.02	0.03
	Min.	7.56	7.37	1.12	0.36	0.42	0.70	1.01	0.80	1.01	1.55	0.90	0.39	1.41	2.20	2.42
	Max.	7.60	7.40	1.15	0.37	0.42	0.71	1.03	0.80	1.05	1.57	0.91	0.39	1.42	2.22	2.45
<i>Gibbosella mantella</i> n. sp.																
Male (N = 1)	Measurements	5.79	5.42	0.88	0.19	0.21	0.53	0.70	0.61	1.14	1.21	0.85	0.27	1.07	1.52	1.96



Table 2.1. (Continued).

Species	Length											Width			
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd
<i>Gibbosella megalfrons</i> n. sp.															
Male (N = 2)	Mean	4.94	4.42	0.75	0.19	0.27	0.37	0.93	1.02	1.01	0.60	0.35	0.78	1.31	1.58
	SD	1.00	0.90	0.14	0.08	0.04	0.04	0.52	0.07	0.12	0.07	0.00	0.11	0.28	0.38
	Range	1.41	1.27	0.20	0.12	0.06	0.06	0.73	0.10	0.18	0.10	0.00	0.15	0.39	0.54
	Min.	4.24	3.78	0.65	0.13	0.24	0.34	0.57	0.97	0.92	0.55	0.35	0.70	1.12	1.31
	Max.	5.65	5.05	0.85	0.25	0.30	0.40	1.30	1.07	1.10	0.65	0.35	0.85	1.51	1.85
<i>Gibbosella nitida</i> n. sp.															
Male (N = 1)	Measurements	5.30	4.75	0.90	0.33	0.35	0.45	0.62	0.85	0.89	0.65	0.35	0.90	1.45	1.61
<i>Gibbosella notoconica</i> n. sp.															
Male (N = 2)	Mean	9.62	9.01	1.37	0.34	0.33	0.64	1.27	1.53	1.90	1.29	0.48	1.35	2.27	3.31
	SD	0.67	0.67	0.09	0.02	0.04	0.01	0.09	0.06	0.19	0.13	0.04	0.08	0.15	0.27
	Range	0.95	0.95	0.13	0.02	0.05	0.02	0.13	0.09	0.26	0.18	0.06	0.11	0.21	0.39
	Min.	9.15	8.53	1.31	0.33	0.31	0.63	1.21	1.49	1.76	1.20	0.45	1.30	2.17	3.11
	Max.	10.09	9.48	1.43	0.35	0.36	0.65	1.33	1.57	2.03	1.39	0.52	1.40	2.38	3.50
<i>Gibbosella pallidacorum</i> n. sp.															
Male (N = 1)	Measurements	6.76	5.78	0.84	0.20	0.23	0.35	0.81	1.04	1.34	0.86	0.30	0.97	1.67	1.89
<i>Gibbosella pallidalata</i> n. sp.															
Male (N = 5)	Mean	5.68	5.18	0.90	0.24	0.29	0.41	0.70	1.07	1.10	0.66	0.36	0.96	1.52	1.83
	SD	0.22	0.18	0.07	0.03	0.04	0.01	0.05	0.03	0.03	0.02	0.01	0.03	0.04	0.07
	Range	0.57	0.46	0.18	0.08	0.10	0.03	0.15	0.09	0.08	0.04	0.03	0.07	0.09	0.15
	Min.	5.36	4.88	0.82	0.20	0.23	0.39	0.63	1.02	1.06	0.63	0.34	0.91	1.47	1.75
	Max.	5.93	5.34	1.00	0.27	0.33	0.42	0.78	1.11	1.14	0.67	0.37	0.98	1.56	1.89
<i>Gibbosella planiscutum</i> n. sp.															
Male (N = 5)	Mean	5.50	5.32	0.88	0.24	0.27	0.52	0.68	0.95	1.08	0.72	0.38	1.02	1.47	1.83
	SD	0.27	0.38	0.08	0.03	0.03	0.02	0.05	0.07	0.07	0.03	0.03	0.06	0.10	0.17
	Range	0.68	0.99	0.20	0.06	0.08	0.05	0.13	0.16	0.16	0.08	0.07	0.15	0.20	0.42
	Min.	5.28	4.91	0.77	0.21	0.24	0.50	0.60	0.85	0.98	0.68	0.35	0.97	1.38	1.67
	Max.	5.96	5.90	0.97	0.27	0.32	0.55	0.73	1.01	1.14	0.76	0.42	1.12	1.58	2.09

Table 2.1. (Continued).

Species	Length											Width			
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sca	Ped	Head	Syn	APr	PPr	Abd
<i>Gibbosella planiscutum</i> n. sp.															
Female (N = 5)	Mean	4.92	4.92	0.86	0.26	0.27	0.71	0.25	0.33	0.71	0.75	0.42	1.07	0.96	1.86
	SD	0.20	0.20	0.01	0.04	0.02	0.01	0.03	0.04	0.04	0.07	0.03	0.03	0.04	0.06
	Range	0.28	0.28	0.02	0.05	0.03	0.02	0.05	0.05	0.06	0.10	0.04	0.04	0.06	0.08
	Min.	4.78	4.78	0.85	0.23	0.25	0.70	0.23	0.30	0.68	0.70	0.41	1.05	0.93	1.82
	Max.	5.06	5.06	0.87	0.29	0.29	0.73	0.28	0.36	0.74	0.80	0.42	1.09	0.99	1.90
<i>Gibbosella quadoeris</i> n. sp.															
Female (N = 5)	Mean	6.46	5.57	0.95	0.25	0.30	0.49	0.78	0.65	1.06	1.22	0.37	1.02	1.61	1.93
	SD	0.22	0.20	0.06	0.04	0.02	0.01	0.04	0.05	0.06	0.08	0.02	0.04	0.08	0.04
	Range	0.56	0.46	0.15	0.11	0.04	0.03	0.09	0.11	0.15	0.21	0.05	0.10	0.21	0.11
	Min.	6.23	5.33	0.88	0.21	0.28	0.47	0.76	0.59	1.00	1.12	0.35	0.97	1.53	1.87
	Max.	6.79	5.78	1.03	0.33	0.32	0.50	0.85	0.70	1.15	1.33	0.39	1.07	1.74	1.99
<i>Gibbosella vangoeris</i> n. sp.															
Male (N = 5)	Mean	8.20	7.25	1.22	0.38	0.36	0.60	0.99	0.87	1.40	1.59	0.42	1.26	2.04	2.60
	SD	0.43	0.30	0.06	0.04	0.02	0.02	0.09	0.04	0.05	0.11	0.04	0.05	0.08	0.13
	Range	1.15	0.81	0.14	0.08	0.05	0.06	0.23	0.11	0.12	0.29	0.11	0.13	0.20	0.36
	Min.	7.55	6.80	1.14	0.34	0.34	0.57	0.85	0.83	1.34	1.48	0.39	1.17	1.91	2.38
	Max.	8.70	7.61	1.28	0.42	0.39	0.63	1.09	0.95	1.45	1.77	0.46	1.31	2.11	2.74
<i>Glymmatophora carolae</i> n. sp.															
Male (N = 4)	Mean	15.77	15.77	2.22	0.80	0.63	1.75	2.58	1.30	2.40	2.94	1.22	3.74	5.32	5.66
	SD	0.84	0.84	0.45	0.24	0.08	0.18	0.14	0.17	0.23	0.31	0.05	0.09	0.14	0.63
	Range	2.01	2.01	0.88	0.46	0.18	0.34	0.35	0.36	0.48	0.68	0.11	0.19	0.31	1.39
	Min.	14.90	14.90	1.82	0.56	0.53	1.57	2.42	1.17	2.26	2.69	1.19	3.67	5.20	5.10
	Max.	16.91	16.91	2.70	1.02	0.71	1.91	2.77	1.53	2.74	3.37	1.30	3.86	5.51	6.49
<i>Glymmatophora crassipes</i> Horváth, 1914															
Male (N = 5)	Mean	17.94	17.94	3.19	1.43	0.52	3.23	1.41	1.23	2.25	2.56	1.27	4.85	4.83	7.85
	SD	1.54	1.54	0.32	0.16	0.07	0.34	0.14	0.13	0.29	0.35	0.09	0.39	0.42	0.99
	Range	4.08	4.08	0.70	0.31	0.19	0.85	0.34	0.30	0.70	0.77	0.45	0.90	0.99	2.40
	Min.	15.52	15.52	2.80	1.29	0.45	2.70	1.28	1.12	1.90	2.13	1.10	4.20	4.11	6.30
	Max.	19.60	19.60	3.50	1.60	0.64	3.55	1.62	1.42	2.60	2.90	1.33	5.10	5.10	8.70

Table 2.1. (Continued).

Species	Length											Width				
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd	
<i>Glymmatophora crassipes</i> Horváth, 1914																
Female (N = 5)	Mean	20.78	20.78	3.36	1.44	0.85	3.57	1.57	1.51	2.36	2.52	2.53	1.43	5.40	5.27	9.29
	SD	2.24	2.24	0.41	0.28	0.23	0.50	0.13	0.12	0.05	0.16	0.30	0.22	0.68	0.80	1.18
	Range	5.60	5.60	1.12	0.72	0.65	1.26	0.30	0.28	0.10	0.30	0.70	0.50	1.56	1.75	2.80
	Min.	18.30	18.30	2.88	1.18	0.55	2.84	1.40	1.37	2.30	2.40	2.20	1.20	4.64	4.45	7.90
	Max.	23.90	23.90	4.00	1.90	1.20	4.10	1.70	1.65	2.40	2.70	2.90	1.70	6.20	6.20	10.70
<i>Maranaspis bidens</i> (Reuter), 1887																
Male (N = 2)	Mean	21.85	21.85	3.25	1.26	0.64	3.93	1.33	1.63	3.33	3.75	2.66	1.53	5.11	5.03	9.21
	SD	0.32	0.32	0.20	0.21	0.02	0.54	0.01	0.03	0.05	0.19	0.08	0.05	0.32	0.31	0.34
	Range	0.45	0.45	0.28	0.30	0.03	0.77	0.01	0.04	0.07	0.27	0.11	0.08	0.45	0.44	0.48
	Min.	21.63	21.63	3.11	1.41	0.65	3.55	1.34	1.61	3.30	3.61	2.72	1.57	4.88	4.81	8.97
	Max.	22.08	22.08	3.39	1.41	0.65	4.32	1.34	1.65	3.37	3.88	2.72	1.57	5.34	5.25	9.45
Female (N = 2)																
	Mean	22.15	22.15	3.65	1.60	1.15	3.74	1.72	1.37	2.90	—	2.78	1.61	4.88	4.14	8.26
	SD	1.20	1.20	0.21	0.14	0.07	0.09	0.04	0.06	0.00	—	0.14	0.03	0.03	0.05	0.07
	Range	1.70	1.70	0.30	0.20	0.10	0.13	0.06	0.08	0.00	—	0.20	0.04	0.04	0.07	0.10
	Min.	21.30	21.30	3.50	1.50	1.10	3.67	1.69	1.33	2.90	—	2.78	1.61	4.88	4.10	8.21
	Max.	23.00	23.00	3.80	1.70	1.20	3.80	1.75	1.41	2.90	—	2.98	1.65	4.92	4.17	8.31
<i>Marojejcoris auranticorinum</i> n. sp.																
Male (N = 4)	Mean	9.35	8.96	1.71	0.57	0.57	0.66	1.16	0.77	1.56	1.82	1.20	0.59	1.66	2.46	2.84
	SD	0.34	0.29	0.07	0.02	0.03	0.10	0.06	0.06	0.07	0.09	0.04	0.06	0.04	0.04	0.14
	Range	0.69	0.64	0.16	0.04	0.07	0.22	0.12	0.14	0.18	0.20	0.08	0.14	0.10	0.10	0.30
	Min.	8.96	8.54	1.64	0.55	0.53	0.54	1.13	0.68	1.47	1.69	1.14	0.53	1.60	2.41	2.70
	Max.	9.65	9.18	1.80	0.59	0.59	0.76	1.25	0.82	1.65	1.88	1.22	0.67	1.69	2.51	3.00
<i>Marojejcoris brevifrons</i> n. sp.																
Male (N = 5)	Mean	7.19	6.58	1.23	0.30	0.34	0.51	0.87	0.66	1.26	1.33	0.97	0.35	1.28	1.88	2.16
	SD	0.96	0.66	0.08	0.04	0.02	0.05	0.07	0.07	0.09	0.11	0.04	0.02	0.11	0.13	0.17
	Range	2.28	1.50	0.21	0.11	0.05	0.13	0.17	0.18	0.22	0.26	0.10	0.06	0.22	0.29	0.41
	Min.	5.63	5.63	1.08	0.24	0.31	0.44	0.79	0.60	1.13	1.18	0.93	0.31	1.15	1.73	1.99
	Max.	7.91	7.13	1.29	0.35	0.36	0.57	0.95	0.77	1.34	1.44	1.02	0.37	1.36	2.02	2.40

Table 2.1. (Continued).

Species	Length											Width			
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd
<i>Marojejcoris francis</i> n. sp.															
Female (N = 2)															
Mean	6.99	6.50	1.15	0.39	0.34	0.50	0.90	0.63	0.93	1.27	0.89	0.44	1.34	1.92	2.19
SD	0.62	0.46	0.04	0.05	0.06	0.01	0.04	0.08	0.09	0.11	0.06	0.00	0.11	0.15	0.16
Range	0.87	0.65	0.06	0.07	0.09	0.02	0.05	0.12	0.12	0.15	0.09	0.01	0.15	0.22	0.22
Min.	6.55	6.18	1.12	0.35	0.30	0.49	0.87	0.57	0.87	1.19	0.84	0.44	1.27	1.81	2.08
Max.	7.42	6.82	1.18	0.42	0.39	0.51	0.93	0.69	1.00	1.34	0.93	0.44	1.42	2.03	2.30
<i>Marojejcoris notadichroa</i> n. sp.															
Male (N = 5)															
Mean	8.46	7.94	1.44	0.48	0.47	0.65	1.03	0.78	1.29	1.46	1.35	0.56	1.56	2.19	2.54
SD	0.45	0.54	0.09	0.03	0.05	0.02	0.08	0.03	0.08	0.05	0.05	0.04	0.11	0.18	0.31
Range	1.01	1.33	0.24	0.08	0.12	0.05	0.22	0.07	0.21	0.13	1.50	0.09	0.25	0.46	0.71
Min.	8.08	7.40	1.33	0.44	0.41	0.62	0.90	0.74	1.22	1.41	1.00	0.53	1.42	1.95	2.18
Max.	9.09	8.73	1.57	0.52	0.53	0.67	1.12	0.81	1.43	1.54	2.51	0.62	1.67	2.41	2.88
<i>Marojejcoris ranomafana</i> n. sp.															
Male (N = 2)															
Mean	6.20	5.46	1.01	0.25	0.38	0.48	0.71	0.55	1.12	1.18	0.70	0.37	1.10	1.56	1.87
SD	0.18	0.06	0.01	0.02	0.00	0.07	0.01	0.07	0.04	0.05	0.03	0.00	0.00	0.01	0.04
Range	0.25	0.08	0.02	0.03	0.00	0.10	0.01	0.10	0.06	0.07	0.04	0.00	0.00	0.01	0.06
Min.	6.08	5.41	1.00	0.24	0.37	0.43	0.70	0.50	1.09	1.15	0.68	0.37	1.10	1.56	1.84
Max.	6.33	5.50	1.02	0.27	0.38	0.52	0.71	0.61	1.15	1.21	0.72	0.37	1.10	1.57	1.90
<i>Tanindrazanus amboasarisicus</i> n. sp.															
Male (N = 4)															
Mean	15.91	15.06	2.74	1.11	0.69	1.19	1.92	1.17	2.29	2.99	2.03	0.75	2.47	3.83	5.03
SD	1.25	0.85	0.27	0.09	0.07	0.10	0.14	0.08	0.09	0.20	0.15	0.11	0.04	0.31	0.36
Range	2.70	1.75	0.60	0.22	0.17	0.20	0.30	0.18	0.20	0.45	0.30	0.25	0.09	0.65	0.80
Min.	14.60	14.30	2.45	1.02	0.62	1.10	1.80	1.05	2.20	2.75	1.90	0.60	2.42	3.50	4.70
Max.	17.30	16.05	3.05	1.24	0.79	1.30	2.10	1.23	2.40	3.20	2.20	0.85	2.51	4.15	5.50
<i>Tanindrazanus andohahela</i> n. sp.															
Male (N = 4)															
Mean	10.83	10.77	1.94	0.73	0.49	0.88	1.35	0.90	1.52	1.99	1.46	0.65	2.01	2.86	3.44
SD	0.42	0.32	0.12	0.10	0.03	0.06	0.04	0.04	0.08	0.08	0.06	0.04	0.10	0.11	0.15
Range	1.03	0.77	0.25	0.22	0.05	0.13	0.10	0.10	0.18	0.18	0.14	0.09	0.19	0.25	0.32
Min.	10.34	10.34	1.78	0.58	0.47	0.79	1.31	0.86	1.41	1.89	1.38	0.60	1.92	2.75	3.22
Max.	11.36	11.10	2.04	0.80	0.53	0.92	1.41	0.96	1.59	2.07	1.52	0.69	2.11	3.00	3.54

Table 2.1. (Continued).

Species	Length											Width			
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd
<i>Tanindrazanus anjzorobeus</i> n. sp.															
Male (N = 3)															
Mean	17.02	16.91	2.99	0.97	1.07	1.01	2.23	1.55	2.32	3.08	2.37	0.74	2.71	4.32	5.72
SD	0.45	0.61	0.33	0.13	0.22	0.12	0.20	0.04	0.04	0.02	0.01	0.02	0.08	0.03	0.11
Range	0.83	1.12	0.59	0.23	0.41	0.22	0.35	0.07	0.08	0.04	0.02	0.04	0.15	0.05	0.22
Min.	16.50	16.20	2.61	0.83	0.81	0.88	2.00	1.51	2.27	3.06	2.36	0.72	2.65	4.30	5.60
Max.	17.33	17.32	3.20	1.06	1.22	1.10	2.35	1.58	2.35	3.10	2.38	0.76	2.80	4.35	5.82
<i>Tanindrazanus antananarivo</i> n. sp.															
Male (N = 5)															
Mean	13.26	12.81	2.20	0.79	0.61	0.87	1.58	1.14	1.90	2.39	1.64	0.65	2.09	3.30	4.27
SD	0.82	0.67	0.17	0.09	0.08	0.12	0.11	0.08	0.09	0.11	0.08	0.02	0.09	0.19	0.39
Range	1.99	1.59	0.43	0.23	0.20	0.32	0.26	0.18	0.22	0.29	0.20	0.06	0.21	0.46	1.00
Min.	12.71	12.41	2.04	0.70	0.48	0.71	1.45	1.07	1.82	2.28	1.58	0.63	2.04	3.17	3.88
Max.	14.69	14.00	2.47	0.93	0.69	1.03	1.71	1.25	2.04	2.56	1.78	0.69	2.25	3.63	4.88
<i>Tanindrazanus bemaraha</i> n. sp.															
Male (N = 2)															
Mean	13.50	13.50	2.48	0.96	0.62	1.09	1.63	1.15	2.12	2.77	1.69	0.80	2.27	3.45	4.40
SD	0.61	0.61	0.17	0.01	0.06	0.05	0.09	0.00	0.00	0.08	0.00	0.05	0.04	0.06	0.07
Range	0.86	0.86	0.23	0.02	0.09	0.06	0.13	0.01	0.01	0.11	0.01	0.07	0.06	0.08	0.09
Min.	13.07	13.07	2.36	0.95	0.57	1.05	1.56	1.14	2.11	2.71	1.68	0.77	2.24	3.41	4.35
Max.	13.93	13.93	2.60	0.97	0.66	1.12	1.69	1.15	2.12	2.83	1.69	0.83	2.30	3.50	4.44
<i>Tanindrazanus brunneus</i> n. sp.															
Male (N = 1)															
Measurements	10.77	10.31	1.91	0.79	0.54	0.75	1.41	1.11	1.90	2.44	1.26	0.70	1.89	2.86	3.54
<i>Tanindrazanus hannajogodae</i> n. sp.															
Male (N = 3)															
Mean	23.33	23.18	3.33	1.35	0.74	1.73	3.34	2.12	3.52	4.51	2.76	1.25	3.83	6.41	7.98
SD	0.75	0.64	0.27	0.12	0.05	0.17	0.12	0.21	0.10	0.09	0.03	0.04	0.12	0.29	0.09
Range	1.37	1.22	0.51	0.22	0.09	0.31	0.24	0.42	0.18	0.18	0.06	0.08	0.24	0.58	0.17
Min.	22.47	22.47	3.02	1.26	0.68	1.63	3.24	1.93	3.46	4.44	2.73	1.21	3.69	6.10	7.91
Max.	23.84	23.69	3.53	1.48	0.77	1.93	3.48	2.35	3.63	4.62	2.79	1.29	3.94	6.68	8.08

Table 2.1. (Continued).

Species	Length											Width				
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd	
<i>Tanindrazanus harinhali</i> n. sp.																
Male (N = 5)	Mean	11.35	11.01	1.77	0.61	0.53	0.89	1.62	0.83	1.72	2.13	1.44	0.70	2.05	3.08	3.60
	SD	0.58	0.59	0.10	0.04	0.04	0.09	0.09	0.11	0.10	0.12	0.09	0.07	0.15	0.19	0.25
	Range	1.61	1.50	0.29	0.10	0.09	0.24	0.25	0.22	0.28	0.31	0.23	0.17	0.36	0.51	0.60
	Min.	10.57	10.39	1.62	0.55	0.50	0.76	1.50	0.70	1.57	1.96	1.34	0.63	1.92	2.87	3.37
	Max.	12.18	11.89	1.91	0.65	0.59	1.00	1.75	0.92	1.85	2.27	1.57	0.80	2.28	3.38	3.97
<i>Tanindrazanus irwini</i> n. sp.																
Male (N = 5)	Mean	21.28	21.05	3.31	1.40	0.73	1.39	2.91	1.65	3.25	4.19	2.54	1.13	3.54	5.57	6.44
	SD	1.31	1.37	0.26	0.21	0.09	0.10	0.25	0.18	0.09	0.10	0.08	0.06	0.17	0.32	0.49
	Range	3.44	3.82	0.62	0.52	0.23	0.24	0.64	0.39	0.22	0.26	0.21	0.14	0.45	0.86	1.26
	Min.	19.32	18.95	2.91	1.08	0.64	1.32	2.54	1.52	3.11	4.05	2.42	1.05	3.28	5.04	5.83
	Max.	22.76	22.76	3.53	1.60	0.87	1.56	3.18	1.90	3.33	4.31	2.64	1.19	3.73	5.90	7.09
<i>Tanindrazanus joffrevillus</i> n. sp.																
Male (N = 1)	Measurements	11.19	11.19	1.99	0.70	0.59	1.03	1.37	1.00	1.82	2.36	1.58	0.83	2.15	2.95	3.84
<i>Tanindrazanus kathrynae</i> n. sp.																
Male (N = 5)	Mean	12.61	12.27	2.25	0.87	0.65	0.89	1.66	1.02	1.69	2.35	1.48	0.73	2.08	3.26	3.86
	SD	0.41	0.39	0.04	0.05	0.02	0.07	0.15	0.12	0.12	0.21	0.06	0.05	0.11	0.17	0.11
	Range	1.08	1.05	0.10	0.14	0.06	0.16	0.40	0.28	0.30	0.50	0.17	0.12	0.29	0.42	0.31
	Min.	12.20	11.62	2.22	0.80	0.62	0.81	1.41	0.87	1.57	2.22	1.41	0.66	1.95	3.01	3.69
	Max.	13.28	12.67	2.32	0.93	0.67	0.97	1.81	1.15	1.88	2.72	1.58	0.78	2.24	3.43	3.99
<i>Tanindrazanus mahafaly</i> n. sp.																
Male (N = 5)	Mean	18.96	18.96	2.72	0.91	0.76	1.32	2.49	1.82	2.64	3.24	2.37	0.98	3.21	5.12	6.08
	SD	0.77	0.77	0.13	0.08	0.05	0.07	0.17	0.16	0.08	0.13	0.19	0.03	0.09	0.18	0.19
	Range	1.89	1.89	0.35	0.19	0.12	0.17	0.40	0.32	0.20	0.32	0.47	0.08	0.22	0.45	0.42
	Min.	18.03	18.03	2.50	0.77	0.68	1.24	2.29	1.64	2.53	3.05	2.24	0.94	3.09	4.86	5.88
	Max.	19.92	19.92	2.85	0.97	0.81	1.40	2.69	1.96	2.72	3.37	2.71	1.02	3.31	5.31	6.30

Table 2.1. (Continued).

Species	Length											Width			
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd
<i>Tanindrazanus marginatus</i> n. sp.															
Male (N = 5)															
Mean	15.53	15.33	2.66	0.97	0.58	0.97	1.96	1.33	2.25	2.80	2.08	0.76	2.43	3.95	5.51
SD	1.81	1.67	0.23	0.06	0.08	0.09	0.25	0.10	0.32	0.33	0.20	0.06	0.20	0.43	0.68
Range	4.01	3.81	0.58	0.16	0.17	0.19	0.53	0.26	0.66	0.69	0.43	0.14	0.46	0.94	1.40
Min.	13.62	13.62	2.42	0.92	0.50	0.89	1.69	1.21	1.90	2.44	1.85	0.69	2.22	3.45	4.78
Max.	17.63	17.43	3.00	1.07	0.68	1.07	2.21	1.47	2.56	3.14	2.29	0.84	2.68	4.39	6.18
<i>Tanindrazanus marojejy</i> n. sp.															
Male (N = 4)															
Mean	18.43	18.22	2.96	1.32	0.73	1.47	2.59	1.25	2.88	3.72	2.05	0.92	3.25	4.97	6.57
SD	0.98	1.19	0.31	0.15	0.17	0.19	0.19	0.06	0.39	0.30	0.10	0.16	0.16	0.10	0.43
Range	2.40	2.67	0.75	0.33	0.41	0.44	0.38	0.12	0.95	0.73	0.22	0.39	0.32	0.21	0.95
Min.	17.20	17.05	2.61	1.12	0.52	1.28	2.42	1.19	2.37	3.32	1.93	0.71	3.10	4.89	6.15
Max.	19.60	19.72	3.36	1.45	0.93	1.72	2.80	1.31	3.32	4.05	2.15	1.10	3.42	5.10	7.10
<i>Tanindrazanus nigripes</i> n. sp.															
Male (N = 5)															
Mean	12.72	12.26	2.20	0.93	0.55	0.90	1.56	1.06	1.77	2.42	1.56	0.74	2.20	3.23	4.00
SD	0.43	0.33	0.07	0.06	0.09	0.09	0.06	0.12	0.09	0.08	0.02	0.01	0.06	0.06	0.16
Range	0.89	0.85	0.18	0.13	0.20	0.20	0.17	0.33	0.20	0.18	0.05	0.03	0.17	0.14	0.36
Min.	12.37	11.78	2.09	0.86	0.45	0.83	1.47	0.90	1.64	2.35	1.53	0.72	2.13	3.16	3.84
Max.	13.26	12.63	2.27	0.99	0.65	1.03	1.64	1.23	1.84	2.54	1.58	0.75	2.30	3.30	4.20
<i>Tanindrazanus notatus</i> n. sp.															
Male (N = 2)															
Mean	11.48	11.20	1.85	0.66	0.45	0.92	1.52	0.92	1.79	2.19	1.44	0.69	2.04	3.03	3.61
SD	1.85	1.46	0.18	0.04	0.05	0.14	0.30	0.00	0.17	0.30	0.15	0.06	0.22	0.46	0.64
Range	2.62	2.06	0.25	0.06	0.07	0.19	0.43	0.00	0.24	0.42	0.21	0.08	0.32	0.65	0.91
Min.	10.17	10.17	1.73	0.63	0.42	0.82	1.31	0.92	1.68	1.98	1.33	0.65	1.88	2.71	3.15
Max.	12.79	12.24	1.98	0.68	0.49	1.01	1.74	0.92	1.91	2.40	1.55	0.73	2.20	3.36	4.06
<i>Tanindrazanus simulans</i> n. sp.															
Male (N = 2)															
Mean	16.10	15.25	2.79	1.36	0.69	1.34	2.11	1.18	2.37	2.68	1.89	0.81	2.43	3.93	4.93
SD	0.14	0.21	0.01	0.04	0.01	0.04	0.12	0.03	0.04	0.04	0.01	0.00	0.05	0.03	0.09
Range	0.20	0.30	0.02	0.06	0.02	0.05	0.17	0.04	0.06	0.05	0.02	0.00	0.07	0.04	0.13
Min.	16.00	15.10	2.78	1.33	0.68	1.31	2.02	1.16	2.34	2.65	1.89	0.81	2.43	3.91	4.86
Max.	16.20	15.40	2.80	1.39	0.70	1.36	2.19	1.20	2.40	2.70	1.91	0.81	2.50	3.95	4.99

Table 2.1. (Continued).

Species	Length											Width				
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd	
<i>Tanindrazanus tenebricus</i> n. sp.																
Male (N = 5)	Mean	16.15	16.12	2.80	0.97	0.74	1.25	2.01	1.35	2.55	3.23	2.34	0.87	2.74	4.13	5.48
	SD	0.64	0.62	0.22	0.18	0.06	0.07	0.16	0.08	0.11	0.16	0.06	0.03	0.10	0.21	0.28
	Range	1.63	1.63	0.56	0.43	0.15	0.18	0.36	0.21	0.29	0.41	0.14	0.07	0.23	0.59	0.63
	Min.	15.40	15.40	2.61	0.85	0.69	1.15	1.82	1.22	2.39	3.01	2.28	0.83	2.61	3.83	5.13
	Max.	17.02	17.02	3.17	1.28	0.84	1.33	2.18	1.43	2.68	3.43	2.42	0.90	2.85	4.41	5.76
<i>Tanindrazanus varicolor</i> n. sp.																
Male (N = 5)	Mean	16.31	16.00	2.53	0.98	0.64	1.06	2.26	1.18	2.31	2.98	2.01	0.94	2.58	4.20	4.95
	SD	0.18	0.08	0.05	0.09	0.07	0.02	0.03	0.04	0.07	0.09	0.04	0.04	0.05	0.09	0.11
	Range	0.41	0.23	0.13	0.19	0.18	0.06	0.06	0.08	0.18	0.23	0.09	0.11	0.12	0.22	0.27
	Min.	16.07	15.89	2.48	0.89	0.57	1.03	2.23	1.14	2.21	2.90	1.97	0.87	2.51	4.06	4.77
	Max.	16.48	16.12	2.61	1.08	0.75	1.09	2.29	1.22	2.39	3.13	2.05	0.98	2.63	4.28	5.04
<i>Tanindrazanus vohiparara</i> n. sp.																
Male (N = 1) Measurements		12.11	11.20	1.96	0.73	0.33	0.68	1.54	0.99	1.68	2.04	1.78	0.59	1.91	3.00	3.65
<i>Toliarus karinae</i> n. sp.																
Male (N = 1) Measurements		12.10	11.70	2.20	1.05	0.60	1.10	1.48	1.05	2.15	1.85	1.50	0.80	2.24	2.98	3.80
<i>Toliarus trichrous</i> n. sp.																
Male (N = 5)	Mean	8.98	8.22	1.55	0.60	0.39	0.69	1.19	0.73	1.23	1.57	1.20	0.64	1.62	2.42	2.57
	SD	0.62	0.76	0.06	0.03	0.04	0.10	0.14	0.09	0.10	0.11	0.12	0.06	0.20	0.27	0.32
	Range	1.58	1.76	0.14	0.08	0.07	0.21	0.30	0.23	0.27	0.31	0.29	0.14	0.47	0.66	0.78
	Min.	8.10	7.22	1.47	0.58	0.36	0.57	1.04	0.60	1.12	1.42	1.07	0.58	1.36	2.06	2.20
	Max.	9.68	8.97	1.61	0.65	0.43	0.78	1.34	0.83	1.39	1.73	1.36	0.72	1.83	2.72	2.98
<i>Toxoptus ambolitantely</i> n. sp.																
Male (N = 5)	Mean	10.51	9.99	1.66	0.61	0.44	0.81	1.27	1.01	1.55	1.82	1.22	0.66	1.93	2.74	3.30
	SD	0.24	0.13	0.06	0.04	0.04	0.06	0.04	0.10	0.05	0.05	0.02	0.01	0.05	0.09	0.05
	Range	0.55	0.35	0.17	0.09	0.10	0.12	0.10	0.22	0.14	0.13	0.05	0.03	0.13	0.24	0.13
	Min.	10.22	9.83	1.55	0.57	0.39	0.76	1.20	0.90	1.50	1.76	1.19	0.64	1.87	2.64	3.23
	Max.	10.78	10.18	1.72	0.65	0.48	0.87	1.30	1.12	1.63	1.89	1.24	0.67	2.00	2.88	3.36



Table 2.1. (Continued).

Species	Length											Width				
	Total	Clyp- -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd	
<i>Toxopus ampitavananima</i> n. sp.																
Male (N = 5)	Mean	11.34	11.20	1.80	0.56	0.42	1.00	1.37	1.16	1.78	2.09	1.64	0.79	2.23	3.17	3.63
	SD	0.62	0.66	0.10	0.05	0.04	0.11	0.10	0.03	0.12	0.21	0.14	0.04	0.14	0.24	0.39
	Range	1.32	1.41	0.25	0.10	0.10	0.26	0.22	0.08	0.29	0.56	0.35	0.10	0.31	0.53	0.92
	Min.	10.71	10.50	1.65	0.52	0.38	0.88	1.26	1.11	1.60	1.78	1.52	0.76	2.11	2.86	3.31
	Max.	12.03	11.91	1.91	0.62	0.48	1.13	1.48	1.20	1.89	2.33	1.87	0.85	2.42	3.39	4.22
<i>Toxopus antsiranana</i> n. sp.																
Male (N = 3)	Mean	12.18	11.81	2.12	0.76	0.47	1.05	1.57	1.10	1.76	2.19	1.58	0.66	2.26	3.24	3.99
	SD	0.61	0.62	0.09	0.07	0.04	0.07	0.12	0.10	0.21	0.26	0.14	0.05	0.10	0.10	0.38
	Range	1.19	1.11	0.17	0.14	0.07	0.14	0.22	0.17	0.38	0.52	0.25	0.10	0.17	0.19	0.66
	Min.	11.51	11.10	2.04	0.70	0.43	0.98	1.43	0.99	1.52	1.91	1.43	0.61	2.15	3.13	3.55
	Max.	12.70	12.21	2.21	0.84	0.51	1.12	1.65	1.16	1.90	2.43	1.68	0.71	2.32	3.32	4.21
<i>Tanindrazanus basalis</i> n. sp.																
Male (N = 5)	Mean	12.39	12.24	1.77	0.56	0.35	1.06	1.61	1.25	1.73	2.13	1.72	0.76	2.54	3.44	4.27
	SD	0.69	0.66	0.12	0.07	0.04	0.04	0.12	0.14	0.07	0.07	0.10	0.03	0.14	0.22	0.19
	Range	1.45	1.52	0.31	0.20	0.11	0.08	0.32	0.30	0.19	0.18	0.23	0.07	0.31	0.51	0.45
	Min.	11.62	11.49	1.57	0.45	0.30	1.02	1.45	1.13	1.65	2.02	1.62	0.73	2.40	3.21	3.97
	Max.	13.07	13.01	1.88	0.64	0.41	1.10	1.77	1.43	1.84	2.20	1.85	0.80	2.70	3.72	4.43
<i>Toxopus brucei</i> n. sp.																
Male (N = 5)	Mean	11.93	11.84	1.73	0.52	0.40	1.18	1.55	1.17	1.70	2.06	1.74	0.88	2.58	3.42	4.17
	SD	0.75	0.80	0.11	0.06	0.04	0.06	0.13	0.17	0.09	0.15	0.14	0.04	0.11	0.27	0.31
	Range	1.64	1.70	0.27	0.15	0.09	0.15	0.28	0.36	0.23	0.41	0.31	0.10	0.26	0.58	0.64
	Min.	11.06	11.00	1.59	0.44	0.37	1.10	1.41	0.99	1.57	1.85	1.59	0.84	2.44	3.14	3.87
	Max.	12.70	12.70	1.86	0.59	0.46	1.24	1.69	1.35	1.80	2.26	1.90	0.94	2.70	3.73	4.51
<i>Toxopus farafangana</i> n. sp.																
Male (N = 5)	Mean	9.89	9.81	1.64	0.46	0.41	0.95	1.18	0.97	1.36	1.71	1.38	0.58	1.91	2.72	3.52
	SD	0.72	0.66	0.17	0.06	0.04	0.09	0.11	0.14	0.09	0.13	0.08	0.04	0.15	0.21	0.27
	Range	1.73	1.61	0.42	0.18	0.10	0.23	0.30	0.37	0.22	0.34	0.21	0.09	0.36	0.50	0.68
	Min.	9.14	9.05	1.47	0.36	0.37	0.86	1.01	0.81	1.27	1.54	1.30	0.54	1.76	2.47	3.16
	Max.	10.87	10.66	1.89	0.54	0.47	1.09	1.30	1.18	1.48	1.88	1.51	0.63	2.12	2.97	3.84

Table 2.1. (Continued).

Species	Length											Width				
	Total	Clyp- -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd	
<i>Toxopus fisheri</i> n. sp.																
Male (N = 5)	Mean	10.95	10.29	1.89	0.66	0.48	0.97	1.34	0.96	1.38	1.95	1.37	0.65	2.08	2.85	3.45
	SD	0.44	0.48	0.09	0.10	0.05	0.07	0.10	0.10	0.16	0.21	0.06	0.05	0.05	0.12	0.13
	Range	1.09	1.26	0.23	0.25	0.12	0.16	0.27	0.27	0.39	0.45	0.15	0.12	0.11	0.25	0.35
	Min.	10.50	9.84	1.76	0.54	0.42	0.89	1.18	0.82	1.27	1.72	1.30	0.61	2.02	2.70	3.26
	Max.	11.59	11.09	1.99	0.79	0.55	1.05	1.45	1.09	1.67	2.17	1.45	0.72	2.13	2.95	3.61
<i>Toxopus griswoldi</i> n. sp.																
Male (N = 5)	Mean	9.61	9.39	1.76	0.59	0.47	0.89	1.24	0.84	1.32	1.66	1.39	0.73	2.05	2.68	3.23
	SD	0.41	0.42	0.08	0.08	0.04	0.06	0.07	0.18	0.07	0.08	0.06	0.03	0.08	0.13	0.28
	Range	1.01	1.02	0.21	0.21	0.09	0.15	0.18	0.48	0.17	0.22	0.16	0.09	0.24	0.32	0.69
	Min.	8.98	8.74	1.65	0.51	0.40	0.82	1.15	0.66	1.25	1.52	1.34	0.69	1.92	2.47	2.84
	Max.	9.99	9.76	1.87	0.71	0.49	0.97	1.34	1.14	1.43	1.74	1.49	0.78	2.16	2.79	3.53
Female (N = 1)	Measurements	11.75	11.75	2.73	1.02	0.76	1.68	0.68	0.47	1.28	1.70	1.77	1.13	2.66	2.60	3.88
<i>Toxopus insignis</i> n. sp.																
Male (N = 5)	Mean	11.58	10.59	1.79	0.58	0.45	0.95	1.44	0.96	1.65	2.16	1.45	0.72	2.13	3.10	3.90
	SD	0.75	0.74	0.13	0.07	0.02	0.08	0.11	0.06	0.11	0.13	0.06	0.04	0.15	0.27	0.37
	Range	1.74	1.67	0.31	0.18	0.04	0.19	0.29	0.13	0.25	0.28	0.14	0.10	0.32	0.58	0.92
	Min.	10.75	9.73	1.68	0.52	0.43	0.86	1.30	0.88	1.49	2.00	1.39	0.66	1.97	2.77	3.44
	Max.	12.49	11.41	1.99	0.70	0.47	1.05	1.60	1.01	1.74	2.27	1.52	0.76	2.29	3.35	4.36
<i>Toxopus itaviana</i> n. sp.																
Male (N = 5)	Mean	11.44	10.42	1.81	0.62	0.45	1.01	1.48	0.99	1.51	2.17	1.46	0.78	2.24	3.06	3.85
	SD	0.28	0.17	0.06	0.05	0.03	0.03	0.06	0.08	0.08	0.16	0.04	0.06	0.07	0.10	0.18
	Range	0.76	0.38	0.15	0.14	0.08	0.08	0.13	0.19	0.22	0.39	0.08	0.13	0.18	0.26	0.46
	Min.	11.09	10.25	1.71	0.54	0.42	0.96	1.41	0.91	1.37	2.05	1.43	0.69	2.17	2.95	3.60
	Max.	11.86	10.63	1.86	0.69	0.49	1.04	1.54	1.10	1.59	2.44	1.51	0.82	2.35	3.21	4.06

Table 2.1. (Continued).

Species	Length											Width				
	Total	Clyp -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd	
<i>Toxopus melobrunneus</i> n. sp.																
Male (N = 5)	Mean	14.66	14.29	2.18	0.72	0.48	0.96	1.84	1.35	2.14	2.77	1.75	0.74	2.49	3.82	4.93
	SD	0.49	0.39	0.13	0.08	0.02	0.09	0.17	0.11	0.06	0.27	0.07	0.03	0.10	0.19	0.20
	Range	1.28	1.00	0.35	0.20	0.06	0.22	0.41	0.23	0.18	0.69	0.15	0.07	0.24	0.50	0.53
	Min.	14.15	13.96	1.99	0.62	0.43	0.85	1.73	1.24	2.06	2.54	1.68	0.70	2.41	3.63	4.62
	Max.	15.43	14.95	2.33	0.82	0.50	1.07	2.13	1.47	2.24	3.23	1.83	0.78	2.65	4.13	5.15
<i>Toxopus miandritsara</i> n. sp.																
Male (N = 2)	Mean	10.05	9.79	1.58	0.54	0.39	0.81	1.20	0.83	1.31	1.64	1.20	0.59	1.81	2.71	3.24
	SD	0.44	0.39	0.01	0.01	0.04	0.00	0.04	0.04	0.07	0.10	0.00	0.03	0.08	0.15	0.01
	Range	0.62	0.55	0.02	0.02	0.06	0.01	0.05	0.06	0.11	0.14	0.00	0.04	0.11	0.22	0.01
	Min.	9.73	9.51	1.57	0.53	0.36	0.80	1.17	0.80	1.25	1.57	1.19	0.58	1.76	2.61	3.23
	Max.	10.36	10.06	1.58	0.55	0.42	0.81	1.23	0.86	1.36	1.72	1.20	0.61	1.87	2.82	3.24
<i>Toxopus namoroka</i> n. sp.																
Male (N = 2)	Mean	11.50	11.50	1.84	0.64	0.44	1.13	1.52	1.24	1.79	2.37	1.68	0.88	2.47	3.30	3.89
	SD	0.08	0.08	0.14	0.04	0.04	0.06	0.02	0.14	0.04	0.11	0.02	0.02	0.03	0.06	0.17
	Range	0.11	0.11	0.20	0.06	0.06	0.09	0.02	0.19	0.05	0.15	0.03	0.02	0.04	0.09	0.25
	Min.	11.44	11.44	1.75	0.61	0.42	1.09	1.51	1.15	1.76	2.30	1.67	0.86	2.45	3.26	3.77
	Max.	11.56	11.56	1.94	0.66	0.47	1.18	1.53	1.34	1.81	2.45	1.69	0.89	2.49	3.35	4.01
<i>Toxopus pallidus</i> n. sp.																
Male (N = 2)	Mean	9.26	8.98	1.40	0.40	0.39	0.74	1.16	0.99	1.37	1.70	1.16	0.56	1.73	2.56	3.02
	SD	0.23	0.17	0.09	0.07	0.01	0.03	0.04	0.06	0.00	0.00	0.00	0.00	0.00	0.06	0.08
	Range	0.32	0.25	0.13	0.10	0.01	0.04	0.06	0.09	0.00	0.00	0.00	0.01	0.00	0.09	0.11
	Min.	9.10	8.85	1.34	0.35	0.39	0.72	1.13	0.95	1.37	1.70	1.16	0.56	1.73	2.52	2.96
	Max.	9.42	9.10	1.47	0.45	0.40	0.76	1.18	1.03	1.37	1.70	1.16	0.56	1.74	2.60	3.07
<i>Toxopus parkeri</i> n. sp.																
Male (N = 5)	Mean	12.51	12.22	1.80	0.61	0.37	1.12	1.63	1.43	1.67	2.15	1.85	0.89	2.73	3.62	4.47
	SD	0.79	0.69	0.09	0.07	0.03	0.06	0.12	0.16	0.12	0.13	0.13	0.05	0.19	0.26	0.33
	Range	2.07	1.82	0.23	0.16	0.06	0.14	0.32	0.38	0.30	0.32	0.32	0.14	0.46	0.63	0.83
	Min.	11.77	11.52	1.69	0.55	0.33	1.04	1.47	1.27	1.57	2.04	1.74	0.84	2.57	3.43	4.09
	Max.	13.84	13.34	1.92	0.71	0.39	1.18	1.79	1.66	1.87	2.36	2.05	0.98	3.03	4.06	4.92

Table 2.1. (Continued).

Species	Length										Width					
	Total	Clyp- -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd	
<i>Toxopus signoretii</i> (Reuter) 1887																
Male (N = 1)	Measurements	12.10	12.20	2.30	0.90	0.62	1.40	1.70	1.46	1.89	2.50	2.15	0.98	2.83	3.75	4.20
<i>Toxopus simulans</i> n. sp.																
Male (N = 4)	Mean	11.96	11.30	1.79	0.55	0.40	0.99	1.43	0.93	1.73	2.06	1.55	0.68	2.17	3.18	3.90
	SD	0.22	0.34	0.08	0.06	0.03	0.04	0.07	0.04	0.01	0.05	0.04	0.03	0.06	0.06	0.20
	Range	0.46	0.74	0.17	0.15	0.07	0.08	0.17	0.08	0.02	0.11	0.08	0.07	0.13	0.16	0.37
	Min.	11.70	10.89	1.69	0.47	0.36	0.95	1.35	0.88	1.72	2.01	1.53	0.66	2.12	3.09	3.70
	Max.	12.16	11.62	1.85	0.62	0.43	1.03	1.51	0.96	1.74	2.11	1.60	0.72	2.24	3.25	4.08
<i>Toxopus steineri</i> n. sp.																
Male (N = 3)	Mean	11.48	11.24	1.89	0.59	0.49	0.88	1.53	1.31	1.69	1.94	1.52	0.76	2.22	3.12	3.57
	SD	0.42	0.31	0.10	0.06	0.04	0.07	0.10	0.12	0.08	0.02	0.02	0.03	0.08	0.03	0.26
	Range	0.76	0.59	0.20	0.12	0.09	0.13	0.20	0.22	0.15	0.04	0.04	0.05	0.16	0.05	0.48
	Min.	11.00	11.00	1.80	0.53	0.45	0.81	1.43	1.23	1.64	1.93	1.50	0.74	2.15	3.10	3.38
	Max.	11.76	11.59	2.00	0.65	0.53	0.95	1.63	1.45	1.79	1.96	1.53	0.80	2.31	3.15	3.87
<i>Toxopus tibialis</i> n. sp.																
Male (N = 3)	Mean	14.80	14.60	2.28	0.64	0.49	1.48	1.92	1.50	2.22	2.73	2.34	1.10	3.08	4.18	4.99
	SD	0.53	0.76	0.16	0.10	0.02	0.03	0.16	0.14	0.02	0.12	0.08	0.06	0.09	0.24	0.24
	Range	1.01	1.50	0.32	0.17	0.03	0.06	0.30	0.28	0.03	0.24	0.17	0.11	0.17	0.47	0.45
	Min.	14.39	13.89	2.10	0.58	0.47	1.44	1.79	1.36	2.20	2.62	2.25	1.04	2.98	3.98	4.72
	Max.	15.40	15.40	2.43	0.75	0.50	1.50	2.09	1.65	2.24	2.86	2.42	1.16	3.14	4.45	5.17
<i>Toxopus toamasina</i> n. sp.																
Male (N = 5)	Mean	12.21	12.02	1.87	0.54	0.41	1.09	1.50	1.16	1.77	2.21	1.83	0.72	2.39	3.30	4.16
	SD	0.35	0.48	0.07	0.04	0.04	0.10	0.10	0.10	0.04	0.06	0.08	0.03	0.11	0.16	0.18
	Range	0.88	1.20	0.20	0.10	0.11	0.24	0.24	0.21	0.11	0.15	0.21	0.08	0.29	0.41	0.47
	Min.	11.69	11.22	1.77	0.49	0.36	1.00	1.32	1.06	1.72	2.15	1.73	0.69	2.20	3.04	3.85
	Max.	12.57	12.41	1.97	0.59	0.46	1.23	1.56	1.27	1.83	2.30	1.94	0.77	2.50	3.44	4.32

Table 2.1. (Continued).

Species	Length											Width			
	Total	Clyp- -Abd	Head	AOc	POc	APr	PPr	Scu	Sea	Ped	Head	Syn	APr	PPr	Abd
<i>Toxopus toliara</i> n. sp.															
Male (N = 5)															
Mean	11.63	11.33	1.99	0.69	0.47	1.02	1.41	1.13	1.43	2.09	1.52	0.74	2.19	3.10	3.85
SD	0.96	0.93	0.19	0.10	0.04	0.07	0.20	0.13	0.17	0.28	0.12	0.05	0.16	0.35	0.48
Range	2.28	2.15	0.47	0.27	0.10	0.17	0.49	0.29	0.43	0.67	0.30	0.12	0.39	0.87	1.19
Min.	10.68	10.39	1.74	0.57	0.41	0.95	1.24	1.02	1.24	1.89	1.40	0.68	1.96	2.74	3.22
Max.	12.96	12.54	2.21	0.84	0.51	1.12	1.74	1.30	1.67	2.57	1.70	0.80	2.35	3.61	4.41
<i>Toxopus vazimba</i> n. sp.															
Male (N = 4)															
Mean	11.99	11.99	2.00	0.65	0.45	1.05	1.62	1.16	1.74	2.25	1.71	0.89	2.50	3.38	3.79
SD	0.59	0.59	0.14	0.06	0.04	0.20	0.11	0.13	0.09	0.10	0.08	0.02	0.13	0.21	0.17
Range	1.44	1.44	0.30	0.15	0.09	0.46	0.26	0.29	0.20	0.24	0.18	0.04	0.30	0.45	0.37
Min.	11.27	11.27	1.79	0.56	0.40	0.77	1.46	1.01	1.66	2.10	1.62	0.87	2.31	3.08	3.63
Max.	12.71	12.71	2.09	0.71	0.49	1.23	1.71	1.30	1.86	2.34	1.80	0.91	2.61	3.52	4.00

Table 2.2. GenBank accession numbers of COI sequences. F – female; M – male; USI – unique specimen identifier; RCW – ethanol specimen collection number.

<b>Species</b>	<b>Sex</b>	<b>USI (UCR_ENT)</b>	<b>RCW</b>	<b>GenBank accession no.</b>
<i>Distirogaster tarsalis</i> (Signoret), 1860	M	00006366	2898	KR606396
<i>Distirogaster</i> n. sp.	M	00088090	3018	KR606395
<i>Distirogaster</i> n. sp.	M	00006369	2928	KR606394
<i>Distirogaster</i> n. sp.	M	00007158	2881	KR606393
<i>Distirogaster</i> n. sp.	F	00007088	2880	KR606392
<i>Gibbosella conisimilis</i> n. sp.	M	00045495	2962	KR606400
<i>Gibbosella conisimilis</i> n. sp.	M	00045427	2996	KR606401
<i>Gibbosella notoconica</i> n. sp.	M	00044813	2908	KR606402
<i>Gibbosella notoconica</i> n. sp.	M	00045558	2913	KR606403
<i>Gibbosella pallidalata</i> n. sp.	M	00045653	2939	KR606404
<i>Gibbosella planiscutum</i> n. sp.	M	00045296	2952	KR606405
<i>Gibbosella quadocris</i> n. sp.	M	00044860	2938	KR606406
<i>Gibbosella quadocris</i> n. sp.	M	00045651	2969	KR606407
<i>Gibbosella quadocris</i> n. sp.	M	00007257	2988	KR606408
<i>Gibbosella quadocris</i> n. sp.	M	00007166	2989	KR606409
<i>Gibbosella</i> sp.	F	00045157	2854	KR606410
<i>Gibbosella</i> sp.	F	00045452	2887	KR606411
<i>Gibbosella</i> sp.	F	00045079	2890	KR606412
<i>Gibbosella</i> sp.	F	00045646	2892	KR606413
<i>Gibbosella</i> sp.	F	00045569	3306	KR606414
<i>Glymmatophora crassipes</i> Horváth, 1914	M	00088087	3020	KR606415
<i>Marojejycoris auranticorium</i> n. sp.	M	00045555	2924	KR606416
<i>Marojejycoris brevifrons</i> n. sp.	M	00006480	2923	KR606417
<i>Marojejycoris ranomafana</i> n. sp.	M	00048057	2944	KR606418
<i>Tanindrazanus bemaraha</i> n. sp.	M	00006473	2964	KR606419
<i>Tanindrazanus harinhali</i> n. sp.	M	00006553	2889	KR606420
<i>Tanindrazanus irwini</i> n. sp.	M	00045339	2925	KR606421
<i>Tanindrazanus joffrevillus</i> n. sp.	M	00007256	2918	KR606422
<i>Tanindrazanus marginatus</i> n. sp.	M	00006887	2902	KR606423
<i>Tanindrazanus nigripes</i> n. sp.	M	00006725	2899	KR606424
<i>Tanindrazanus nigripes</i> n. sp.	M	00045465	2961	KR606425
<i>Tanindrazanus nigripes</i> n. sp.	M	00045328	2990	KR606426
<i>Tanindrazanus nigripes</i> n. sp.	M	00045228	2991	KR606427
<i>Tanindrazanus nigripes</i> n. sp.	M	00006705	2993	KR606428
<i>Tanindrazanus notatus</i> n. sp.	M	00044868	3312	KR606429
<i>Tanindrazanus tenebricus</i> n. sp.	M	00006723	2931	KR606430
<i>Tanindrazanus varicolor</i> n. sp.	M	00006250	2921	KR606431
<i>Tanindrazanus varicolor</i> n. sp.	M	00006485	2922	KR606432

Table 2.2. (Continued).

<b>Species</b>	<b>Sex</b>	<b>USI (UCR_ENT)</b>	<b>RCW</b>	<b>GenBank accession no.</b>
<i>Tanindrazanus varicolor</i> n. sp.	M	00006482	2949	KR606433
<i>Toxopus ampitavananima</i> n. sp.	M	00006732	3002	KR606434
<i>Toxopus antsiranana</i> n. sp.	M	00007189	2955	KR606435
<i>Toxopus antsiranana</i> n. sp.	M	00007191	2956	KR606436
<i>Toxopus basalis</i> n. sp.	M	00045525	2907	KR606437
<i>Toxopus brucei</i> n. sp.	M	00045338	2901	KR606438
<i>Toxopus brucei</i> n. sp.	M	00045286	2912	KR606439
<i>Toxopus brucei</i> n. sp.	M	00007133	3008	KR606440
<i>Toxopus brucei</i> n. sp.	M	00007043	3009	KR606441
<i>Toxopus brucei</i> n. sp.	M	00044971	3011	KR606442
<i>Toxopus farafangana</i> n. sp.	M	00045332	2884	KR606443
<i>Toxopus farafangana</i> n. sp.	M	00006729	2906	KR606444
<i>Toxopus farafangana</i> n. sp.	M	00044965	3006	KR606445
<i>Toxopus fisheri</i> n. sp.	M	00045431	2910	KR606446
<i>Toxopus griswoldi</i> n. sp.	M	00006435	2916	KR606448
<i>Toxopus griswoldi</i> n. sp.	F	00045042	2894	KR606447
<i>Toxopus italaviana</i> n. sp.	M	00006428	2934	KR606449
<i>Toxopus melobrunneus</i> n. sp.	M	00045243	2994	KR606450
<i>Toxopus mianद्रitsara</i> n. sp.	M	00044852	2919	KR606451
<i>Toxopus mianद्रitsara</i> n. sp.	M	00006148	3007	KR606452
<i>Toxopus namoroka</i> n. sp.	M	00045251	2917	KR606453
<i>Toxopus pallidus</i> n. sp.	M	00045432	2957	KR606454
<i>Toxopus parkeri</i> n. sp.	M	00006973	3010	KR606455
<i>Toxopus simulans</i> n. sp.	M	00045124	2980	KR606456
<i>Toxopus toamasina</i> n. sp.	M	00007056	2951	KR606457
<i>Toxopus toliara</i> n. sp.	M	00006589	2933	KR606458
<i>Toxopus toliara</i> n. sp.	M	00006468	2966	KR606459
<i>Toxopus toliara</i> n. sp.	M	00006688	2997	KR606460
<i>Toxopus toliara</i> n. sp.	M	00006580	3001	KR606461
<i>Toxopus vazimba</i> n. sp.	M	00006472	2953	KR606462
Ectrichodiinae sp.	F	00045353	2891	KR606397
Ectrichodiinae sp.	F	00044794	2893	KR606398
Ectrichodiinae sp.	F	00045654	2895	KR606399

### **Chapter 3: Phylogenetics and biogeography of the endemic Madagascan millipede assassin bugs (Hemiptera: Reduviidae: Ectrichodiinae)**

#### **Abstract**

For at least the past 80 my, Madagascar, a major biodiversity hotspot, has been isolated from all other landmasses. This long-term isolation, along with geologic and climatic factors within Madagascar and throughout the Indian Ocean, has undoubtedly influenced the evolution of the island's biota. However, few systematic analyses incorporating modern divergence dating and biogeographic analyses have focused on Madagascan insects. The diverse Madagascan millipede assassin bugs (Heteroptera: Reduviidae: Ectrichodiinae) offer an opportunity to contribute to a limited body of insect-related research that explores Madagascar's historical biogeography. A molecular dataset (COI mtDNA and 18S, 28S D2 and D3-D5 rDNAs) for 56 taxa (39 ingroup) and a combined morphological (145 characters) and molecular dataset for 110 taxa (93 ingroup) are analyzed with maximum likelihood (ML) and parsimony approaches. Based on the molecular ML phylogeny, divergence times were estimated using fossil and secondary calibrations and biogeographic analyses performed using DIVA, DEC, and DEC+j models to determine the role and patterns of vicariance and dispersal in the origin of Madagascan Ectrichodiinae. Results indicate that Ectrichodiinae in Madagascar do not form a monophyletic group, different clades are closely related to Afrotropical and Oriental lineages, and have colonized the island via transoceanic dispersal at least twice from the Oriental region and once from the Afrotropical region in the last ~68 my. Additionally, the DEC+j and DIVA models infer a single out-of-Madagascar dispersal event to the Afrotropical region. Oceanic and geologic factors that may have facilitated dispersal between these three regions are discussed. Results of the combined analyses are used to explore character support for Madagascan taxa and inform taxonomic



diagnoses. Our results are congruent with the small but growing body of biogeographic research supporting Cenozoic transoceanic dispersal for Madagascan invertebrates to and from Oriental and Afrotropical regions.

## **Introduction**

Madagascar, the fourth largest island on Earth, is one of the world's biodiversity hotspots with more than 13,000 plant, 900 vertebrate, and 5,800 invertebrate species known (Goodman & Benstead, 2005; Phillipson et al., 2006). For non-invasive species, ~86% of invertebrates, ~50% of birds, more than 95% of other vertebrates, and more than 90% of vascular plants are endemic to the island (Goodman & Benstead, 2005; Yoder & Nowak, 2006; Phillipson et al., 2006; Buerki et al., 2013). Numerous phylogenetic studies have provided evidence that a large proportion of Madagascar's biota is closely related to African lineages (~37% of plants, ~30% of vertebrates, ~58% of invertebrates [see Yoder & Nowak, 2006]). However, a significant proportion of the biota is closely related to Oriental taxa (see Vences, 2004; Yoder & Nowak, 2006; Warren et al., 2010); about 32% of plants, 32% of vertebrates, and 17% of invertebrates are sister to Oriental lineages (Yoder & Nowak, 2006). This extant, unique biodiversity is, in the words of Ganzhorn et al. (2014), "rooted in the plate tectonics of the Mesozoic" (i.e., long isolation period), as well as the island's geography, climate, and late colonization by humans (Goodman & Benstead, 2003; Scales, 2014).

Madagascar was once part of Gondwana, but ~160 mya, the Gondwanan landmass began to split with Indo-Madagascar separating from Africa (Plummer & Belle, 1995; Gnos et al., 1997). By at least 80 mya, Madagascar separated from India and has been isolated ~400 km from Africa's southeastern coast since (Storey et al., 1995; Torsvik et al., 2000; Seward et al., 2004; Ali & Aitchison, 2008). Many plant and vertebrate studies have investigated historical

biogeographic patterns for various Madagascan taxa. Although some systematic studies support influences of Gondwanan vicariance for the presence of some Madagascan lineages (e.g., ranid frogs [Bossuyt et al., 2006], boid snakes [Noonan & Chippindale, 2006], chameleons [Okajima & Kumazawa, 2010]), many recent studies based on divergence dating analyses support post-Gondwanan transoceanic dispersal from Afrotropical and Oriental regions (e.g., endemic mammalian lineages [Poux et al., 2005], Chrysophylloideae [Bartish et al., 2011], gerrhosaurid lizards [Raselimanana et al., 2009]). Comparatively fewer studies have investigated biogeographic histories of Madagascan insects. Several Madagascan insects have been hypothesized to be relicts of Gondwanan vicariance, e.g., Diplatyidae (Dermaptera) (Popham, 2000), Notonemourinae (Plecoptera) (Illies, 1965; Paulian & Viette, 2003), some Blephariceridae (Diptera) (Paulian & Viette, 2003), and Sialidae (Liu et al., 2015), but there is an increasing number of systematic studies that support Cenozoic transoceanic dispersal from the Afrotropical and Oriental regions. For example, allodapine bees (Fuller et al., 2005; Schwarz et al., 2006; Chenoweth & Schwarz, 2011) and scarabid beetles (Wirta et al., 2008; Wirta et al., 2010; Sole et al., 2011) colonized Madagascar multiple times in the last 65 my. Fungus-growing termites in Madagascar originated from a single colonization event ~7–11 mya (Nobre et al., 2010). Other studies on carpenter bees (Rehan et al., 2010), mayflies (Monaghan et al., 2005), pierid butterflies (Nazari et al., 2011), and dytiscid beetles (Bukontaite et al., 2015) have also supported Cenozoic colonization.

Millipede assassin bugs (Heteroptera: Reduviidae: Ectrichodiinae) are a diverse group of specialized millipede predators with 736 species in 121 genera that are distributed in circumtropical and some temperate regions (Maldonado, 1990; Dougherty, 1995; Carpintero & Maldonado, 1996; Forthman & Weirauch, 2012; Forthman et al., in press). Recently, Forthman et al. (in press) published a monograph of Madagascan millipede assassin bugs (excluding the genus

*Distirogaster* Horváth) in which 63 new species and three new genera were described, increasing the known Madagascan millipede assassin bug fauna by six-fold. All species and six of the eight genera found on Madagascar are endemic; of the 37 species of *Glymmatophora* Stål and 39 species of *Maraenaspis* Karsch, two and one, respectively, are endemic to Madagascar with the remainder distributed in Africa. This diverse Madagascan millipede assassin bug fauna thus presents an opportunity to investigate and contribute to a limited yet rapidly growing body of research exploring historical influences that have shaped Madagascar's terrestrial insect fauna. However, such an evolutionary investigation has been hindered by a lack of phylogenetic hypotheses; relationships between Ectrichodiinae genera have not been tested cladistically beyond a genus-level phylogeny of the New World fauna based on morphological data (Dougherty, 1995). A recent systematic study of the Reduviidae indicates that Ectrichodiinae originated 57–80 mya and, therefore, after the Madagascar-India split (Hwang & Weirauch, 2012). Thus, dispersal is likely responsible for the current distribution of millipede assassin bugs worldwide, but it is unknown what factors (e.g., ocean currents, land bridges, and stepping-stone islands) may have facilitated dispersal between Madagascar and other biogeographic regions.

Given the recent taxonomic treatment, Upper Cretaceous age, and worldwide distribution of the subfamily, as well as the general restriction of genera to particular biogeographic regions, millipede assassin bugs are an excellent model for biogeographic studies of the Madagascan fauna. Here, we present molecular phylogenetic results that include representatives of six of the eight Madagascan Ectrichodiinae genera and provide a framework for testing the number, timing, and origin of Madagascan lineages. Based on these results, we perform divergence dating and biogeographic analyses to investigate the geographic and oceanic influences that could have facilitated long-distance dispersal between Madagascar and other biogeographic areas. We also present results of a combined morphological and molecular phylogenetic analysis on a large

sample of Madagascan Ectrichodiinae species that is used to formalize taxonomic decisions and inform diagnoses in Forthman et al.'s (in press) Madagascan Ectrichodiinae monograph.

## **Material and Methods**

### *Taxon sampling, vouchering, and databasing*

A total of 110 terminal taxa were examined, comprising 93 ingroup (Ectrichodiinae) and 17 outgroup taxa (seven Reduviidae subfamilies). Ingroup sampling comprised all eight genera and 67 species of Madagascan Ectrichodiinae. Due to the lack of DNA quality material, only 56 terminal taxa were sequenced (39 ingroups, including 14 Madagascan species in six genera; sequence data for the Madagascan genus *Toliarus* Forthman, Chłond, & Weirauch and the species *Maraenaspis bidens* [Reuter] are unavailable). Each taxon is represented by a mounted primary specimen voucher (molecular vouchers are listed in Table 3.1). For specimens sampled for molecular data, a hind leg was removed for DNA extraction; once extraction was completed, the leg was card mounted and associated with the pinned specimen. Each molecular voucher is associated with a unique specimen identifier (USI) label to connect it with specimen information and, where possible, images in the Planetary Biodiversity Inventory database (<http://www.research.amnh.org/pbi/locality/index.php>) and Heteroptera Species Pages (<http://research.amnh.org/pbi/heteropteraspeciespage/>) maintained by the American Museum of Natural History. All USI labels are comprised of the prefix AMNH\_PBI or UCR\_ENT followed by an 8-digit number. Sequenced specimens are also associated with an RCW number, which is the ethanol specimen collection number.

Specimens examined for this study are deposited in the following institutions: **AMNH**, American Museum of Natural History, New York, USA; **BMNH**, Natural History Museum, London, United Kingdom; **CAS**, California Academy of Sciences, California, USA; **FCAP**,

Universidade Federal do Pará, Pará, Brazil; **MNHN**, Muséum National d'Histoire Naturelle, Paris, France; **MRAC**, Musée Royal de l'Afrique Centrale, Tervuren, Belgium; **MTEC**, Montana State University, Montana, USA; **SU**, Department of Zoology, University of Silesia, Poland; **UCR**, University of California, Riverside Entomological Research Museum, California, USA; **USNM**, National Museum of Natural History, Washington D.C., USA.

*Morphological methods, terminology, and abbreviations*

Extreme sexual dimorphism (Fig. 3.1) is common among millipede assassin bugs and poses a significant problem when analyzing evolutionary relationships. As such, male ectrichodiine specimens were targeted for morphological character coding. However, two terminal taxa, *Maraenaspis coccinea* Horváth and *Racelda* sp., are represented by apterous females. The male of *Maraenaspis coccinea* has, to our knowledge, yet to be discovered and available specimens of *Racelda* were restricted to the one female at the time of this project. Despite the possibility of extreme dimorphism, some morphological features (e.g., antennal segmentation, relative lengths of labial segments, scutellar apical processes, etc.) remain similar among males and females within ectrichodiine species and genera. As such, these sexually static morphological features are inferred and coded in the analysis for taxa only represented by females.

External morphology and genitalic characters were examined using Nikon SMZ1000 and SMZ1500 dissecting microscopes. For males, genitalia (abdominal segment 8, pygophore, and phallus) were dissected, cleared in heated 10% potassium hydroxide (KOH) for 3–8 minutes, washed with distilled water and subsequently with 100% ethanol (EtOH), stained with Chlorazol Black E in 70% EtOH solution to give contrast to membranous areas, examined in glycerol, and permanently stored in genitalic capsules pinned to the specimen. A total of 145 morphological characters were coded in the Descriptive Language for Taxonomy (DELTA) program (Dallwitz,

1980; Dallwitz et al., 1999) using a modified morphological matrix from Weirauch (2008) and Weirauch (2010). In general, terminology follows a subset of terms used by Dougherty (1995), Weirauch (2008), and Forero & Weirauch (2012). Terminology for wing venation follows Hill (2014) and Weirauch (2008), although homology concepts are currently being reviewed across Paraneoptera (Dávid Rédei, pers. comm.). The morphological characters and character states used in this study are given in Table 3.2. The morphological matrix is provided in Supplementary Material (SM) 3.1 in nexus format.

#### *Molecular markers, PCR, sequencing, and alignment*

One mitochondrial (COI) and three nuclear (18S, 28S D2, and 28S D3–D5 rDNAs) gene regions were sampled. Sequences from Weirauch & Munro (2009) and Hwang & Weirauch (2012) were used for outgroup terminals and some Ectrichodiinae taxa (Table 3.1). To sequence remaining ectrichodiine taxa, primer sequences for 18S (18SF, 18SR) and 28S D3–D5 (D3Fa, D5Ra) were obtained from Weirauch & Munro (2009); 28S D2 (D2Fa, D2Ra) from Forero et al. (2013); and COI (C1-J-2183F) from Simon et al. (1994) and (C1-N-2609R) from Damgaard & Sperling (2001). DNA was extracted from a hind leg for each specimen using QIAGEN DNeasy Blood and Tissue Kit protocols. PCR was performed using either GE Healthcare Life Sciences PuReTaq-Ready-To-Go-PCR-Beads or EmeraldAmp GT PCR Master Mix and Fisher Scientific or BioRad T100 Thermal Cyclers. Gene regions were amplified following Weirauch & Munro (2009), except the initial denaturation was set at 5 minutes, and for the 28S gene regions, the annealing temperature was set to 50°C. Gel electrophoresis with SyberSafe gel stain and a UV illuminator were used to check amplification results. PCR products were cleaned using Bionline SureClean and sequenced on an Applied Biosystems 3730xl DNA Sequencer at UCR's Institute for Integrative Genome Biology. Sequences were assembled and edited using Sequencher v4.8

and are available on GenBank (Table 3.1). Each gene region was treated as a separate partition and aligned in MAFFT (Katoh & Standley, 2013) using the G-INS-i algorithm. Gene regions were concatenated in SequenceMatrix v1.7.8. (Vaidya et al., 2011) and exported with external gaps coded as question marks. The concatenated, aligned molecular matrix is provided in nexus format in SM 3.2.

#### *Molecular phylogenetic analyses*

A maximum likelihood (ML) analysis was performed using the molecular dataset and RAxML-HPC2 on XSEDE v8.0.24 (Stamatakis, 2014) on the Cyberinfrastructure for Phylogenetic Research Science Gateway v3.1 (CIPRES; [http://www.phylo.org/sub\\_sections/portal/](http://www.phylo.org/sub_sections/portal/)). Each gene partition was analyzed under a GTR+ $\Gamma$ +I model, which is the best fit model for each partition as determined by MEGA v6.06 (Tamura et al., 2013). Rapid bootstrap analysis (BS) was performed for 1,000 iterations and followed by a ML best tree search. Aside from estimating the proportion of invariable sites for each partition, default settings were used.

The molecular matrix was also subjected to parsimony analyses with New Technology search and equal (EW) and implied weights (IW) in TNT v.1.1 (Goloboff et al., 2008). Uninformative characters (1,990) were inactivated prior to analysis. For IW, three concavity constant values ( $k$ ) were used: 3, 6, and 10. Internal gaps were treated as a fifth state, while external gaps were treated as missing. Default settings were used for sectorial search, drift, and tree fusing. The initial driven search level was set at 100 with level checked every 3 hits, initial addition sequences = 14, find minimum length 100 times, and random seed = 4,325. Standard BS resampling with absolute frequencies was performed using 500 replicates and New Technology

search (initial driven search = 38, check level every 3 hits, initial addition sequences = 7, find minimum length 10 times).

#### *Combined morphological and molecular phylogenetic analyses*

Morphological and molecular datasets were concatenated using Mesquite v2.75 (Maddison & Maddison, 2011) and subjected to ML (RAxML-HPC v.8.1.15; Stamatakis, 2014) and EW parsimony analyses (New Technology search in TNT). For the ML analysis, each gene partition was analyzed under a GTR+ $\Gamma$  model, while the morphological partition was analyzed with a Mkv model (Lewis, 2001). Rapid BS was performed for 1,000 iterations and followed by a ML best tree search. Our equal weights parsimony analysis used the same search and bootstrap approaches applied to our molecular dataset (see previous section). Unambiguous character optimizations were examined on the strict consensus tree in WinClada v1.00.08 (Nixon, 2002) (SM 3.3).

#### *Divergence dating estimation*

Divergence dating estimates were generated on the molecular ML best tree using the CIPRES web server's BEAST2 on XSEDE (Bouckaert et al., 2014). Each gene region was assigned an unlinked site model (GTR+ $\Gamma$ +I) and linked to the same tree model (Yule). The number of gamma rate categories for each site model was set to 4 and the proportion of invariant sites set to 0.5, with the option to estimate the value of both parameters and the gamma shape. All other site model parameters and estimation options remained at default settings. Following Hwang & Weirauch (2012), a linked clock model was assigned to 28S D2 and 28S D3–D5 to reflect their single identity. As such, three unlinked relaxed lognormal clock models were assigned for 18S, 28S, and COI. Initially, all non-calibration priors were not modified from default settings, however low posterior, prior, and mutation rate effective sample size (ESS)



values were extremely low (<50) after performing four independent runs in BEAST. To improve ESS values for these parameters, the default Jeffrey's prior was replaced with a log normal prior for the mutation rates (A. Rambaut & A. Drummond, <https://groups.google.com/forum/#!forum/beast-users>).

Given that there are no described Ectrichodiinae fossils, two outgroup fossils (*Triatoma dominicana* Poinar and *Apicrenus fossilis* Maldonado, Santiago-Blay, & Poinar) and two secondary calibrations (root node and Ectrichodiinae+Tribelocephalinae node) were used from a recent cladistic analysis of Reduviidae (Hwang & Weirauch, 2012). Fossils were assigned to nodes following Hwang & Weirauch (2012), using an apomorphy-based approach (Parham et al., 2011). For fossil-based node calibrations, a prior lognormal distribution was assigned with the minimum age corresponding to the minimum age of the fossil and with the 95% confidence interval encompassing the range of estimated dates recovered by Hwang & Weirauch (2012). For secondary calibrations, a prior normal distribution was assigned with the 95% confidence interval corresponding to the estimated 95% highest posterior density (HPD) credibility interval from Hwang & Weirauch (2012). Calibration nodes, means, standard deviations or sigma values, and offsets (in real space) are given in SM 3.4. Four Markov Chain Monte Carlo (MCMC) chains were ran for 200 million generations, with parameters and trees logged every 20,000 generations. Each chain was assessed for convergence in Tracer v1.6 and combined after discarding 5% burn-in. Combined ESS values for each parameter were  $\geq 200$ . LogCombiner v2.1.3 was used to discard 5% of trees as burn-in and combine the remaining trees from each run. Median heights were annotated using TreeAnnotator v.2.1.2.

### *Biogeographic analyses*

Ancestral ranges were reconstructed on the maximum clade credibility tree from the BEAST analysis using DIVA (Dispersal-Vicariance Analysis) in RASP v3.02 (Ronquist, 1997; Yu et al., 2010, 2015) and DEC (dispersal-extinction-cladogenesis) (Ree et al., 2005; Ree & Smith, 2008) and DEC+j (Matzke, 2014) models in the BioGeoBEARS v0.2.1 package (Matzke, 2013) in R v3.2.0 (R Core Team, 2015). Distribution ranges of Ectrichodiinae were divided into four areas: Neotropical, Afrotropical, Madagascar, and Oriental. Outgroup taxa were trimmed due to limited sampling of other assassin bug subfamilies.

The parsimony, event-based DIVA method was developed by Ronquist (1997) and reconstructs ancestral distributions based on a simple biogeographic model and a three-dimensional cost matrix. No penalty is allotted when speciation is the result of vicariance, but dispersal and extinction have a penalty of one per unit area added to or deleted from a distribution (Ronquist, 1997). To reduce the acquisition of multiple areas at nodes near the root, the maximum number of areas for each node was two.

The DEC models geographic range evolution by stochastic dispersal and local extinction events along branches and estimates the likelihood of ancestral ranges at cladogenesis (Ree et al., 2005; Ree & Smith, 2008). This model has two free parameters specifying the dispersal rate ( $d$ ; range expansion) and extinction rate ( $e$ ; range contraction) along branches (Ree et al., 2005; Ree & Smith, 2008; Matzke, 2014). The DEC model also has a fixed cladogenesis model that assumes that one daughter lineage will always inherit one subarea of a widespread ancestral range (Matzke, 2014). The DEC+j model has an additional free parameter  $j$  that enables one daughter lineage to disperse to a new range outside the ancestral range, thus modeling founder-event speciation events (Matzke, 2014). For both DEC and DEC+j models, the range constraint matrix was set at default with the maximum number of areas for each node set to two. Five time scales

were constructed in the dispersal matrix to reflect changes in ocean currents in the Mozambique channel and the emergence of Indian Ocean islands: 1) 0–15 mya, 2) 15–23 mya, 3) 23–34 mya, 4) 34–56 mya, and 5) 56–66 mya. Dispersal rates between areas were assigned as 0.1 (very low; oppositely directed ocean circulation, lack of islands and land bridges, and areas not connected), 0.25 (low; ocean circulation in dispersal direction or presence of islands or land bridges), 0.5 (moderate; ocean circulation in dispersal direction and presence of islands or land bridges), 0.75 (high; areas closely adjacent or directly connected), and 1.0 (same area) based on paleoceanographic and paleogeographic reconstructions (Sanmartín et al., 2001; Heydt & Dijkstra, 2006; Ali & Huber, 2010; Lomolino, 2010; Pyron, 2014). Dispersal rates assigned for each time scale are given in SM 3.5. The DEC model is nested within the DEC+j model, and, as a result, we compare the two models using a Likelihood Ratio Test (LTR).

## Results

### *Molecular phylogenetic analyses*

The first published molecular ML best tree (Ln score =  $-25,426.19528$ ) of millipede assassin bugs is shown in Fig. 3.2. Many nodes received moderate (BS = 70–89) to high support (BS = 90–100), but several deeper nodes were poorly supported (BS < 70). Ectrichodiinae is monophyletic with high support and recovered as the sister group to Tribelocephalinae, which is congruent with previous morphological and molecular cladistic analyses (Weirauch, 2008; Weirauch & Munro, 2009; Hwang & Weirauch, 2012). Two major ectrichodiine clades are recovered with low support: a predominantly Neotropical clade (node 1), which includes the Afrotropical genus *Santosia* Stål, and an Old World clade (node 2) that includes Afrotropical, Oriental, and Madagascan taxa.

Four Madagascan Ectrichodiinae lineages are recovered with low to high support. A representative of the Madagascan genus *Gibbosella* Chłond is nested within a clade of Oriental taxa with moderate support, but relationships within this clade, especially the *Gibbosella* and *Caecina* Stål sister group relationship, are poorly supported. The genera *Tanindrazanus* Forthman, Chłond, & Weirauch, *Toxopus* Bergroth, and *Marojejycoris* Forthman, Chłond, & Weirauch is a poorly supported clade. Relationships within this clade are largely weakly supported. *Tanindrazanus* is recovered as the sister group of *Marojejycoris*, and together are closely related to *Toxopus*. *Distirogaster* also forms a clade, but is weakly supported as the sister group to a clade comprising Afrotropical, Madagascan, and Oriental taxa. The Madagascan species *Glymmatophora crassipes* Horváth is highly supported as the sister to the Afrotropical genus *Ectrichodia* Lepeletier & Serville.

A number of unresolved relationships are recovered throughout the EW parsimony phylogeny (SM 3.6), e.g., near the root of the phylogeny (i.e., most outgroups and Ectrichodiinae+Tribelocephalinae), between *Tanindrazanus*+*Marojejycoris* and two other clades (*Toxopus* and *Toxopus*+*Distirogaster*), among others. Ectrichodiinae is also rendered paraphyletic with respect to Tribelocephalinae. The ectrichodiine genus *Vilius* Stål is sister to Tribelocephalinae genera, which is a questionable result. Due to the lack of resolution in our EW phylogeny, we used IW under three concavity constant values that resulted in one ( $k = 6$  or  $10$ ) or three ( $k = 3$ ) most parsimonious trees (SM 3.6). Some of the relationships recovered from these IW analyses are similar with the molecular ML phylogeny. However, differences are observed (e.g., non-monophyly of *Toxopus* and Ectrichodiinae), some of which are questionable and unlikely based on morphology (e.g., *Vilius* nested with Tribelocephalinae, *Maraenaspis coccinea* sister to *Caecina* sp.). Some IW results consistently differed between weighting schemes, e.g., the phylogenetic positions of *Marojejycoris*, *Toxopus toamasina*, and *Toxopus griswoldi*. Despite

topological differences between EW and IW parsimony trees, these results provide evidence for the non-monophyly of Madagascan Ectrichodiinae.

#### *Combined morphological and molecular phylogenetic analyses*

The TNT consensus tree of 938 parsimonious trees is shown in Fig. 3.3 (outgroups not shown but same as in Fig. 3.2) and was used to inform taxonomic decisions in Forthman et al. (in press). As in the molecular (SM 3.2) and combined ML analyses, Ectrichodiinae are monophyletic and recovered as the sister to Tribelocephalinae with high support (BS = 99). A predominantly Neotropical clade (node 1) is also recovered with low support, but unlike the ML analysis, this clade includes the Oriental genus *Vilius* along with *Santosia*. An Old World clade (node 2) is recovered with several topological differences from the ML tree. *Mendis* Stål+*Bannania* Hsiao+*Neolibavius* Miller is weakly supported as the sister to the remaining taxa, followed by a weakly supported *Microstemmatoides* Putschkov+*Synavecoris* Villiers+nr *Bannania* clade. A clade (node 3) comprised of *Ectrichodia*, *Glymmatophora*, *Cleptria* Stål, *Maraenaspis*, and *Ectrychotes* Burmeister is recovered as sister to a predominantly Madagascan clade (node 4) with low support. Relationships within the latter clade are supported with BS<50, with *Marojejycoris* sister to *Gibbosella*+*Caecina*, *Marojejycoris*+*Gibbosella*+*Caecina* sister to *Distirogaster*+*Toliarus*+*Tanindrazanus*+*Toxopus* (node 6), *Distirogaster* sister to *Toliarus*+*Tanindrazanus*+*Toxopus* (node 7), and *Toliarus* sister to *Tanindrazanus*+*Toxopus*. Character optimization results for the parsimony tree are given in SM 3.3.

Our combined ML result (SM 3.7) is largely congruent with the combined parsimony topology with the following exceptions: 1) *Maraenaspis bidens* is sister to *Racelda* sp., 2) *Gibbosella* is the sister to the *Caecina*+*Mendis*+*Neolibavius*+*Bannania*+nr *Bannania* clade, and 3) *Distirogaster* is the sister group to the *Glymmatophora*+*Ectrichodia*+*Centraspis*+

*Ectrychotes+Microstemmatoides+Cleptria+Maraenaspis coccinea* clade. The ML result supports the taxonomic decisions of Forthman et al. (in press) and produces minor character optimization differences in comparison to the parsimony tree for taxa and clades of interest (data not shown).

#### *Divergence dating estimation*

Results of the BEAST molecular divergence dating analysis are shown in Fig. 3.4, with HPD intervals shown in SM 3.8. Based on the analysis, Ectrichodiinae diverged from Tribelocephalinae approximately 68.53 mya (95% HPD = 57.61–79.39 mya), which is congruent with Hwang & Weirauch (2012). Within Ectrichodiinae, the Neotropical Ectrichodiinae+*Santosia* clade diverged from the Old World clade about 63.51 mya (95% HPD = 52.39–74.44 mya). Within the Old World clade, all Madagascan Ectrichodiinae lineages and genera diverged from their respective sister groups within the last 46 my: 1) *Gibbosella quadocris* Forthman, Chlond, & Weirauch (median = 27.33 mya, 95% HPD = 12.22–42.12 mya), 2) *Marojejycoris*+*Tanindrazanus*+*Toxopus* (median = 35.55 mya, 95% HPD = 25.34–45.60 mya), 3) *Distirogaster* (median = 32.86 mya, 95% HPD = 23.48–42.59 mya), and 4) *Glymmatophora crassipes* (median = 19.24 mya, 95% HPD = 11.46–27.85 mya).

#### *Biogeographic analyses*

Ancestral range reconstruction results based on the DEC and DEC+j models are reported and shown in Fig. 3.4. Results of the ancestral area reconstruction using the DIVA model are shown in SM 3.9 and are most similar to the highest probability ranges reconstructed by DEC+j. The likelihood,  $d$ ,  $e$ , and  $j$  parameter values for the DEC and DEC+j models are as follows: (1) DEC: LnL = -39.09,  $d = 0.0056$ ,  $e = 0.0006$ ,  $j = 0$  and (2) DEC+j: LnL = -27.10,  $d = 1e^{-12}$ ,  $e = 1e^{-12}$ ,  $j = 0.1238$ . The DEC+j model performed significantly better than the DEC (LRT  $X^2 = 23.97$ , df = 1,

$p = 9.80e^{-7}$ ), with the  $j$  parameter (or founder speciation effect) by far the most significant contributor to the current distribution.

In analyses with all three biogeographic models, a Neotropical and Oriental distribution is recovered for the ancestral node of Ectrichodiinae. The ancestral areas for the Neotropical Ectrichodiinae+*Santosia* clade and the Old World clade are inferred as Neotropical and Oriental, respectively, in all analyses. In the DEC+j analysis, the ancestral range of *Gibbosella quadocris* and *Caecina* sp. is recovered as Oriental with high probability, indicating the ancestor of *Gibbosella* dispersed to Madagascar from this region. An Oriental distribution is also reconstructed for the node that includes *Gibbosella* and *Ectrychotes* with subsequent dispersal to Madagascar for the node containing *Marojejycoris* and *Ectrychotes*. The ancestor of the clade containing *Centraspis* Schaum and *Ectrychotes* is reconstructed to have dispersed to the Afrotropical region from Madagascar, with a re-colonization from Africa to Madagascar by *Glymmatophora crassipes*. Results from DIVA are similar to DEC+j highest probability results, but differ in the following with respect to the Madagascan fauna: 1) the node containing *Gibbosella* and *Ectrychotes* is inferred as Madagascar+Oriental, 2) the ancestral range of *Gibbosella quadocris* and *Caecina* sp. is Madagascar+Oriental, 3) an Afrotropical+Madagascar distribution is recovered for the clade containing *Distirogaster* and *Ectrychotes*, and 4) the ancestral range of *Glymmatophora crassipes*+*Ectrichodia* is Afrotropical+Madagascar.

Reconstructions from the DEC analyses differ from DEC+j results for some nodes and by the presence of more widespread ancestral ranges. In the DEC analysis, a Madagascar+Oriental ancestral distribution is recovered for *Gibbosella quadocris*+*Caecina* sp. with high probability, as well as for the node containing *Marojejycoris* and *Ectrychotes* and the node containing *Distirogaster* and *Ectrychotes*. An Afrotropical+Oriental ancestral range is recovered for the

clade including *Centraspis* and *Ectrychotes*, with the ancestral node of *Glymmatophora crassipes*+*Ectrichodia* having an Afrotropical+Madagascar distribution.

## Discussion

### *Phylogeny of Madagascan Ectrichodiinae*

Results of our molecular ML phylogeny are used to investigate the number of Madagascan Ectrichodiinae lineages, as well as to estimate temporal divergences and biogeographic history. Our combined morphological and molecular phylogeny was used by Forthman et al. (in press) to formalize taxonomic decisions and inform diagnostic features for Madagascan millipede assassin bugs. Both molecular ML and combined parsimony phylogenetic analyses recovered some similar relationships. Both phylogenetic hypotheses support a paraphyletic New World fauna, with respect to *Santosia* in the molecular ML phylogeny or *Santosia*+*Vilius* in the combined parsimony result. This predominately New World clade is recovered as sister to a larger clade of Afrotropical, Oriental, and Madagascan Ectrichodiinae with relatively weak support. Madagascan lineages are recovered with close relationships to Oriental and Afrotropical taxa in both analyses, which is congruent with Vences' (2004) and Yoder & Nowak's (2006) conclusions that a large proportion of the Madagascan biota exhibits relationships with Afrotropical and Oriental taxa. Species of *Glymmatophora* share a close relationship with *Ectrichodia* species. *Gibbosella* species are recovered as sister to *Caecina*, albeit with low support in both phylogenies. Although the molecular and combined phylogenies differ in their phylogenetic placement of *Marojejycoris*, the genera *Tanindrazanus* and *Toxopus* retain a close relationship in both analyses.

Despite recovering similar relationships, the two phylogenies show some degree of discordance in higher-level relationships within Ectrichodiinae, a result that may be due to the increase in missing molecular data in the combined analysis. Unlike the molecular phylogeny,



most Madagascan taxa form a clade in the combined phylogeny (node 4, including *Caecina*) with low support. The phylogenetic placement of *Marojejycoris* and *Distirogaster* differ; in the ML phylogeny, *Marojejycoris* is recovered as sister to *Tanindrazanus* and *Distirogaster* as sister to a clade containing *Ectrichodia* and *Ectrychotes*, whereas the two genera are included within node 4 of the combined analysis. In general, more nodes are recovered with  $BS \geq 50$  in the molecular phylogeny compared to the combined phylogeny. Regardless of the discordance between the two phylogenetic hypotheses, the combined phylogeny has permitted exploration of potential character support for taxa and been useful in determining diagnostic features for Forthman et al.'s (in press) taxonomic monograph.

#### *Temporal divergence and biogeography*

Based on our temporal divergence estimates and ancestral range reconstructions, millipede assassin bugs colonized Madagascar via transoceanic dispersal more than once within the last ~68 my. The DEC+j model outperformed the DEC model based on LRT and infers two colonization events from the Oriental region to Madagascar, once between 25–57 mya and once within the last 42 my. Subsequently, a single colonization event from Madagascar to the Afrotropical region is inferred to have occurred around the Eocene-Oligocene boundary (21–43 mya) with a back-colonization event (ancestral node of *Glymmatophora*+*Ectrichodia*) from the Afrotropical region to Madagascar within the last ~28 my. These results are congruent with the DIVA results (SM 3.9). Although the DEC results differ from the DEC+j in terms of the ancestral ranges reconstructed for several nodes, in general we find similar patterns: two dispersal events into Madagascar from the Oriental region and one dispersal from Africa to Madagascar. However, an out-of-Madagascar dispersal event to the Afrotropical region is not recovered in the DEC analysis; upon diverging from *Distirogaster*, the branch leading to the node containing *Centraspis*

and *Ectrychotes* has an Oriental distribution, with a range expansion into the Afrotropical region at that node.

Several hypotheses have been proposed for dispersal events from the Oriental region to Madagascar and may explain the two colonization events reconstructed from the DEC+j model. While many researchers reconstruct an isolated India in the Late Cretaceous (~65 mya) (see Yoder & Nowak, 2006; Ali & Aitchison, 2008), Ali & Aitchison (2008) proposed a paleogeographic model in which India became progressively isolated but remained connected to Madagascar via the Seychelles-Mascarene Plateau. This connection was extended in the Palaeogene by the development of volcanic islands (Ali & Aitchison, 2008; Zhou et al., 2012) and would have facilitated dispersal as India moved north. Warren et al. (2010) reviewed geologic evidence and also proposed that the rise and fall of sea levels exposed islands between India and Madagascar over the last 65 my, thus reducing the distance of open ocean to traverse for potential colonizers. Warren et al. (2010) further proposed climatic influences that would facilitate long-distance dispersal (e.g., winter monsoon winds blowing from India towards Madagascar facilitating aerial dispersal).

Several studies have supported “out-of-Madagascar” dispersal events to surrounding islands and continents for plants and vertebrates (e.g., Jansa et al., 1999; Wikström et al., 2010; Harmon et al., 2008), as well as for several groups of insects. Torres et al. (2001) concluded that satyrine butterflies in the subtribe Mycalesina have a complex biogeographical history that involves dispersal from Madagascar to Africa. The mayfly genus *Cloeodes* Traver is inferred to have a Madagascan origin followed by dispersal into Africa (Monaghan et al., 2005). Zakharov et al. (2004) inferred out-of-Madagascar dispersal for species of the butterfly genus *Papilio* Linnaeus to Oriental and Afrotropical regions within the last 10 my. Recently, the out-of-Madagascar biogeographical hypothesis has been supported by systematic analyses of Madagascan diving

beetles (Dytiscidae), with results indicating dispersal to Oriental and Afrotropical regions around the Oligocene and Miocene (Bukontaite et al., 2015).

Our DEC+j results support an out-of-Madagascar dispersal event to Afrotropical regions about 21–43 mya. Westward dispersal could not have been facilitated by ocean currents as these currents had an eastward movement during the Eocene and reversed direction only ~15 mya (Ali & Huber, 2010). McCall (1997) hypothesized that a land bridge known as the Davie Ridge once connected Africa and Madagascar during the mid-Eocene to early Miocene (i.e., 45–26 mya). However, this land bridge has not been supported by some studies (e.g., Leclaire et al., 1989; Bassias, 1992; Rogers et al., 2000), although some suggest that isolated islands may have been present (Rogers et al., 2000; de Wit, 2003). Thus, it is possible that islands in the Mozambique Channel have facilitated westward aerial dispersal during this time. Female specimens for extant Madagascan species of *Distirogaster* (5 spp.), *Toxopus* (2 spp.), and *Glymmatophora* (1 sp.) are apterous, which makes present-day aerial dispersal unlikely. However, it is unknown if the ancestors of these taxa were apterous or capable of flight, thus enabling long-distance aerial dispersal. Female specimens for the majority of Madagascan species, as well as many Afrotropical and Oriental taxa, remain unknown and hinder our ability to reconstruct ancestral states. Future discoveries of female specimens will allow us to investigate the evolution of winglessness for each sex and to subsequently refine hypotheses on how ancestral species may have dispersed in and out of Madagascar.

Based on our DEC+j analysis, the presence of islands and the eastward ocean currents of the Mozambique Channel within the last ~28 my may have facilitated the re-colonization of Madagascar from Africa. The divergence time estimate of *Glymmatophora crassipes* (95% HPD = 11.46–27.85 mya) is consistent with this hypothesis. However, aerial dispersal cannot be ruled out as ocean currents reversed direction ~15 mya, which would hinder eastward dispersal via

rafting. *Glymmatophora crassipes* is apterous in both sexes, but some African species in this genus are macropterous, and it is uncertain when and where the ancestor of *G. crassipes* has lost the capability of flight. Apterous and macropterous *Glymmatophora* species will need to be targeted for future analyses to determine if ocean currents or aerial dispersal are responsible for extant Madagascan *Glymmatophora*. Ancestral state reconstructions on the evolution of winglessness across *Glymmatophora* may provide further insight into possible avenues for the colonization of Madagascar.

Five Ectrichodiinae species in three genera — *Mascaregnasa* Distant, *Quinssyana* Distant, and *Rochonia* Distant — are known from another Indian Ocean island, the Seychelles (Maldonado, 1990). Each species is only known from the male holotype and, thus, were not included in our molecular or morphological analyses. We hypothesize, based on morphological similarities examined in habitus images of the holotypes, that *Quinssyana* is closely related to the Madagascan genus *Gibbosella* and the African genus *Synavecoris*, while *Toxopus* is closely related to *Rochonia*. *Mascaregnasa* is very different from other millipede assassin bug genera, and we are unable to hypothesize affinities with other taxa. Future inclusion of these taxa in systematic analyses will be critical for testing the role of the Seychelles as a stepping-stone island, which has been postulated for other taxa (e.g., *Tachycnemis* frogs [Vences et al., 2003a, b], *Nephilengys* hermit spiders [Kuntner & Agnarsson, 2011], and baetid mayflies [Monaghan et al., 2005]). Millipede assassin bugs from other Indian Ocean islands are currently unknown, but future taxonomic surveys on these islands may result in material that would also benefit future biogeographic investigations for Madagascan Ectrichodiinae.

*Combined parsimony analysis to inform taxonomic diagnoses of Madagascan Ectrichodiinae*

Character optimizations of the combined parsimony analysis were used as the basis for diagnoses in the taxonomic monograph of Madagascan Ectrichodiinae (Forthman et al., in press). A detailed discussion of important morphological characters for selected nodes shown in Fig. 3.3 is provided in SM 3.10. Here, we only discuss some examples of male diagnostic features for endemic Madagascan genera that are also included in Forthman et al. (in press). Species of *Marojejycoris* are distinguished by features such as a complete pronotal transverse suture (57–1; Fig. 3.3H), the laterally visible MGE (71–1; Fig. 3.3K), and the fore wing lacking the distal part of M beyond the M+Cu junction (104–1; Fig. 3.3M). Diagnostic features for *Gibbosella* include the 8-segmented antennae (38–3; except 6-segmented in *G. pallidalata* Forthman, Chłond, & Weirauch), dorsally directed paramedian scutellar processes (62–1; except in *G. planiscutum* Forthman, Chłond, & Weirauch) (Fig. 3.3I), and transversely bicolored dorsal laterotergites (105–1) with posterior protuberances (107–1, 108–1, 109–1, 110–1; shared with *Caecina*). A number of features are diagnostic for *Distirogaster*, e.g., the metallic coloration (2–1), 8-segmented antennae (38–3), posterior tubercles on dorsal laterotergites III–VI (107–1, 108–1, 109–1, 110–1), and paramedian carinae on abdominal sternites (120–1; Fig. 3.3N). The small (1–0) red and black (2–2) body and a punctate posterior pronotal lobe (51–2) distinguish *Toliarus*. Although *Tanindrazanus* is not supported by any synapomorphic characters in our analysis, it is consistently recovered as a clade in molecular and combined analyses. An incomplete transverse suture between the meso- and metasterna (66–1), a laterally expanded dorsal laterotergite II (106–1), and the transversely bicolored dorsal laterotergites (105–1) are treated as diagnostic characters for *Toxopus*.

## Conclusion

Madagascar's unique biodiversity is, at least partially, the result of long-term geologic isolation, transoceanic dispersal events followed by speciation, local geography, and climate change. The recently revised and diverse millipede assassin bug fauna of Madagascar have presented an opportunity to contribute to this growing knowledge of Madagascan invertebrates by investigating its historical biogeography. Phylogenetic hypotheses generated from molecular and combined morphological and molecular cladistic analyses are similar in some respects but discordant at higher-level relationships. Overall, the molecular phylogeny received higher BS support compared to the combined phylogeny. Regardless, results from the combined phylogeny have been important for informing taxonomic diagnoses in Forthman et al. (in press).

Based on our molecular dataset, millipede assassin bugs are shown to have colonized Madagascar within the last ~68 my. Given this relatively recent age, transoceanic dispersal rather than vicariance is responsible for the Madagascan Ectrichodiinae fauna we see today. Millipede assassin bugs colonized Madagascar twice from the Oriental region and once from the Afrotropical region. However, DEC+j and DIVA models reconstruct an out-of-Madagascan dispersal event to the Afrotropical region, whereas the DEC model does not. Temporal divergence estimates and biogeographic results indicate that dispersal from the Oriental region to Madagascar may have been facilitated by the Seychelles-Mascarene Plateau and volcanic islands in the Indian Ocean over the last ~65 my. Factors facilitating dispersal from the Afrotropical region to Madagascar are more difficult to determine, but aerial dispersal via stepping-stone islands in the Mozambique Channel is the most probable hypothesis when accounting for ancient oceanographic reconstructions (i.e., ocean currents had an eastward direction). Further testing of the geologic and oceanic factors facilitating dispersal will require a larger sample of Afrotropical millipede assassin bugs, as well as Seychellois species. Furthermore, apterous females are known

for several Madagascan species, making aerial dispersal impossible. However, it remains to be investigated if the ancestors of these taxa possessed an apterous condition or were capable of flight. Our results are congruent with a small, yet, largely growing body of biogeographic studies for Madagascan invertebrates; Madagascan millipede assassin bugs have a complex biogeographic history, with Cenozoic transoceanic dispersal between Oriental, Afrotropical, and Madagascan regions solely responsible for the endemic fauna we currently find present on the island.

## References

- Ali, J.R., Aitchison, J.C. 2008. Gondwana to Asia: plate tectonics, paleogeography and the biological connectivity of the Indian sub-continent from the Middle Jurassic through latest Eocene (166–35 Ma). *Earth-Science Reviews* 88: 145–166.
- Ali, J.R., Huber, M. 2010. Mammalian biodiversity on Madagascar controlled by ocean currents. *Nature* 463: 653–656.
- Álvarez-Padilla, F., Ubick, D., Griswold, C.E. 2012. *Noideattella* and *Tolegnaro*, two new genera of goblin spiders from Madagascar, with comments on the gamasomorphoid and silhouettelloid oonopids (Araneae, Oonopidae). *American Museum Novitates* 3745: 1–76.
- Bartish, I.V., Antonelli, A., Richardson, J.E., Swenson, U. 2011. Vicariance or long-distance dispersal: historical biogeography of the pantropical subfamily Chrysophylloideae (Sapotaceae). *Journal of Biogeography* 38: 177–190.
- Bassias, Y. 1992. Petrological and geochemical investigation of rocks from the Davie Fracture Zone (Mozambique Channel) and some tectonic implications. *Journal of African Earth Sciences* 15: 321–339.
- Bossuyt, F., Brown, R.M., Hillis, D.M., Cannatella, D.C., Milinkovitch, M.C. 2006. Phylogeny and biogeography of a cosmopolitan frog radiation: Late Cretaceous diversification resulted in continent-scale endemism in the family Ranidae. *Systematic Biology* 55: 579–594.
- Bouckaert, R., Heled, J., Kühnert, D., Vaughan, T., Wu, C.-H., Xie, D., Suchard, M.A., Rambaut, A., Drummond, A.J. 2014. BEAST 2: a software platform for Bayesian evolutionary analysis. *PLoS Computational Biology* 10: e1003537.
- Buerki S., Devey, D.S., Callmander, M.W., Phillipson, P.B., Forest, F. 2013. Spatio-temporal history of the endemic genera of Madagascar. *Botanical Journal of the Linnean Society* 171: 304–329.
- Bukontaite, R., Ranarilalaitiana, T., Randriamihaja, J.H., Bergsten, J. 2015. In or out-of-Madagascar?—colonization patterns for large-bodied diving beetles (Coleoptera: Dytiscidae). *PLoS ONE* 10: e0120777.
- Carpintero, D.J., Maldonado, J. 1996. Diagnostic characters and key to the genera of American Ectrichodiinae (Heteroptera, Reduviidae). *Caribbean Journal of Science* 32: 125–141.
- Chenoweth, L.B., Schwarz, M.P. 2011. Biogeographical origins and diversification of the exoneurine allodapine bees of Australia (Hymenoptera, Apidae). *Journal of Biogeography* 38: 1471–1483.
- Dallwitz, M.J. 1980. A general system for coding taxonomic descriptions. *Taxon* 29: 41–46.
- Dallwitz, M.J., Paine, T.A., Zurcher, E.J. 1999. User's guide to the DELTA Editor. Accessible at <http://delta-intkey.com>.



- Damgaard, J., Andersen, N.M., Cheng, L., Sperling, F.A.H. 2000. Phylogeny of sea skaters, *Halobates* Eschscholtz (Hemiptera, Gerridae), based on mtDNA sequence and morphology. *Zoological Journal of the Linnean Society* 130: 511–526.
- Dougherty, V. 1995. A review of the New World Ectrichodiinae genera (Hemiptera: Reduviidae). *Transactions of the American Entomological Society* 121: 173–225.
- Forero, D., Berniker, L., Weirauch, C., 2013. Phylogeny and character evolution in the bee-assassins (Insecta, Heteroptera: Reduviidae). *Molecular Phylogenetics and Evolution* 66: 283–302.
- Forero, D., Weirauch, C. 2012. Comparative genitalic morphology in the New World resin bugs Apiomerini (Hemiptera, Heteroptera, Reduviidae, Harpactorinae). *Deutsche Entomologische Zeitschrift* 59: 5–41.
- Forthman, M., Chłond, D., Weirauch, C. (in press). Taxonomic monograph of the endemic millipede assassin bug fauna of Madagascar (Hemiptera: Reduviidae: Ectrichodiinae). *Bulletin of the American Museum of Natural History*.
- Forthman, M., Weirauch, C. 2012. Toxic associations: a review of the predatory behaviors of millipede assassin bugs (Hemiptera: Reduviidae: Ectrichodiinae). *European Journal of Entomology* 109: 147–153.
- Fuller, S., Schwarz, M., Tierney, S. 2005. Phylogenetics of the allodapine bee genus *Braunsapis*: historical biogeography and long-range dispersal over water. *Journal of Biogeography* 32: 2135–2144.
- Ganzhorn, J.U., Wilmé, L., Mercier, J.-L. 2014. Explaining Madagascar's biodiversity. In Scales, I.R. (ed.): *Conservation and Environmental Management in Madagascar*. Routledge, NY, pp. 17–43.
- Gnos, E., Immenhauser, A., Peters, T. 1997. Late Cretaceous/early Tertiary convergence between the Indian and Arabian plates recorded in ophiolites and related sediments. *Tectonophysics* 271: 1–19.
- Goloboff, P.A., Farris, J.S., Nixon, K.C. 2008. TNT, a free program for phylogenetic analysis. *Cladistics* 24: 774–786.
- Goodman, S.M., Benstead, J.P. 2003. *The Natural History of Madagascar*. University of Chicago Press, Illinois.
- Goodman, S.M., Benstead, J.P. 2005. Updated estimates of biotic diversity and endemism for Madagascar. *Oryx* 39: 73–77.
- Harmon, L.J., Melville, J., Larson, A., Losos, J.B. 2008. The role of geography and ecological opportunity in the diversification of day geckos (*Phelsuma*). *Systematic Biology* 57: 562–573.

- von der Heydt, A., Dijkstra, H.A. 2006. Effect of ocean gateways on the global ocean circulation in the late Oligocene and early Miocene. *Paleoceanography* 21: PA1011.
- Hill, L. 2014. Revision of *Silhouettanus* with description of nine new species (Hemiptera: Heteroptera: Schizopteridae). *Zootaxa* 3815: 353–385.
- Hwang, W.S., Weirauch, C. 2012. Evolutionary history of assassin bugs (Insecta: Hemiptera: Reduviidae): insights from divergence dating and ancestral state reconstruction. *PLoS ONE* 7: e45523.
- Illies, J. 1965. Phylogeny and zoogeography of the Plecoptera. *Annual Review of Entomology* 10: 117–140.
- Jansa, S.A., Goodman, S.M., Tucker, P.K. 1999. Molecular phylogeny and biogeography of the native rodents of Madagascar (Muridae: Nesomyinae): a test of the single-origin hypothesis. *Cladistics* 270: 253–270.
- Katoh, K., Standley, D.M. 2013. MAFFT multiple sequence alignment software version 7: improvements in performance and usability. *Molecular Biology and Evolution* 30: 772–780.
- Krishnankutty, S.M., Dietrich, C.H. 2011. Taxonomic revision and phylogeny of an endemic leafhopper genus *Nesocerus* (Hemiptera: Cicadellidae: Idiocerinae) from Madagascar. *Zoological Journal of the Linnean Society* 162: 499–543.
- Kuntner, M., Agnarsson, I. 2011. Biogeography and diversification of hermit spiders on Indian Ocean islands (Nephilidae: *Nephilengys*). *Molecular Phylogenetics and Evolution* 59: 477–488.
- Leclaire, L., Bassias, Y., Clocchiatti, M., Ségoufin, J. 1898. La Ridge de Davie dans le Canal de Mozambique: approche stratigraphique et géodynamique. *Comptes Rendus de l'Académie des Sciences - Series II* 308: 1077–1082.
- Lewis, P.O. 2001. A likelihood approach to estimating phylogeny from discrete morphological character data. *Systematic Biology* 50: 913–925.
- Liu, X., Hayashi, F., Yang, D. 2015. Phylogeny of the family Sialidae (Insecta: Megaloptera) inferred from morphological data, with implications for generic classification and historical biogeography. *Cladistics* 31: 18–49.
- Lomolino, M.V., Riddle, B.R., Whittaker, R.J., Brown, J.H. 2010. *Biogeography*. Sinauer Associates, Maryland.
- Maddison, W.P., Maddison, D.R., 2011. Mesquite: a modular system for evolutionary analysis. Version 2.75. Available: <http://mesquiteproject.org>.
- Maldonado, J. 1990. Systematic catalogue of the Reduviidae of the world (Insecta: Heteroptera). *Caribbean Journal of Science, Special Edition*: 1–694.

- Matzke, N.J. 2013. Probabilistic historical biogeography: new models for founder-event speciation, imperfect detection, and fossils allow improved accuracy and model-testing. *Frontiers of Biogeography* 5: 242–248.
- Matzke, N.J. 2014. Model selection in historical biogeography reveals that founder-event speciation is a crucial process in island clades. *Systematic Biology* 63: 951–970.
- McCall, R.A. 1997. Implications of recent geological investigations of the Mozambique Channel for the mammalian colonization of Madagascar. *Proceedings of the Royal Society of London B* 264: 663–665.
- Monaghan, M.T., Gattolliat, J.-L., Sartori, M., Elouard, J.-M., James, H., Derleth, P., Glaizot, O., de Moor, F., Vogler, A.P. 2005. Trans-oceanic and endemic origins of the small minnow mayflies (Ephemeroptera, Beatidae) of Madagascar. *Proceedings of the Royal Society B* 272: 1829–1836.
- Mugrabi, D.F., Azevedo, C.O. 2010. Insecta, Hymenoptera, Bethyridae: range extension and filling gaps in Madagascar. *Check List* 6: 062–063.
- Nazari, V., Larsen, T.B., Lees, D.C., Brattström, O., Bouyer, T., van de Poel, G., Hebert, P.D.N. 2011. Phylogenetic systematics of *Colotis* and associated genera (Lepidoptera: Pieridae): evolutionary and taxonomic implications. *Journal of Zoological Systematics and Evolutionary Research* 49: 204–215.
- Nixon, K.C. 2002. WinClada ver. 1.00.08. Published by the author, Ithaca, NY.
- Nobre, T., Eggleton, P., Aanen, D.K. 2010. Vertical transmission as the key to the colonization of Madagascar by fungus-growing termites? *Proceedings of the Royal Society B* 277: 359–365.
- Noonan, B.P., Chippindale, P.T. 2006. Vicariant origin of Malagasy reptiles supports Late Cretaceous Antarctic land bridge. *American Naturalist* 168: 730–741.
- Okajima, Y., Kumazawa, Y. 2010. Mitochondrial genomes of acrodont lizards: timing of gene rearrangements and phylogenetic and biogeographic implications. *BMC Evolutionary Biology* 10: 141.
- Parham, J.F., Donoghue, P.C.J., Bell, C.J., Calway, T.D., Head, J.J., Holroyd, P.A., Inoue, J.G., Irmis, R.B., Joyce, W.G., Ksepka, D.T., Patané, J.S.L., Smith, N.D., Tarver, J.E., van Tuinen, M., Yang, Z., Angielczyk, K.D., Greenwood, J.M., Hipsley, C.A., Jacobs, L., Makovicky, P.J., Müller, J., Smith, K.T., Theodor, J.M., Warnock, R.C.M. 2011. Best practices for justifying fossil calibrations. *Systematic Biology* 61: 1–14.
- Paulian, R., Viette, P. 2003. An introduction to terrestrial and freshwater invertebrates. In Goodman, S.M., Benstead, J.P. (eds.): *The Natural History of Madagascar*. University of Chicago Press, IL, pp. 503–511.

- Phillipson, P.B., Schatz, G.E., Lowry II, P.P., Labat, J.-N. 2006. A catalogue of the vascular plants of Madagascar. In Ghazanfar, S.A., Beentje, H.J. (eds.): *Taxonomy and Ecology of African Plants: Their Conservation and Sustainable Use, Proceedings XVIIIth AETFAT Congress*. Royal Botanic Gardens, Kew, pp. 613–627.
- Plummer, P.S., Belle, E.R. Mesozoic tectono-stratigraphic evolution of the Seychelles microcontinent. *Sedimentary Geology* 96: 73–91.
- Popham, E.J. 2000. The geographical distribution of the Dermaptera (Insecta) with reference to continental drift. *Journal of Natural History* 34: 2007–2027.
- Poux, C., Madsen, O., Marquard, E., Vieites, D.R., de Jong, W.W., Vences, M. 2005. Asynchronous colonization of Madagascar by the four endemic clades of primates, tenrecs, carnivores, and rodents as inferred from nuclear genes. *Systematic Biology* 54: 719–730.
- Pyron, R.A. 2014. Biogeographic analysis reveals ancient continental vicariance and recent oceanic dispersal in Amphibians. *Systematic Biology* 63: 779–797.
- R Core Team. 2015. R: a language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria.
- Raselimanana, A.P., Noonan, B., Karanth, K.P., Gauthier, J., Yoder, A.D. 2009. Phylogeny and evolution of Malagasy plated lizards. *Molecular Phylogenetics and Evolution* 50: 336–344.
- Ree, R.H., Moore, B.R., Webb, C.O., Donoghue, M.J. 2005. A likelihood framework for inferring the evolution of geographic range on phylogenetic trees. *Evolution* 59: 2299–2311.
- Ree, R.H., Smith, S.A. 2008. Maximum likelihood inference of geographic range evolution by dispersal, local extinction, and cladogenesis. *Systematic Biology* 57: 4–14.
- Rehan, S.M., Chapman, T.W., Craigie, A.I., Richards, M.H., Cooper, S.J.B., Schwarz, M.P. 2010. Molecular phylogeny of the small carpenter bees (Hymenoptera: Apidae: Ceratinini) indicates early and rapid global dispersal. *Molecular Phylogenetics and Evolution* 55: 1042–1054.
- Rogers, R.R., Hartman, J.H., Krause, D.W. 2000. Stratigraphic analysis of Upper Cretaceous Rocks in the Mahajanga Basin, Northwestern Madagascar: implications for ancient and modern faunas. *Journal of Geology* 108: 275–301.
- Ronquist, F. 1997. Dispersal-vicariance analysis: a new approach to the quantification of historical biogeography. *Systematic Biology* 46: 195–203.
- Sanmartín, I., Enghoff, H., Ronquist, F. 2001. Patterns of animal dispersal, vicariance and diversification in the Holarctic. *Biological Journal of the Linnean Society* 73: 345–390.
- Scales, I.R. 2014. Conservation as the crossroads: biological diversity, environmental change and natural resource use in Madagascar. In Scales, I.R. (ed.): *Conservation and Environmental Management in Madagascar*. Routledge, NY, pp. 1–13.

- Schwarz, M.P., Fuller, S., Tierney, S.M., Cooper, S.J.B. 2006. Molecular phylogenetics of the exoneurine allodapine bees reveal an ancient and puzzling dispersal from Africa to Australia. *Systematic Biology* 55: 31–45.
- Seward, D., Grujic, D., Schreurs, G. 2004. An insight into the breakup of Gondwana: identifying events through low-temperature thermochronology from the basement rocks of Madagascar. *Tectonics* 23: TC3007.
- Simon, C., Frati, F., Beckenbach, A., Crespi, B., Liu, H., Flook, P. 1994. Evolution, weighting, and phylogenetic utility of mitochondrial gene sequences and a compilation of conserved polymerase chain reaction primers. *Annals of the Entomological Society of America* 87: 651–701.
- Sole, C.L., Wirta, H., Forgie, S.A., Scholtz, C.H. 2011. Origin of Madagascan Scarabaeini dung beetles (Coleoptera: Scarabaeidae): dispersal from Africa. *Insect Systematics and Evolution* 42: 29–40.
- Stamatakis, A. 2014. RAxML version 8: a tool for phylogenetic analysis and post-analysis of large phylogenies. *Bioinformatics* 30: 1312–1313.
- Storey, M., Mahoney, J.J., Saunders, A.D., Duncan, R.A., Kelly, S.P., Coffin, M.F. 1995. Timing of hot spot-related volcanism and the breakup of Madagascar and India. *Science* 267: 852–855.
- Tamura, K., Stecher, G., Peterson, D., Filipski, A., Kumar, S. 2013. MEGA6: Molecular Evolutionary Genetics Analysis version 6.0. *Molecular Biology and Evolution* 30: 2725–2729.
- Torres, E., Lees, D.C., Vane-Wright, R.I., Kremen, C., Leonard, J.A., Wayne, R.K. 2001. Examining monophyly in a large radiation of Madagascan butterflies (Lepidoptera: Satyrinae: Mycalesina) based on mitochondrial DNA data. *Molecular Phylogenetics and Evolution* 20: 460–473.
- Torsvik, T.H., Tucker, R.D., Ashwal, L.D., Carter, L.M., Jamtveit, B., Vidyadharan, K.T., Venkataramana, P. 2000. Late Cretaceous India-Madagascar fit and timing of break-up related magmatism. *Terra Nova* 12: 220–224.
- Vaidya, G., Lohman, D.J., Meier, R. 2011. SequenceMatrix: concatenation software for the fast assembly of multi-gene datasets with character set and codon information. *Cladistics* 27: 171–180.
- Vences, M., Kosuch, J., Glaw, F., Bohme, W., Veith, M. 2003a. Molecular phylogeny of hyperoliid treefrogs: biogeographic origin of Malagasy and Seychellean taxa and re-analysis of familial parphyly. *Journal of Zoological Systematics and Evolutionary Research* 41: 205–215.

- Vences, M., Vieites, D.R., Glaw, F., Brinkmann, H., Kosuch, J., Veith, M., Meyer, A. 2003b. Multiple overseas dispersal in amphibians. *Proceedings of the Royal Society* 270: 2435–2442.
- Vences, M. 2004. Origin of Madagascar's extant fauna: a perspective from amphibians, reptiles and other non-flying vertebrates. *Italian Journal of Zoology* 71, S2: 217–228.
- Warren, B.H., Strasberg, D., Bruggemann, J.H., Prys-Jones, R.P., Thébaud, C. 2010. Why does the biota of the Madagascar region have such a strong Asiatic flavour? *Cladistics* 26: 526–538.
- Weirauch, C. 2008. Cladistic analysis of Reduviidae (Heteroptera: Cimicomorpha) based on morphological characters. *Systematic Entomology* 33: 229–274.
- Weirauch, C. 2010. *Tribelocodia ashei*, new genus and new species of Reduviidae (Insecta: Hemiptera), has implications on character evolution in Ectrichodiinae and Tribelocephalinae. *Insect Systematics and Evolution* 41: 103–122.
- Weirauch, C., Munro, J.B. 2009. Molecular phylogeny of the assassin bugs (Hemiptera: Reduviidae), based on mitochondrial and nuclear ribosomal genes. *Molecular Phylogenetics and Evolution* 53: 287–299.
- Wikström, N., Avino, M., Razafimandimbison, S.G., Bremer, B. 2010. Historical biogeography of the coffee family (Rubiaceae, Gentianales) in Madagascar: case studies from the tribes Knoxiaceae, Naucleaceae, Paederiaceae and Vangueriaceae. *Journal of Biogeography* 37: 1094–1113.
- Wirta, H., Orsini, L., Hanski, I. 2008. An old adaptive radiation of forest dung beetles in Madagascar. *Molecular Phylogenetics and Evolution* 47: 1076–1089.
- Wirta, H., Viljanen, H., Orsini, L., Montreuil, O., Hanski, I. 2010. Three parallel radiations of Canthonini dung beetles in Madagascar. *Molecular Phylogenetics and Evolution* 57: 710–727.
- de Wit, M.J. 2003. Madagascar: heads it's a continent, tails it's an island. *Annual Review of Earth and Planetary Sciences* 31: 213–248.
- Yoder, A.D., Nowak, M.D. 2006. Has vicariance or dispersal been the predominant biogeographic force in Madagascar? Only time will tell. *Annual Review of Ecology, Evolution, and Systematics* 37: 405–431.
- Yu, Y., Harris, A.J., He, X. 2010. S-DIVA (Statistical Dispersal-Vicariance Analysis): a tool for inferring biogeographic histories. *Molecular Phylogenetics and Evolution* 56: 848–850.
- Yu, Y., Harris, A.J., Blair, C., He, X. 2015. RASP (Reconstruct Ancestral State in Phylogenies): a tool for historical biogeography. *Molecular Phylogenetics and Evolution* 87: 46–49.

- Zakharov, E.V., Smith, C.R., Lees, D.C., Cameron, A., Vane-Wright, R.I., Sperling, F.A.H. 2004. Independent gene phylogenies and morphology demonstrate a Malagasy origin for a wide-ranging group of swallowtail butterflies. *Evolution* 58: 2763–2782.
- Zhou, L., Su, Y.C.F., Thomas, D.C., Saunders, R.M.K. 2012. ‘Out-of-Africa’ dispersal of tropical floras during the Miocene climatic optimum: evidence from *Uvaria* (Annonaceae). *Journal of Biogeography* 39: 322–335.

Figure 3.1. Examples and criteria of limited (top) and extreme (bottom) sexual dimorphism in Ectrichodiinae. Arrows in top panel indicate distal margin of wings.



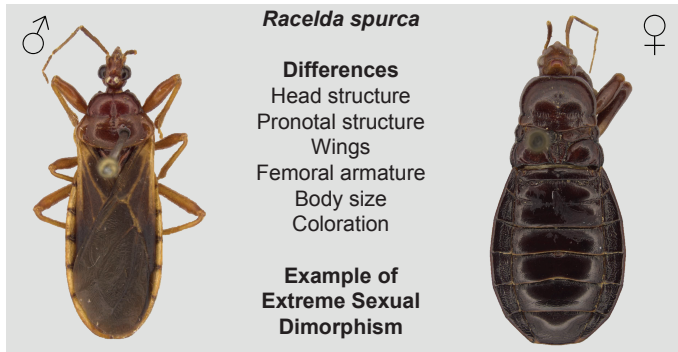
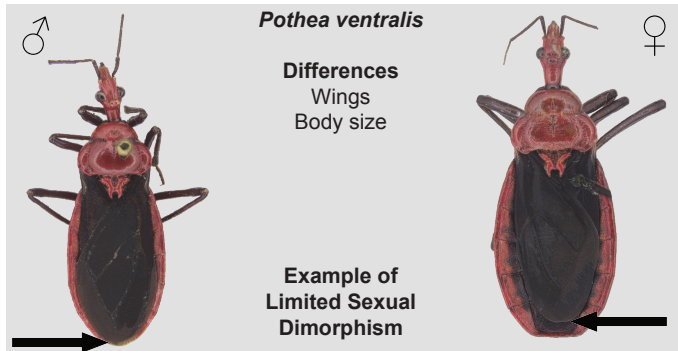


Figure 3.2. Best tree based on ML analysis of 56 taxa, four G-INS-I aligned gene partitions (COI, 18S, 28S D2, 28S D3–D5), and GTR+ $\Gamma$ +I model of sequence evolution (final Ln = -25,426.19528). Terminal branches are colored by occurrence of a taxon in a given biogeographic region, which corresponds to the map legend. Numbers in gray circles refer to nodes discussed in the text. Bootstrap (BS) values  $\geq 50$  are reported below branches, with the exception that BS values are reported next to the node of *Microstemmatoides atrocyanea* and *Centraspis ducalis*, as well as the node of the clade containing *Microstemmatoides atrocyanea* and *Ectrichodia crux*.

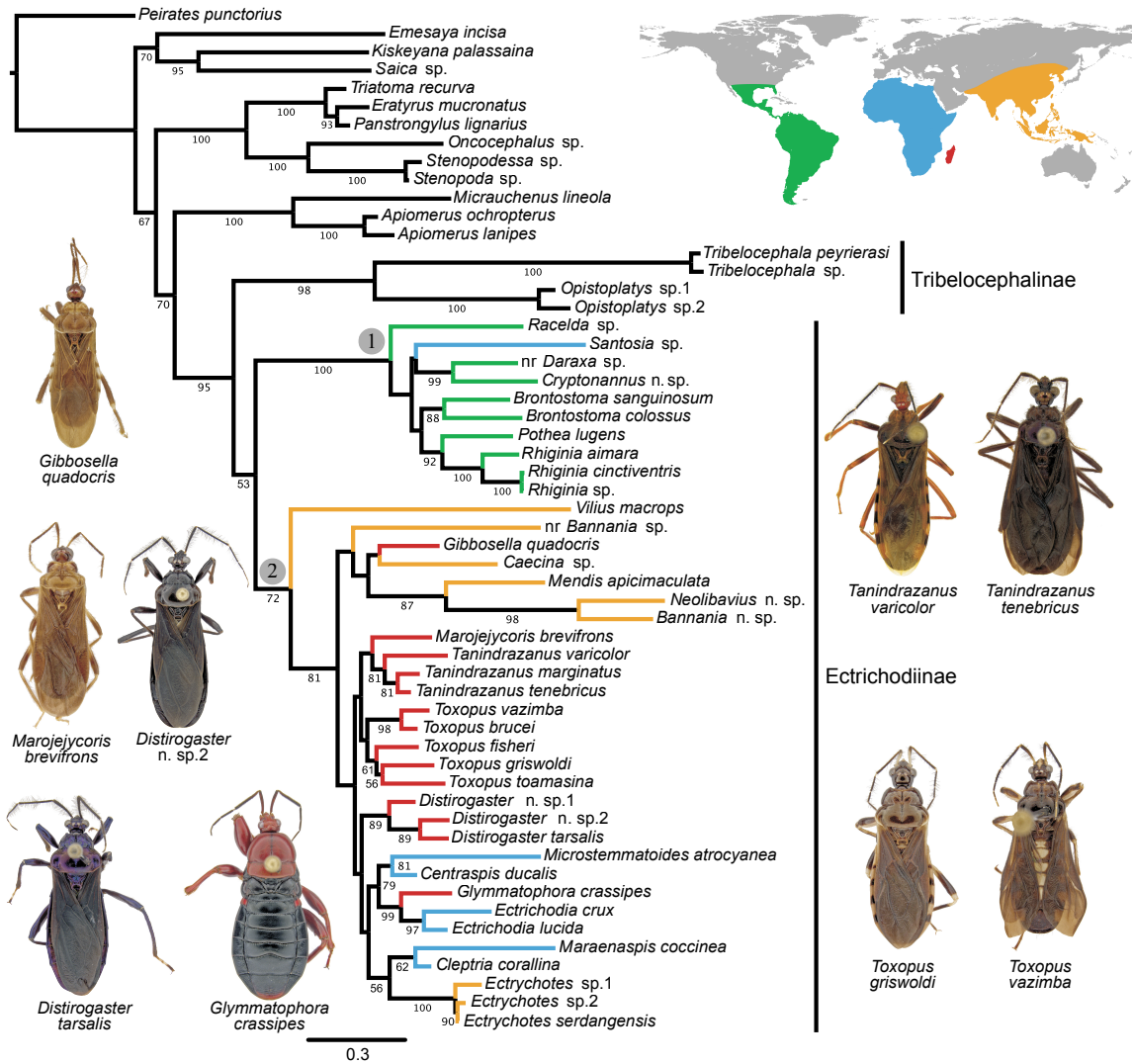


Figure 3.3. Strict consensus tree from 938 parsimonious trees from a TNT analysis of 110 taxa (outgroups removed from figure) and a combined morphological and molecular dataset (length = 7,250 steps; RI = 0.587; CI = 0.356). Terminal branches are colored by taxon distribution, which corresponds to the map legend. Numbers in gray circles refer to nodes discussed in the text and SM10. Bootstrap values  $\geq 50$  are reported below branches. Examples of morphological characters that are discussed in the text are given: A.–D. Dorsal (A., B.) and lateral (C., D.) head morphology: A. *Glymmatophora crassipes*, B. *Toxopus tibialis*, C. *Maraenaspis bidens*, D. *Tanindrazanus kathrynae*; E. Lateral antennal morphology of *Toxopus insignis*; Dorsal (F.–H.), lateral (I.), and ventral (J.): F. *Maraenaspis bidens*, G. *Toxopus italaviana*, H. *Marojejycoris notadichroa*, I. *Gibbosella conisimilis*, J. *Tanindrazanus irwini*; K. Metathoracic gland evaporatorium of *Tanindrazanus varicolor* in lateral view; L. Ventral mid leg morphology of *Glymmatophora crassipes*. M. Hemelytral morphology of *Gibbosella planiscutum*; N. Ventral abdominal morphology of *Distirogaster tarsalis*. A., C.–G., and I.–M. modified from Forthman et al. (in press). Character numbers and character state codings are listed in Table 3.2. Abbreviations used in figures: 1A, first anal vein; aa, antennal articulation; ain, antennal insertion site; ap, antennal pseudoarticulation; apl, anterior pronotal lobe; app, anterolateral pronotal projection; asp, anterior femoral subapical protuberance; bf, basiflagellomere; cl, clypeus; co, corium; cp, corial pterostigma; Cu, cubitus; df, distiflagellomere; e, compound eye; exM, extension of M beyond M+Cu distal junction; fmp, femoral medial protuberance; g, gula; is, interocular sulcus; L2, labial segment II (first visible segment); L3, labial segment III (second visible segment); M, media; mc, metacoxa; mf, mid femur; mge, metathoracic gland evaporatorium; mms, transverse suture between meso- and metasterna; mss, mesosternum; mt, mid trochanter; mts, metasternum; p, pedicel; pa, papillae; pc, sternal paramedian carinae; pcd, postclypeal depression; ppl, posterior pronotal lobe; psp, posterior femoral subapical protuberance; pts, pronotal transverse suture; R, radius; s, scape; s2, sternite II (fused sternites I and II); s3, sternite III; sc, scutellum; sis, sternal intersegmental suture; sl, synthlipsis; sld, sternal medial longitudinal depression.

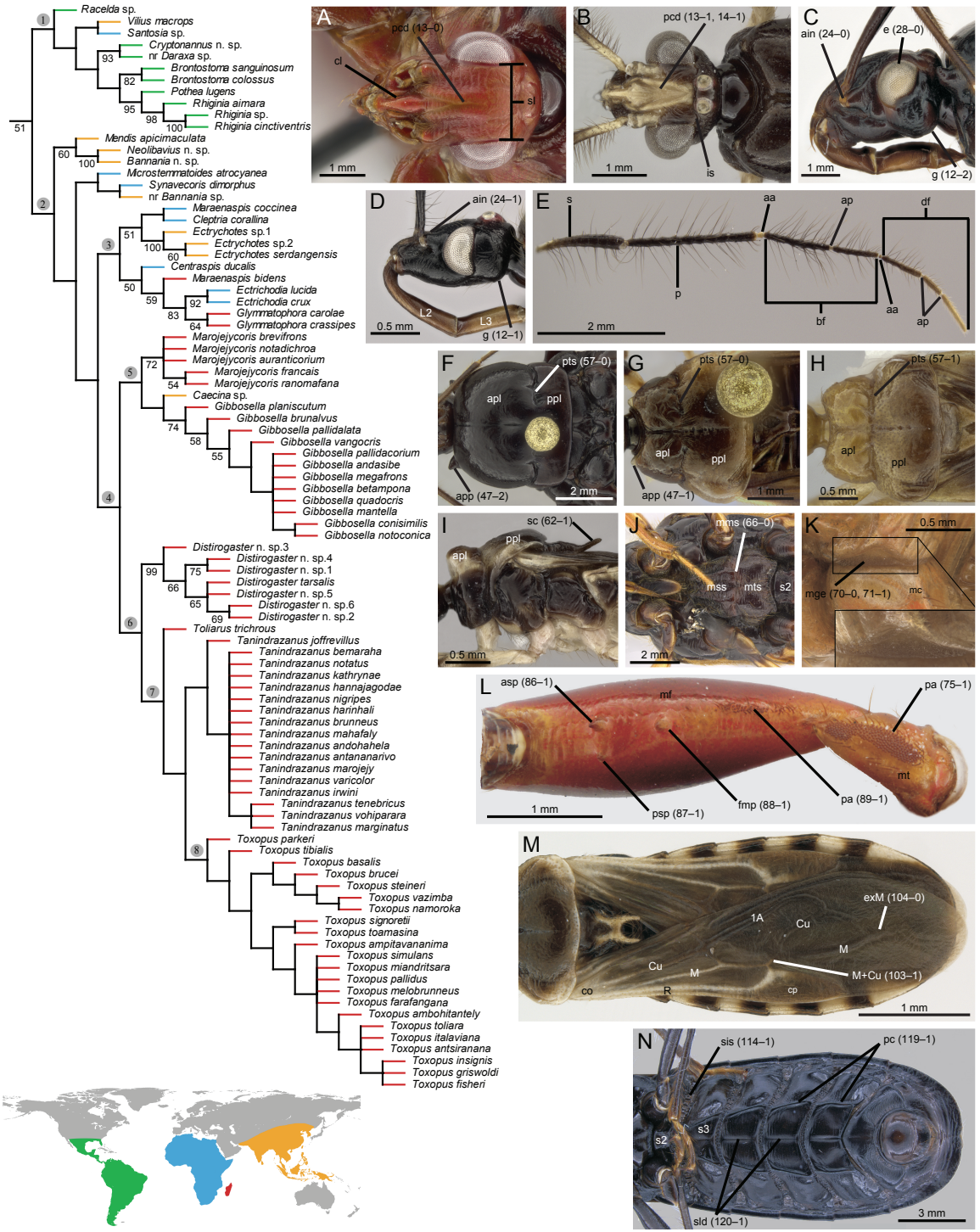


Figure 3.4. Ancestral range reconstructions from DEC (left) and DEC+j (right) models for select nodes. Pie charts indicate the relative likelihoods of each reconstructed distribution, with geographical distributions color coded according to the legend at the top of the figure. The DEC+j model performed significantly better than the DEC: DEC+j LnL =  $-27.10$ ; DEC LnL =  $-39.09$ ; LRT p-value =  $9.80e^{-7}$ . Select dispersal events contributing to Madagascar's Ectrichodiinae diversity is shown for each model. Dispersal events unique to each model are indicated by dashed arrows.

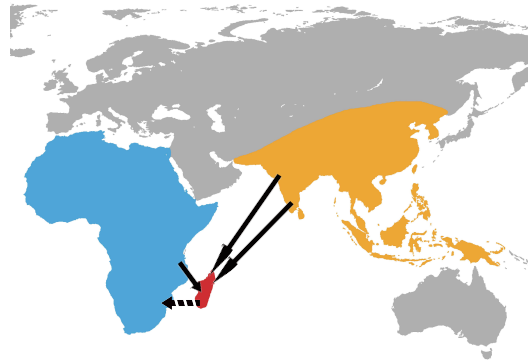
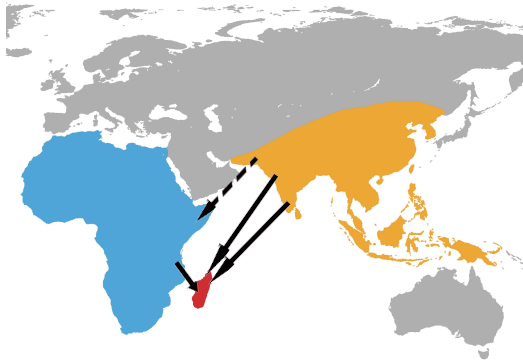
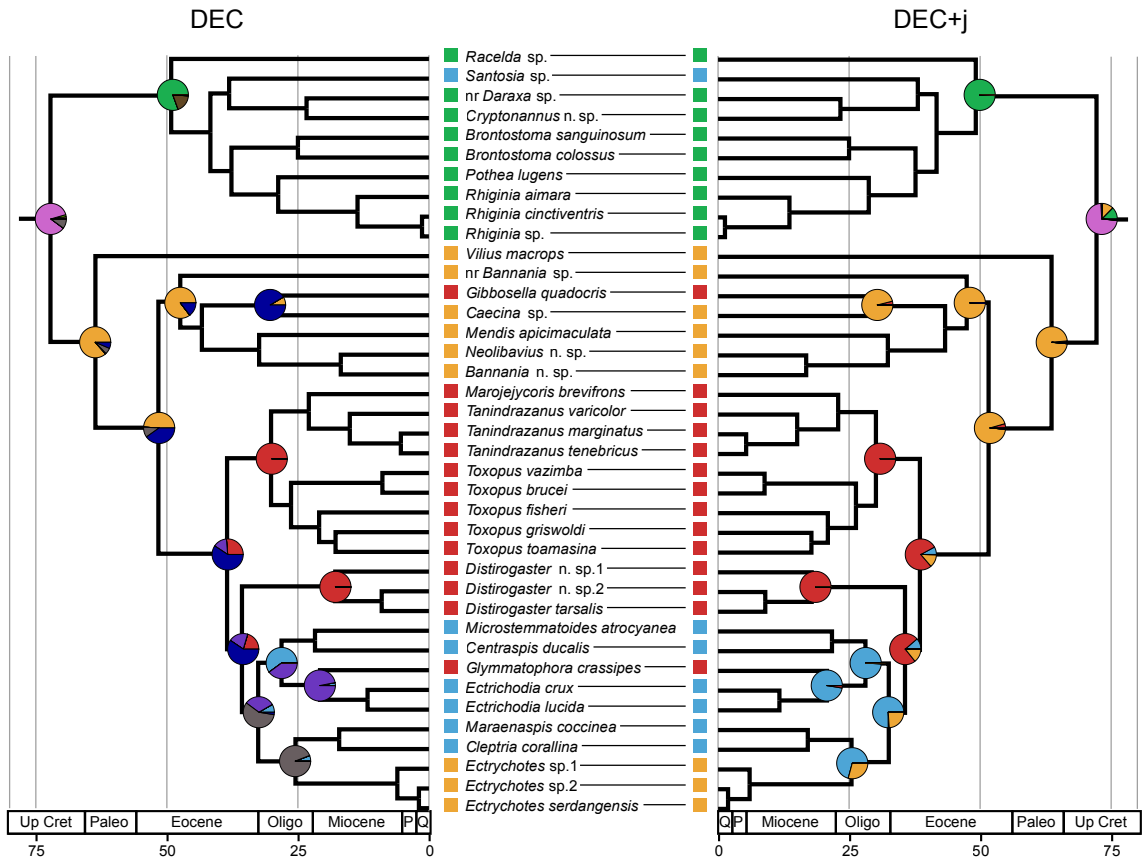
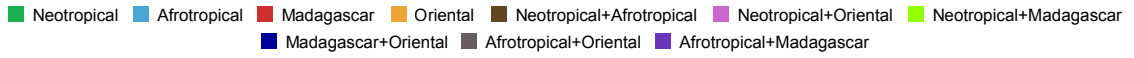


Table 3.1. Taxon sampling, USI codes, ethanol molecular specimen collection numbers (RCW), depositories, and GenBank accession numbers for sequenced specimens. Asterisks indicate sequences retrieved from GenBank.

Subfamily	Taxon	USI	RCW	Depository	GenBank accession no.				
					COI	18S	28S D2	28D3-D5	
Ectrichodimae	<i>Bannania</i> n. sp.	UCR_ENT 00119027	3064	UCR	KT221890	KT221910	KT221939	KT221968	
	nr <i>Bannania</i> sp.	UCR_ENT 00004465	609	UCR	KT221908	KT221937	KT221966	KT221996	
	<i>Brontostoma colossus</i>	UCR_ENT 00119017	3049	UCR	KT221891	KT221911	KT221940	KT221969	
	<i>Brontostoma sanguinosum</i>	UCR_ENT 00002617	1243	UCR	KT221892	KT221912		KT221970	
	<i>Caecina</i> sp.	UCR_ENT 00004440	2636	UCR	KT221893	KT221913	KT221941	KT221971	
	<i>Centraspis ducalis</i>	UCR_ENT 00119021	3014	UCR	KT221894	KT221914	KT221942	KT221972	
	<i>Cleptria corallina</i>	AMNH_PBI 00218770	14	UCR	FJ230462*	FJ230543*		FJ230621*, FJ230700*	
	<i>Cryptonannus</i> n. sp.	UCR_ENT 00002757	1433	UCR	KT221895	KT221915		KT221973	
	nr <i>Daraxa</i> sp.	UCR_ENT 00119016	3076	UCR	KT221909	KT221938	KT221967	KT221997	
	<i>Distirogaster tarsalis</i>	UCR_ENT 00006366	2898	CAS	KR606396	KT221918	KT221945	KT221976	
<i>Distirogaster</i> n. sp.1	UCR_ENT 00007158	2881	UCR	KR606393	KT221916	KT221943	KT221974		
<i>Distirogaster</i> n. sp.2	UCR_ENT 00088090	3018	UCR	KR606395	KT221917	KT221944	KT221975		
<i>Ectrichodia crux</i>	UCR_ENT 00119022	3026	UCR	KT221896	KT221919	KT221946	KT221977		
<i>Ectrichodia lucida</i>	AMNH_PBI 00218769	13	UCR		FJ230461*	FJ230542*	FJ230620*, FJ230699*		
<i>Ectrychotes serdangensis</i>	AMNH_PBI 00218830	76	UCR		FJ230479*	FJ230560*	FJ230638*, FJ230717*		
<i>Ectrychotes</i> sp.1	UCR_ENT 00119028	571	UCR	KT221897	KT221920	KT221947	KT221978		
<i>Ectrychotes</i> sp.2	AMNH_PBI 00218932	188	UCR	JQ942322*	FJ230503*	FJ230584*	FJ230661*, FJ230740*		
<i>Gibbosella quadocris</i>	UCR_ENT 00044860	2938	UCR	KR606406					
<i>Glymmatophora crassipes</i>	UCR_ENT 00088087	3020	UCR	KR606415	KT221921	KT221948	KT221979		
<i>Maraenaspis coccinea</i>	AMNH_PBI 00218772	16	UCR		FJ230463*	FJ230544*			
<i>Marojejcoris brevifrons</i>	UCR_ENT 00006480	2923	CAS	KR606417	KT221922		KT221980		



Table 3.1. (Continued).

Subfamily	Taxon	USI	RCW	Depository	COI	GenBank accession no.		
						18S	28S D2	28D3-D5
	<i>Mendis apicimaculata</i>	UCR_ENT 00119026	2647	UCR	KT221898	KT221923	KT221949	KT221981
	<i>Microstemmaatoides atrocyanea</i>	UCR_ENT 00119029	3029	UCR	KT221899	KT221924	KT221950	KT221982
	<i>Neolibavius</i> n. sp.	UCR_ENT 00003965	1830	UCR	KT221900	KT221925	KT221951	KT221983
	<i>Pothea lugens</i>	UCR_ENT 00119018	3304	UCR	KT221902	KT221927	KT221953	
	<i>Racelda</i> sp.	AMNH_PBI 00218801	41	UCR		FJ230472*	FJ230553*	FJ230631*, FJ230710*
	<i>Rhiginia aimara</i>	UCR_ENT 00119019	3077	UCR	KT221903	KT221928	KT221954	
	<i>Rhiginia cinciventris</i>	AMNH_PBI 00218891	139	UCR	JQ942333*	FJ230490*	FJ230571*	FJ230648*, FJ230727*
	<i>Rhiginia</i> sp.	UCR_ENT 00119020	3047	UCR	KT221904	KT221929	KT221955	KT221985
	<i>Santostia</i> sp.	UCR_ENT 00004138	2046	UCR	KT221905	KT221930	KT221956	KT221986
	<i>Tanindrazanus marginatus</i>	UCR_ENT 00006887	2902	CAS	KR606423	KT221931	KT221957	KT221987
	<i>Tanindrazanus tenebricus</i>	UCR_ENT 00006723	2931	CAS	KR606430	KT221932	KT221958	KT221988
	<i>Tanindrazanus varicolor</i>	UCR_ENT 00006482	2949	UCR	KR606433	KT221933	KT221959	KT221989
	<i>Toxopus brucei</i>	UCR_ENT 00045338	2901	UCR	KR606438		KT221960	KT221990
	<i>Toxopus fisheri</i>	UCR_ENT 00045431	2910	CAS	KR606446		KT221961	KT221991
	<i>Toxopus griswoldi</i>	UCR_ENT 00006435	2916	CAS	KR606448	KT221934	KT221962	KT221992
	<i>Toxopus toamasina</i>	UCR_ENT 00007056	2951	SU	KR606457		KT221963	
	<i>Toxopus vazimba</i>	UCR_ENT 00006472	2953	CAS	KR606462	KT221935	KT221964	KT221993
	<i>Vilius macrops</i>	UCR_ENT 00119025	3068	UCR	KT221907	KT221936		KT221995
Emesinae	<i>Emesaya incisa</i>	AMNH_PBI 00219017	282	UCR	JQ942323*	FJ230515*	FJ230598*	FJ230672*, FJ230751*
Harpactorinae	<i>Apiomerus lanipes</i>	AMNH_PBI 00219016	281	UCR	JQ942298*	FJ230514*	FJ230597*	FJ230671*, FJ230750*

Table 3.1. (Continued).

Subfamily	Taxon	USI	RCW	Depository	COI	GenBank accession no.		
						18S	28S D2	28D3-D5
Peiratinae	<i>Apiomerus ochropterus</i>	AMNH_PBI 00218777	22	UCR		FJ230466*	FJ230548*	FJ230625*, FJ230704*
	<i>Micrauchenus lineola</i>	AMNH_PBI 00218790	35	UCR	JQ942329*	FJ230471*	FJ230552*	FJ230630*, FJ230709*
	<i>Peirates punctorius</i>	AMNH_PBI 00218960	216	UCR		FJ230508*	FJ230590*	FJ230666*, FJ230745*
Saicinae	<i>Kiskeyana palassaina</i>	AMNH_PBI 00190561	10	USNM		FJ230460*	FJ230541*	FJ230619*, FJ230698*
Stenopodainae	<i>Saica</i> sp.	AMNH_PBI 00218796	42	UCR		FJ230473*	FJ230554*	FJ230632*, FJ230711*
	<i>Oncocephalus</i> sp.	UCR_ENT 00000182	79	UCR		FJ230481*	FJ230562*	FJ230640*, FJ230719*
	<i>Stenopoda</i> sp.	AMNH_PBI 00218904	154	UCR		FJ230493*	FJ230574*	FJ230651*, FJ230730*
Triatominae	<i>Stenopodessa</i> sp.	UCR_ENT 00000078	398	FCAP		FJ230532*	FJ230611*	FJ230688*, FJ230767*
	<i>Eratyrus micronatus</i>	UCR_ENT 00052168	1811	UCR		JQ897555*	JQ897635*	JQ897711*
	<i>Panstrongylus lignarius</i>	UCR_ENT 00052166	1813	UCR		JQ897584*	JQ897656*	JQ897741*
	<i>Triatoma recurva</i>	AMNH_PBI 00218913	170	UCR		FJ230496*	FJ230577*	FJ230654*, FJ230733*
Tribelocephalinae	<i>Opistoplatys</i> sp.1	UCR_ENT 00119024	3056	UCR	KT221901	KT221926	KT221952	KT221984
	<i>Opistoplatys</i> sp.2	UCR_ENT 00052187	1592	UCR		JQ897612*	JQ897682*	JQ897767*
	<i>Tribelocephala peyrierasi</i>	AMNH_PBI 00219033	287	CAS		FJ230521*	FJ230601*	
	<i>Tribelocephala</i> sp.	UCR_ENT 00119023	3048	UCR	KT221906		KT221965	KT221994

Table 3.2. Morphological characters and character state codings.

Character no.	Character	Character states
<b>GENERAL</b>		
0	Wing form	(0) Macropterous; (1) Brachypterous; (2) Apterous; (3) Micropterous
1	Body length	(0) Small; (1) Medium; (2) Large
2	General coloration	(0) Dull, ranging from pale to dark brown/black; (1) Metallic; (2) Red and black, sometimes with yellow or white markings; (3) Red and dull; (4) Orange and black; (5) Red and metallic
3	Bulbous hairs on integument	(0) Present; (1) Absent
4	Vestiture	(0) Very dense; (1) Moderately dense; (2) Sparse; (3) Glabrous
<b>HEAD</b>		
5	Relative head length in dorsal view	(0) As long as wide; (1) Longer than wide; (2) Wider than long
6	Ventral surface of head	(0) Flat; (1) Anteromedially depressed; (2) Medially depressed; (3) Paramedially depressed
7	Depth of ventral head depression	(0) Shallow; (1) Deep
8	Ventral tubercles on head	(0) Absent; (1) Present
9	Ventrolateral tubercle on gula	(0) Absent; (1) Present
10	Anteocular length	(0) Shorter than postocular; (1) As long as postocular; (2) Longer than postocular
11	Postocular-neck constriction	(0) Absent, gradually transition; (1) Present, distinctly constricted
12	Gula shape in lateral view	(0) Flat, conforming to rounded head shape; (1) Moderately swollen ventrolaterally, not produced beyond ventral head margin; (2) Greatly swollen ventrolaterally, produced beyond ventral head margin
13	Postclypeal depression	(0) Flat; (1) Depressed
14	Width of postclypeal depression	(0) Narrow; (1) Broad
15	Depth of postclypeal depression	(0) Shallow; (1) Deep
16	Extension of postclypeal depression	(0) Posterior margin of clypeus to middle of synthlipsis; (1) Posterior margin of clypeus to interocular sulcus
17	Maxillary plate in lateral view	(0) Reaching or nearly reaching dorsal clypeal margin; (1) Not reaching dorsal clypeal margin
18	Clypeal process	(0) Absent; (1) Present

Table 3.2. (Continued).

Character no.	Character	Character states
19	Shape of clypeus	(0) Without dorsal projection; (1) With dorsal projection
20	Clypeal apex relative to labral base	(0) Not elevated; (1) Elevated
21	Antennifer armature	(0) Unarmed; (1) With dorsolateral tubercle
22	Antennal shield	(0) Absent, not expanded; (1) Present, expanded
23	Antennal insertion site in lateral view	(0) Not concealed by antennal shield; (1) Concealed by antennal shield
24	Point of antennal insertion	(0) Sublateral or lateral; (1) Dorsal
25	Presence of ocellar lens	(0) Present; (1) Absent
26	Orientation of ocellar lens	(0) Medially; (1) Laterally
27	Ocellar lens size	(0) Large; (1) Small
28	Eye height in lateral view	(0) Less than or half of head height; (1) More than half of head height
29	Eye shape	(0) Adpressed; (1) Globbose
30	Anterior eye margin	(0) Convex; (1) Concave
31	Posterior eye margin	(0) Concave; (1) Straight
32	Dorsal eye margin	(0) Not attaining dorsal head surface; (1) Attaining dorsal head surface
33	Ventral eye margin	(0) Not attaining ventral head surface; (1) Attaining ventral head surface
34	Synthlipsis width	(0) About width of eye; (1) About 1.5 times width of eye; (2) About 2 times width of eye; (3) About 2.5 times width of eye; (4) About 3 times or more width of eye
35	Scapus length	(0) Not surpassing clypeal apex; (1) Surpassing clypeal apex
36	Pedicel curvature	(0) Straight; (1) Curved
37	Scapo-pedicellar articulation	(0) Slightly bent; (1) Strongly bent, pedicel and flagellomeres point posteriad
38	Antennal segmentation	(0) 4-segmented; (1) 6-segmented; (2) 7-segmented; (3) 8-segmented; (4) >8 segments
39	Labrum structure	(0) Subdivided by transverse membrane; (1) Completely sclerotized
40	Ventral margin of labial segment III	(0) Straight; (1) Convex; (2) Concave

Table 3.2. (Continued).

Character no.	Character	Character states
41	Labial segment III length	(0) Longer than labial segment II; (1) Shorter than labial segment II; (2) Subequal to labial segment II
<b>THORAX</b>		
42	Pronotal length	(0) Wider than long; (1) As long as wide; (2) Longer than wide
43	Pronotal collar	(0) Distinct in dorsal view; (1) Indistinct in dorsal view
44	Anterior pronotal margin shape	(0) Medially concaved; (1) Straight or nearly straight
45	Anterior pronotal lobe shape	(0) Flat; (1) Dorsally elevated, conical
46	Anterior pronotal lobe structure	(0) Rugose; (1) Smooth; (2) Tuberculate
47	Anterolateral armature of pronotum	(0) Absent; (1) Small tubercle; (2) Strongly projecting process
48	Lateral carinae on anterior pronotal lobe	(0) Absent; (1) Present
49	Anterior pronotal lobe length	(0) Longer than posterior lobe; (1) Shorter than posterior lobe; (2) As long as posterior lobe
50	Anterior pronotal lobe width	(0) More than half as wide but not as wide as posterior lobe; (1) As wide as posterior lobe
51	Posterior pronotal lobe structure	(0) Striated; (1) Smooth; (2) Punctate; (3) Tuberculate; (4) Longitudinal ridges
52	Posterior pronotal lobe lateral depressions	(0) Present; (1) Absent
53	Posterior pronotal lobe lateral depression structure	(0) Not foveate; (1) Foveate
54	Extension of pronotal longitudinal depression on anterior lobe	(0) Reaching or nearly reaching anterior margin; (1) Not reaching anterior margin; (2) Absent
55	Presence of pronotal longitudinal depression on posterior lobe	(0) Absent; (1) Present
56	Structure of pronotal longitudinal depression on posterior lobe	(0) Not foveate; (1) Foveate
57	Pronotal transverse suture	(0) Incomplete, divided by paramedian ridges; (1) Complete, not divided by paramedian ridges
58	Pronotal transverse suture structure	(0) Not foveate; (1) Foveate

Table 3.2. (Continued).

Character no.	Character	Character states
59	Scutellar processes	(0) None, scutellum triangular; (1) Two paramedian apical processes; (2) Two paramedian and one medial apical processes; (3) Two paramedian and two lateral processes
60	Distance between paramedian scutellar processes	(0) Narrow; (1) Broad
61	Length of paramedian scutellar processes	(0) Short; (1) Long
62	Orientation of paramedian scutellar processes	(0) Horizontally directed; (1) Dorsally directed
63	Scutellar disc surface	(0) Medially depressed; (1) Flat
64	Prosternal stridulatory process length	(0) Short; (1) Long, surpassing posterior margin of fore coxal cavity
65	Shape of apex of prosternal stridulatory process	(0) Acute; (1) Rounded
66	Transverse suture between meso- and metasterna	(0) Complete; (1) Incomplete, at least partially
67	Mesosternal surface	(0) Medially and laterally depressed; (1) Medially depressed; (2) Flat
68	Metasternal surface	(0) Medially depressed; (1) Flat or convex
69	Presence of metathoracic gland	(0) Absent; (1) Present
70	Presence of metathoracic gland evaporatorium	(0) Present; (1) Absent
71	Metathoracic gland evaporatorium size	(0) Small, not visible in lateral view; (1) Large, visible in lateral view
<b>LEG</b>		
72	Shape of fore coxa	(0) Globular; (1) Elongate
73	Shape of fore femur	(0) Slender; (1) Incrassate
74	Presence of papillae on fore trochanter	(0) Absent; (1) Present
75	Presence of papillae on mid trochanter	(0) Absent; (1) Present
76	Presence of papillae on hind trochanter	(0) Absent; (1) Present
77	Presence of anteroventral row of spines/tubercles on fore femur	(0) Absent; (1) Present
78	Presence of posteroventral row of spines/tubercles on fore femur	(0) Absent; (1) Present
79	Presence of anterior subapical tubercle on fore femur	(0) Absent; (1) Present
80	Presence of posterior subapical tubercle on fore femur	(0) Absent; (1) Present
81	Presence of medial tubercle on fore femur	(0) Absent; (1) Present

Table 3.2. (Continued).

Character no.	Character	Character states
82	Presence of carina on fore femur	(0) Absent; (1) Ventrally entirely carinate; (2) Ventrally carinate basally
83	Presence of papillae on fore femur	(0) Absent; (1) Present
84	Presence of anteroventral row of spines/tubercles on mid femur	(0) Absent; (1) Present
85	Presence of posteroventral row of spines/tubercles on mid femur	(0) Absent; (1) Present
86	Presence of anterior subapical tubercle on mid femur	(0) Absent; (1) Present
87	Presence of posterior subapical tubercle on mid femur	(0) Absent; (1) Present
88	Presence of medial tubercle on mid femur	(0) Absent; (1) Present
89	Presence of papillae on mid femur	(0) Absent; (1) Present
90	Presence of anteroventral row of spines/tubercles on hind femur	(0) Absent; (1) Present
91	Presence of anterior subapical tubercle on hind femur	(0) Absent; (1) Present
92	Presence of posterior subapical tubercle on hind femur	(0) Absent; (1) Present
93	Presence of subapical medial tubercle on hind femur	(0) Absent; (1) Present
94	Presence of medial tubercle on hind femur	(0) Absent; (1) Present
95	Presence of fossula spongiosa on fore tibia	(0) Present; (1) Absent
96	Presence of fossula spongiosa on mid tibia	(0) Present; (1) Absent
97	Protibial groove	(0) Absent; (1) Present
<b>FOREWING</b>		
98	Extension of corium on fore wing	(0) Restricted to areas adjacent to basal wing veins and with pterostigma-like appearance; (1) Well-developed corium
99	Vestiture of corium on fore wing	(0) Long, simple setae; (1) Glabrous
100	Basal part of M and Cu on fore wing	(0) Separate veins; (1) Forming one vein, at least partially
101	Presence of cubital cell on fore wing	(0) Absent; (1) Present
102	Shape of cubital cell on fore wing	(0) Quadrate; (1) Hexagonal
103	Distal part of M and Cu on fore wing	(0) Separate veins; (1) Fused basally
104	Extension of distal part of M on fore wing	(0) Extends beyond distal M+Cu junction; (1) No extension beyond distal M+Cu junction

Table 3.2. (Continued).

Character no.	Character	Character states
<b>ABDOMEN</b>		
105	Color pattern of dorsal laterotergites	(0) Concolor; (1) Transversely bicolor; (2) Longitudinally bicolor
106	Expansion of dorsal laterotergite II	(0) Not expanded; (1) Laterally expanded; (2) Posterior tubercle
107	Dorsal laterotergite III armature	(0) Unarmed; (1) Posterior tubercle
108	Dorsal laterotergite IV armature	(0) Unarmed; (1) Posterior tubercle
109	Dorsal laterotergite V armature	(0) Unarmed; (1) Posterior tubercle
110	Dorsal laterotergite VI armature	(0) Unarmed; (1) Posterior tubercle
111	Connection of dorsal laterotergite and mediotergite	(0) Separated by membrane; (1) Sclerotized
112	Connection of ventral laterotergites and mediotergites III-VII	(0) Separated by membrane; (1) Sclerotized
113	Ventral connexival suture on sternite II	(0) Absent; (1) Present
114	Carinulation of intersegmental suture between abdominal sternites II and III	(0) Not carinulate; (1) Entirely carinulate
115	Carinulation of intersegmental suture between abdominal sternites III and IV	(0) Not carinulate; (1) Entirely carinulate; (2) Laterally carinulate
116	Carinulation of intersegmental suture between abdominal sternites IV and V	(0) Not carinulate; (1) Entirely carinulate; (2) Laterally carinulate
117	Carinulation of intersegmental suture between abdominal sternites V and VI	(0) Not carinulate; (1) Entirely carinulate; (2) Laterally carinulate
118	Carinulation of intersegmental suture between abdominal sternites VI and VII	(0) Not carinulate; (1) Entirely carinulate; (2) Laterally carinulate
119	Presence of paramedian carinae on abdominal sternites	(0) Absent; (1) Present
120	Abdominal sternite surface	(0) Flat; (1) Medially depressed; (2) Convex or keeled
121	Spiracle shape	(0) Ovoid or elliptical; (1) Circular
<b>MALE GENITALIA</b>		
122	Anteroposterior thickness of median pygophore process	(0) Flat; (1) Thickened, at least partially
123	Lateral thickness of median pygophore process	(0) Flat; (1) Thickened, at least partially
124	Lateral shape of median pygophore process	(0) Subtriangular; (1) Blade-like; (2) Hook-like; (3) Knob-like; (4) Rod-like
125	Shape of median pygophore process in caudal view	(0) Subtriangular; (1) Subquadrate; (2) Rod-like; (3) Spade-like



Table 3.2. (Continued).

Character no.	Character	Character states
126	Apex of median pygophore process	(0) Entire; (1) Medially notched, bifid
127	Process on ponticulus basilaris	(0) Absent; (1) Present
128	Median process on ductifer	(0) Absent; (1) Present
129	Length of basal plate extension	(0) Shorter than basal plate; (1) As long as basal plate; (2) Longer than basal plate
130	Posterior margin of dorsal phallothecal sclerite	(0) Rounded; (1) Medially notched
131	Armature of dorsal phallothecal sclerite	(0) Unarmed; (1) Armed with denticle-like processes
132	Presence of vesica on endosoma	(0) Absent; (1) Present
133	Sclerotization of posterior endosomal margin	(0) Membranous; (1) Entirely sclerotized; (2) Paramedian sclerites; (3) Slender sclerotized processes; (4) Sclerotized lateral margin; (5) Sclerotized lateral hooks adjacent to fan-like lobes and with a medial dorsally projecting spine
134	Medial sclerotization of posterior endosomal region	(0) Membranous; (1) Slightly sclerotized; (2) Distinctly sclerite
135	Lateral sclerotization of posterior endosomal region	(0) Membranous; (1) Sclerotized
136	Medial sclerotization of medial endosomal region	(0) Membranous; (1) Slightly sclerotized; (2) Paramedian sclerites
137	Lateral sclerotization of medial endosomal region	(0) Membranous; (1) Sclerotized
138	Medial sclerotization of anterior endosomal region	(0) Membranous; (1) Sclerotized
139	Lateral sclerotization of anterior endosomal region	(0) Membranous; (1) Sclerotized
<b>GLANDS</b>		
140	Presence of Brindley's gland	(0) Present; (1) Absent
141	Dorsal abdominal scent gland (DAG) I in adults	(0) Absent; (1) Present
142	DAG II in adults	(0) Absent; (1) Present
143	DAG III in adults	(0) Absent; (1) Present
144	Size of DAG III relative to DAGs I and II, external view	(0) Similar size; (1) Larger

Supplementary material 3.1. Morphological character matrix in nexus format for cladistic analysis.

```
#NEXUS
BEGIN TAXA;
  TITLE Taxa;
  DIMENSIONS NTAX=110;
  TAXLABELS
    Apiomerus_lanipes Apiomerus_ochropterus Bannania_n_sp
  Brontostoma_colossus Brontostoma_sanguinosum Caecina_sp Centraspis_ducalis
  Cleptria_corallina Cryptonannus_n_sp Distirogaster_n_sp1 Distirogaster_n_sp2
  Distirogaster_tarsalis Ectrichodia_crux Ectrichodia_lucida Ectrychotes_serdangensis
  Ectrychotes_sp1 Ectrychotes_sp2 Emesaya_incisa Eratyrus_mucronatus Gibbosella_quadocris
  Glymmatophora_crassipes Kiskeyana_palassaina Maraenaspis_coccinea
  Marojejycoris_brevifrons Mendis_apicimaculata Micrauchenus_lineola
  Microstemmatoides_atrocyanea Neolibavivus_n_sp nr_Bannania_sp nr_Daraxa_sp
  Oncocephalus_sp Opisthoplatys_sp1 Opisthoplatys_sp2 Panstrongylus_lignarius
  Peirates_punctorius Pothea_lugens Racelda_sp Rhiginia_aimara Rhiginia_cinctiventris
  Rhiginia_sp Saica_sp Santosia_sp Stenopoda_sp Stenopodessa_sp Tanindrazanus_marginatus
  Tanindrazanus_tenebricus Tanindrazanus_varicolor Toxopus_brucei Toxopus_fisheri
  Toxopus_grisworldi Toxopus_toamasina Toxopus_vazimba Triatoma_recurva
  Tribelocephala_peyrierasi Tribelocephala_sp Vilius_macrops Synavecoris_dimorphus
  Maraenaspis_bidens Glymmatophora_carolae Distirogaster_n_sp3 Distirogaster_n_sp4
  Distirogaster_n_sp5 Distirogaster_n_sp6 Toliarius_trichrous Tanindrazanus_joffrevillus
  Tanindrazanus_irwini Tanindrazanus_marojejy Tanindrazanus_mahafaly
  Tanindrazanus_harinjali Tanindrazanus_hannajagoda Tanindrazanus_kathrynae
  Tanindrazanus_notatus Tanindrazanus_antananarivo Tanindrazanus_nigripes
  Tanindrazanus_bemara Tanindrazanus_andohahela Tanindrazanus_brunneus
  Tanindrazanus_vohiparara Toxopus_signoretii Toxopus_parkeri Toxopus_tibialis
  Toxopus_basalis Toxopus_steineri Toxopus_namoroka Toxopus_ampitavananima
  Toxopus_italviana Toxopus_pallidus Toxopus_simulans Toxopus_insignis
  Toxopus_ambohitantely Toxopus_antsiranana Toxopus_toliara Toxopus_farafangana
  Toxopus_miandritsara Toxopus_melobrunneus Marojejycoris_auranticorium
  Marojejycoris_notadichroa Marojejycoris_francais Marojejycoris_ranomafana
  Gibbosella_vangocris Gibbosella_planiscutum Gibbosella_pallidalata Gibbosella_brunalvus
  Gibbosella_conisimilis Gibbosella_notoconica Gibbosella_pallidacorium Gibbosella_andasibe
  Gibbosella_betampona Gibbosella_megafrons Gibbosella_mantella
  ;
END;

BEGIN CHARACTERS;
  TITLE Character_Matrix;
  DIMENSIONS NCHAR=145;
  FORMAT DATATYPE = STANDARD GAP = - MISSING = ? SYMBOLS = " 0 1 2 3
4 5";
  MATRIX
```

*Apiomerus\_lanipes* 0201110-001000---000000-  
1011110010210001000010?111010000-100---10012101-  
000000000000000000000011110010001000001100000001111011001000001010101110  
*Apiomerus\_ochropterus* 0201110-000000---000000-  
1011110010110001000010?111010000-100---10012101-  
0000000000000000000000111100100010000011000000011?????????????????01110  
*Bannania\_n\_sp* 0031100-101100---100000-  
1001110000211010110010111101011110121000111001000000010-000010-0001----0000000-  
00021110000111111101110000000110101000111  
*Brontostoma\_colossus* 0221310-002110---100000-  
000011001001103011001011010000111011000011000101011100000001000001000000001100  
-001000000001200002011010000100000200000111  
*Brontostoma\_sanguinosum* 0221310-002110---000000-  
000011000001103012011011020100111011000011000101011100000001000001000000001100  
-0000000000001000002001110000100000200000111  
*Caecina\_sp* 0001110-002100---100000-  
1000110000111010011??122120201111?0110001100010000000000000000000000000000001100000-  
100211110001111101111000002000010100000111  
*Centraspis\_ducalis* 0241300-002110---  
100001000001100100110100200001101000001010310001110110001110000000100000100000  
0001100-000100000001122001010200000000200100000111  
*Cleptria\_corallina* 0101210-  
002111000100001000001100100110301210001111010011100110001110010001000000000000  
0000000000000000-000100000001111201011000000000110100000111  
*Cryptonannus\_n\_sp* 0001300-002100---100000-  
1001110010010030120110100101002100111000111001000100000000110000000000011011?0  
-001100000001111200111000000000200000000111  
*Distirogaster\_n\_sp1*  
020132200021110101000110000011001101103012000012110000011001100011000100000000  
000010000000000000000100-000211110001111211010200000110000100000111  
*Distirogaster\_n\_sp2*  
020131200021110000000110000011000011103012000012110000011001100011000100011100  
0000010000000000000000100-000211110001111211010200001010000100000111  
*Distirogaster\_tarsalis*  
011132200021110000000110000011000011103012000011110000011001100011100100011100  
000010000000000000000100-000211110001111211010400001010000100000111  
*Ectrichodia\_crux* 0201310-002120---  
101001110010100002110101211101000100011100110001100110001110001110100111101100  
0001110-0021000000011112000????????????????????00111  
*Ectrichodia\_lucida* 0221310-002120---  
1010011100101000021101012111010001000111001100011000100010000011100001110?????0  
001110-0?21000000011112000????????????????????00111  
*Ectrychotes\_serdangensis* 0121210-001110---  
100001010011100002110301111101002010011101210001100010001111000002100000100010  
0000100-000100000001111200011020001000000100000111  
*Ectrychotes\_sp1* 0121210-002110---  
100001010011100001110301111101002010011101210001100010001111000001100000100010  
0000100-001100000001111200011030001000000100000111

Ectrychotes\_sp2 0121210-002110---  
 000001010011100002110301111101002010011101210001100010001111000002100000100000  
 0000100-000100000001111201011000001000000100000111  
 Emesaya\_incisa 1201110-001000---100000-11--0101003110010121101000011-  
 20---0---10112101-1000011--000000000000001100100---  
 00000000100000001????????????????0000-  
 Eratyrus\_mucronatus 02212110002100---100100-  
 001011001?0010000010002211001-010000---00112111-00000?????00001100000000001000-  
 000000000010000000111400111001--000010000-  
 Gibbosella\_quadocris  
 000121100011010001000010100111000011103012000011010100011101111011?00100010000  
 000000000000000000000000-101211110001111001101010001000000100000111  
 Glymmatophora\_crassipes 2121310-002110---100001101--  
 0100003110101111101000111-11000110011110010001111001010100111101101000-----  
 01000000011112000103000000000011000000111  
 Kiskeyana\_palassaina ?031210---110????110000-  
 1????1????1000111?0?110?????????0---101?2101-  
 10?????????0?0000?00000110????????1000000100000001?????????????????0000-  
 Maraenaspis\_coccinea ?151210-  
 00211????10000??0????1????1103012?1?0101?????????1??1111011000111100000100000  
 10000000??????10000000011112000?????????????????00111  
 Marojejycoris\_brevifrons  
 000111100011111000000010100011001101101011000010010100011101100011000101010000  
 00000000000000000000100000-010000000001111201101??000000000100000111  
 Mendis\_apicimaculata 01013010012100---  
 101011000011100002110201110101001010111101110001110010101110000002100000100000  
 0001110-00210000001111101011000000100000101000111  
 Micrauchenus\_lineola 0121110-001100---000000-  
 1011110000210001000010?111010000-100---10112101-  
 0000000000000000000000001111001000000001100000001110?0000000000000001110  
 Microstemmatoides\_atrocyanea0111310-002120---  
 10100110001110000211010001110110101001100111000111001000100000001000000000000  
 0000110-000100000001111001111001000000000100000111  
 Neolibavius\_n\_sp 0001100-101100---100000-  
 1001110000211010110010111101011110121000111001000100010-000010-0001----0000000-  
 0?12110000111111101110000000110100000111  
 nr\_Bannania\_sp 0021100-001110---100000-  
 1001110010211010120010100101010110111000110101000100000000000000000000000100  
 -100211100001111101111410001000000101000111  
 nr\_Daraxa\_sp 0001300-002110---100000-  
 100111001021102011001010110000011011100011100100011100000011000001000000001110  
 -001000000000000002101120000100110000000111  
 Oncocephalus\_sp 0101110-002100---000010-  
 1000110011111012211102010031-10-100---10012111-0100010-  
 000000000000000110001110010000010100000021?????????????????0110-  
 Opistoplatys\_sp1 000001?002000---100100-11--  
 101011011141021?100001010000-100---000???11-0000000000000000000000001100000-  
 01000000001000000?11020000000000000011110

*Opisthoplatys\_sp2* 0100010-002000---100100-11--1010110111?1011?100001011-  
00-100---000??11-00000000000000000000001100000-  
000000000010000000????????????????????11110  
*Panstrongylus\_lignarius* 02012110002100---100100-  
001011000?0000000001022110000010000---00012111-000000011000001100000000001100-  
001000000010000000111000111001--000010000-  
*Peirates\_punctorius* 3101110-002110---000000-11--0100002100011020101110011-  
110100---11012111-0100011--10011--0000000000-0----  
00000000100000001????????????????????01110  
*Pothea\_lugens* 0121310-002000---101000-  
000111000021103001011010020101111011000011000100000000000000000000000000101100  
-002000000001000000011011000100020000000111  
*Racelda\_sp* ?101310-00211????100000-  
0????1????1103011?1?0100????????????1??0111001?0111100000100000100000000??????0  
0000000012222001????????????????????00111  
*Rhignia\_aimara* 0121310-002110---101000-  
100101000021103002001001010001111011000011100100000000000000000000000000000000001100  
-00200000000100000001110000100020000000111  
*Rhignia\_cinctiventris* 0121310-002110---101000-  
000011000021003002001011010000111011000011000100000000000020000000000000000000000000001100  
-002000000001000000001110000100021010100111  
*Rhignia\_sp* 0121310-002110---101000-  
000011000021003002001011010000111011000011000100000000000020000000000000000000000000001100  
-002000000001000000001110000100021010100111  
*Saica\_sp* 0121110-000100---100000-11--0101004100011121112000031-10-  
100---10112101-1000011--0000000000000001100010-  
0?00000000100000001????????????????????0000-  
*Santosia\_sp* 0121310-002100---100000-  
00011100001100310100101001010101101100001100010100000000000000000000000000000000000011?0  
-001000000001000000001120000000401011000111  
*Stenopoda\_sp* 0201110-102100---000010-10011101001111012111002010041-  
10-000---10012111-0100010-  
0000000000000000100011100000000101000000211100000100001000010110-  
*Stenopodessa\_sp* 0201110-102100---000010-100111010021110121111020100(3  
4)1-10-000---10012111-0100010-  
00000000000000001000111000211110100000021????????????????????0110-  
*Tanindrazanus\_marginatus*  
010111100021111100001010100011001101102002000011010000011101100011000101000000  
00112201001000002000000100000111  
*Tanindrazanus\_tenebricus*  
010111100021111100001010100011001001102002000011010000011001100011000101000000  
00112201001000002000000100000111  
*Tanindrazanus\_varicolor*  
014111100021111100001010100011000011102002000011010000011001100011000101000000  
0000000000000000000000000000-00(0 1)0000000111201001000000000000100000111  
*Toxopus\_brucei*  
010120210021111111000010100011000021102002000011010100011001100011100101010000  
0000000000000000000000000000-001100000001122001001000000000000100000111

*Toxopus\_fisheri* 0101210-  
00211100010000100001110000211020020000110100000110011000111001010100000000000  
0000000000000000-001100000001222001001020000000300000000111  
*Toxopus\_griswoldi*  
010121100021110001000010000111000021102002000011010000011001100011100101010000  
0000000000000000000000000000-00110000000122201001020000000300000000111  
*Toxopus\_toamasina*  
010110210021111111000010100011001101102002000011010000011001100011100101010000  
0000000000000000000000000000-001100000001220001001000001000000100000111  
*Toxopus\_vazimba*  
010111210021111111000010100011000021102002000011010100011001100011100101010000  
0000000000000000000000000000-00110000000111201001000000000000100000111  
*Triatoma\_recurva* 02312110002100---100100-  
001011000?00100000001002110000010000---00012111-000000011000001100000001101000-  
0020000000100000001????????????????0000-  
*Tribelocephala\_peyrierasi* 010001??002100---110000-01--  
101000011131020?0000010?0?0101?0---000??11-00000000000000000000000000001100010-  
00000000001000000101102000000031111111110  
*Tribelocephala\_sp* 010001??002100---110000-01--  
101000011131020?0000010?0?0101?0---000??11-00000000000000000000000000001100010-  
00000000001000000101102000000031111111110  
*Vilius\_macrops* 01213030012110---  
101001000001100001110000000101011010021110310001100111-  
000000000000000000000000000000100-00000000000111101111000000000501000100111  
*Synavecoris\_dimorphus* 00013010001110---100000-  
1001110000211010100010101101000110011000111001000100010-11000000000000000000110-  
1112000000011100111????????????????00111  
*Maraenaspis\_bidens* 2211310-002120---100001001--  
0100002110301211101200111-11001110001110010001111000000100000100000000-----  
0100000001222200010200000000100000000111  
*Glymmatophora\_carolae* 0121320-001120---  
100001100011100002110101101001001010011000110011110010001000001012000111001101  
0000110-001100000001111200010300000100011000000111  
*Distirogaster\_n\_sp3*  
011131200021110000000010000111000041103012000011110000011001100011000100010000  
0000000000000000000000100-000211110001111211010200001010000100000111  
*Distirogaster\_n\_sp4*  
021132200021110000000110000011001101103012000011110000011001100011000100010000  
0000100000000000000000100-000211110001111211010200001010000100000111  
*Distirogaster\_n\_sp5*  
02113220002111000000011000011100001110?012000011110000011001100011?00100011100  
0000110000000000000000100-000211110001111211010400001010000100000111  
*Distirogaster\_n\_sp6*  
02013020002111000000011000001100001110?012000012110100011001100011000100011100  
0000010000000000000000100-000211110001111211010200000010000100000111  
*Toliarus\_trichrous*  
002111100021110101000010100111000021101002000010010200011001100011000101010000  
0000000000000000000000000000-00000000000111201001000001000000100000111

Tanindrazanus\_joffrevillus 0101110-  
0021111000000101000110000211020020000100101000110011000110001010100000000000  
0000000000000000-000000000001111001001100001000000100000111

Tanindrazanus\_irwini 0141110-  
00211111000010101000110000111020000000101100000110011000110001010000000000000  
0000000000000000-00(0 1)000000001122001001000001000000100000111

Tanindrazanus\_marojejy 0121110-  
00211110000010101000110000111020000000100100000111011000110001010000000000000  
0000000000000000-000000000001222001001000000000000100000111

Tanindrazanus\_mahafaly  
020111100021111000001010100011000011102000000010010000011001100011000101010000  
0000000000000000000000000000-000000000001000001001000001000000100000111

Tanindrazanus\_harinjali 01011110002111100000101010001100001110(1  
2)00200001001000001100110001100010101000000000000000000000000000000000-  
00000000000011112010010000000000000100000111

Tanindrazanus\_hannajagoda 0221110-  
00211111000010101000110000111020000000101100000110011000110001010100000000000  
0000000000000000-000000000001220001001000001000000100000111

Tanindrazanus\_kathrynae  
01211110002111110000101010001100002110200200001101000001110110001100010100000  
0000000000000000000000000000-000000000001112001001000002000000100000111

Tanindrazanus\_notatus  
010111100021111100001010100011000011102002000010010000011101100011000101010000  
0000000000000000000000000000-000000000001111201001000000000000100000111

Tanindrazanus\_antananarivo  
01311110002111110000101010001100000110200200001101000001110110001100010100000  
0000000000000000000000000000-000000000001112001001000001000000100000111

Tanindrazanus\_nigripes  
010111100021111100001010100011000011102002000011010000011001100011000101010000  
0000000000000000000000000000-000000000001111001001000001000000100000111

Tanindrazanus\_bemarahaha  
01011110002111110000101010001100001110200100001111000001110110001100010100000  
0000000000000000000000000000-000000000001111201001000001000000100000111

Tanindrazanus\_andohahela  
010111100021111100001010100011000011102001000010010000011001100011000101010000  
0000000000000000000000000000-000000000001111201001000001000000100000111

Tanindrazanus\_brunneus 0101110-  
00211100100010101001110000311020010000100100000111011000110001010000000000000  
0000000000000000-000000000001111201001100001000000100000111

Tanindrazanus\_vohiparara  
01011110002111110000001010001100110110200200001101010001110110001100010100000  
0000000000000000000000000000-000000000001110001001000000000000100000111

Toxopus\_signoretii  
0101202?002111111100001010001100100110?0?2000011010101011001100011100101010000  
000000000000000000000000-0011000000012220010????????????????????00111

Toxopus\_parkeri  
010110200021111011000010100011000021102002000011010000011001100011100101010000  
0000000000000000000000000000-001100000001111001001000001000000100000111

*Toxopus tibialis*  
010110110021111111000010100011000011102002000011010000011001100011100101010000  
00000000000000000000000000000000-001100000001111001001000001000000100000111

*Toxopus basalis*  
010120210021111111000010100011000111102002000011010000011001100011100101010000  
00000000000000000000000000000000-001100000001122001001000000000000100000111

*Toxopus steineri*  
010111210021110111000010100011000021102012000011010100011001100011100101010000  
00000000000000000000000000000000-001100000001122001001000000000000100000111

*Toxopus namoroka*  
010121210021110011000010100011000021102002000011010100011001100011100101010000  
00000000000000000000000000000000-001100000001111201001000000000000100000111

*Toxopus ampitavananima*  
010111210021111111000010100011000011102002000010010000011001100011100101010000  
00000000000000000000000000000000-001100000001222001001000001000000100000111

*Toxopus italaviana*  
010111100021111001000010000011000021102012000011010000011001100011100101010000  
00000000000000000000000000000000-00110000000122201001020000000300000000111

*Toxopus pallidus*  
010111100011110111000010100011000011102002000011010000011001100011100101010000  
00000000000000000000000000000000-001100000001000001101000001000??????00111

*Toxopus simulans*  
010111210021110111000010100011000011102012000011010000011001100011100101010000  
00000000000000000000000000000000-001100000001000001001000001000000100000111

*Toxopus insignis* 0101210-  
002111000100001000011100002100200200001101000001100110001110010101000000000000  
00000000000000000000000000000000-001100000001122001001020000000300000000111

*Toxopus ambohitantely*  
010111100021110111000010000111000021102012000011010000011001100011100101010000  
00000000000000000000000000000000-001100000001220001001000001000000100000111

*Toxopus antsiranana*  
010111100021110001000010000111000011102002000011010000011001100011100101010000  
00000000000000000000000000000000-001100000001200001001020000000300000000111

*Toxopus toliara*  
010111100021110001000010000011000021102002000011010100011001100011100101010000  
00000000000000000000000000000000-001100000001122001001020000000300000000111

*Toxopus farafangana*  
010111100011111101000010100011000011102012000011010100011001100011100101010000  
00000000000000000000000000000000-001100000001000001001000000000000100000111

*Toxopus miandrtsara*  
010111200021111111000010000011000021102012000011010100011001100011100101010000  
00000000000000000000000000000000-001100000001000001001000001000000100000111

*Toxopus melobrunneus*  
010111210021111111000010100011000011102002000011010100011101100011100101010000  
00000000000000000000000000000000-001100000001000001001000001000000100000111

*Marojejycoris auranticorium*  
000111100011111000000010100111000021101011000010010100011101100011000101010000  
00000000000000000000000000000000-01000000000111201101120000000000100000111



Marojejycoris\_notadichroa  
 00011110001111110000010100011000021101011000010010100011101100011000101010000  
 00000000000000000000000000000000-010000000001111201101120000000000100000111  
 Marojejycoris\_francais  
 000111100011111000000010100011000021102011000010010100011101100011000101010000  
 00000000000000000000100000-010000000001122001101??000000000100000111  
 Marojejycoris\_ranomafana  
 000111100011111000000010100111000021102011000010010000011101100011000101000000  
 00000000000000000000000000-010000000001122001101??001000??????00111  
 Gibbosella\_vangoeris  
 000121100011011001000010000111000011103011000011010100011101111011000100000000  
 0000000000000000000000000000-101211110001111002101130?01000000100000111  
 Gibbosella\_planiscutum  
 000111100011111001000010100111000021103011000011010100011001100011100100010000  
 0000000000000000000000000000-101211110001111201101000000000000100000111  
 Gibbosella\_pallidalata  
 000121100011110001000010000111000021101011000010010100011101111011000100010000  
 00000000000000000000100000-101211110001111002101100000000000100000111  
 Gibbosella\_brunalvus 00011110001110---  
 100001000011100002110301100001101010001100111101100010001000000000000000000000000  
 0000000-101211110101122011101000000000000100000111  
 Gibbosella\_conisimilis  
 000121100011011001000010000111000011103002000110010100011101111011?21100000000  
 0000000000000000000000000000-101211110001222002101000001000000100000111  
 Gibbosella\_notoconica  
 000121100011010101000010000011001101103002000110010100011101111011020100000000  
 00000000000000000000000000-101211110001122002101000001000000100000111  
 Gibbosella\_pallidacorium  
 0001201000110101010000101000110000011030120000111101000111011??0111001000??00??  
 ??0000000000000000000000-101211110001120002101000001000000100000111  
 Gibbosella\_andasibe 00012110000100---  
 100001010011100003110301200001101010001110110101100010000000000000000000000000000  
 0100000-101211110001100002101000000000000100000111  
 Gibbosella\_betampona  
 000120100011010001000010100011000101103011000011010100011101111011?20100010000  
 00000000000000000000100000-101211110001122000101000000000000100000111  
 Gibbosella\_megafrons  
 000121100001011001000010100111000041103012000011010100011101111011?00100010000  
 00000000000000000000??00000-101211110001111002101000??000000100000111  
 Gibbosella\_mantella  
 000120100011010101000010000011001101103011000011010100011101111011020100010000  
 0000000000000000000000000000-101211110001122002101000001000000100000111  
 ;  
 END;

Supplementary material 3.2. Molecular character matrix in nexus format for cladistic analysis.

```
#NEXUS
BEGIN TAXA;
  TITLE Taxa;
  DIMENSIONS NTAX=56;
  TAXLABELS
    Apiomerus_lanipes Apiomerus_ochropterus Bannania_n_sp
  Brontostoma_colossus Brontostoma_sanguinosum Caecina_sp Centraspis_ducalis
  Cleptria_corallina Cryptonannus_n_sp Distirogaster_n_sp1 Distirogaster_n_sp2
  Distirogaster_tarsalis Ectrichodia_crux Ectrichodia_lucida Ectrychotes_serdangensis
  Ectrychotes_sp1 Ectrychotes_sp2 Emesaya_incisa Eratyrus_mucronatus Gibbosella_quadocris
  Glymmatophora_crassipes Kiskeyana_palassaina Maraenaspis_coccinea
  Marojejycoris_brevifrons Mendis_apicimaculata Micrauchenus_lineola
  Microstemmatoides_atrocyanea Neolibavivus_n_sp nr_Bannania_sp nr_Daraxa_sp
  Oncocephalus_sp Opisthoplatys_sp1 Opisthoplatys_sp2 Panstrongylus_lignarius
  Peirates_punctorius Pothea_lugens Racelda_sp Rhiginia_aimara Rhiginia_cinctiventris
  Rhiginia_sp Saica_sp Santosia_sp Stenopoda_sp Stenopodessa_sp Tanindrazanus_marginatus
  Tanindrazanus_tenebricus Tanindrazanus_varicolor Toxopus_brucei Toxopus_fisheri
  Toxopus_grisworldi Toxopus_toamasina Toxopus_vazimba Triatoma_recurva
  Tribelocephala_peyrierasi Tribelocephala_sp Vilius_macrops
  ;
END;

BEGIN CHARACTERS;
  TITLE Character_Matrix;
  DIMENSIONS NCHAR=3054;
  FORMAT DATATYPE = DNA GAP = - MISSING = ?;
  MATRIX
    Apiomerus_lanipes
  GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAA
  CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-
  CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--
  GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT
  TAAAAAGCTCGTAGTTGGTTCTGCGTGCCACGCTGTCGGTTCGCCGCCTGTCGGTGTA
  ACTGGCATGTCGTGGCATGTCCTGTCGGTGGTGAACAGAAT-----
  ----CCCCGGTA----C-GATGTAGGCTT----TATTGCCGAAG-T-CTGTATCG-----TGC-
  GTGATTCTGTTTGCCGA---
  TCTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG
  AACAAATTAGAGTGCTTAAAGCAGGCT-
  AAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAACAGGACCTTAGTTCTATT
  TTGTTGGTTTTAGGGATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTGCGTA
  TTGCGACGTTAGAGGTGAAATTCTTGATCGTCGCAAGACGCACTAGAGCGAAAGCA
  TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA
  GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTTCT
  TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCGGGGG
  AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG
  TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG
```

GAAGGATTGACAGATTGATAGCTCTTTCTTGATTTCAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGAAGGTGGTAAACTCCATCTAAGGCTAA  
ATATAACCACGAGACCGATAGCGAACAAAGTACCGTGAGGGAAAGTTGAAAAGAACT  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAAAAGACCTGAA--  
TATACGAAAGGGGAGATTACCGTCTTACCATAT---ATTCTGCTTTTAAAGTGC-----  
ATTGAGATGGTCATTT-GCCCTGCTCAG----TGTA--AACTGT-TAGTAG-  
TGGGTTTGGTCGCTCGCATTTAATAT-----  
GTAAGAATAATGGTGGACCGCATTTCTCCCTTAGT-AATACGTTGCGACCTGTT-  
AAAT-----AATATTCTAAG-TA----TAGTTAAAGTGCCTGTTT-AAAAGTATTT-----  
-----ACTGAAA-----GGTA---TGCTTTTCTACAGCTTTAA--CTAAATAT-----A--CT-  
GGATATTT-----AT---ATTTTAAACAGTAAAAAAATGACGGTGTGAGCCACTTAAAA-  
---TTGCTA--ATTTATTAGTAATT----GATAGTG-----TTATTGTCC-GTTTACAA-  
GTGGGTT-----GTTTTTTAGT-A-TAATCGTA--CTTAAGTCCGTTTAT-  
ACTTTACTCCTA-CTGTTTGT-AAAC-TGTG--ACTGT--AACAGTGCTT---  
ATTTTAAACACCTGTCTGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT?????????  
?????????CTTGAAACACGGACCAAGGAG-----GCAAGTCGTTGGGTG--  
AATAAAACCCAAAGGCAAAATGAAAGTAAAGAGTTGAGCATAATTTA-----  
AAAAAAA-----ATTTTTTTTTTA-----  
TGCTCCTCTTAAGGGAAGATAG--AAATTG-CTTTAATAATAG-----  
TTGTACTT-----TATAAC---TA-----TT-CGTTAATGTAAAT-  
TTCTTGCACTCCCAGGGCGCTTG-TTTAACGC-  
TTGCGTGGCAATGCGCACCTAGAGCGTACACGCTGATACCCGAAAGATGGTGAACATA  
TGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGA  
CGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCT  
AGTAGCTGGTTCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTTAGAT-----  
-----AAAGAATCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ATCCCCGGCTTTCTTAAAA-----TTGAAGCCGTGGGAAT--GATCAG-----  
-  
ATGCCTGGTAAGCAGAACTGGCGCTGTGG?????????CTGAAGTTTATATTCTTATTCT  
ACCAGGATTTGGCTTAATTTCCCATATCATTGCTATAGAAACAGGTAAAAATGAGGC  
ATTTGGAGCCCTAGGAATAATTTATGCTATATTAGCAATTGGATTATTAGGATTCATT  
GTATGAGCTCACCACATATTTACTGTAGGTATAGACGTAGATACACGAGCTTACTTT  
ACTTCAGCTACGATAATTATTGCTGTTTCTACTGGAATTAATCTTTAGATGACTGG  
CTACCCTGCATGGCAGAATATTAACCTTACACACCAAGAATCTTATGAGCTTTGGGAT  
TTGTATTCTTATTACTATTGGTGGGTAAACAGGTGTAGTTCTAGCCAATTCAAGTAT  
TGATATTGTTTTGCATGATACTTACTATGTAGTTGCACATTT???????

*Apiomerus\_ochropterus*

????????????AAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAATAACGATA  
CGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCACGCTGTCGGTTCGCCGCTGTCGGTGTA  
ACTGGCATGTCGTGGCATGTCCTGTCGGTGGTGAACAGAAT-----  
---CCCCGGTA---C-GATGTAGGCTT----TATTGCCGAAG-T-CTGTATCG-----TGC-  
GTGATTCTGTTTGCCGA---  
TCTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG

AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAACAGGACCTTAGTTCTATT  
TTGTTGGTTTTAGGGATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCA????????????????  
??CGTGTTGCTTGATAGTGCAGCCCT  
AAGAAGGTGGTAAACTCCATCTAAGGCTAAATATAACCACGAGACCGATAGCGAAC  
AAGTACCGTGAGGGAAAGTTGAAAAGAAGCTTTGAAGAGAGAGTTCAAGAGTACGCT  
GAAACCGCCTCAGGGGTAAACGGAAAAGACCTGAA--  
TATACGAAAGGGGAGATTCACGTCTTACCATAT---ATTCTGCTTTTAAGTGC-----  
ATTGAGATGGTCATTC-GCCCTGCTCAG----TGTA--AACTGT-TAGTAG-  
TGGGTTTGGTCGCTCGCATTTAATAT-----  
GTAAGAATAATGGTGGACCGCATTCTCCCTTAGT-AAGACGTTGCGACCTGTT-  
AAAT-----AATATTCTAAG-TA----TAGTTAAAGTGCCTGTTT-AAAAGTATTT-----  
-----ACTGAAA-----GGTA---TGCTTTTTTACAGCTTTAA--CTA-ATAT-----A--CT-  
GGATATTT-----AT---ATTTTAAACAGT-AAAAAATGACGGTGTGAGCCACTTAAAA--  
--TTGCTA---ATTTATTTAGTAATT----AATAGTG-----TTATTGTCC-GTTTACAA-  
GTGGGTTG----TTTTTTTTTAGT-A-TAATCGTA--CTTAAGTCCGTTTAT-  
ACTTTACTCCTA-CTGTTTGT-AAAC-TGTG--ACTGT--AACAGTGCTT-----  
ATAACACCTGTCGGCAGCGATCCAGGTTTCCAGGCTCTTATCCGGTCCGT????????????  
????????CTTGAACACGACCAAGGAGTTTATCGTGTACGCAAGTCATTGGGTG--  
AATAAAACCCAAAGGCGAAATGAAAGTAAAGAGTTGAGCATAATTTA-----  
AAAAAAT-----TTTTTTTCTTTA-----  
TGTTCCCTCTTAAGGGAAGATAG--AAATTG-CTTTAATAATAG-TTTTAAA-----  
CTTGTTT-----TATAAC---TA-----TTTGTTAATGTTAAT-  
TTCTCGCACTCCAGGGCGCTTG-TTTAACGC-  
TTGCGTGGCAATGCGCACCTAGAGCGTACACGCTGATACCCGAAAGATGGTGAACATA  
TGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGA  
CGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCT  
AGTAGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTTAGAT-----  
-----AAAGAATCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTGC-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ATCCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGAATAGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??  
??  
??  
??  
??  
??  
?????

Bannania\_n\_sp

??  
TCGGAATGAGTACACTTTAAAT-CTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAATTTGTTGCGGT

TAAAAAGCTCGTAGTTGGTTCTGCGTTTCGCACTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGAATGTCCTGTCGGTGGTGAATAAAAT-----  
----TCCTGGTG----C-TAACCTAGGCT-TT--TATAGCTGAAG-T-CGGTGCCG-----TGG-  
TTGATTTTATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCAAGTAGGCCGACACATTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAAGATCTGCCTAAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGTCAGCCAGCGATCCGCCAATGTTTCTT  
TAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGA  
AGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGG????????????????  
??  
??  
??  
GTCAGCCCTAATTAGGTGGTAAACTCCATCTAAG  
GCTAAATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGAAAAGTTGAAAA  
GAACTTTGAAGAGAGAGTTCAATAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA-  
TATTTGAAAGAAGAGATTCACGTCTTAGCACTT---TGTTTACTTTTAAGTAC----  
CTTTGAGATGGTAATTCAGTTTTATCCAG----TGAA--TACTG--TGGTAA-  
GAGGTTTTGCCGCTCGTACTTTATT---  
AAGTAAGCTGTAGTGCTGGACCGCACTTCTCTTTTAGT-AGGACGTTGCAACCTGTC-  
AGAT-----AATACTCTAAG-TA----TGATTAAAGTGTCTCTGT-AAATGTATTAAGA-----  
-----CTTGTTTT-----TATT---ATATCTACAACAGCTTTTA--TCAATTT-----A--  
CT-GGGTATTT-----A---TTTTTTGACAGT-  
ATCAACTGATGGTGTAGAGCCGTTGATTA---CTAAGC-----  
GAGCAGCCG-----TTTTACAA-GTTAGGT-----GTATATAAGTT-A-  
ATGTCTAA--CATCTGTTAGTCTAA-CTTTATCACCTA-CTGTTTGC-AAAA-TGTG--  
CCTTGC-ATTAGTGCCT-----  
ACAACACCGGTTGGCAGCGATTGAGGTTTCTTATCCGACCCGTCTTGAAACA?  
??  
AAAGTAAAGGATAAAAAGATAAAAG-----AAGTTAA-----  
GCAATTTTCATCTTTGAT-----TCTTTATCCTAGGGAAGATGG--ATGTCT-  
TGTTTTAAAAG-----CTTTAG-----  
TTTTCTAAACATAGAT-CTCCCGCATTCCCAGGGCGCTTG--  
CACTTGCTTTGCAAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATG  
GTGAACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGC  
GATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCG  
AACCATCTAGTAGCTGGTCCCTCTGAAGTTCCCTTAGGATAGCTTGCCTCGATT--  
-----  
TATAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCTGGCTTTCTTAAAGAAAAAATTTTTTTTTTTTGAAGTCTTGGGATTAGGAT-  
-----  
CA????????????????TTTTATATTCTAATTCTACCTGGATTGGACTTATTTACATATTAT  
CGCTATAGAAACCGGGAAAAATGAAGCCTTCGGCTCATTGGGGATAATTTATGCTAT  
ACTAACAATTGGTTTATTAGGATTCATTGTTTGGAGCCCATCATATATTCACAGTAGGG  
ATAGATGTAGACACCCGAGCCTACTTTACCTCTGCAACTATAATTATTGCTGTACCTA

CTGGAATTAATAATTTTCAGTTGATTAGCTACTCTTCACGGTAGCGTATTGAACCTTTC  
CCCTAGATTATTATGAGCATTGGGATTTCGTTTCTTATTCACTATTGGAGGATTAACC  
GGAGTTATTTAGCAAACCTCAAGAATTGATATTGTACTTCATGATACATACTATGTTG  
TAGCCCATTTCC????

Brontostoma\_colossus

??CCGTAA  
TCGGAATGAGTACACTTAAAT-CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTGCGGCATGTCTGTCGGTGGTGAATGGAAT-----  
---CCCCGGTG---C-TTCTTGGGCTT-CT---TTAGCCTAAG-T-ATGTGCCG-----TAC--  
GGATGCCATTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTAATTGATCAAGAACGAAAGTTAGAGGCTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGG????????????????  
??  
????????????????????????????????????AGAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTA  
AGGCTAAATACAACTACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAA  
AAGAAGTTGAAGAGAGAGTTCAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGAAGTGA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATAT--TGTTTAGCTTTTAAATGT---  
TCAAGAGATGTTTATTC-GCCCTATTAG---TGCA--AACTG--TAATAG-  
AGGGTTTATTCGCTCGCATTTAATA---  
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGATCTGTC-  
AGTT-----AATACTCTAAG-TA-----TGGTTAAAGTGTCTGTAT-AAAGAAAATTTAGT-----  
-----TTCGGCT-----TTTT--TTTTTTTTACAGCTTTAA--TCA-TTTT----ATA--  
CT-GAGTATTT-----TA--AACTTATGACAGT-  
ATCAATTGATGGTGTGAGCCACTTTATAAC-TTTTTCT-----AAAGTG---  
-----TCTTTAGCCGGTTTACAA-GTAGGT-----TAGTAGTT-A-  
ATTTGCGGCTTTTCGGTCCGTATAA-CTAGGCCACTTA-CTGTTAGT-AAAC-TGTG--  
CCTA--GACTGTGCTT-----  
ATAACACTGATCAGCAGCGATCCAG????????????????????????????????????  
??ACCCAAAGGCATAATAAAAGTAA  
AGAGTAATTTAATTATAA-----AAGTGTTAG-----  
TACTTTTTTTTA-----TTAAATTCTTAAAGGAAGATAG--ATTAA-  
TTTTTTAATAG-AAAAAGAGTGA-----TATTTTTTATTT-----CTCTTT-----  
----TTTTTTTTTAGAAAT-ATCTTTCATTCCTAGGGCGCTTG-  
TTTTCTCGCATTGCGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAAGTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCGCGAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT--  
-----

GAAACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCCAAATAGATGAG-  
AACCTCGGCT??  
??  
TATATATTTAATCTTACCCGGATTGGTATTA  
TTTCACACTCTATCAGTATAGAGACCGGAAAAAATGAAACCTTTGGAGCACTAGGAA  
TAATCTATGCTATAATTGCCATTGGATTACTAGGATTTATTGTATGAGCACACCATAT  
ATTTACTGTAGGAATAGATGTAGATACACGAGCATATTTCACTTCAGCAACAATAAT  
TATTGCCGTACCCACAGGAATTAATAATTTTAGATGACTTGCAACCTTACACGGCAA  
AAACATCGAGTTCACGCCAAGTATATTATGAACATTAGGATTCGTTTTCTATTACC  
ATTGGAGGATTAAGTGGTATCATTCTAGCAAAATTCAGGAATTGATATTATTCTACATG  
ACACTTATTATGTAGTAGCCCATTTCC????

*Brontostoma sanguinosum*

??  
GTAA  
TCGGAATGAGTACACTTTAAAT-CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGGAAT-----  
---CCCCGGTG---C-TGCCTGGGCTT-CT---TTAGCCCAAG-T-ATGTGCCG-----TAC--  
GGACGCCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTGCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAG??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
TACGCAAG  
TCATTGGGTA--  
ATTA AAAACCCAAAGGCGCAATAAAAGTAAAGAGCAATTTAAATAAGT-----  
ATTAGTAC-----TTTTTT-----  
TTAAATTCTTAAAGGAAGACAG--ATTTT-AACTGCT-----TT-----  
----ATAAGAAAATT-----CTTTTTAAAGGAGAA-  
ATCTTGCATTCTAGGGCGCTTG-  
TTTCTCGCATTGCGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC

GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT--

-----  
AAAACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCCAAATAGATGAG-  
AACCTCGGCTTTCTTAAAA-----

TTGAAGCCGTAGGATTTGGATCAGAGTGC??  
????????????????TATATTCTAATTCTACCTGGATTTCGGAATTATTTCTCACACTATTGCAA  
TAGAAACCGGTAAAAATGAAGCATTGGAGCACTAGGTATAATTTATGCAATAATTG  
CTATTGGATTATTAGGATTCATTGTATGAGCCCATCATATATTTACAGTAGGAATAGA  
TGTAGATACACGTGCATACTTCACCTCAGCAACTATAATCATTGCTGTTCCCTACAGGA  
ATTAATACTTTCAGATGACTAGCCACATTACACGGAAGTAATTAACCTTTACCCCA  
AGTATAATGTGAACTTTAGGGTTTGTATTCTTATTACCATTGGAGGATTAACCTGGAG  
TCATTTTAGCAAATTCAGTATTGACATTATTTACATGATACCTACTATGTAGTAGC  
TCACTTCC?????

Caecina\_sp

??  
??GGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCGGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGCCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAAT-----  
---TCCTGGTA---C-TGCGTAGGCTT-TA--AATAGCTGAAG-T-TTGTGCCG-----TGG--  
TAATTTCAATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACATTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATCTCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCC????????  
??  
??  
??  
??  
??  
AAATACAACCACGAGTCCGATAGCGAA  
CAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACG-  
TGAAACCG-CTCAGGGGTAAACGGAGAAGATCTGAA--  
TGTTGAAAGAGAGATTACGTTCTTACCAGTT--GCTCTACTTTGAAATAC----  
TTTTGAGATGGTAATTC-GCCCTATCCAG----TGTA--TACTG--TGTTGG-  
TGGGTTTTGCGCTCGTATTTCAAT----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTATC-  
AGAC-----AATACTCTAAGTTT-----TAATTAAGCGCCTGTGT-ATTGGTTGTATATTT--  
-----ATTTA-----TATA-AT---CTTATACAGCTTTAA--TAAATTTT-----A--  
CT-GGGTATTT-----A---TTTTCTGATAGT-  
AACGACTGATGGTGTAGAGCCACTTTACT-----ATATAATG----  
TAAAAGTGTG-----CGATTTTTTGGGC-GTTTACAA-GTGAGTT----  
TAGGCTTTTTAGTT-A-ATGTCGAA--CTTTGTTTCGTCTAA-CTATGCCGCTTA-  
CTGTTTGT-AAAC-TGCG--CCTTG--GACAGTGCTT-----

ACAACACCGGTCGGCAGCGATTGAGGTT??  
??





TTAAAGTGTT-----CAATTTATTTGCC-GTTTACAA-GTGGGTT-----TAGGTATTAGTT-  
A-ATGTGCGAA--CTTTTGTTCGTCTAA-CTAAGCCGCCTA-CTGTTTGT-AAAC-TGTGC-  
CCTTG--AACAGTGCTT-----  
ACAACACCGGTCGGCAGCGATC??  
??  
AAAGATAAAAG-----ATAAATTTT-----CTAATTTTCATCTTTGATTC---  
-----TCTCTCTCTCAAGGGAAGATGG--AAATCT-TATTTTGAATAA-TATTTAG-----  
-----TATTT-----ATATTAAG-TAA-----GTA-TTTAAAATATAGA-  
TTCCCGCATTCCAGGGCGCTTGTTCCTCGCTTTCGCGAGTCTATGCGCACCTAGAGC  
GTACACGATGATACCCGAAAGATGGTGAACCTATGCCTGGCCAGGACGAAGCCAGGG  
GAAACCCTGGTGGAGGTCGCGCAGCGATTCTGACGTGCAAATCGATCGTCAGAGCTGG  
GTATAGGGGCGAAAGACCAATCGAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCC  
CTCAGGATAGCTTGCCTCGGTT-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTC??  
??  
TATTGCAATAGAAACAGGTA AAAATGAAGCCTTTGGGTCATTAGGTATAATTTATGC  
TATATTAACAATTGGATTATTAGGATTTATTGTATGAGCTCATCATATATTCACAGTA  
GGAATAGATGTTGACACTCGAGCTTACTTTACCTCAGCCACTATAATTATTGCTGTTC  
CTACAGGAATTA AAAATTTTATGTTGATTAGCTACTTTACATGGAAGAATTTTATTATT  
CACCCCTAGACTTTTATGAGCACTGGGATTCGTATTTCTATTCACTATTGGGGGATTA  
ACAGGAGTAATTCTAGCAAATCAAGAATTGATATTATTCTACATGA????????????  
????????????????

Cleptria\_corallina

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGAGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCTGTCGGTGGTGAATGAAAT-----  
---CCCCGGTG---C-TATGTAGGCTT-TT--ATTAGCTGAAG-T-ACGTGCCG-----TTG--  
TGACTTCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTTCAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGA  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--

TATCCGAAAGGGGAGATTCACGTCTTACCACTT---GCTCGACTTTGAAGTGC----  
TTTTGAGATGGTAATTT-GCCCTATCCAG----TTTT--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTATA----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA----TGGTTAAAGTGCCTGTCT-TTAGGTTATAAAATT---  
-----TATTT-----TATA-AT---CTATTTACAGCTTTAA--TCAATTTT-----A--CT-  
GGGTATTT-----A---TTTTCTGACAGT-AACGATTGATGGTGTAGAGCCACTTTT-----  
-----ATAAAGTGTC-----CTTTTTTTTTTGCC-GTTTACAACGTGGGTT-----  
-TAGGTATTAGTT-A-ATGTCGAA--CTTTGTTCGTCTAA-CTAACCCGCCTA-  
CTGTTTGT-AAAC-TGTG--CCTCG--GACAGTGCTT-----  
ACAACACCGGTGCGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT????????????  
????????CTTGAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTG-  
ATACAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGCAAGATGGAGG-----  
TTGA-----TTTTTCAATCTAGAT-----  
TCTTGCTCTCAAGGGAAGATGG--ATATCT-TGTTTTAAATTA-----  
TATTT-----ATATTAGG-TA-----TA-TTTAAATTTAGA-  
TTCCCGCATTCCAGGGCGCTTG-  
TTTCTCGCTTTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT

-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCAACCTATTCTCAAACCTCCAAATAGATGAG-  
ACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGCGCTGTGG??  
??  
??  
??  
??  
??  
????  
???

Cryptonannus\_n\_sp

??CGGGACTCTATTGAGG-  
CCC GTAATCGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-  
TTGGAGGGCAAGTCT--  
GGT GCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGGAAT-----  
---CCCCAGGA---C-TGCTTAGGCTT-TT---ATAGCTTATG-T-ATGTGCTG-----GAA--  
GGATTCCATTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGCTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA

ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGA????????????????  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
????CAAGTCATTGGGTA--

ATTAACCAAGCGTAATAAAAGTAAAGAGTAATTTAAATAAGT-----  
AGTAATA-----CTTT-----  
TTAAATTCTTAAAGGAAGATAG--ATTTT-ATAGTTT-----CTTTG---  
-----TTGAAATTAAT-ATCTAGCATTCTAGGGCGCTTG-  
TTTCTCGCATTGCGAGTATATGCGCACCTAAAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCGCTCATT--  
-----

GTAAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCTCGGCTTTCTTAAAA-----  
TTGAAGCCGTAGGACTCGGATCAGAGTGCAA-----  
AG?????????GAAGTATATATTTAATCTTACCGGATTCGGAATTATTTACATGTA  
ATTGCAAGAGAAACAGGTAAAAATGAGGCCCTTCGGGGCCTTAGGAATGATTTATGCC  
ATACTAGCAATTGGACTCTTGGGATTCATTGTATGAGCCATCACATATTCACAGTTG  
GAATAGACGTAGATACACGTGCATATTTTACATCAGCAACTATAATTATTGCCGTAC  
CTACAGGAATCAAATCTTCAGATGGTTAGCTACCCTTACGGTAGTAATATTTTATT  
TACTCCGAGAATTTTGTGAGCCCTGGGGTTTCGTATTCCTGTTACAATCGGAGGTCTA  
ACCGGAATTATTTAGCTAACTCAAGTATTGACATTACCCTGCATGATACCTATTATG  
TAGTTGCCCACTTCCAT???

Distirogaster\_n\_sp1

??  
TCGGAATGAGTACACTTTAAAT-CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCCGCCCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCTGTCGGTGGTGAATGGAAT-----  
---CCCTGGTG---C-TACGTAGGCTT-TT--ATTAGCTGAAG-T-TTGTACCG-----TGG--  
TGACTTCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA

TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAAT????????????????????????  
??  
??  
??  
CTAAGGCTAAATACAACC  
ACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAG  
AGAGTTCAAGAGTACG-TGAAACCG-CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTACCACTT---GCTCGACTTTAAAGTGC---  
TCTTGAGATGGTAATTC-GCCCTATCCAG---TGCA--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTATT---  
AGTAAGGCGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA----TAATTAAGTGCCTGTCT-AAAGGTTATAAATTT---  
-----TTGTT-----TATT-AC---CTTTTTACAGCTTTAA--TTAATTTT-----A--CT-  
GGGTATTT-----AT---TTTTCTGACAGT-AACGACTGATGGTGTAGAGCCACTTAAAC---  
-ATT-----ATATAATG---TAAAAGTGT-----TCATTTGCC-GTTTACAA-  
GTGGGTTA--GGTATTTATTAGTT-A-ATGTCGAA--CTTTTGTTCTGTCTAA-  
CTAAGCCGCCTA-CTGTTTGT-AAAC-TGTG--CCTTG--AACAGTGCTT-----  
ACAACACCGGTGCGCAGCGA????????????????????????????????????  
??  
ACGCAAGTCATTGGGTG-  
ATACAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGAAAAGATAGAAG-----  
AAGAAA-----TTCATTTTCATCTTTGAT-----  
TCTCTCTCTCAAGGGAAGATGG--ATATCA-TATTTTAAATAT-TATTTGG-----  
TATTT-----ATATTAAG-TA-----TATTTTAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTCTGCGAGTTTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTGC-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTAGGATCAGAGTGCC????????????????????????  
?????????AAGTATACATTTTGTATTTACCCGGATTTGGTTTAAATTTCCCACATTATTGCT  
ATAGAAACAGGAAAAAATGAAGTTTTTGGGACATTAGGTATAATTTATGCCATACTA  
ACTATCGGTTTATTAGGATTCATTGTTTGTGAGCACATCATATTTTACAGTAGGTATAG  
ATGTAGATACACGAGCCTACTTTACATCAGCAACAATAATCATCGCTGTACCTACAG  
GTATTAATAATTTTCAGATGATTAGCCACCCTACATGGAAGAATTTTAAATTTTACC  
CAGCTTGTTATGAGCACTAGGATTCGTTTTCTATTTACAATTGGGGGGTTAACGGGA  
GTAGTTTTAGCAAATTCAAGAATTGATATTATCTTACATGATACATATTATGTAGTAG  
CACA?????????

Distirogaster\_n\_sp2

??  
?????????AAGTATACATTTTGTATTTACCCGGATTTGGTTTAAATTTCCCACATTATTGCT  
??TGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTGCGGTGT  
ACTGGCATGTCGCGGCATGTCTGTGCGGTGGTGAATGGAAT-----

---CCCCGGTG---C-TACGTAGGCTT-TT--ATTAGCTGAAG-T-TTGTACTG-----TGG--  
TGAATTCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAA??  
??  
??  
??  
GCGAA  
CAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACG-  
TGAAACCG-CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTACCACTT---GCTCGACTTTAAAGTGC----  
TCTTGAGATGGTAATTC-GTCCTATCCAG---TGCA--TACTG--TGATAG-  
AGGGTTTTATCGCTCGCATTTTATT----  
AGTAAGGCGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----TATACTCTAAG-TA-----TAATTAAGTGCCTGTCT-AAAAAGGTTTTATTTA-  
-----TTTAT-----TACC-TT---TTTTTTACAGCTTTAA--TTAATTTT-----A--CT-  
GGGTATTT-----AT---TTTTCTGACAGT-AACGACTGATGGTGTAGAGCCACTTAAAA---  
-----CATTACATAATG---TAAAAGTGT-----TTACTTGCC-GTTTACAA-  
GTGGGTTAG-GTATTTTTATTAGTT-A-ATGTCGAA--CTTTTGTTTCGTCTAA-  
CTAATCCGCCTA-CTGTTTGT-AAAC-TGTG--  
CC??  
??  
??  
GCGGAATGAAA  
GTAAAGAGTAGAAAGATAAAAG-----AAGGAAA-----  
TTAATTTTCATCTTTGAT-----TCTCTCTCTCAAGGGAAGATGG--ATATCT-  
TATTTTAAATAA-TATTTAG-----TATTT-----ATATTAAG-TA-----  
TA-TTTAAAATTTAGA-TTCCCGCATTCCAGGGCGCTTG-  
TTTCTCGCTCTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGTT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTCT??  
??  
GAGGTATACATTTCTTATTCTACCAGGGTTTGGAT  
TAATCTCACATATCATTGCAATAGAAACCGGAAAAAATGAAGTTTTTGGGTCATTAG  
GGATAATTTACGCAATACTCACTATTGGATTACTAGGCTTTATTGTTTGGAGCCCATCA  
CATATTCACAGTAGGGATAGATGTTGATACCCGAGCCTATTTTACATCAGCAACAAT  
AATCATTGCTGTACCCACAGGTATTAATAATCTTTAGATGATTAGCTACTCTGCATGGT  
AGCATTTTAACCTTCTCCCCAGATTATTATGAGCTTTAGGATTTGTTTTCTTATTAC  
AATCGGAGGACTAACAGGGGTAGTTTTAGCAAACCTCAAGAATTGATATTATTTACA  
TGATACATACTATGTAGTAGCCATT?????????

Distirogaster tarsalis

??CCGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCCGCCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGGAAT-----  
----CCCCGGTG----C-TACGTAGGCTT-TT--ATTAGCTGAAG-T-TTGTACCG-----TGG--  
TGAATTCATTTGCCGA---  
TTTCTCCTACTCCGGTGTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
AACAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCCGGGGGCATTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCGAAGGCGATCA  
GATACCGCCCTAGTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCTT  
TTAATGACTCGGCGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTCCCGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACG????????????????????????  
??  
??  
??  
??  
??  
??  
TATGACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGCTT  
TGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTACCCTT---GCTCGACTTTAAAGTGC----  
TCTTGAGATGGTAATTC-GCCCTATCCAG----TGCA--TACTG--TGATAG-  
AGGGTTTTATCGCTCGCATTTTATT----  
AGTAAGGCGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA----TAATTAAGTGCCTGTCT-AAAAAGGTTTTATTA---  
-----ATTAA-----TTAC-CT---TTTTTACAGCTTTAA--TTAATTTT-----A--CT-  
GGGTATT-----AT---TTTTCTGACAGT-AACGACTGATGGTGTAGAGCCACTTTAAA---  
-ACAT-----TACATAATG---TAAAGTGT-----TCATTTGCC-GTTTACAA-  
GTGGGTTAG-GTATTTTATTAGTT-A-ATGTCGAA--CCTTTGTTTCGTCTAA-  
CTAATCCGCCTA-CTGTTTGT-AAAC-TGTG--CCTTG--AACAGTGCTT-----  
ACAACACCGGTGCGGACGATCCAGGTTTCCAGGCTCTTATCC????????????????  
??  
ACCCAAAGGCGTAATGAA  
AGTAAAGAGTAGAAAGATAGAAG-----AAGGAA-----  
YYHHHH?TCATCTTTGAT-----TCTCTCTCAAGGGAAGATGG--ATA??-  
??TTTTAAATAA-TATTTAG-----TATTT-----ATATTAAG-TA-----  
TA-TTTAAAATTTAGA-TTCCCGCATTCCAGGGCGCTTG-  
TTTCTCGCTTTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCGCTCGGT  
-----  
GTAAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCH?????TTCTCAAACCTCAAATAGATGAG-ACCCCCGGCTTTCTTAAA-  
-----  
TGAAGCCG??  
GTATATATTTTAAATTCTGCCTGGATTTGGTTTTAATTTACATATATTGCTATAGAAAC





CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT

-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTCT??  
??  
AGTATATATTCTTATTTTACCAGGATTCGGCTT  
AATCTCCCATATTATTGCTATAGAGACAGGGAAAAATGAAGCATTGGATCACTTGG  
ATAAATTTATGCTATATTAACCTATTGGATTATTAGGATTCATCGTTGAGCTCATCAT  
ATATTTACAGTAGGAATAGACGTTGATACACGAGCTTATTTTACATCAGCTACTATA  
ATTATTGCTGTACCTACAGGAATTAATAATTTTTCAGATGATTAGCAACCTTACATGGA  
AGTATTTTAAGATTTTCACCAAGATTACTTTGAGCATTAGGCTTCGTGTTCCCTTTTAC  
AATCGGTGGATTAACCGGAGTAATTTGGCTAATCAAGAATTGATATTATTTTACAT  
GACTTATTATGTTGTTGCACACTTCCA???

*Ectrichodia lucida*

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGAGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAAT-----  
---CCCCGGTG---C-TACGTAGGCTT-TT--ATTAGCTGAAG-T-ATGTGCCG-----TGG--  
TGATTTCAATTTACCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACATTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTTCTTGATTCAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGT  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTACGTCTTACCACTT---GTTCAACTTTAAAGTGC----  
TTTTGAGATGGTAATTC-GCCCTATCCAG----TGTA--TACTG--TGATAG-  
AGGGTTTTATTGCTCGCATTTTATTT--  
AAGTAAGGTACAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA-----TGGTTAAAGTGCCTATTT-AAATGTTATAAATTT---  
-----TATT-----TATG-AC---ATTTTTATAGCTTTAA--TCAATTCT-----A--CT-  
GGGTTTTT-----TTTTCTGACAGT-AACGACTGATGGTGTAGAGCCACTTATTA-----  
-----CATTTTATAATGT--TTAAAAGTGTG-----TAAATTGCC-GTTTACAA-  
GTGGGTT-----AGGTATTAGTTAA-ATGTGCGAA--CTTATGTTTCGTCTAA-  
CTAAGCCGCTAGCTGTTTGT-AAAC-TGTG--CCTTA--GACAGTGCTT-----

ACAACACCGGTCAGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT????????????  
????????CTTGAAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTG-  
ATACAAAACCCAAAGGCGTAATGAAAGTAAAGAGTGAGTAGAGAAAGA--  
TTTTAAAAGGATAAGAATT-----TTTATTTTAACTTTGATTT-----  
CTTTATCTCTCTCAAGGGAAGATGG--AAATCT-TATTTTAGATTAATATTTAG-----  
----TATATTTTTTT-----ATATTAAG-TA-----AA-  
TTTAAAATGTAGATTTCTCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTTTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCTGATCAGAGTGCCAAGTGGGCAATTTTGGTAAGCAGAAC  
TGGCGCTGTGG??  
??  
??  
??  
??  
??  
????

*Ectrychotes\_serdangensis*

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGAGAGGTAGTGACAAAAAATAA  
CGATACGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCCAGTCT--  
GGTGCCAGCAGCCGCGGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCTGTCGGTGGTGAATGAAGT-----  
---CCCCGGTA---C-TGCGTAGGCTT-TT--ATTAGCTGAAG-TTGTTTATCG-----TGG--  
TGACTTCATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTGAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACT  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATTCGAAAGAGGAGATTACGTTTACCCTT---GCTCGACTTTAAAGTGC----  
TCTTGAGATGGTAATTC-GCCCTATTAG----TATT--TACTG--TGGTAG-

AGGGTTTTACCGCTCGCATTTTATA----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAA-TA----TGATTAAAGTGACTGTCT-AGAGGTTATAGAATT--  
-----TATTT-----TATA-AT---CTTTTTACAGCTTTAA--TTAATTTT-----A--  
TT-GAGTATTT-----A---TTTTCTGACAGT-AACGACTGATGGTGTAGAGCCACTTTA----  
-----ATT---TTAAAGTGTT-----CATTTGTATTGCC-GTTTACAA-  
GTGGGTT-----TTGATATTAGTT-A-ATGTTGAA--  
CTTTTGTTCGTCTAACCTAAGTCGCCTA-CTGTTTGT-AAAC-TGTG--CCTTG--  
AACAGTGCTT-----  
ACAACACCGGTCAGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT????????????  
????????CTTGAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTG-  
ATACAAAACCTAAAGGCGAAATGAAAGTAAAGAGTAGTAAGATAGAGG-----  
AAGAAAT-----TTATTTTTCATCCTTGAT-----  
TCTTGCTCTCAAGGAAGATGG--TTATCT-TATTTA-----  
-----AATTAT-AT-----TA-TTTAAGATGTAGA-TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTTTGCGAGTTTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACATATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCTCTCAGGATAGCTTGCCTCGATT  
-----

GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCCAAATAGATGAG-  
ACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGCGCTGTGG??  
??  
??  
??  
??  
??  
??  
???

Ectrychotes\_sp1

??  
CCGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCCAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATAAAGT-----  
---CCCCGTA---C-TGCGTAGGCTA-TT--ATTAGCTGAAGTT-GTTTATCG-----TGG--  
TGACTTTATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGG????????????????  
??

??AAACTCCATCTAAGGCTAAAT  
ACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGCTTT  
GAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGAAGCTGAA--  
TATTCGAAAGAGGAGATTACGCTTACCATTT---GCTCGACTTTGAAGTGC----  
TTTTGAGATGGTAATTC-GCCCTATTCAG----TATT--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTATA----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----ATACTCTAAA-TA----TGATTAAGTGACTGTCT-AGAGGTTATAGAATT--  
-----TATTT-----TATA-AT---CTTTTACAGCTTTAA--TTAATTTT-----A--  
TT-GAGTATTT-----A---TTTTCTGACAGT-AACGACTGATGGTGTAGAGCCACTTTA----  
-----ATT---TTAAAGTGTT-----CATTTGTATTGCC-GTTTACAA-  
GTGGGTT-----TAGATATTAGTT-A-ATGTTGAA--CTTTGTTCGTCTAA-  
CTAAGTCGCTA-CTGTTTGT-AAAC-TGTG--CCTTG--AACAGTGCTT-----  
ACAACACCGGTCAGCAGCGATCCAGGTTTCAGTCTC????????????????????????????????  
??  
ATACAAAACCTAAAGGCGAAATGAAAGTAAAGAGTAGTAAGATAGAGG-----  
AAGAAAT-----TTATTTTTCATCCTTGAT-----  
TCTTACTCTCAAGGGAAGATGG--TTATCT-TATTTTAAATTA-----  
TATT-----TAATTAT-AT-----TA-TTTAAGATGTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTTTGCGAGTTTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCTCTGAAGTTTCTCTCAGGATAGCTTGCGCTCGATT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAA-----

Ectrychotes\_sp2

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGAGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGGCCAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCCTGTCGGTGGTGAATGAAGT-----  
---CCCCGGTA---C-TGCGTAGGCTT-TT--ATTAGCTGAAG-TTGTTTATCG-----TGG--  
TGACTTCATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG

AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTGAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGA  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATTGAAAGAGGAGATTCACGTCTTACCCTT---GCTCGACTTTAAATTGC----  
TTTTGAGATGGTAATTC-GCCCTATTAG----TATT--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTATA----  
TGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAA-TA----TGATTAAGTGACTGTCT-AGAGGTTATAGAATT--  
-----TATTT-----TATA-AT---CTTTTACAGCTTTAA--TTAATTTT-----A--  
TT-GAGTATTT-----A---TTTTCTGACAGT-AACGACTGATGGTGTAGAGCCACTTTA----  
-----ATT---TTAAAGTGTT-----CATTTGTATTGCC-GTTTACAA-  
GTGGGTT-----TTGATATTGGTT-A-ATGTTGAA--CTTTGTTCGTCTAA-  
CTAAGTCGCCTA-CTGTTTGT-AAAC-TGTG--CCTTG--AACAGTGCTT-----  
ACAACACCGGTCAGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT?????????????  
????????CTTGAAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTG-  
ATACAAAACCTAAAGGCGAAATGAAAGTAAAGAGTAGTAAGATAGAGG-----  
AAGAAAT-----TTATTTTTCATCCTTGAT-----  
TCTTGCTCTCAAGGAAGATGG--TTATCT-TATTTTAAATTA-----  
TATT-----TAATTAT-AT-----TA-TTTAAGATGTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTTTGCGAGTTTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCTCTCAGGATAGCTTGCCTCGATT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTCTTAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGCGCTGTGG????????????????CATTTTAATCCTTCCGGGATTTGGTTAATTTCC  
CATATTATTTCAATAGAAACAGGAAAAAGAGAAGCATTGTTGCTTTAGGTATAATT  
TATGCAATAATTACTATCGGACTTTTAGGATTTATTGTATGGGCACACCATATATTCA  
CTGTAGGTATAGATGTTGATACACGAGCATATTTTACCTCAGCTACTATAATTATTGC  
AGTTCTACAGGAATTAATAATTTTAGATGATTAGCAACTTTACATGGAAGAATATT  
AACTTTCTCTCTAGACTTTTATGAGCATTAGGCTTCGTATTCTTTTACTATTGGAG  
GATTAACAGGAGTAATTTAGCAAATTCAGGATTTGATATTGTACTACACGATACTT  
ACTATGTAGTAGCACATTT???????

Emesaya\_incisa

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCGCCGCTGTTCGGTGT  
ACTGGCATGTTCGCGGCATGTCCTGTTCGGTGGTGAACGGGAT-----  
----CCCCGGTG---C-TAACTGAGACT----TCTTGTCAAAG-TAATGTGCCG-----TGT-  
GTGATGTTCGTTTGCCGA---  
TCTCTCTACTCCGGTGCTCTTAAACGAGTGTTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAATAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCCGACG-  
TAGAGGTGAAATTCTTGGATCGTCGCAAGACGCCTAGAGCGAAAGCATTGTTGCCAAG  
TATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCGAAGGCGATCAGATACCGCC  
CTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCTATAATGACT  
CGGCGGGGAGCTTCTACTCGGGAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGT  
TGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAGTGGAGCCTG  
CGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTG  
ACAGATTGATAGCTCTTTCTTGATTGAGTGGGTAGTGGTGCATGGCCGTTCCGTGTTG  
CTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCAC  
AAGACCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAG  
AGTTCAAGAGTACG-TGAAACCG-TTCAGGGGTAAACGGAAAAGATCTGAA--  
ATTCCGAAAGGGGAGATTCACGTCTTACCATTT---ACTTATCTTTTTTATGTT---  
TTAAGAGATGGCAAGTC-GCCCTATCCAG----TGCA--AACTG--TGGTAG-  
AGGGTCTGGTTCGCTCGCATATTAAG-----  
GTTTTGGTAGTGGTGGACCGCACTTCTCCCTCAGT-AGGACGTTGCGACCTGTC-  
ATTT-----TACATTCTAAG-TGCGTTTGTGTTAAAGTGCCTGTAT-TTAGATAAT-----  
-----GTCGCAA-----GATA---TTATCTTTTACAGCTTTAA--CAATATGC-----A--  
CC-GAGTGTTA-----AT---TTTTATGACAGT-  
AACAAATGATGGTGTGAGCCACTTAAAA---TTGTTGTCATTATTGATGATAATG----  
GAGAGTGT-----CTTAGGCCG-GTTTGCAA-GTGGGCT-----GTTTTATAGTT-A-  
TATTCCGA--CTTTGTCCGTCTTG-CTATTGCTCTCA-CTGTTGGT-AGAC-TGTG--  
CCTCT--GACAGTGCTT-----  
ACAACATCAGTCAGCAGCGATCAAGGTTTCAGATTCTTATCCGGTCCGT?????????????  
??????CTTGAAACACGGACCAAGGAGTTTATCGTGTACGCGAGTCATTGGGTA--  
AAGAAAACCCAAAGGCGAAATGAAAGTAAATAGTGGGAAATTGATTT-----  
-----TATTTCTATATTATCAG-----  
TTTCCCCTATAAGGGAAGATGG--AAAATATATCAGTAATGG-TGT-----  
-----GATAATATCATA-----TTGTTGTTGATAACAAGT-  
TTCCCGCACTCCCGGGGCGCTTG-  
TTTCTCGCTTTGCGAGTCACTGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACATATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-

AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAACCCCCGGCTTTCTTAA  
AA-----  
ATGAAGCCCGTGGGATTCGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGCGCTGTGG????????????TACATTCTAATTTACCAGGATTCGGATTAATTTC  
CCACATCATCTCTATAGAAAGAGGTAAAAAGGAAGCCTTCGGGTCATTAGGTATAAT  
TTACGCAATATTATCAATTGGAATTCTAGGATTCATTGTATGAGCACATCATATATTC  
ACCGTAGGTATAGATGTAGATACACGAGCATACTTTACATCAGCAACAATAATTATT  
GCTGTTCTACAGGAATTAAGGTCTTCAGTTGATTAGCAACAATAACAGGATCAGTT  
ATAACATTCTCTCCAGTATACTTTGAGCACTAGGGTTCGTATTCTATTTACAATAG  
GAGGGTTAACAGGAGTAATTCTAGCTAACTCAAGAATTGACATTGTACTACATGATA  
CATACTATGTAGTAGCCCATTT??????

*Eratyrus mucronatus*

??AT  
CGGAATGAGTACACTTTAAAT-CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCCTGTCGGTGTA  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAACGGAGT-----  
----CCACAATG---C-TACATAGGCTT----TATTGCCGATG-T-TAGCGTCG-----TGT-  
GTACTTCCGTTGGCCGA---  
TCTCTCTACTTCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAAACAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGAGCTTCCACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGT??  
???  
???  
???  
ATACAACCACGAGACCGATA  
GCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGAG  
TACG-TGAAACCG-CTCAGGGGTAACGGAAAAGACCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCAGTT---GCCCACTTTTAAATTA-----  
TTTGAGATGGCCTCTC-GCCCTATTCAG----TGTAACAGCTG--TGGTAG-  
TGGGTTTAGTCGCTCGCATTAAATT-----  
GGAAGGGTGTGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACCTGTC-  
AAAT-----AATATCCTAAG-TA---TTTGGTTAATTTGTCTGTTT-TTAGTTAAT-----  
-----GCTGAAA-----GGTA--TTACCTTTAACAGTTTATAG--CCGCTTAT---ATA--  
CT-GGAT-----ATAATTGACAGT-AATGAATCATGGTGTGAGCCACTTGAAA---  
-TTATGT--ATTTATATATAATG-----GAAAGTGTC-----CTTGTGGCT-  
GTTTGCAA-GTGGGT-----GGTAATTAGTT-T-TAGTCGGA--TTTTGGTCCGTCTAG-  
CTATTCCGCCTA-CTGTTGGT-AAAC-TGTT--CCTTG--GACAGTGCTT-----  
ATAACACCGGTCGGCAGCGATTCAGTTTCAGGCTCTTATCCGACCCGTCTTGAAACA  
CG????????????????CACGGACCAAGGAGTTAGCGGTACGCAAGTCATTGGGTG--  
TATAAACTCAAAGGCGAAATGAAAGTAAAGAGTAGAGATTTGTTAA-----  
GAAG-----TTACTTTTTTACA-----  
ATCTCCTCTTTAGGGAAGATGG--AAATAT-CATACAGC-----  
-----ATTGTATGAAGAT-TTCCCGCATTCCAGGGCGCTTG-





GTAGATACACGAGCTTATTTTACATCAGCAACTATAATTATTGCTGTACCCACAGGA  
ATTAAAATTTTTCAGTTGACTTGCCACCCTTCAAGGAAGAGTACTAAACCTTACCCCA  
GAACACTTTGAGCCCTAGGATTTGTATTTCTATTCACAATTGGAGGATTAACAGGAG  
TTATCCTAGCAAACCTCAAGAATCGATATCGTACTCCATGATACTTATTATGTAGTAGC  
ACATTTTCATT??

Glymmatophora\_crassipes

??CCGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAAT-----  
---TCCCGGTG---C-TACGTAGGCTT-TT--ATTAGCTGAAG-T-ATGTGCCG-----TGG--  
TGATTTCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAAATGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCCGCCAGCGATCCGCCGATGTTCTT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGGTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACT??  
??  
??  
??GC  
GAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGAGTA  
CG-TGAAACCG-CTCAGGGGTAAACGGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTACCACTT--GCTCGACTTTAAAGTGC---  
TTTTGAGATGGTAATTC-GCCCTATCCAG---TGTA--TACTG--TGGTAG-  
AGGGTTTTATTGCTCGCATTTTATT--  
AAGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA----TGGTTAAAGTGTCTATCT-AAAGGTTATAAATTTA-  
-----TCTAAGTT-----TATT-AC--CTTTTATAGCTTTAA--TCAATTAT---  
ACA--CT-GGGTGTTT-----GT---TTTTCTGACAGT-  
AACGATTGATGGTGTAGAGCCACTTGAAC----ATTATGA-TAATTTTATTATAATG----  
TAAAAGTGTC-----CATGTGTTTCC-GTTTACAA-GTGGGT-----TAGGTATTAGTT-  
A-ATGTGAA--CTTTTGTCTGCTAA-CTAAGCCGCTA-CTGTTTGT-AAAC-  
TGTGC??  
??  
GTAAT  
GAAAGTAAAGAGAGTGTTAGAGGAGGGGAAAGAAAAGGATAAAAGTTTT-----  
-----TATACTTTAACTTTGATTTCTCTCTCTCTCTCTCTCTCTCAAGGGAAGATGG--  
AAATCT-TATTTAAGTGA-TGT-----TA-----AA-  
TTTAAATGTTGA-TTCCCGCATTCCCAGGGCGCTTG-  
TTTTCTCGCTTTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATAACCGAAAGAT  
GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGAAACCCTGGTGGAGGTCCGACG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCGCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-

AAACGACCTCAACCTATTCTCAAACCTCCAAATAGATGAG-  
ACCCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCTGATCAGAGTGCCAAGTGGGCCAATTTTGGTAA?????????  
????????????????????TTATATTTTAATTTTACCAGGATTTGGTTTAATTTACATATT  
ATCGCAATAGAAACAGGAAAAACGAGGCATTTGGATCACTAGGAATAATTTACGC  
TATACTAACTATTGGTCTCTTAGGGTTTATTGTATGAGCACACCATATATTTACAGTA  
GGAATAGATGTAGATACTCGAGCTTACTTTACATCAGCCACTATAATTATTGCTGTAC  
CTACTGGAATTAATAATCTTTAGATGATTAGCCACTTTACATGGAAGAATTTAATATT  
CACACCTAGACTATTATGAGCATTAGGATTTGTATTTTATTACAATTGGAGGATTA  
ACTGGAATTATTCTTGCTAATTCAAGAATTGATATTATTCTTCATGACACTTACTATG  
TAGTAGCCCACTTCCA???

Kiskeyana\_palassaina

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCACGCTGTCGGTTCGCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAACGGGGT-----  
----CCCCGGTG---C-AACTTTGGCTT-TTCATAAAGTCAGAG-C-CTGTGCCG-----  
TGT-GTGTCCCCGTTTGCCGA---  
TCTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAATAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTGCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTCATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTGAGTGGGTAGTGGTGCATGGCCG  
TTCCGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATACAACCACGAGACCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACT  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAAAAGACCTGAA--  
TTTCCGAAAGGGGAGATTCACGTCTTACCGCTC---GCTCGATTTTTTTATGT----  
TTAAGAGATGGTCACTC-GCCCTATCTAG---TGCA--AACTG--TGATAG-  
AGGGTGTGGTTCGCTCGCATTTTTGA-----  
ATTAGAGCAGTGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACCTGTC-  
ATAT-----AATAATCTAAG-TA---TTGGTTAAAGTGCCCTGTAT-AAAGGTTGA-----  
-----TCGCAA-----GGT---TAACTTTTTACAGCTTTAA--CCGTTTTT----ATA--CC-  
GATTATTT-----TTGTTTGACAGT-AACGATTGATGGTGTGCGAGCCACTTGATA----  
TTATTA-----GCAATAATAATG-----GAAAGTG-----TCTTTGTCC-GTTCGCAA-  
GTGGGTA-----AGGTAATAGAT-G-TTGTGCAATTTTATGATTCGGCTCG-  
CTATTCTCCTA-CTGTTGGT-GAAA-TGTG--TCTCA--GACAGTGCTC-----  
ACAACACCGGTCGGCAGCGATCCAGGTTTCAAGGCTTATCCGGTCCGT?????????????  
????????CTTGAACACGGACCAAGGAGTTTATCGTGTACGCGAGTCATTGGGTA--  
AAATAAACCCAAAGGCCGAAATGAAAGTAAAGAGTGGACAATTGTTGA-----

AAAAAT-----TTATTTTTCTTCC-----  
AATTTCCCTCGAGGGAAGATAG--AAAGCT-AGTGTCTTTGAT-----  
GAGTTTTCTTT-----CGGGTTAATTTA-----TCATTGGCATCATAGCT-  
TTCTCGCATTCCCGGGGCGCTTG-  
TTTCTCGCTTTGCGGAGACTATGCGCACCTAGAGCGTACACGCTGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTCTTAAAA-----  
GTGAAGCCGTGGGACTAGGATCAGAGTGCCAAGTGGGCCAATTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??  
??  
??  
??  
??  
????

*Maraenaspis\_coccinea*

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGAGAGGTAGTGACAAAAAATAA  
CGATACGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAAT-----  
---CCTCGGTG---C-TACCTAGGCTT-TT--ATTAGCTGAAG-T-ACGTGCCG-----TTG--  
TGACTTCATTTGCCGA---  
TTTTTCCTACTTCGGTGCTCTTAAACGAGTGTCGAAGTAGGCCGACACATTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATCTCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCCCGTTCTATT  
TTATTGGTTTTAGGAATACGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCCTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTTCAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACT  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTACAGTCTTACCCTT---GCTCAACTTTGAAGTGC----  
TCTTGAGATGGTAATTT-GCCCTATCCAG---TTTA--TACTG--TGATAG-  
AGGCTTTTACCGCTCGCATTTTATA----  
AGTAAGGTGCAGTGGTGGACCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTATC-  
AGAT-----AATACTCTAAG-TT-----AGATTAAAGTGCCTGTAT-ATAGGTTTTTAAATT-

-----AATTTTTT-----TCTA-AT---CTATTTACAGCTTTAA--TCAAATTTA---  
TTA--CT-GGGTATTG-----AT--CTTTACTGATAGT-  
AATGACTGATGGTGTGAGCCACTTTTTT---TCT-----TTACTGAAC----  
TATAAGTGTC-----CTATGTGGTGGCC-GTTTACAA-GTGGGTT---  
CTAGGTTATTAGTT-A-ATGTCGAA--CTTTGTTCGTCTAA-CTAAGCCGCCTA-  
CTGTTAGTTAAAC-TGTG--CCTCG--GACAGTGCTT-----  
ACAACACCGGTTCGGCAGCGATCCAGGC-----

TCTTATCCGGTCCGT??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
????????????????

Marojeycoris\_brevifrons

??  
??TGAGTTACACTTTAAATCCCTTAAACAAGGATCCATTTGGAGGGCAAGTTCCTGGTC  
GCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGTTAA  
AAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCGCCGCCTGTCGGTGTACT  
GGCATGTGCGGCGCATGTCCTGTCGGTGGTGAATGAAAT-----  
CCCCGGTG---C-TACGTAGGCTT-TT--ATTAGCTGAAG-T-TTGTGCCG-----TGG--  
TGACTTCATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGCGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCTCT  
TTAATGACTCGGC??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
????????????????

??  
??TACGCAAGTCATTGGGTG-  
ATATAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGAAGGATAAAAG-----  
AAGGAAA-----TTAATTTTCATCTTTGAT-----  
TCTTTCTCTCAAGGGAAGATGG--ATATCTTTATTTTAAAT-----  
-----TA-----TA-TTTAAAATTTAGA-TTCCCGCATTCCAGGGCGCTTG-  
GTTCTCGCTATGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGATT  
-----  
TTAAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGGCTTTCTTAAA-----  
TTGAAGCCGTGGGATTAGGATCAGAGTGCCAAGT??  
??  
??  
??  
??  
??  
??

Mendis apicimaculata

??  
????????????????????AT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCCCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTAAATAAAAT-----  
---TCTCCGCGTACT-TACCTAGGCTT-TT--ATTAGCTGAAG-T-TTGTGTTG-----  
TGG--TGACTTTATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTCGAAGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAAGATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCGGGGG  
AAGTATGGTTGCAAAGCTGA??  
??  
??  
??CCATCTAAGGCTAAATACAACCACGA  
GTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAG  
TTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGGAATATCTCGAAAGAGGAGATTACAGTCTTA  
CCATTT---GCTCTACGTTGAAGTAC----CTTTGAGATGGTAATTC-GCCCTATCCTG----  
TGCA--TACTG--TGATAG-TGGTTTTACCGCTCGTATTTTCATT----  
AGTAAGCTGCAGTGGTGGACCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACCCTAAG-TA----TAATTAAGTGCCTGTGTAAATATAATAAAAAT-  
-----TGTTT-----TATT--CAATTTATAACAGCTTTAA--TAAATTT-----A--  
CT-GGGTATTT-----A---TTTTCTGACAGT-



AAGTACCGGTAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACG-  
TGAAACCG-CTCAGGGGTAAACGGAAAAGACCTGAA--  
TATCCGAAAGGGGAGATTACGTCTTGCCATAT---ATTCTCCTGTTAAGTGC-----  
ATTGAGATGGTCATTT-GTCCCGCCAG---TGTA--AACTG--TGGTGG-  
TGGGTTTGGTCGCTCGCATTTAAAT-----  
GTAAGAATAATGGTGGACCGCATTCTCCCTTAGT-AAGACGTTGCGACCTGTT-  
AAAT-----AATACTCTAAG-TA---TAAGTTAAAGTGCCTGTTT-AAGGTTATA-----  
-----CTTTTAT-----GGTA---TGCTTTATTCAGCTTAA--CTAATAAT-----A--CT-  
GGGTATTT-----AT---ATATTTAACAGT-  
AACAAATGACGGTGCTGAGCCACTTGCTTAAGATTGCTG--  
AATTTATTTAGTAATTTTGTGTAAGTGT-----TTATTGTCC-GTTCACAA-  
GTGGGGT-----GTTATATAGT-A-TAGTCGTAC-  
TTTTAGTGAGTCTATGCTATATCTCCTA-CTGTTGGTGAAGC-TGTG--ACTCT--AACTG-  
GCTT-----  
AAAACACCTGTCGGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT?????????????  
??????CTTGAAACACGGACCAAGGAGTTTATCGTGTACGCGAGTCATTGGGTG--  
AATAAAACCCAAAGGCGAAATGAAAGTAAAGAGTTGAGCATATTTGA-----  
TTAGAA-----TTTATTCTTTTCTTATTA-----  
TGTTCTCTTGAGGGAAGATAA--AAATTG-CGTTAATATTTA-----  
ATTGTATTAATT-----TACAAT-----TGATTTAATGTGAAT-  
TTTTCGCACTCCAGGGCGCTTG-TTTAACGC-  
TTGCGTGACAATGCGCACCTAGAGCGTACACGCTGATACCCGAAAGATGGTGAACATA  
TGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGA  
CGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCT  
AGTAGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGTAT-----  
-----CTTAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAG-  
AGCCCCGGCTTTCTTAAAA-----  
TGAAGCCGTGGGATTAGGATCAGAGTGCCAAGTGGGCAATTTTGGTAAGCAGAACT  
GGCGCTGTGG?????????CCGAAGTCTACATTTTAAATCCTCCCTGGGTTTGGTCTAATTT  
CACATATTATTAGTATAGAAACAGGAAAGGCAGAAGCCTTTGGCTCATTAGGAATAA  
TTTACGCAATACTTGCCATTGGCCTCCTGGGATTTATTGTATGAGCACACCATATATT  
TACTGTCGGAATAGATGTAGATACACGAGCATATTTTACTTCAGCCACTATAATTATT  
GCTGTCCCTACAGGAATCAAATCTTTAGCTGACTAGCAACCTTACATGGCTGTATA  
ATCAACTATTACCTAGTATTTTATGGGCTCTAGGCTTTGTATTTCTATTTACTATGGG  
GGGATTAACGGGAGTAATCTTAGCCAACCTCATCAATTGATATCATTCTCCACGATAC  
TTACTATGTTGTAGCCCACTTCCACT??  
Microstemmatoides atrocyanea  
??CCGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCCGCCCTGTCGGTGT  
ACTGGCATGTTCGCGGCATGTCCTGTCGGTGGTGAATGAAAT-----  
----CCCCGGTA----C-TACGTAGGCTT-TT--ATTAGCTGAAG-T-ATGTGCCG-----TGG--  
TGATTTCAATTTACCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT

TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAGCTTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGG?????????????  
??  
??  
??  
CCAC  
GAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAG  
AGTTCAAGAGTACG-TGAAACCG-CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTGCCACAT---GCTCGACTTTAGAGTGCT---  
TTTTGAGATGATAATTC-GCCCTATCCAG----TGTA--TACTGTGTATAGG-  
AGTGTTTTATTGCTCGCACTTTCTTTTAAAGTAGGATGCTGTGGTGGATCGCACTTCT  
CCTTTAGT-AGGACGTTGCGACCTGTC-AGAT-----ATACTCTAAA-TA-----  
TGGTTTTAGTGCCTGTGC-TAAGGCTATTGGAAAAAGTGTCTACATTTTTTTTTAA----  
-----TATA---ACTTTTTACAGCTTTAA--CCAATTCT-----A--TTGGGTATTT-----  
TGT---TTTCTTGACAGT-AACGACTGATGGTGTAGAGCCACTTTTAA---GCATTT-----  
TCAAAGAAATG---TAAAAAGTGTT-----TTTTTTTTATTTGCC-GTTTACAA-  
GTGGGCT---TCAAGGTATTAGTT-A-ATGTCGAA--CTTTGTTCGTCTAA-  
CTAAGCCGCCTA-CTGTTGT-AAAC-TGTGC-ACTTA--AACAGTTCTT-----  
ACAACACCGGTGCGGCGGCGATCCAG????????????????????????????????  
??  
ACCCAAAGGCGTAATGAAAGTAA  
AGAGTAGAAAAGAGAGAA-----  
GATGAAAAGGTAAATTATATATTTTTTTTGTCTTTGATATTCTCTCTC-----  
TCTTTCTCTCTAGGGAAGATGG--AAATCT-TATTTTAAATAT-----  
ATACAATATCT-----TGTAT-----TATATTAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTTTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACATATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCCCGGCTTTCT??  
??  
AAGTATATATTTTAAATTTTACCAGGATTTGGGTT  
AATCTCACATATTATTTCAATAGAACTGGCAAAGAGAAGCCTTTGGCTCCTTAGG  
ATAAATTTATGCAATGTTAACTATTGGACTACTAGGATTCATCGTTTGAGCTCACCAT  
ATATTTACAGTAGGAATAGATGTTGATACACGAGCTTATTTTACTTCTGCAACAATA  
TTATTGCAGTACCTACAGGGATTAATAATCTTTAGATGATTAGCCACCTTACATGGCA  
GACCTTTATCTTTCTCCCTAGTCTTTTATGAAGATTAGGATTCGTATTTTTATTTACA  
ATTGGAGGATTAAGTGAATTATTTTAGCTAATTCAAGAATTGATATTATTCTTCATG  
ATACTTATTATGTTGTTGCCA???????????

Neolibavius\_n\_sp

??  
CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAATTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTTCGCACTGTCCGGTTCACCGCCTGTCCGGTGT



ACTGGCATGTCGCGGAATGTCCTGTCGGTGGTGAATAAAGT-----  
----TCCTGGTG----C-TACCTAGGCTT-TT--TATAGCTGAAG-T-GTGTGCCG-----TGG--  
TGACTTTATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAAGTAGGCCGACACGTTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAAGATCTGCCTAAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACAGGCGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATTCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTTCT  
TTAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAAT????????????????????????  
??  
??  
??  
TGGTAAACTCCATCTAAGGCTAAATAT  
AACCACGAGTCCGATAGTAAACAAGTACCGTGAGGAAAAGTTGAAAAGAAGCTTGA  
AGAGAGAGTTCAATAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA-  
TATCTCGAAAGAAGAGATTCAAGTCTTAGCATTT--TGTTTACTTTTAAGTAC----  
CTTTGAGATGGTAATTCAGTATTATCCAG----TAAA--TACTG--TGGTAA-  
GAAGTTTTGCCGCTCGTACTTTATTTT-  
AAGTAAGCTGTAATGCTGGACCGCACTTCTCTTTTAGTAAGGACGTTGCAACCTGTC-  
AGAT-----AATACTCTAAG-TA----TGATTAAGTGTCTCTGT-AAATGTAATTTAAAA---  
-----CTTGTTTT-----AATATTATATCTTGCAGTAGCTTTTA--TCAATTTTA---  
CTA--CT-GGGTGTTT-----A---TATTCTGACAGT-ATCAACT-  
ATGGTGTAGAGCCACTAAAAA---TTAAGT-----AAGCATCCG-----  
-----TTTTACAA-GTTTGGT-----AATTATAAGTT-A-ACGTCTAA--  
CTTTTGTTAGTCTAA-CTTTATTACCTA-CTGTTTGT-AAAA-TGTA--  
ACTTGCTTTTAGTGCTC-----  
ATCACACCGGTTGGCAGCGATTTCAGGTTTCAGATTCTTATCCG????????????????  
??  
TAAAGGATAAAAAGATAGAAG-----AAGTTAA-----  
GCAATTTTCATCTTTGAT-----TCTTTTATCCTAGGGAAGATGG--ATATCT-  
TGTTTTAAAAA-----AAATTTTTTT-----  
TTTTTTAAACATAGAT-CTCCCGCATTCCCAGGGCGCTTG--  
GACTTGCTTTGCAAGTCTATGCGCACCTAGAGGGTACACGATGACACCCGAAAGATG  
GTGAACTATGCCTGGCCAGGCCGAAGCCAGGGGAAACCCTGGGGGAGGTCCGCAGC  
GATTCTGACGTGCAAATCGATCGTTAGAGCTGGGTATAGGGGGGAAAGACCAATCG  
AACCATCTAGTAGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCGCTCGATT--  
-----  
TTAAAGAGTTTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCAGGCTTTCTTAAAGAGAGAAAAAAGAAAATTTTT-----  
-----  
AATTTGGGCACCCAGAGGTCTATATTCTAATTCTACCGGGGTTTGGCCTAATTTTACA  
TATTATCGCCATAGAGACAGGGAAAAATGAAGCCTTTGGTTCCTTGGGAATAATTTA  
TGCTATGCTAACCATTGGCCTTTTAGGATTTATTGTATGAGCACATCATATATTTACT  
GTAGGAATAGATGTAGATACACGAGCCTACTTTACTTCAGCAACAATAATTATTGCT  
GTACCCACCGGAATTTAAATTTTTAGATGGCTAGCAACGCTACATGGAAGAGTTCTA



AGCCCCGGCTTTCTTAAAA-----  
TTGAAGCCGCGGGATTTCGGATCAGAGTGCC??  
?????????AAGTTTATATTCTCATTTTACCTGGATTGGATTAATTTCCCATATTGTATCC  
ATAGAAACAGGTAAAACAGAAGCCTTCGGTTCTTTAGGAATAATTTATGCTATACTA  
ACTATTGGCTTATTAGGATTCATTGTATGAGCTCATCATATATTTACAGTAGGTATAG  
ATGTAGATACACGTGCTTATTTTACATCTGCTACTATAATTATTGCTGTTCCACAGG  
AATTAATAATTTTAGATGATTAGCCACCCTTCATGGAAGAATCCTTAATTTCTCCCCT  
AGTTTACTTTGAGCATTGGGTTTTATCTTCTTATTTACAATTGGAGGGTTAACAGGAA  
TTATTCTAGCAAATTCAAGAATTGATATTGTACTTCACGACACCTACTATGTAGTAGC  
CCATTTTCATT??

nr\_Daraxa\_sp

??  
CCGTAA  
TCGGAATGAGTACACTTAAAT-CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGGAAT-----  
---CCCCAGGA---C-TTCTTAGGCTT-C---ATAGCTTATG-T-ATGTTCTG-----TAA--  
GGACTCCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGG????????????????????  
??  
??AGAGTGCAGCCCTAAGTAGGTGGTAACTCCATCTA  
AGGCTAAATATAACTACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAA  
AAGAACTTTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGAAGTAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATAT--TGTTTAACTTTTAAATGT----  
ATATGAGATGTTTATTC-GCCCTATTGAG----TGCA--AACTG--TAATAG-  
AGGGTTTATTCGCTCGCATTTAATT----  
TAGTAAAACAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGATCTGTT-  
AGAT-----AATACACTAAG-TA----TGATTAAAGTGTCTGTAT-TGAAGAAAGGAA-----  
-----ATTTTTT-----TCTT--TCTATTTTACAGCTTTAA--TCAATTAT----ATA--  
CT-GTGTATTT-----T--ATTTACTGACAGT-ATCAATTGATGGTGTGAGCCACTTA-----  
--TTTTTC-----AAGTG-----TCTTTAGCCTGTTTACAA-GTAGGTT---  
-----TATTAGTT-A-ATTTCCGGGCTTTTTGGTCCGTATAA-CTAGGCCACTTA-  
CTGTTAGT-TAAC-TGTG--CCTTA--GACTGTGCTT-----  
ATAACACTGATCAGCAGCGATCC??  
????????????????GTTTATTGTGTACGCAAGTCATTGGGTA--  
ATTAACCCAAAGGCGTAATAAAAGTAAAGAGTAATTTAAATAAA-----  
GGTAGCAAT-----ACTTTTTTTTTT-----  
TTAAATTCTTAAAGGAAGATAG--ATTGTT-----ATTACATTT----  
----TTTATAAAA-----GAATTTTATAAT-ATCTTGCAATTCCTAGGGCGCTTG-

TCTCTCGCATTGCGAGTATATGCGCACCTATAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCTCTGAAGTTTCCCTCAGGATAGCTTGCGTTCATT---

-----  
GAAACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCTCGGCTTTCTTAAAAC??  
??GAAGTATATATTCTCATTTTACCTGGATTTCG  
AATTATTTCCCATATTATTGCAATAGAAACAGGTA AAAATGAAGCTTTCGGAACACT  
TGGAATAATTATGCAATATTGGCTATTGGATTATTAGGGTTTATTGTATGAGCTCAT  
CATATATTTACCGTAGGAATAGATGTAGATACACGAGCTTATTTACATCAGCTACC  
ATAATTATTGCTGTACCTACAGGAATCAAAATTTTCAGATGGTTAGCTACTCTACATG  
GTAGAGATATTATTTTACTCCGAGTATCTTATGATCTTTTGGATTTGTATTTTATTC  
ACTATAGGGGGGATAACTGGAATTATTTTAGCTAACTCTAGTATTGATGTTACTTTAC  
ATGATACTTACTACGTAGTTGCCATTTTC?????

Oncocephalus\_sp

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGAGGTTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAACGGGAT-----  
----CCACGGTA----C-TGCATAGGCTT--C--TTTAGTCGATG-T-TAGTGTG-----TGT-  
GTATTTCCGTTTGCCGA---  
TCTCTCCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTAGTGCATGGAATGATAAAACAGGACCTCAGTTCTATT  
TTGTTGGTTTTTCGGAATATGAGGTAATGATCAATATGGACTGGCGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTTCAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATACAACCACGAGACCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACT  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCCTACTAGTT--TGCTTTACATTTAAATGT-----  
ATTGAGATGGCTTCTC-GCCCTATTTAG---TGTA AAAACTA--TGATAG-  
TGGGTTTAGTCACTCGCATTTAAATT-----  
GTCAGGGCAGTAGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACCTGTC-  
AAATAAT---TTTATCCTAAG-TA---TAGGCTAAATTGTCTGTTT-ATCGGTTTT-----  
-----ATCTTGC-----GATA---TTTCCTTTAACAGTTTTAG--CCATTAAT----ATG--  
CT-GGAT-----ATAATTGACAGT-AACAAATAATGGTGTAGGGCCACTATTTA---  
-----ATAGTG-----TCTATGGCT-ATTTGCAA-GTGAGTT----

GTATATTTTTAGTT-T-TCGTCGGA--TGTTTGTCCGTCAAA-CTATTACGCTTA-  
CTGTTGGT-AAAT-AGTA--CAAAA--  
GACAGTGCCTAATAATAAAACACCAGTCGGCAGCGATTCAGGTTTCAGGCTCTTATCC  
GGTCCGT????????????????????CTTGAAACACGGACCAAGGAG-----  
-----CTTTTAT---  
-----AATCTCTCTTAAGGGAAGATGG--AAATAT-AACAATACA-----  
-----ATTTGTTTGTAAAAT-  
TTCCCGCACTCCCAGGGCGCTTG-  
TATCTCGCTTTGCGAATCGATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT

GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGATGCCCCGGCTTTCTTAA  
AA-----GTGAAGCCGTGGGA-----  
TGTAAGCAGAACTGGCGCTGTGG????????????????????????????????????  
??  
??  
??  
??  
??  
????????????????????????

Opisthoplatys\_sp1

??  
CCGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTAAACAAGGATCTA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTTGCCGCCTGTCGGTGTA  
ACTGGCATGTCGCGGTATGTCCTGTCGGTGGTGAATGTAATG-----  
---GCAACGGTG---CCAGGGTATGCTT-----TATCAAAAG-T-  
ATACGCGGGTAACTTGTGT-TCAATTACATTTGCCGT---  
TTTTTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAGAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGG????????????????  
??  
??  
??  
TGAAGAGTGCAGCCCTAATTAGGTGGTAACTCCATCTA  
AGGCTAAATACAACCAGAGACCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAA  
AAGA ACTTTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
TTCAGGGGTAAACGGAAAAGACCTGAA--  
TATTCGAAAGGGGAGATTCACGTCTTACCGCCT---  
GTTTAACTATTGAATGTTATTGAGATGTTTCTTT-GCCCTGTCTAG--TTTATA--  
TACTG--TGATGG-TGGGTTGAAACGCTCGCATTTAATTA----  
GTA AAAACAGCGGTGGACCGCACTTCTCCCTCAGT-

AGGACGTTGCGATTTGTCAAAAA-----AATATTCTAAG-CA----  
TCCATTAAGTGTCTGTAT-TTGTGAAAGGTGTAG---GGGAATTTATTCTTTTACA-----  
-----TTTT---TTTCCCTATACAGCTTTAA--TAAATAAA-----TG--CT-GAATA-----  
TATTTAACAGT-AACAAATGATGGTGTGAGCCACTTTTA-----  
TATAAAGTG-----TCTTGCCA-ATTTACAA-GTGGGGTA-----TGAACATTAGTT-  
A-ATGTCGGTC-TTAATGTCCGTCTAA-CTGGTTCCTA-CTGTTGGT-AAAT-  
TGTTTGCTTTG--GACAGCGCTA-----  
ACAGCACCGATCAGCAGCGATCAA??  
??GCATAATAAAAGTAAAA  
GGTGTTTTGGATGATGGGGG-----AAGATTATT-----  
TTTTTTTTTTTTTCTTCATTTAT-----CCTTAATCCTAAGGGAAGATGA--  
TAATTA-CTTTTTCTCAA-----TTTTTTT-----TTGAG-----  
TAAAAAGATAAACAAT-TGTCCGCATTCCTGGGGCGCTTG--  
TTCTTGCTTTGCAAGTCTATGCGCACCTAGAGCGTACACGATAATACCCGAAAGATG  
GTGAACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGC  
GATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCG  
AACCATCTAGTAGCTGGTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGTT--  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCCCGGCTTTCT??  
??  
GTTAATTTACATATTATTGCAATAGAAACAGGTAAAAATGAAGCTTTTGGGGCAT  
TAGGGATAATTTATGCTATATTAGCTATTGGATTATTGGGATTTATTGTTTGAGCACA  
CCATATATTCACAATTGGCATAGATGTAGATACCCGAGCCTACTTTACATCAGCAAC  
TATAATTATTGCTGTTCCACAGGGATTAAAATCTTTAGATGATTAGCAACTTTACAT  
GGTAGAGTAATTCAACTCTCCCCAAGAACATTATGGGCCTTAGGTTTTGTTTTCTAT  
TACTATGGGAGGTTAAACGGGTATTATTTAGCCAATTCCAGTATTGATATTATTT  
ACATGATACATATTATGTAGTCGCTCACT?????????  
Opisthoplatys\_sp2  
??CGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTAAACAAGGATCTA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTTGCCGCTGTGCGGTGA  
ACTGGCATGTCGCGGTATGTCCTGTCGGTGGTGAATGTAATG-----  
---GCAACGGTG---CCAGGGTATGCTT-----TATCAAAG-T-  
ATACGCGGGTAACTTGTGT-TCAATTACATTTGCCGT---  
TTTTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAGAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACT????????????????????????????????????  
??  
??  
??ACCACGAGACCG

ATAGCGAACAAAGTACCGTGAGGGAAAGTTGAAAAGAAGCTTTGAAGAGAGAGTTCAA  
GAGTACG-TGAAACCG-TTCAGGGGTAAACGGAAAAGACCTGAA--  
TATTCGAAAGGGGAGATTACAGTCTTACCGCCT---  
GTTTAACTATTGAATGTTATTATTGAGATGTTTCTTC-GCCCTGTCCAG--TTTATA--  
TACTG--TGATGG-TGGGTTGAAACGCTCGCATTTAATTA-----  
GTA AAAACAGCGGTGGACCGCACTTCTCCCTCAGT-  
AGGACGTTGCGATTTGTCAAAAA-----AATATTCTAAG-CA----  
TCCGTTAAAGTGTCTGTAT-TTGTGAAAGGTGTAG---GGGAATTTTTTCTTTTACAT-----  
-----TTTT--TTCCCTATACAGCTTTAA--TAAATAAA----TG--CT-GAATA-----  
TATTTTAAACAGT-AACAAATGATGGTGTGAGCCACTTTACA-----  
AAAAGTG-----TCTTGCCA-ATTTACAA-GTGGGGTA----TGAACATTAGTT-A-  
ATGTCGGTC-TTAATGTCCGTCTAA-CTGGTTCTCCTA-CTGTTGGT-AAAT-  
TGTATGCTTTG--GACAGCGCTA-----  
ACAGCACCGATCGGCAGCGATCAAGGTTTCAGGCTCTTATCCGACCCGTCTTGAAAC??  
??  
AGTAAAAGGTGTTTTGGATGATGGGGG-----GGAAAATT-----  
TTTTTYTTCTTCATTTAT-----CCTTAATCCTAAGGGAAGATGA--TAATTA-  
CTTTTTTTTCC-----CCTGATAA-----AGGGGA-----  
GAGAAAAGAAAACGAT-TGTCCGCATTCCCGGGGCGCTTG--  
TTCTTGCTTTGCAAGTCTATGCGCACCTAGAGCGTACACGATAATACCCGAAAGATG  
GTGAACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGC  
GATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCG  
AACCATCTAGTAGCTGGTCCCTCTGAAGTTCCCTCAGGATAGCTTGCGCTCGGTT--  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCCCGGCTTTCTTAAAA-----  
TT??  
??  
??  
??  
??  
??  
??  
??  
Panstrongylus\_lignarius  
??  
CGTAATCGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-  
TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCCGCCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAACGGAGT-----  
----CCACAATG----C-TACATAGGCTT----TATAGCCGATG-T-TAGCATCG-----TGT-  
GTACTTCCGTTGGCCGA---  
TCTCTCTACTTCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACAGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAAACAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA

GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCTT  
TTAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAA????????????????????????  
??  
??  
??  
ATACAACCACGA  
GACCGATAGCGAACAGTACCGTGAGGGAAAGTTGAAAAGA ACTTTGAAGAGAGAG  
TTCAAGAGTACG-TGAAACCG-CTCAGGGGTAAACGGAAAAGACCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCAGTT--GCTCAACTTTTAAATGA----  
TTTGAGATGGCCTCTC-GCCCTATTAG---TGTAACAGCTG--TGGTAG-  
TGGGTTTGGTCGCTCGCATTTAAATT----  
GGAAGGGTAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACCTGTC-  
AAAT-----AATATTCTAAG-TA---TTTGGCTAATTTGTCTGTTT-TTAGGCAGT-----  
-----ACTGAAA-----GGTA---TTTCTTTAACAGTTTTAG--CCGTTTAA---ATA--  
CT-GGAT-----AAAATTGACAGT-AACGAATCATGGTGTGAGCCACTTGAAA--  
-TTGTAT--ATTTATATGTATAATG----GAAAGTGTC-----CTTGCGGCT-  
GTTTGCAA-GTGGGTT-----GGTAATTAGTT-T-TAGTCGGA--TTTTGGTCCGTCTAG-  
CTATTCCGCCTA-CTGTTGGT-AAAC-TGTT--CCTTG--GACAGTGCTT-----  
ATAACACCGGTCGGCAGCGATTAGGTTTCCAGGCTCTTATCCGACCCGTCTTGAACA  
CG????????????????CACGGACCAAGGAGTTTACGCGTGTACGCAAGTCATTGGGTG--  
TATAAAACCCAAAGGCGAAATGAAAGTAAAGAGTAGAGATTTGTAAA-----  
GAAG-----TTTACTTTTTTACA-----  
ATCTTCTCTTTAGGGAAGATGG--AAATAT-TATACAGT-----  
-----AATGTATGAAAAT-TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTTGCTTTGCGAGTCGATGCGCACCTAGAGCGTACACGCTGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ATCCCCGGCTTTCTTAAAA-----  
GTGAAGCCGTGGGAATCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAA  
CT-----  
G??  
??  
??  
??  
??  
??  
Peirates\_punctorius  
GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTTTTGGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCCTGTCGGTGGTGAACGGGAT-----  
----CCCCGGTG---C-TGTTGAGGCCG----TCAGGTCAAAG-C-ATGTACCG-----TGT-  
GTGTTTCCGTTTGCCTGA--  
TCTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG



AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAACAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGAGCTTCTACTCGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTGAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATATAACCACGAGACCGATAGCAAACAAGTACCGTGAGGGAAAGTTGAAAAGA  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAAAAGACCTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCACTT---GCTCAGCTTTTAAATAT-----  
TATGAGATGGCCATTC-GCCCTGTCCAG---TGTA--AACTG--TGATGG-  
TGGGTTTGTTCGCTCGTATTTATTAT-----  
GCAAGGGTGGTGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACCCGTC-  
AAAC-----AATGCTCTAAG-TA---TTTGTTTTAGTGTCTGTAT-AAGTGTATA-----  
-----CCGTTA-----GGTATATTTATCTTGTACAGCTTTAA--CAATTTAT----ATA--  
CT-GGGTAT-----TTGTTAGACGGT-AACAAACGATGGTGTGCGAGCCACTTTAAT--  
--TTTTT-----AAAGTGT-----CCACCGGCT-GTTTGCAA-GTGAGTT--  
---TGGTTAATAAAT-T-AGTCGGGC--TTTTGGTCCGTCTAA-TTATTCCACTTA-  
CTGTTGGC-GGAC-ATAG--CCTCG--GACAGTGCTT-----  
ACAACACCAGTCGGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT????????????  
????????CTTGAAACACGGACCAAGGAGTTTATCGTGTACGCAAGTCATTGGGTT--  
AGCTAAACCCAAAGGCGAAATGAAAGTAAATAGTGTTAA-----  
-----TAGTGGTTTTCCATTT-----TTTACACTTAAGGGAAGATAG--  
AATCTA-TTTGTAGATTTT-----AAATTTTTTT-----ATAAAAAA-TT-----  
-----TAGTTTACAAAATAGT-TTCTCGCACTCCCTGGGCGCTTG-  
TTCTCGTTTTTTGCGAGTCAATGCGCACCCAGAGCGTACACGCTGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGAT  
A-----  
AAGAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCCCGGCTTTCTTAAAA-----  
CTGAAGCCGTGGGACTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??  
??  
??  
??  
??  
??  
?????

Pothea\_lugens

??  
TCGGAATGAGTACACTTTAAAT-CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--



CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATAAGAT-----  
----CCCCGGTG---CC-TGCGTAGGCTT-CT---ATAGCTGAAG-TATGGTGCCG-----TAC-  
-GGATGTTATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGTAATAAGCGAAAGCA  
TTTGCCAAGTATGTCTAATTGATCAAGAACGAAAGTTAGAGGCTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTAATTTGACTCAACACGGGAAATCTCACCAGACCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTCAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAA  
ATATAACTACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGA  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATAT--TGTTAACTTTTAAATGTA---  
TTCTGAGATGTTTATTC-GCCCTATTCAG----TGCA--AACTG--TAATAG-  
AGGGTTTATTCGCTCGCATTTTATT----  
TAGTAAGAACAGTGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGATCTGTC-  
AGAT-----AATACTCCAAG-TA----TGGTTAAAGTGTCTGTAT-  
TTAAAATAATCTATCTTGGGG-----TTTACTCTAGGGTAGGTAAAATTA---  
TTTTTTCTTACAGCTTTAA--TCAATTTATTAATA--CT-GAGTATTT-----  
ATATTGATAGT-ATCAATTGATGGTGTGAGCCACTTAAACAA-----  
AAAAAGTG-----TCTTTAGCC-GTTTACAA-GTAGGTT-----TAATAGTT-  
GTTTTTCGGGTTTTTCGATCCGTATAA-CTAGAACACTTA-CTGTTTGT-AAAC-TGTG--  
CCTTA--GACTGTGCCTTA----  
TAAACACTGATCAGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT????????????  
??????CTTGAAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTA--  
ATTA  
AAACCCAAAGGCGTAATAAAAAGTAAAGAGTAATTTAATTAAT-----  
-----ATTTTCTTATATTTAA-----TTCTATTCTTAAAGGAAGATAG--  
ATTTTT-TTTCTACTAT-----ATCTGTTTT-----TTTACTAAA-----  
---ATAGTGTATAGTAGAA-ATCTAGCATTCTAGGGCGCTTG-  
TCTCTCGCTTTGCGAGTATATGCGCACCTTGAGCGTACACGATGATACCCGAAAGAT  
GGTGA  
ACTATGCCTGGCCAGGATGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGGTTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCTCTGAAGTTTCCTCAGGATAGCTTGCCTCATT  
-----  
GAACAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAA  
ACTCCAAATAGATGAG-  
AACATCGGCTTTCTTAAAA-----  
TGAAGCCGAGAAAAATGGATCAGAGTGCAAAGTGGGCCAATTTTGGTAAGCAGAAC

TGGCGCTGTGG??  
??  
??  
??  
??  
????

Rhiginia\_aimara

??  
CCGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGTATGTCCTGTCGGTGGTGAATGGAAT-----  
---CCCCGGTG---C-TGCCTAGGCTT-TT---ATAGCTGAAG-C-ACGTGCCG-----TAC--  
GGACTCCATTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGC??  
??  
????????????????AGAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACA  
ACTACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAA  
GAGAGAGTTCAAGAGTACG-TGAAACCG-CTCAGGGGTAAACGGAGAAGAAGTAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATAATATGTTTAACTTTTAAATGT---  
ACAAGAGATGTTTATTC-GCCCTATTCAG---TGCA--AACTG--TAATAG-  
AGGGTTTATTCGCTCGCATTTAATT---  
TAGTAAAACAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGATCTATC-  
AGAT-----AATACTCTAAG-CA----TGATTAAAGTGTCTGTAT-TAGAATTATTAGTT-----  
-----TTTAAATT-----AATT---TTTTCTTTTACAGCTTTAG--TCAATTAT---ATG--  
CT-GAGTATTTGTTGTTGT---TTCTGCTGATAGT-  
ATCAAATGATGGTGTGAGCCACTTATTT---TTATTT-----AAAGTG-----  
-----TCTTTAGCC-GTTTACAA-GTAGGTT-----TAGTAGTT-A-  
ATTTCCGGGCTTTTCGGTCCGTATAA-CTAGGACACTTA-CTGTTTGT-AAAC-TGTG--  
CCTA--GACTGTGCTT-----  
ATAACACTGATCGGCAGCGATTTCAG??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
TGAAGTTT

ACATTTTAATTTTACCAGGATTTGGGCTTATTTACATATTATTGCTATAGAAACAGG  
AAAAAATGAAGCTTTCGGAGCCTTAGGTATAATTTACGCTATATTAGCAATCGGACT  
TTTAGGATTTATTGTATGAGCCCACCACATATTTACAGTAGGAATAGATGTTGACACT  
CGAGCTTATTTTACATCAGCAACAATAATTATTGCCGTACCTACAGGAATTTAAATTT  
TTAGTTGATTAGCCACTTTACATGGTAGACCCATTACATTTACTCCCAGAATTATATG  
AGCATTAGGATTTATTTTTCTATTCACTATAGGAGGATTAAGTGGTGTAAATTTTAGCA  
AATTCAGAATTGATATTATTCTTCATGATACATATTATGTAGTAGCACATTTTCATT  
A?

*Rhiginia cinctiventris*

?????????AAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAATAACGATA  
CGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCCGTTCCGCCCTGTCCGGTGT  
ACTGGCATGTCCGCGTATGTCCTGTCCGGTGGTGAATGGGAT-----  
---CCCCGGTG---C-TGCCTAGGCTT-TT---ATAGTCGAAG-C-ACGTGCCG-----TAC--  
GGATGCCATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAAGGAATGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCA?????????????????  
??CGTGTTGCTTGATAGTGCAGCCCT  
AAGTAGGTGGTAAACTCCATCTAAGGCTAAATACTACGAGTCCGATAGCGAAC  
AAGTACCGTGAGGGAAAGTTGAAAAGAAGTTGAAAGAGAGAGTTCAAGAGTACG-  
TGAAACCG-CTCAGGGGTAAACGGAGAAGAACTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATGATATGTTAACTTTTAAATGT----  
ACAAGAGATGTTTATTC-GCCCTATTCAG----TGCA--AACTG--TAATAG-  
AGGGTTTATTCGCTCGCATTTAATT----  
TAGTAAAACAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGATCTATC-  
AGAT-----AATACTCTAAG-CA----TGATTAAGTGTCTGTAT-TAGAATTATTAGTTT---  
-----TTTGGCT-----AATC---TTTTCTTTTACAGCTTTAA--TCAATTAT----ATG--  
CT-GAGTATTT-----T---ACTTTTTGATAGT-ATCAAATGATTGTGTTGAGCCACTTA-----  
--TTATTT-----AAAGTG-----TCTTTAGCC-GTTTACAA-GTAGGTT---  
-----TAGTAGTT-A-ATTTCCGGGCTTTTCGGTCCGTATAA-CTAGGACACTTA-  
CTGTTTGT-AAAC-TGTG--CCTTA--GACTGTGCTT-----  
ATAACACTGATCAGCAGCGATTCAGGTTTCTTATCCGGTCCGT?????????????  
???????CTTGAACACGGACCAAGGAGTTTAT?GTGTACGCAAGTCATTGGGTA--  
ATTAACCCAAAGGCGTAATAAAAGTAAAAAGTGATTAAAAA-----  
-----TATTT-----TTAAATCTTAAAGGAAGAAAG--ATTTTT-  
TAAAATT-----TTACTACG-----AGTAAAAAAAAT-----TTT-  
-----AAAAA-ATCTTGCATTCTAGGGCGCTTG-  
TCTCTCGCATTGCGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT

```

GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCGCTCATT--
-----
GAAGCGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-
AACCTCGGCTTTCTTAAAA-----
TGAAGCCGTAGGATTCGGATCAGAGGCCAAGGG-----
CCTGGTAAGCAGA AACTGGCGCTGTGG?????????TGAAGTCTACATTTTAATTCTACC
CGGATTCGGACTAATCTCACACATCATTGCTATAGAAACAGGAAAAAATGAAGCCTT
TGGAGCTCTCGGAATAATCTATGCAATATTAGCAATTGGTCTTCTAGGATTCATTGTA
TGGGCCACCATATATTTACAGTAGGTATAGACGTAGATACTCGAGCCTATTTCACTT
CAGCAACAATAATTATTGCAGTACCCACAGGAATTAATAATTTTCAGTTGATTAGCCA
CTTACACGGAAGTCCCATCACTTTCACCCCTAGCATTCTTTGAGCACTAGGATTTGT
ATTCTTATTCACAATTGGGGGATTA ACTGGAGTTATCTTAGCAAATTCTAGAATTGAT
ATTATCCTTACGATACTACTACGTAGTAGCTCACTTC?????
    Rhinia_sp
?????????????????????????????????????????????????????????????CCGTAA
TCGGAATGAGTACACTTTAAAT-CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT
ACTGGCATGTCGCGGTATGTCCTGTCGGTGGTGAATGGGAT-----
---CCCCGGTG---C-TGCCTAGGCTT-TT---ATAGTCGAAG-C-ACGTGCCG-----TAC--
GGATGCCATTGCGCA--
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG
AACAAATTAGAGTGCTTAAAGCAGGCT-
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGAAGGCGATCA
GATACCGCCTAGTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA
ATAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGG????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
AAGGCTAAATACA ACTACGAGTCCGATAGCGAACCAAGTACCGTGAGGGAAAGTTGA
AAAGA ACTTTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-
CTCAGGGGTAAACGGAGAAGAACTGAA--
TTTTCGAAAGGGGAGATTCACGTCTTACCATGATATGTTAACTTTTAAATGT----
ACAAGAGATGTTTTATTC-GCCCTATTCAG---TGCA--AACTG--TAATAG-
AGGGTTTATTCGCTCGCATTTAATT----
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGATCTATC-
AGAT-----AATACTCTAAG-CA-----TGATTAAAGTGTCTGTAT-TAGAATTATTAGTTT---
-----TTGGCT-----AATC---TTTTCTTTTACAGCTTTAA--TCAATTAT----ATG--
CT-GAGTATTT-----T---ACTTTTTGATAGT-ATCAAATGATTGTGTTGAGCCACTTA----
--TTATTT-----AAAGTG-----TCTTTAGCC-GTTTACAA-GTAGGTT--
-----TAGTAGTT-A-ATTTTCGGGCTTTTTCGGTCCGTATAA-CTAGGACACTTA-
CTGTTTTGT-AAAC-TGTG--CCTTA--GACTGTGCTT-----

```

ATAACACTGATCAGCAGCGATTTCAGGTTTCAGTCTCTTATCCGACCCGTCTTGAACA  
 C???GGCGTAA  
 TAAAAGTAAAAAGTGATTTAAAAA-----  
 TATTT-----TTAAATTCCTTAAAGGAAGAAAG--ATTTTT-TAAAATT-----  
 -----TTTACTAYTA---GRRTAAAAAAAAT-----TTT-----AAAAA-  
 ATCTTGCATTCCCTAGGGCGCTTG-  
 TCTCTCGCATTGCGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
 GGTGAACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
 CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
 GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCATT--  
 -----  
 GAAGCGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
 AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
 AACCTCGGCTTTCTTAAAA-----TTGAAGCCGAAGGATTCCGGATCAGA-----  
 -----  
 AT?????????????????CTACATTTAATTCTACCCGGATTGACTAATCTCACACATCAT  
 TGCTATAGAAACAGGAAAAAATGAAGCCTTTGGAGCTCTCGGAATAATCTATGCAAT  
 ATTAGCAATTGGTCTTCTAGGATTCATTGTATGGGCCACCATATATTTACAGTAGGT  
 ATAGACGTAGATACTCGAGCCTATTTCACTTCAGCAACAATAATTATGTCAGTACCC  
 ACAGGAATTTAAATTTTCAGTTGATTAGCCACTTTACACGGAAGTCCCATCACTTTTA  
 CCCCTAGCATTCTTTGAGCACTAGGATTTGTATTCTTATTACAATTGGAGGATTAAC  
 TGGAGTTATCTTAGCAAATTCTAGAATTGATATTATCCTTCACGATACCTACTACGTA  
 GTAGCTCACTTCCA???

Saica\_sp

??????????AAATTACCCACTCCCGGCACGGGAGGTAGTGACAAAAATAACGATA  
 CGGGACTCTTTTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
 CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
 GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
 TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTGCGGTGA  
 ACTGGCATGTCGCGGATGTCTGTGCGGTGGTGAACGGGG-----  
 ----CACCGTA---C-TGCTTTGGCTT-TT---TAAGTCGATT-C-TAGTGCCG-----  
 GGCTGTTCCCTGTTTGCCGA--  
 TCTCTCCTACTTCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
 AACAAATTAGAGTGCTTAAAGCAGGT-  
 GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAAATAGGACCTCGGTTCTATT  
 TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGTCATTGTA  
 TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
 TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
 GATACCGCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTTCT  
 TTAATGACTCGGCGGGGAGCTTCTACTCGGGAAACCAAAGCTTTTTGGGTTCCGGGG  
 AAGTATGGTTGCAAAGCTGAAACCTTAAAGGAATGACGGAAGGGCACCACCAGGAG  
 TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAATCTCA????????????????????  
 ???  
 AAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCACGAGACCGATAGCGAAC  
 AAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACG-  
 TGAAACCG-CTCAGGGGTAAACGGAAAAGACCTGAA--  
 TTTTCGAAAGGGGAGATTCACGTCTTACCGCTT-AAATTTTCTTTTATGTAT----  
 AAAGAGATGGTCACTC-GCCCTATTCAG---TGCA--AACTG--TAATAG-

AGGGTTTGGTCGCTCGCATTAAAG-----  
ATTTGTTAGTGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACCTGTC-  
AAATAA----AATGCTCTAAG-TA----TTTTTAAATTGTCTGTAC-AATGGTTAT-----  
-----ACTGTAA-----AAGG--TTTTCCATATACAGTTTTAA--ATAATTTT----ATA--  
CT-GAGTA-----TATTTTGACAGT-AACAAACGATGGTGTGAGCCACTTATTA--  
-TTTTTTT-----AATGTAAGTG-----TCTTTGGCC-GTTTGCAA-  
GTGGGTA-----AGGTAATAGTT-T-TAGTCGGA--CTATTGTCCGTCTAG-  
TTATTCCTCCTA-CTGTTGGT-AAAC-TGTG--CCTTA--GACTGTGCTT-----  
ACCACACCAGTCGGCAGCGATCCAGGTTTCCAGGCTCTTATCCGGTCCGT????????????  
?????CTTGAAACACGGACCAAGGAGTTTAGCGTGTACGCAAGTCATTGGGTA--  
TTTAAACCCAAAGGCGAAATGAAAGTAAAGAGTAGTAAATTGTTGC-----  
AGAAATT-----TATTTTTGTATAT-----  
AATTTCTTTAGGGAAGATGG--AAAGTT-AATCATAAATTGATTATTGA-----  
TAATTTTTTATC-----TATAAT-----TGTTTTTGATATAGCT-  
TTCCCGCATTCCCGGGGCGCTTG-  
CATCTCGCTTTGCGAGTCAGTGCACCTAGAGCGTACACGCTGATACCCGAAAGAT  
GGTGAACATATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT

-----  
GTGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCGAAAACGAC  
CTCAACCTATTCTCAAACCTCAAATAGATGAG-AACCCCGGCTTCTTGAAAA-----  
-----

ATGAAGCCGTGGGATTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??  
??  
??  
??  
??  
????  
???

Santosia\_sp

??AT  
CGGAATGAGTACACTTTAAAT-CCTTTAACCAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGTCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGGATGGAAT-----  
---CCACGTTA---C-TGCTTTAGTTTACT--ATAAGCTGAAG-T-ATGTATCG-----TTC--  
GGGCGCCATTTGCCGA--  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAAT????????????????????????  
??



??  
??  
??  
??  
GTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACG-  
TGAAACCG-ATCAGGGGTAAACGGAGAAGATCTGAA--  
TTTTCGAAAGGGGAGATTACGTCTTACCATAT--TGTTAACTTATAAAATGTA---  
TTAAGAGATGTTTATTC-GCCCTATTTAG----TGCA--AACTG--TAATAG-  
AGGGTTTATTCGCTCGCATTATTTT---  
TAGTAAAACAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGATCTGTC-  
AGTT-----AATACTCGAAG-TA----TGATTAATAATGTCTGTAT-TTATGTAATTTTTTTT----  
-----TTTTTA-----AATT---ATATTTTTTACAGCTTTAA--TCAATTTA---ATA-  
CT-GAGTATTT-----T---AATATTGATAGT-  
AACAAATGATAGTGTGAGCCCTTTAAA-----AAAAAGTGTC-----  
-----TAGCC-GTTTACAA-GTATGTTTT-TAGGCTTATTTAGCT-A-  
ATTTCCGGTTTTATGATCCGTATAG-CAAGGCCACATA-CTGTTGGT-AGATGTGCT--  
CCTA--  
GA??  
??  
??  
??  
AGTAAAGAG  
TAATTTAATTTAGT-----AGTAATA-----CTTTAA-----  
TTAACTTCTTAAAGGAAGAAAG--ATTTA-ATTTT-----TTTTTTC-  
-----ATTAAAAGTTT-----TATACTTTTAAATGAAA-  
GTCTTGCATTCTAGGGCGCTTG-  
TCTCTCGCGTTGCGAGTATATGCGCACCTAAAGCGTACACGATGATACCCGAAAGAT  
GGTGAAGTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCATT-  
-----  
TTGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAG-----  
-----  
AA??  
TTGCAATAGAAACAGGAAAAAATGAAGCATTGGGGCTTTAGGTATAATTTATGCAA  
TGCTATCTATTGGTCTTTTAGGATTCATTGTTTGGAGCTCATCACAATTTACAATCGG  
AATAGATGTAGATACTCGAGCTTATTTACATCTGCTACTATGATTATTGCTGTCCCA  
ACAGGAATTAATAATTTTAGTTGACTTGTACTCTTCATGGAAGTAAAATTATATTCA  
CCCAAGTATAATATGGGCTTTAGGTTTTGTTTTTTGTTACTGTTGGCGGTTAACA  
GGCGTAATTTAGCAAATCAAGAATTGATATTATTTACATGACACCTATTATGTAG  
TAGCCATTTT?????  
Stenopoda\_sp  
GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCTGTCGGTGGTGAACGGAGT-----  
----CCACGATG---C-TGCATAGGCTT-CC--TTAGCCGATG-T-TAGGGTCG-----TGT-  
GTATTTCCGTTTGCCGA-----  
TCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTGAA  
CAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAACAGGACCTCAGTCTATT

TTGTTGGTTTTAGAAATATGAGGTAATGATCAATAAGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGTCAGCCAGCGATCCGCCGATGTTTCAT  
TTAATGACTCGGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCA????????????????  
??  
AAGCAGGTGGTAAACTCCATCTAAGGCTAAATAAAACCACGAGGCCGATAGCGAAC  
AAGTACCGTGAGGGAAAGTTGAAAAGAAGCTTTGAAGAGAGAGTTCAAGAGTACG-  
TGAAACCG-TTCAGGGGTAAACGGAAAAGACCTGAA--  
TATCCGAAAGGGGAGATTCACGTCCTACTAATT--CATCCTACATTTGAATGA----  
TTTGAGATGGCCCCTC-GCCCTATTTAG---TGTAACAGCTA--TAATAG-  
TGGGTTTGGTCGCTCTCATTTAAATA----  
GTTAGGATGGTGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCAACCTGTC-  
AAGC-ATTATAATATCCTAAG-CA----TTGGCTAAATTGTCTGTTT-GAAGGTTTA-----  
-----ATCGCAA-----GATA---TTTCCTTTAACAGTTTTAG--CTATTTTT---ATG-  
-CT-GGAT-----ATAATTGACAGT-ATCAAATGATGGTGTGAGCCACTTAAAA--  
--TT-----AAGTGT-----CTTATGGCT-ATTTGCAA-GTGAGTT---  
GGTTTTTTTTAGTT-T-GAGTCGGA--TTTTAATCCGTCCAA-CTATTCATCTTA-  
CTGTTGGT-AAAT-AGTA--CCTTA--GACTGTGCTT-----  
ATAACACCGGTTGGCAGCGATTCAGGTTTCAGGCTCTTATCCGGTCCGT????????????  
??????CTTGAAACACGGACCAAGGAGTTTATCGTGTACGCAAGTCATTGGGCG--  
TTTAAAACCCAAAGGCGAAATGAAAGTAAAGAGTTGAAATTTGTAAA-----  
AAAAA-----TTTTTTTTTTTAC-----  
AAACTCTCTTAAGGGAAGATGG--AAATAT-CGCATATATTTA-----  
-----TA-TGTGTGGAAATAT-TTCTCGCACTCCCAGGGCGCTTG-  
TATCTCGCTTAGCGAATCAGTGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ATCCCCGGCTTTCTTAAAA-----  
GTGAAGCCGTGGGGCTATGATCAAAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??  
??  
??  
??  
??  
??  
????  
???

Stenopodessa\_sp

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAA  
CGATACGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGTGTA

ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAACGGAGT-----  
----CCACGATG---C-TGCATAGGCTT-CC--TTTAGCCGATG-T-TAGGGTTCG-----TGT-  
GTATTTCCGTTTGCCGA-----  
TCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTGAA  
CAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAACAGGACCTCAGTTCTATT  
TTGTTGGTTTTAGAAATATGAGGTAATGATCAATAAGGACTGGCGGGGGCATTTCGTA  
TTGCGACG-  
TAGAGGTGAAATTCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCATTGCGCAAG  
TATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCAGATACCGCC  
CTAGTTCTAACCATAAACGATGTCAGCCAGCGATCCGCCGATGTTCAATTAATGACTC  
GGCGGGGAGCTTCTACCCGGGAAACCAAAGCTTTTGGGTTCGGGGGAAAGTATGGTT  
GCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAGTGGAGCCTGC  
GGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTGA  
CAGATTGATAGCTCTTTCTTGATTGAGTGGGTAGTGGTGCATGGCCGTTCCGTGTTGC  
TTGATAGTGCAGCCCTAAGCAGGTGGTAAACTCCATCTAAGGCTAAATAAAACCACG  
AGGCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGA  
GTTCAAGAGTACG-TGAAACCG-TTCAGGGGTAAACGGAAAAGACCTGAA--  
TATCCGAAAGGGGAGATTACGTCCTACTAATT--CATCCTACATTTGAATGA----  
TTTGAGATGGCCCATC-GCCCTATTTAG----TGTATCAGCTA--TAATAG-  
TGGGTATGGTCGCTCTCATTAAATT-----  
GTTAGGATGGTGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCAACCTGTC-  
AAGCAATTATAATATCCTAAG-CA---TTGGCTAAATTGTCTGTTT-GAAGGTTTA-----  
-----ATCGCAA-----GATA---TTCCTTTAACAGTTTTAG--CTATTTTT---  
ATG--CT-GGAT-----ATAATTGACAGT-  
ATCAAATGATGGTGTGAGCCACTTAAAA---TT-----AAGTGT-----  
---CTTATGGCT-ATTTGCAA-GTGAGTT---GGTTTTTTTAGTT-T-GAGTCGGA--  
TTTTAATCCGTCCAA-CTATTCATCTTA-CTGTTGGT-AAAT-AGTAC-CCTA--  
GACTGTGCTT-----  
ATAACACCGGTTGGCAGCGATTGAGGTTGAGGCTCTTATCCGGTCCGT?????????????  
??????CTTGAAACACGGACCAAGGAGTTTAGCGTGTACGCAAGTCATTGGGCG--  
TTTAAACCCAAAGGCGAAATGAAAGTAAAGAGTTGAAATTTGTAAA-----  
AGAAAATTT-----TTTTTTTTTTTAC-----  
AAACTCTCTTAAGGGAAGATGG--AAATAT-CACATATATCTT-----  
-----TA-----TA-TGTGTGAAAATAT-TTCTCGCACTCCAGGGCGCTTG-  
TATCTCGCTTAGCGAATCAGTGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ATCCCCGGCTTTCTAAAAG-----GAAGCCGGG-----  
TGTAAGCAGAACTGGCGCTGTGG????????????????????????????????  
??  
??  
??  
??

??  
????????????????????????????

Tanindrazanus\_marginatus

??CCGTAA  
TCGGAATGAGTACACTTTAAAT-CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAAT-----  
---CCCTGGTG---CTTACGTAGGCTT-TT--ATTAGCTGAAG-T-TTGTGCCG-----TGG--  
TGACTTCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAAGCTTTTGGGTTCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACG????????????????????  
??  
??  
??CTAAGGCTAAATACAAC  
CACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGA  
GAGAGTTCAAGAGTACG-TGAAACCG-CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCGCTT---GCACGACTTTGAAGTGC----  
TTTTGAGATGGTAATTC-GCCCTATCCAG----TGAA--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTATT---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA----TAGTTAAAGTGCCTGTCA-AAAGGTTATAAAATT--  
-----TATTT-----TATA-AC---CATTTTACAGCTTTAA--CTAATTTT-----A--  
CT-GGGTATTT-----A---TTTTCTGACAGT-  
AACGACTGATGGTGTAGAGCCACTTAAAA----ATTA-----TTTATAATT---  
TAAAAGTGC-----TTTTGGTTGCC-GTTTACAA-GTGGGT----TTTAGGTATTAGTT-  
A-ATGTCGAA--CTTTTGTTCGTCTAA-CTAAGCCGCTA-CTGTTTGT-AAAC-TGTG--  
CCTA--GACAGTGCTT-----  
ACAACACCGGTCGGCAGCGATCCAGG????????????????????????????????????  
????????????????????????????????GTACGCAAGTCATTGGGTG-  
ATATAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGAAGGATAGAAG-----  
AAGAAAA-----TTAATTTTCATCTTTGAT-----  
TCTTTCTCTCAAGGGAAGATGGATATATCT-TATTTTAAATAA-TATTTAG-----  
--TATTT-----ATATTAAG-TA-----TA-TTTAAAATTTAGA-  
TTCCCGCATTCCAGGGCGCTTG-  
TATCTCGCTCTGCGAGATTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTGC-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-

ACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTAGGATCAGAGTGCCAA????????????????????????????????  
????????????AGTATATATTCTAATTCTACCAGGATTTGGATTAATTTACACATTATTG  
CAACAGAACTGGCAAGAGAGAAGCATTGGATCATTGGGTATAATTTACGCTATAA  
TACTATCGGACTTCTTGGATTTATTGTATGAGCTCACCATATGTTCACTGTTGGGAT  
AGATGTTGACACTCGAGCCTACTTCACCTCAGCAACTATAATTATTGCCGTACCCACA  
GGAATTAATAATCTTCAGATGACTAGCAACATTACATGGAAGAGTAATAACCTTTTCT  
CCTAGAATACTATGAGCCTTAGGATTTGTGTTCCCTGTTCACTATTGGAGGCTTAACAG  
GAGTCATGCTAGCTAATTCCAGAATTGATATTATCTTACACGATACATACTATGTAGT  
AGCACATTTTC?????

*Tanindrazanus tenebricus*

??  
??  
GCCAGCAGCCGCGGTAATT  
CCAGCTCCAATAGCGTATATTAAGTTGTTGCGGTTAAAAAGCTCGTAGTTGGTTCTG  
CGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGTACTGGCATGTCGCGGCATGTCCT  
GTCGGTGGTGAATGAAAT-----CCCCGGTG----  
CTTACGTAGGCTT-TT--ATTAGCTGAAG-T-TTGTGCCG-----TGG--  
TGACTTCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAG????????????????????????????????  
??  
??  
AGTCCG  
ATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAA  
GAGTACG-TGAAACCG-CTCAGGGGTAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCGCTT---GCACGACTTTGAAGTGC----  
TTTTGAGATGGTAATTC-GTCCTATCCAG----TGAA--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTATT----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA----TAGTTAAAGTGCCTGTCA-AAAGGTTATAAAATT--  
-----TATTT-----TATA-AC---CATTTTACAGCTTTAA--CTAATTTT-----A--  
CT-GGGTATTT-----A---TTTTCTGACAGT-  
AACGACTGATGGTGTAGAGCCACTTAAAA---ATTA-----TTTATAATT----  
TAAAAGTGTG-----TTTTGGTTGCC-GTTTACAA-GTGGGTT----TTAGGTATTAGTT-  
A-ATGTCGAA--CTTTTGTTCGTCTAA-CTAAGCCGCCTA-CTGTTTGT-AAAC-TGTG--  
CCTTA--GACAGTGCTT-----  
ACAACACCGGTCGGCAGCGATCCAGGTTTCAGGCTCTTATCCGACCCGTCTTGAACA  
CGGACCAAGGAGA????????????????????????????????ACGCAAGTCATTGGGTG-  
ATATAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGAAGGATAGAAG-----  
AAGAAAA-----TTAATTTTCATCTTTGAT-----  
TCTTTCTCTCAAGGGAAGATGGATATATCT-TATTTTAAATAA-TATTTAG-----  
--TATTT-----ATATTAAG-TA-----TA-TTAAAATTTAGA-

TTCCCGCATTCCCAGGGCGCTTT-  
TATCTCGCTCTGCGAGATTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCGCTCGGTT

-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCCCGGCTTTCTTAAAA-----

TTGAAGCCGTGGGATTAGGATCAGAGTGCCAA????????????????????????????  
????????????GTATATATTCTGATTTTACCAGGATTTGGACTCATTTCACATATTATTGC  
TACAGAAACCGGTAAAAGAGAAGCGTTTGGGTCACTAGGAATAATTTATGCAATAAT  
TACTATTGGACTTTTAGGATTCATTGTCTGAGCTCATCATATATTCACTGTGGGAATA  
GATGTTGATACTCGAGCCTATTTCACTTCAGCAACTATAATCATTGCTGTACCTACAG  
GAATTA AAAATCTTCAGATGATTAGCAACATTACATGGAAGAACCATGACATTTTCCC  
CAAGAATATTATGAGCATTAGGATTTGTATTCTTATTACAATTGGAGGTTTAAACAG  
GAGTAATACTAGCCAACCTCAAGAATTGATATTGTTCTTCACGATACATACTATGTAGT  
AGCACATTTTCAC???

*Tanindranus varicolor*

??  
??  
??  
??  
GTTTCGCCGCCTGTCGGT  
GTTACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAAT-----  
-----CCCTGGTG---CTTACGTAGGCTT-TT--AATAGCTGAAG-T-TTGTGCCG-----  
TGG--TGAATTCATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCT????????????????????????????????????  
??  
??  
??  
??  
??  
GTTAGGTGGTAAACTTCAT  
CTAAGGCTAAATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTT  
GAAAAGA ACTTTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCATTT---GAATGGCTTTAAAATGC----  
TTTTGAGATGGGAATTC-GCCCTATCTAG----TGAA--TACTG--TGATAG-  
AGGGTTTTACCGCTCGCATTTTATT----  
AGTAAGGTTTCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA-----TAGTTAAAGTGCCTGTTT-AAAGATTATAAAATT---  
-----TATTT-----TATA-AC---CATTTTACAGCTTTAA--TTAATTTT-----A--CT-  
GGGTATTT-----A---TTTTCTGACAGT-AACGAATGATGGTGTAGAGCCACTTAAAA---  
-ATTA-----TTTATAATT--TTATAAGTGTC---CAATTTTTTTTGGATTGCC-  
GTTTACAA-GTGGGTT---TTTAGGTATTAGTT-A-ATGTCAAA--CTTTTGTTCGTCTAA-  
CTAAGCCGCTA-CTGTTTGT-AAAC-TGTG--CCTTG--GACAGTGCTT-----

ACAACACCGGTCAGCAGCGATCCAGGTTTCAGGCTCTTATCCG????????????????????  
??  
??  
??  
??  
??  
??  
??  
TATTTAG-----TATTT-----  
ATATTAAG-TA-----TA-TTTAAAATTTAGA-TTCCCGCATTCCCAGGGCGCTTG-  
TATCTCGCTTTGCGAGATTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACCTATGCCTGGCCAGGACGAAGCCAGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCGCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ACCTCCGGCTTTCTTAAA-----  
TTGAAGCCGTGGGATTTGGATCAGAGTGCCAAGTGGGCCAATTTTGTAAGCAGAAC  
TGGC-----  
GA????????CCAGAAGTTTACATTTAATTTTACCAGGATTGGGAATTATTTTCGCACAT  
TATTGCAATAGAAACCGGTAAAATGAAGCATTTCGGCTCATTGGGGATAATCTATGC  
ATAATCACTATTGGATTATTGGGGTTTATTGTCTGAGCTCATCACATTTACTGTA  
GGGATAGATGTAGATACTCGAGCCTACTTCACTTCGGCAACTATAATTATCGCTGTA  
CCAACAGGAATCAAATCTTTAGATGATTAGCAACATTACAAGGCAGTATCATATTA  
TTTTCTCCTAGAATGATATGAGCCCTTGCTTCGTATTTTATTACAATTGGAGGAT  
TAACAGGTGTCATACTCGCTAATTCAAGAATTGATATTATCCTCCATGATACTTATTA  
CGTAGTAGCACATTTCCA???

*Toxopus\_brucei*

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
CGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGT  
TGAAAAGA ACTTTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTACGTCTTACCACTT---GCTCGACTTTGAAGTGC----  
TCTTGAGATGGTAATTC-GCCCTATCCAG----TGCA--TACTG--TGATAG-  
AGGGTTTTACCGCTCGCATTTTATA---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGGA-----ATACTCTAAG-TA-----TAGTTAAAGTGCCTGTT-AAATGTTATAAAATT---  
-----TATATTT-----TATA-ACTTTATTTTACAGCTTTAA--TTAATTT-----A--  
CT-GGGTATTT-----TTTTTTGTTCTGACAGT-  
AACGACTGATGGTGTAGAGCCACTTAAAC----ATTATT-----TTTTATAATG----  
TAAAAGTGTGCCT-TATTCTTTTTTTGCT-GTTTACAA-GTGGGT-----

TAAGGTATTAGTT-A-ATGTCGAA--CTTTTGTTTCGTCTAA-CTAAGCCGCCTA-  
CTGTTTGT-AAAC-TGTGC-CCTTG--  
GACAGTGC??  
????????????????????????????????ACGCAAGTCATTGGGTG-  
ATAGAAAACCCAAAGGCGTAATGAAAGTAAAGAGTATAAAGATAGAAG-----  
AAGAAAA-----TTCATTTTCATCTTTGAT-----  
TCTTCTCTCAAGGGAAGATGG--ATATCT-TATTTTAAATAA-TATTTAG-----  
TATTT-----TTATTAAG-TA-----TT-TTTAAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-  
TATCTCGCTCTGCGAGTCTACGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCGCTCAAT  
-----  
GTGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGAT??  
????????TATATATTCTCATTTTACCAGGATTTGGCTTAATTTACATATTATCGCAACAG  
AGACTGGTAAAAGTGAAGCTTTTGGATCATTAGGAATAATTTATGCAATATTAACAA  
TTGGATTATTAGGATTTATTGTTTGAGCACACCATAATTTACTGTAGGTATAGATGT  
AGACACCCGAGCATATTTTACCTCTGCAACCATAATTATTGCAGTACCTACAGGAAT  
TAAAATTTTAGATGACTAGCCACTTTCATGGAAGATCTACAAAATTCTCACCAAG  
AATACTATGAGCATTAGGATTCGTATTCTTTTACTATTGGTGGATTAACGGGAGTA  
GTACTTGCAAACCTCAAGAATTGATATTATTTTACATGACACTTATTATGTAGTAGCCC  
ACTTCCA???

*Toxopus fisheri*

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
????????????????????AAGGCTAAATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGG  
GAAAGTTGAAAAGAACCTTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAACCGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCACTT---GCTCGACTTTGAAGTGC----  
TCTTGAGATGGTAATTC-GCCCTATCCAG---TGCA--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTATT---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AAAT-----AATACTCTAAG-TA-----TAGTTAAAGTGTCTGTCT-AAAGGTTATAAAAATT---  
-----TATTT-----TATA-ACATTTATTAACAGCTTTAATTTTAAATTT-----A-



-CT-GGGTATTT-----ATT-TTTTTTTGACAGT-  
AACAAATGATGGTGTAGAGCCACTTAAAC----ATTA-----TTTATAATG----  
TAAAAGTGTGTC-----CATTTTTGCC-GTTTACAA-  
GTTGGTTTAAGGTATTTTTTTAGTT-ATTAGTCGAA--CTTTTGTTCGTCTAA-  
CTAAGCCGCCTA-CTGTTTGT-AAAC-TGTG--CCTTG--GACAGTGCTT-----  
ACAACACCGTTCGGCAGCGATCCAGG????????????????????????????????  
????????????AGGAGTTTATTGTGTACGCAAGTCATTGGGTG-  
ATAGAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGAAAGATAGAAG-----  
AAGAAA-----TTAATTTTCATCTTTGAT-----  
TCTTTCTCTCAAGGGAAGATGG--ATATCT-TATTTTAAATAA-TATTTAG-----  
TATTT-----ATATTGAG-TA-----TA-TTTAAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-  
TATCTCGCTCTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCCCGGCTTTCTTAAAA-----  
TGAAGCCGT??  
??TACATTTTAAATTCTACCCGGATTCGGATTAATTTCCCATATTATCGCAATAGAAACA  
GGA AAAAATGAAGCATTGGATCTCTAGGGATAATTTATGCAATACTTACTATTGGA  
CTATTAGGATTTATTGTTTGGGCCATCATATATTTACAGTTGGAATAGATGTAGATA  
CCCGAGCTTATTTCACTTCAGCAACAATAATTATTGCCGTACCTACCGGTATTAGGAT  
TTTTAGTTGACTAGCTACAATACATGGAAGTATTATAACTTTCTCACCCAGAATATTA  
TGAGCATTAGGATTCGTATTTTTATTACAATTGGAGGATTAACAGGAGTTATTTTG  
CAA ACTCTAGAATTGATATTATTTACATGATACTTATTATGTAGTTGCACACTTCCA  
T???

*Toxopus griswoldi*

??  
??  
CAATAGCGTATATTAAGTTGTTGCGGTTAAAAGCTCGTAGTTGGTTCTGCGTGCC  
GCGCTGTGCGTTCGCCGCCTGTCGGTGTACTGGCATGTCGCGGCATGTCCTGTGCGT  
GGTGAATGAAAT-----CCCCGGTG----C-TACGTAGGCTT-TT--  
ATTAGCTGAAG-T-TTGTGCCG-----TGG--TGA CTTCATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTGCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGAGCTTCTA????????????????????????????????  
??  
??  
??  
??  
??  
CTGAGGGGAAAGTTGAAAAGA ACTTTGAAGAGAGAGTTCAAGAGTACG-

TGAAACCG-CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTACGTCTTACCCTT---GCTCGACTTTGAAGTGC----  
TCTTGAGATGGTAATTC-GTCCCTATCCAG----TGAA--TACTG--TGGTAG-  
AGGGTTTTACCGCTCGCATTTTTTTT----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGAT-----AATACTCTAAG-TA-----TAGTTAAAGAGTCTGTTT-AAAGGTTATAAAATT---  
-----TATTT-----TGTA-ACTTTTTTTTTACAGCTTTGA--TTAATTTT-----A--  
CT-GGGTATTT-----TT--TATTCTGACAGT-  
ATCGACTGATGGTGTAGAGCCACTTAAAC----ATTA-----TTTATAATG---  
TTTAAAGCGTC-----CATTGTTGCC-GTTTACAA-GTGGGTAA--  
GGTATTAATTAGTT-A-ATGTCGAA--CTTTGTTCTGCTCTAA-CTAAGCCCCCTA-  
CTGTTTGT-AAAC-TGTG--CCTTG--GACAGTTCTT-----  
ACAACACCGGTCAGCAGCGATCCAGG????????????????????????????????  
??  
AGTAGAAAGATAGAAG-----AAGAAA-----  
TTTATTTTCATCTTTGAT-----TCTTTCTCTCAAGGGAAGATGG--ATATC--  
TAATTTAAATTA-----TATTT-----ATTATTAAGTTA-----TA-  
TTTAAATTTAGA-TTCCCGCATTCCCAGGGCGCTTG-  
TATTTGCTTTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGA ACTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCCAAATAGATGAG-  
AACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGC-----  
G????????????????????TTCTTATTTTACCGGGATTTGGATTAATTTACATATTATTT  
AATAGAACTGGGAAAAACGAAGCGTTTGGATCACTTGGATAATCTACGCCATAAT  
TACAATTGGATTATTAGGATTCATTGTATGAGCGCATCATATATTTACAGTAGGAATA  
GATGTAGATACCCGAGCTTATTTTACATCAGCAACTATAATCATTGCAGTACCAACA  
GGTATTAATAATCTTTAGATGATTAGCAACCATAACCGGTAGTGTTACTATATCTTCAC  
CTAGAATACTTTGAGCATTAGGATTTGTATTCTTATTACCATTGGGGGATTAACAGG  
AGTTGTATTAGCCAACCTCAAGAATTGATATTATTTTACACGATACATATTATGTAGTA  
GCCCACT????????

Toxopus toamasina

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??



??  
??  
AAGTTGAAAAGAAGTTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTACGCTTACCCTT---GCTCGACTTTGAAGTGC---  
TCTTGAGATGGTAATTC-GCCCTATCCAG---TGCA--TACTG--TGATAG-  
AGGGTTTTACCCTCGCATTATA---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-AGGACGTTGCGACCTGTC-  
AGGA-----AATACTCTAAG-TA----TAGTTAAAGTGCCTGTTT-AAATGTTATAAAAATT---  
-----TATATTT-----TATA-ACTTTATTTTACAGCTTTAA--TTAATTTT-----A--  
CT-GGGTATTT---TTTTTTTTTTTTTTGACAGT-  
AACAACTGAGGGGGGAGAGCCCCTTAAAA---ATTTTT-----TTTTTAAAG---  
TAAAAAGGGTTCGCTTATTTTTTTTTTTTTGTC-GTTTACAA-GTGGGT---  
TTAAGGTATTAGTT-A-ATGTCGAA--CTTTGTTCGTCTAA-CTAAGCCGCTA-  
CTGTTTGT-AAAC-TGTGC-CCTG--GACAGTGCTT-----  
ACTACACCGTTCGCGAGCGAT??  
??  
GTACGCAAGTCATTGGGTG-  
ATAGAAAACCAAAGGCGTAATGAAAGTAAAGAGTATAAAGATAGAAG-----  
AAGAAA-----TTCATTTTCATCTTTGAT-----  
TCTTCTCTCAAGGGAAGATGG--ATATCT-TATTTTAAATAA-TATTTAG-----  
TATTT-----TTATTAAG-TA-----TG-TTTAAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTT-  
TATCTCGCTCTGCGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGAT  
GGTGAAGTATGCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCGAG  
CGATTCTGACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT  
-----  
GTGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
AACCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAAG??  
??  
AGAAGTATATAATTTTACCAGGATTCGGATTAATTTACATATTATC  
GCAACAGAGACTGGTAAAAATGAAGCTTTCGGATCCCTGGGAATAATCTATGCAATA  
TTAACAATTGGATTGTTAGGATTTATTGTATGAGCACACCACATATTCACTGTAGGAA  
TAGACGTGGATACACGAGCATATTTACCTCTGCAACAATAATTATTGCAGTACCCA  
CAGGAATTAATAATTTTTCAGATGACTAGCCACTCTCCATGGAAGAATAACAATTCT  
CCCCAAGAATGTTGTGAGCACTAGGATTCGTATTTCTATTTACTATTGGAGGATTAAC  
AGGAGTGGTTCTCGCAAACCTCAAGAATTGATATCATTTTACATGATACTTATTATGT  
GTAGCCCATTTCT?????

*Triatoma recurva*

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAA  
CGATACGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGTA  
ACTGGCATGTCGCGGCATGTCTGTGCGGTGGTGAACGGAGT-----  
----CCACAATG---C-TACATAGGCTT-----TATAGCCGATG-T-TAGCGTCG-----TGT-  
GTACTTCCGTTGGCCGA---

TCTCTCTACTTCGGTGCTCTTAAACGAGTGTCTGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATCTCTGCCTGAATAGTGGTGCATGGAATGATAAAACAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCT  
TTAATGACTCGGCGGGGAGCTTCCACCCGGGAAACCAAAGCTTTTGGGTTCGGGGG  
AAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGAG  
TGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATTG  
GAAGGATTGACAGATTGATAGCTCTTCTTGATTCAAGTGGGTAGTGGTGCATGGCCG  
TTCCGTGTTGCTTGATAGTGCAGCCCTAAGCAGGTGGTAACTCCATCTAAGGCTAA  
ATACAACCACGAGACCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGT  
TTGAAGAGAGAGTTCAAGAGTACG-TGAAACCG-  
CTCAGGGGTAAACGGAAAAGACCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCAGTT--GCTCAACTTTTAAATGA-----  
TTTGAGATGGCCTCTC-GCCCTATCCAG----TGTAAGTCTG--TGGTAG-  
TGGGTTTGGTCGCTCGCATTAAATT-----  
GGAAGGGTAATGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACCTGTC-  
AAAT-----AATATCCTAAG-TA---TATGGCTAATTTGTCTGTTT-TTAGGCAAT-----  
-----ACCGCGA-----GGTA---TTTCCTTCAACAGTTTTAG--CCGTTTAT----  
ATAT?CT-GGAT-----ATAATTGACAGT-  
AACGAATCATGGTGTGAGCCACTTGAAA---TTGTAT--ATTTATATGTATAATG----  
GAAAGTGTG-----CTTGTGGCT-GTTTGCAA-GTGGGT-----GGTTAATTAGTT-T-  
TAGTCGGA--TTTTGGTCCGTCTAG-CTATTCCGCCTA-CTGTTGGT-AAAC-TGTT--  
CCTTG--GACAGTGCTT-----  
ATAACACCGGTTCGGCAGCGATTGAGGTTTCCGCTCTTATCCGGTCCGT????????????  
?????CTTGAAACACGGACCAAGGAGTTTAGCGTGTACGCAAGTCAATTGGGTG--  
TATAAAACCAAAGGCGAAATGAAAGTAAAGAGTTGAGATTTGTAGA-----  
AAAG-----TTTACTTTTTTACA-----  
ATTTTCTCTTTAGGGAAGATGG--GAATAT-CATGCAGT-----  
-----AATGTATGAAGAT-TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTTGCTTTGCGAGTCTATGCGCACCTAGAGCGTACACGCTGATACCCGAAAGAT  
GGTGAAGTATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAG  
CGATTCTGACGTGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATC  
GAACCATCTAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT  
-----  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAG-  
ATCCCCGGCTTTCTTAAAA-----  
GTGAAGCCGTGGGAATCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??  
??  
??  
??  
??  
????  
???

Tribelocephala\_peyrerasi

GCAGCAGGCACGCAAATTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAA  
CGATCGGGGACTCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCAATAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTTCCGCCTGTCGGTGT  
ACTGGCATGTCTCGGCATGTCTGTCGGTGGTGAATGTAATCTGGCCAATAGGTTA  
AGGGTTAGTAGTTCGATGTGCGCTAATGCG----C-TACACGGGCTT-TT--  
CCCCTTCTAACTTGTTGGTTTT-----TTT--  
TGATTATGTTTACCGATTTTTTTTCTACTCCGGTGCTCTTAAACGAGTGTCTGAAGTA  
GGCCGACACGTTCACTTTGAACAAATTAGAGTGCTTAAAGCAGGCTGGAAATATCTG  
CTGAATAGTGGTGCATGGAATGATAAAATAGGACCTCGGTTCTATTTTATTGGTTTT  
AGGAATATGAGGTAATGATCAATATGGACTGGCAGGGGGCATTTCGTATTGCGACGTTA  
GAGGTGAAATTCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCATTGCGCAAGTA  
TGTCTTAATTGATCAAAAACGAAAGTTAGAGGTTCCAA-  
GCGATCAGATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGA  
TGTTCCTTAATGACTCGGCGGGGAGCTTTTACCCGGGAAACCAAAGCTTTTGGGTTCC  
CGGGGGAAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCCAC  
CAGGAGTGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCA????????????  
??CGTGTTGCTTGATAGTGC  
AGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATAGTACCACGAGACCGATAG  
CGAACAAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGAGT  
ACG-TGAAACCG-CTCAGGGGTAACGGAAAAGACCTGAA--  
TATTCGAAAGGGGAGATTCACGTCTTACCGCAT---GTTTAACTTTTTATCTT----  
TTATGAGATGATTTTTC-GCCCTGTCCAGTTTTTTTTA--TACTG--  
TGATGGTTGGGTTAAATCGCTCGTATATAATTA-----  
GTAAAACAGCGGTGGACCGCACTTCTCCCTTAGT-AGGACGTTGCGACTCGTG-  
GGACAAT-ATATTATTCTAAG-CT----ATATTAAGTGTCCTTTAT-ATATTGTATGTAGC-  
-----ATTGTTA-----TTTTA---TAATATATAGCTATAA--TAAATAAA-----  
TG--CT-GAATA-----ATTTCCGACAGT-  
AACGAATGATAGTGTTGAGCCGCTAATTT---TTTTCT-----TTAGCG-----  
-----TTTTGACCA-ATTTACAA-GTGGGGTA----KGAATAATTATT-A-ATGTCGAA--  
CTTTTGTTTCGTCTAA-TATGTTCTCCTA-CTGTTGGT-AAAT-TGTTTGGTTTCA--  
ACAAGGCTT-----  
ATAACTGATCAGCAGCGATCCAGTTTCCAGGCTCTTATCCGGTCCGT????????????  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??







GGAATAGATGTAGATACCCGAGCTTACTTCACCTCTGCAACAATAATCATTGCTGTC  
CCCACAGGTATTTAAAATCTTCAGATGACTCGCCACCTTACACGGCAGTATAATTATA  
CTTTCACCAAGAATTATATGAGCATTAGGATTCGTATTCTTACCATAGGAGGAT  
TAACTGGAGTAATTCTAGCTAATTCAAGAATCGACATTATCCTTCACGATACTTACTA  
TGTCGTGGCACACTT???????

;  
END;

BEGIN SETS;

CHARSET 18S\_G\_INS\_i\_aligned = 1 - 1024;  
CHARSET 28S\_D2\_G\_INS\_i\_aligned = 1025 - 1855;  
CHARSET 28S\_D3\_5\_G\_INS\_i\_aligned = 1856 - 2628;  
CHARSET COI\_G\_INS\_i\_aligned = 2629 - 3054;

TAXSET LENGTH\_ATLEAST\_400 = 1 - 56;  
TAXSET LENGTH\_ATLEAST\_600 = 1 - 19 21 - 56;  
TAXSET LENGTH\_ATLEAST\_800 = 1 - 19 21 - 56;  
TAXSET LENGTH\_ATLEAST\_1000 = 1 - 19 21 - 56;  
TAXSET LENGTH\_ATLEAST\_1200 = 1 - 19 21 - 50 52 - 56;  
TAXSET LENGTH\_ATLEAST\_1400 = 1 - 19 21 - 50 52 - 56;  
TAXSET LENGTH\_ATLEAST\_1600 = 1 - 19 21 - 23 25 - 50 52 - 56;  
TAXSET LENGTH\_ATLEAST\_2000 = 1 - 4 6 - 8 10 - 19 21 - 22 25 - 35 37 39 -

47 50 52 - 53;

TAXSET LENGTH\_ATLEAST\_3000 = 1 17 - 18;

TAXSET CHARSETS\_ATLEAST\_4 = 1 3 - 4 6 - 7 10 - 13 16 - 18 21 25 - 30 32 39  
- 40 42 45 - 47 50 52;

TAXSET CHARSETS\_ATLEAST\_3 = 1 - 19 21 - 22 25 - 50 52 - 53 55 - 56;

TAXSET CHARSETS\_ATLEAST\_2 = 1 - 19 21 - 56;

TAXSET TAXA\_HAVING\_18S\_G\_INS\_i\_aligned = 1 - 19 21 - 47 50 52 - 54 56;

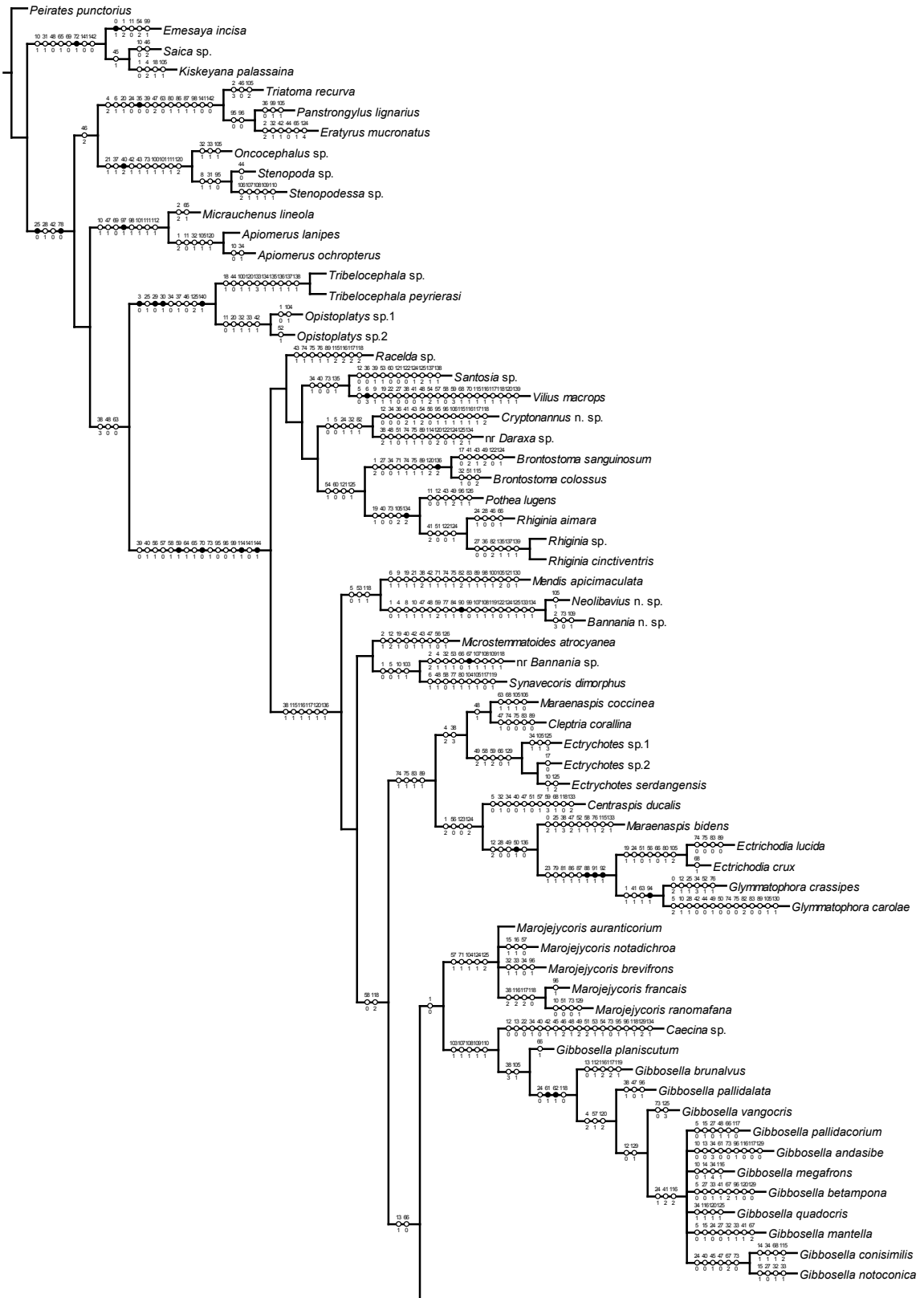
TAXSET TAXA\_HAVING\_28S\_D2\_G\_INS\_i\_aligned = 1 - 4 6 - 8 10 - 19 21 - 23  
25 - 55;

TAXSET TAXA\_HAVING\_28S\_D3\_5\_G\_INS\_i\_aligned = 1 - 19 21 - 22 24 - 35 37  
39 - 50 52 - 53 55 - 56;

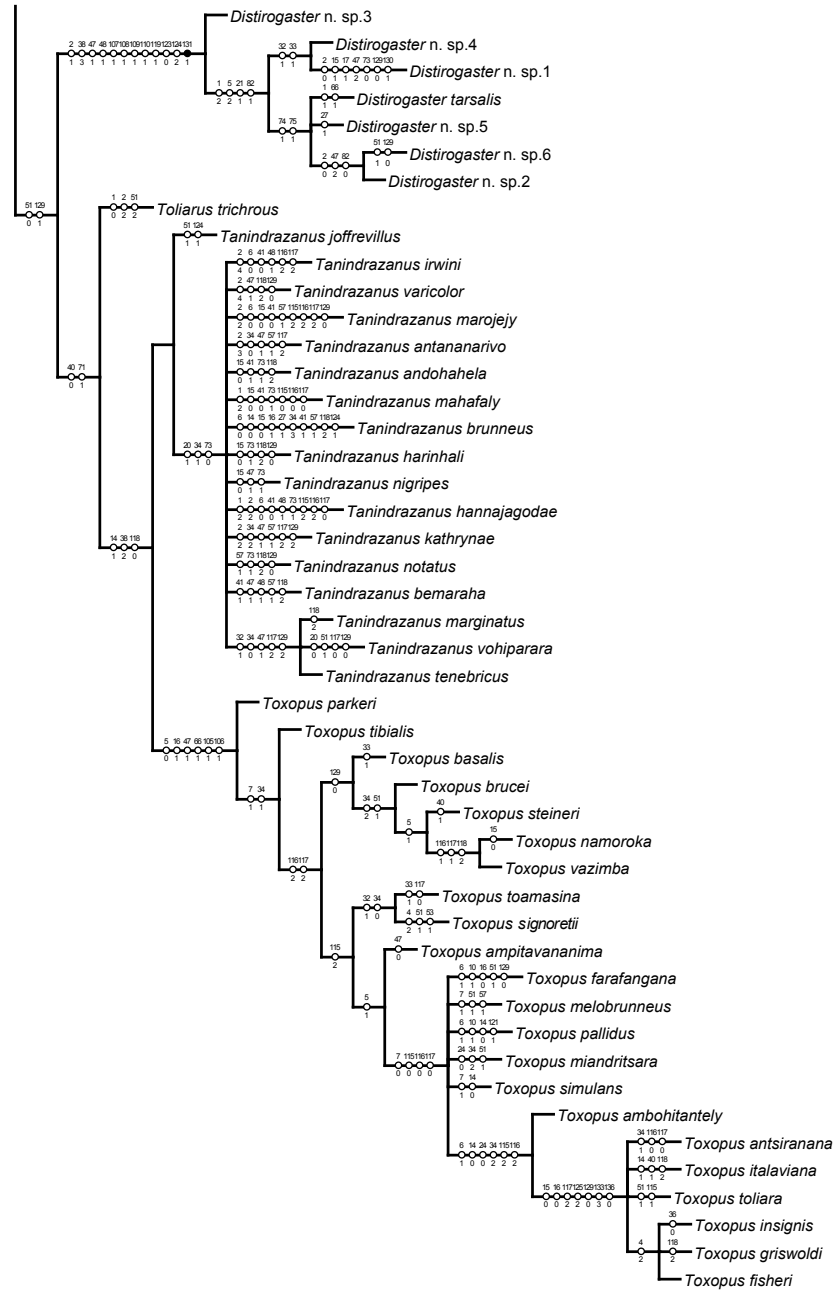
TAXSET TAXA\_HAVING\_COI\_G\_INS\_i\_aligned = 1 3 - 7 9 - 13 16 - 18 20 - 21  
25 - 30 32 36 38 - 40 42 45 - 52 55 - 56;

END;

Supplementary material 3.3. Unambiguous character optimizations on the strict consensus tree from parsimony analysis of 110 taxa and a combined morphological and molecular dataset.



Supplementary material 3.3. (Continued).



Supplementary Material 3.4. Fossil information and calibration nodes, means, standard deviations (SD) or sigma values, and offsets (in real space).

<b>Fossil taxon</b>	<b>Locality</b>	<b>Taphonomy</b>	<b>Age (mya)</b>	<b>References</b>	<b>Calibration node</b>	<b>Mean</b>	<b>SD</b>	<b>Sigma</b>	<b>Offset</b>
<i>Apicrenus fossilis</i>	Dominican	Amber	15–20	Maldonado et al. (1993); Iturralde-Vinent & MacPhee (1996)	Harpactorinae: Apiomerini	20	0.446		15
<i>Triatoma dominicana</i>	Dominican	Amber	15–20	Poinar (2005); Iturralde-Vinent & MacPhee (1996)	Triatominae	20	0.092		15
				Hwang & Weirauch (2012)	Root	96.85		9.39	
				Hwang & Weirauch (2012)	Ectrichodiinae + Tribelocephalinae	67.47		7.36	

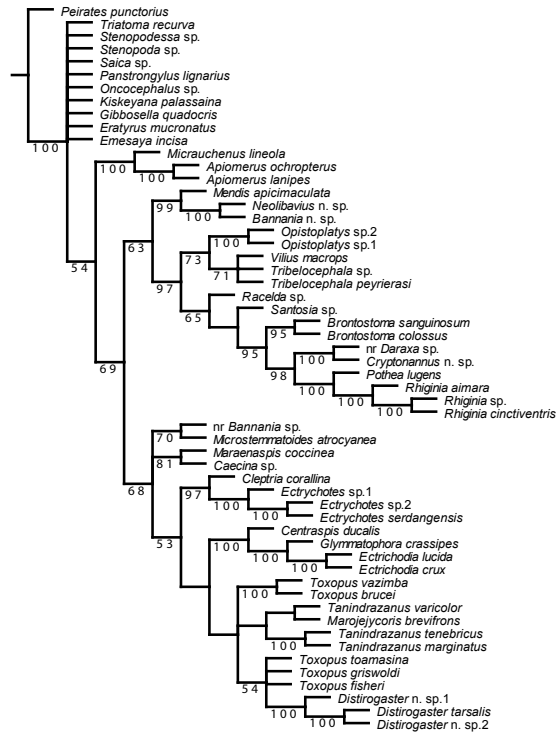
Supplementary material 3.5. DEC and DEC+j time-stratified dispersal rate matrix. N = Neotropical, A = Afrotropical, M = Madagascar, O = Oriental.

Time constraint (mya)		Rate matrix			
0–15		N	A	M	O
	N	1.00	0.25	0.10	0.25
	A	0.25	1.00	0.10	0.75
	M	0.10	0.50	1.00	0.25
	O	0.25	0.75	0.50	1.00
15–23		N	A	M	O
	N	1.00	0.25	0.10	0.25
	A	0.25	1.00	0.50	0.75
	M	0.10	0.10	1.00	0.25
	O	0.25	0.75	0.50	1.00
23–34		N	A	M	O
	N	1.00	0.25	0.10	0.25
	A	0.25	1.00	0.50	0.50
	M	0.10	0.10	1.00	0.25
	O	0.25	0.50	0.50	1.00
34–56		N	A	M	O
	N	1.00	0.25	0.10	0.25
	A	0.25	1.00	0.50	0.10
	M	0.10	0.10	1.00	0.25
	O	0.25	0.25	0.50	1.00
56–66		N	A	M	O
	N	1.00	0.25	0.10	0.25
	A	0.25	1.00	0.50	0.25
	M	0.10	0.10	1.00	0.25
	O	0.25	0.25	0.50	1.00

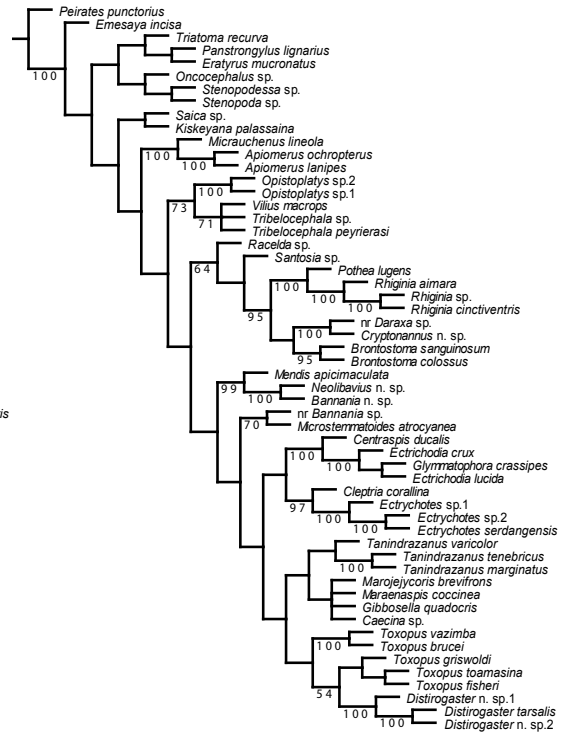
Supplementary material 3.6. Equal (EW) and implied weights (IW) molecular parsimony results for 56 taxa. A. Strict consensus tree from 147 parsimonious trees from an EW analysis (length = 6,232 steps; RI = 0.563; CI = 0.336). B. Strict consensus tree from 3 parsimonious trees from an IW analysis with  $k = 3$  (length = 470.59899 steps; RI = 0.560; CI = 0.335). C. & D. Single parsimonious trees from IW analyses when  $k = 6$  (C.) (length = 340.36981 steps; RI = 0.560; CI = 0.335) and  $k = 10$  (D.) (length = 252.28554 steps; RI = 0.561; CI = 0.336). For A.–D., bootstrap values  $\geq 50$  are reported below branches.



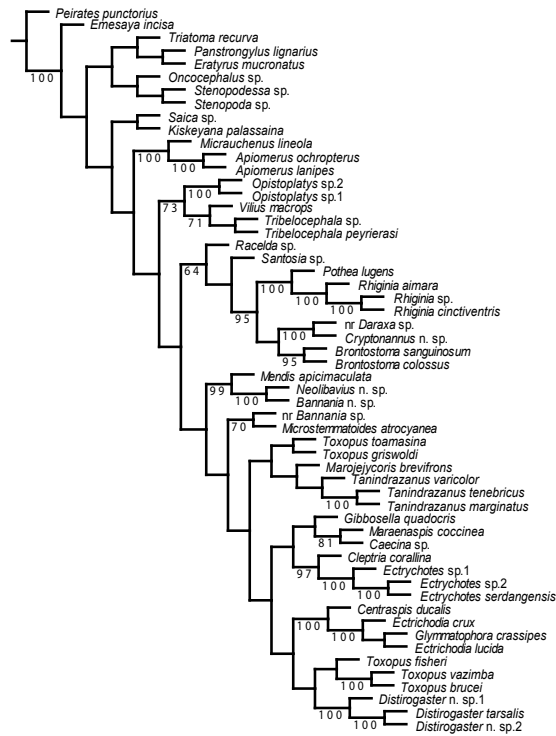
A. EW



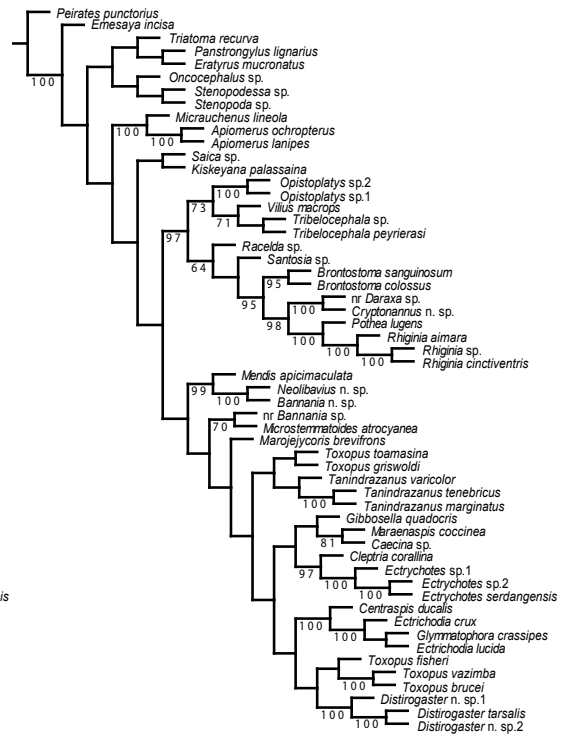
B. IW (k = 3)



C. IW (k = 6)



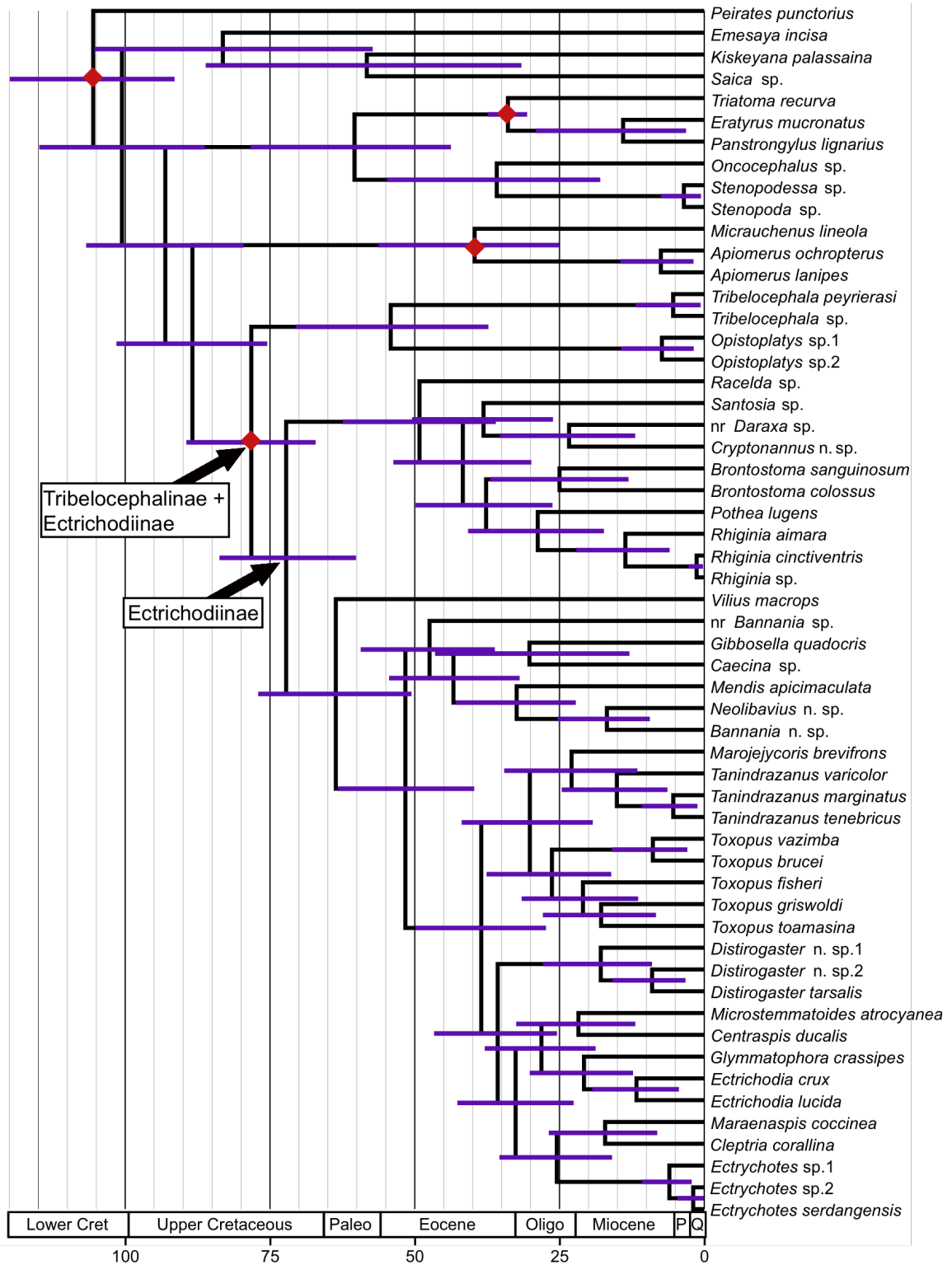
D. IW (k = 10)



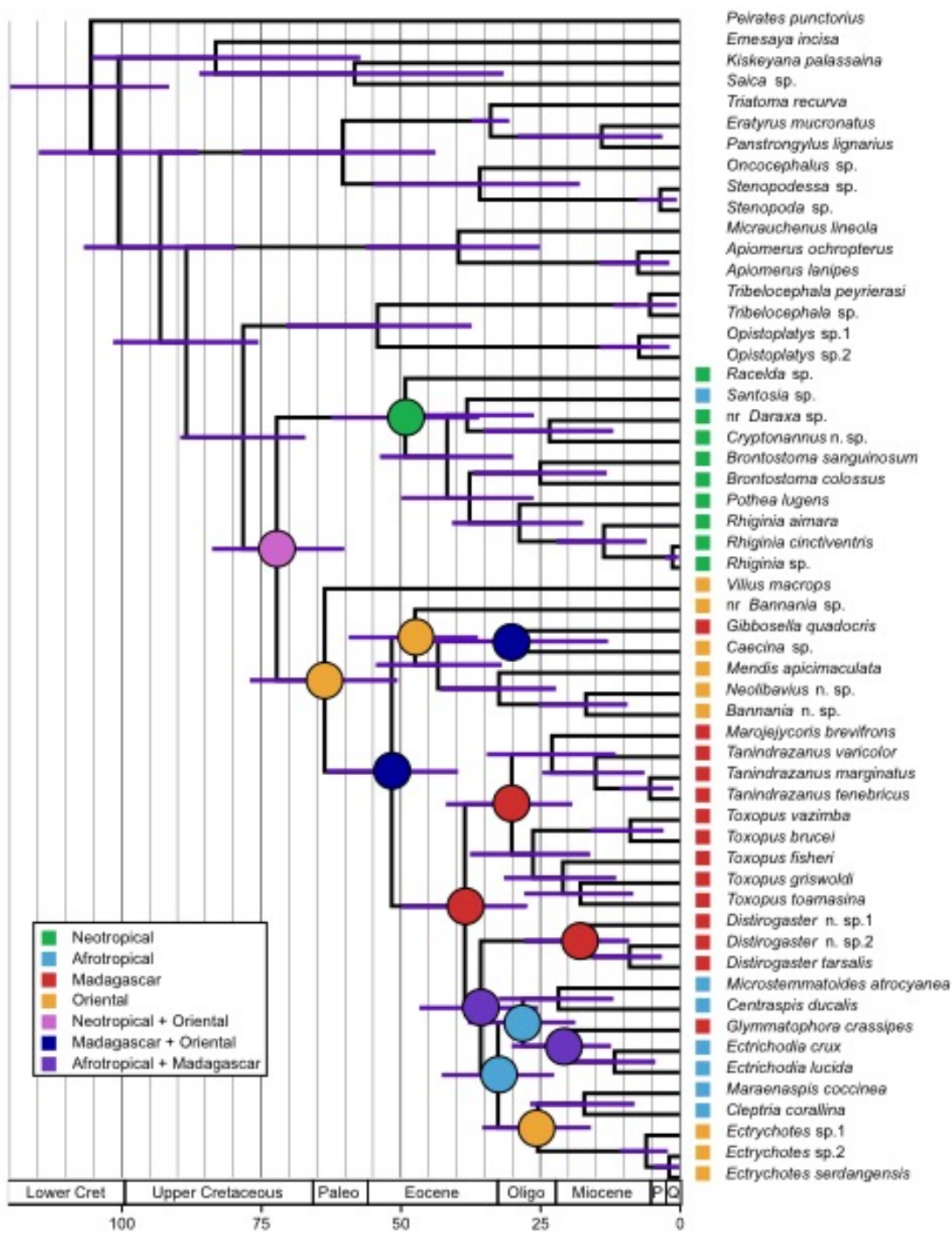
Supplementary material 3.7. Best tree based on ML analysis of 110 taxa, GTR+ $\Gamma$ +I model of sequence evolution for each G-INS-I aligned gene partitions (COI, 18S, 28S D2, 28S D3–D5), and Mkv model of morphological evolution (final Ln = -31,224.182191). Outgroups not shown. Bootstrap values  $\geq 50$  are reported below branches.



Supplementary material 3.8. Divergence time estimates based on a BEAST analysis using the ML best tree (Fig. 3.2) and the same molecular dataset (COI, 18S, 28S D2, 28S D3–D5). Each gene partition was assigned an unlinked site model (GTR+ $\Gamma$ +I) and were linked to the same tree model. A linked clock model was assigned to 28S D2 and 28S D3–D5 to reflect their single identity. Three unlinked relaxed lognormal clock models were assigned for 18S, 28S, and COI. Placement of two outgroup fossil and two secondary calibrations are indicated by red diamonds. Purple node bars indicated 95% HPD credibility intervals.



Supplementary material 3.9. Ancestral range reconstructions from the DIVA model for select nodes. Geographical distributions are color coded according to the legend.



Supplementary material 3.10. Discussion of morphological support for major lineages recovered in the combined morphological and molecular phylogenetic analysis.

Phylogenetic hypotheses generated from the combined cladistic analysis are discussed in order of the node labels shown in Fig. 3.3 and with emphasis on Madagascan taxa. Important morphological characters are discussed for selected nodes, with character numbers and states shown in Table 3.2. Results of unambiguous character optimizations on the parsimony consensus tree are shown in SM 3.3. Character support for Madagascan genera has informed taxonomic diagnoses in Forthman et al.'s (in press) monograph.

#### Monophyly of Ectrichodiinae

The subfamily Ectrichodiinae is recovered as monophyletic and sister to the Tribelocephalinae with high support (BS = 99). Based on unambiguous character optimization, four synapomorphies support Ectrichodiinae monophyly: scutellum with two paramedian apical processes (59–1), metathoracic gland evaporatorium (MGE) present (70–0), carinulate intersegmental suture between sternites II and III (114–1; Fig. 3.3N), and the larger size of dorsal abdominal gland (DAG) III relative to DAGs I and II (144–1). The scutellum is further modified within Ectrichodiinae: an additional medial apical process (59–2) evolved twice independently in *Neolibavivus*+*Bannania* (rudimentary in *Bannania*) and *Ectrychotes*, and four (59–3) evolved independently in *Vilius* and *Centraspis*. Twelve homoplasious characters further support the monophyly of millipede assassin bugs (SM 3.3).



#### Node 1: Neotropical Ectrichodiinae+Santosia+Vilius clade

A predominately Neotropical clade, including *Santosia* and *Vilius*, is recovered but not supported by any morphological synapomorphies. One terminal taxon, *Racelda* sp., is coded from an apterous female as a result of the inability to identify the species. Thus, the male representative of this taxon is unknown, resulting in some missing morphological data in the matrix. Dougherty (1995) investigated the phylogenetic relationships for 16 New World genera. Her analysis used a ground-plan approach with character polarity determined ad hoc through comparison with other reduviid taxa. Although that analysis excluded Old World Ectrichodiinae, Dougherty hypothesized a non-monophyletic New World based on shared features between Old World taxa and the New World genus *Ectrichodiella* Fracker and Bruner. Our results are congruent with Dougherty's hypothesis in that the New World fauna is not monophyletic in both molecular and combined analyses. However, our results should be tested further with a larger sample of Ectrichodiinae genera, including the genus *Ectrichodiella* from the Greater Antilles.

#### Node 2: Old World Ectrichodiinae clade

This clade of Afrotropical, Oriental, and Madagascan ectrichodiines receives weak support from six homoplastic characters, e.g., the carinulate intersegmental sutures between sternites III and VI (115–1, 116–1, 117–1) and the medially depressed abdominal sternites (120–1; Fig. 3.3N). In some taxa (e.g., *Tanindrazanus mahafaly* Forthman, Chłond, & Weirauch and *Maraenaspis bidens*), the intersegmental suture carination patterns are restricted to lateral areas or are entirely absent. In some species of *Gibbosella* and *Glymmatophora*, the sternites are convex or flat. This clade is further supported by the presence of 6-segmented antennae (38–1), which is further subdivided in many taxa (e.g., 7-segmented as in Fig. 3.3E). The medial region

of the endosoma is slightly sclerotized (136–1) in most species, but is membranous (136–0) in some examined species of *Glymmatophora*, *Maraenaspis*, and *Toxopus*.

Node 3: *Maraenaspis*+*Cleptria*+*Ectrychotes*+*Centraspis*+*Ectrichodia*+*Glymmatophora* clade

This clade is supported by the presence of small papillae on the fore and mid trochanters (74–1, 75–1) and femora (83–1, 89–1; Fig. 3.3L). Within this clade, the Madagascan species *Maraenaspis bidens* does not form a sister relationship with the Afrotropical *Maraenaspis coccinea*, potentially an artifact of missing data. Sexually static morphological features were inferred from a female of *M. coccinea* since males are currently unknown. Furthermore, the genus may not be monophyletic and, to date, has not undergone thorough taxonomic revision. In our analysis, *M. bidens* is recovered as the sister to *Ectrichodia*+*Glymmatophora*. This relationship is supported by one synapomorphy: the anterior pronotal lobe is as wide as the posterior lobe (50–1; Fig. 3.3F). The *Maraenaspis bidens*+*Ectrichodia*+*Glymmatophora* clade is further supported by four homoplastic characters, such as the ventrolaterally swollen gula produced beyond the ventral head margin (12–2; Fig. 3.3C) (slightly reduced in *Glymmatophora crassipes*), the eye less than half of head height in lateral view (28–0; Fig. 3.3C), the anterior pronotal lobe longer than posterior lobe (59–0; Fig. 3.3F), and the membranous medial endosomal region (136–0).

Madagascan species of *Glymmatophora* form a sister group relationship to the Afrotropical *Ectrichodia* species by several homoplastic features and the following synapomorphies: presence of a medial tubercle on the mid femur (88–1; Fig. 3.3L) and an anterior and posterior subapical tubercle on the hind femur (91–1, 92–1; Fig. 3.3L). The Madagascan *Glymmatophora* are recovered as closely related and are supported by one synapomorphic character: the presence of a medial tubercle on the hind femur (94–1; Fig. 3.3L). The medium body size (1–1), longer labial

segment II relative to segment III (41–1; e.g., Fig. 3.3D), and flat scutellar disc (63–1) provide additional support for this relationship.

Node 4: *Marojejycoris*+*Caecina*+*Gibbosella*+*Distirogaster*+*Toliarus*+*Tanindrazanus*+*Toxopus* clade

The majority of Madagascan species, including a species of *Caecina* from Thailand, forms a weakly supported clade in our analysis. Two homoplastic characters support this large clade: the depressed postclypeus (13–1; Fig. 3.3B) and a complete transverse suture between the meso- and metasterna (66–0; Fig. 3.3J). Within this clade, the postclypeus is not depressed (13–0; Fig. 3.3A) in *Caecina* sp. and some *Gibbosella* species. Species of *Toxopus*, as well as some *Gibbosella* and *Distirogaster* species, have an incomplete transverse suture (66–1) between the meso- and metasterna.

Node 5: *Marojejycoris*+*Caecina*+*Gibbosella* clade

The *Marojejycoris*+*Caecina*+*Gibbosella* clade is weakly supported by one homoplasious character, the small body (1–0). The Madagascan genus *Marojejycoris* is highly supported by five homoplastic characters, such as a complete pronotal transverse suture (57–1; Fig. 3.3H), the MGE visible in lateral view (71–1; Fig. 3.3K), the distal part of M not extending beyond the M+Cu junction on the fore wing (104–1; see Fig. 3.3M for 104–0), and the shape of the medial pygophore process (124–1, 125–2). The Madagascan genus *Gibbosella* is recovered as sister to *Caecina* by the distal part of M and Cu basally fused on the fore wing (103–1; Fig. 3.3M) and presence of a posterior tubercle on dorsal laterotergites III–VI (107–1, 108–1, 109–1, 110–1). *Gibbosella* forms a clade supported by an 8-segmented antenna (38–3) (6-segmented in *G. pallidalata* Forthman, Chlond, & Weirauch) and the transversely bicolor dorsal laterotergites

(105–1). Two synapomorphic characters support *Gibbosella*, with the exclusion of *G. planiscutum* Forthman, Chłond, & Weirauch sister to the remaining *Gibbosella*: the long (61–1) and dorsally directed (62–1) paramedian scutellar processes (Fig. 3.3I). The sublateral antennal insertion (24–0; Fig. 3.3C) and lack of carinulae on the intersegmental suture between abdominal sternites VI and VII (118–0) are also optimized as homoplastic along this branch. Some species in *Gibbosella* exhibit a dorsal antennal insertion (24–1; Fig. 3.3D).

Node 6: *Distirogaster*+*Toliarus*+*Tanindrazanus*+*Toxopus* node

Two homoplasious characters are optimized on the branch of this clade: a striated posterior pronotal lobe (51–0) and the basal plate extension as long as the basal plate (129–1). Some taxa within this clade have smooth (51–1) or punctate (51–2) posterior pronotal lobes and a basal plate extension that is shorter (131–0) or longer (131–2) than the basal plate. Within this clade, the Madagascan genus *Distirogaster* is recovered as the sister group to other members of the clade. The presence of denticle-like processes on the dorsal phallothecal sclerite (131–1) is optimized as a synapomorphy of *Distirogaster*. *Distirogaster* is further supported by 11 homoplasious characters: the metallic coloration (2–1), 8-segmented antenna (38–3), relatively small anterolateral pronotal tubercles (47–1; Fig. 3.3G), laterally carinate anterior pronotal lobe (48–1), posterior tubercles on dorsal laterotergites III–VI (107–1, 108–1, 109–1, 110–1), paramedian carinae on abdominal sternites (120–1; Fig. 3.3N), and shape of the medial pygophore process (123–0, 124–2). One species of *Distirogaster* has a reversal to dull coloration (2–0), and large anterolateral projections on the pronotum (47–2; Fig. 3.3F) have evolved twice independently in the genus.

Node 7: *Toliarus*+*Tanindrazanus*+*Toxopus* clade

The straight ventral margin of labial segment III (40–0; Fig. 3.3D) and the laterally visible MGE (71–1; Fig. 3.3K) are homoplastic characters optimized on the branch of the Madagascan *Toliarus*+*Tanindrazanus*+*Toxopus* clade. However, the ventral margin of labial segment III is convex (42–1; Fig. 3.3C) rather than straight in some *Toxopus* species. The genus *Toliarus* is sister to *Tanindrazanus*+*Toxopus* and possesses the following homoplastic features: a small body (1–0), red and black coloration (2–2), and a punctate posterior pronotal lobe (51–2).

Node 8: *Tanindrazanus*+*Toxopus* clade

The broad postclypeal depression (14–1; Fig. 3.3B), 7-segmented antennae (38–2; Fig. 3.3E), and smooth intersegmental suture between sternites VI and VII (118–0) are homoplastic characters supporting this clade. The postclypeal depression is narrowed (14–0) in *Tanindrazanus brunneus* Forthman, Chłond, & Weirauch, *Toxopus namoroka* Forthman, Chłond, & Weirauch, and *Toxopus steineri* Forthman, Chłond, & Weirauch. Approximately half of the examined specimens of *Tanindrazanus harinhali* Forthman, Chłond, & Weirauch have 6-segmented antennae rather than 7; such polymorphism is also observed in individual specimens between the right and left antenna. Although the genus *Tanindrazanus* is not supported by any synapomorphic characters, it is consistently recovered as a clade both molecular and combined analyses. *Toxopus* is supported by six homoplastic features: a head as long as it is wide (5–0), a postclypeal depression extending from the posterior clypeal margin to the interocular sulcus (16–1) as in Fig. 3.3B, the small anterolateral tubercles on the pronotum (47–1; Fig. 3.3G), an incomplete transverse suture between the meso- and metasterna (66–1), the transversely bicolor dorsal laterotergites (105–1), and a laterally expanded dorsal laterotergite II (106–1). In 13 *Toxopus* species, i.e., the clade including *Toxopus ampitavananima* Forthman, Chłond, & Weirauch and

*Toxopus fisheri* Forthman, Chłond, & Weirauch, the head is longer than wide (5–1). The postclypeal depression only extends to the middle of the synthlipsis (16–0) in *Toxopus farafangana* Forthman, Chłond, & Weirauch and the clade including *Toxopus antsiranana* Forthman, Chłond, & Weirauch and *Toxopus fisheri*. The anteromedial pronotal tubercle is absent (47–0) in *Toxopus ampitavananima*.

**Chapter 4: Millipede assassin bugs (Heteroptera: Reduviidae: Ectrichodiinae) show off: evolution of aposematic coloration and extreme sexual dimorphism**

**Abstract**

Aposematic coloration has been shown to be positively associated with body size in certain groups of organisms. Amongst the studies using insects as model systems (Lepidoptera; Hemiptera) to investigate this phenomenon, larval or adult gregariousness may have confounded body size. Millipede assassin bugs (Heteroptera: Reduviidae: Ectrichodiinae) display a diversity of color patterns across a range of body sizes, are typically solitary as adults, and are thus an excellent model for investigating a possible correlation. Some species also exhibit drastic sexual dimorphism with major changes in female morphology that has not been explored in a phylogenetic context. Here, a comprehensive phylogeny is constructed for 152 Ectrichodiinae and Tribelocephalinae taxa using morphological and molecular data. Based on the combined maximum likelihood phylogeny, ancestral states for coloration and sexual dimorphism were reconstructed, and phylogenetic logistic regression was employed to test for a coloration-body size correlation. The Ectrichodiinae+Tribelocephalinae ancestor is reconstructed as cryptically colored with limited sexual dimorphism. Multiple origins of aposematic and cryptic coloration and extreme sexual dimorphism are reconstructed. Prevalence of aposematic coloration is positively associated with male body length when regressions are performed on ultrametric trees but not on the phylogram. Thus, our results support the hypothesis that selection on Ectrichodiinae body size may influence evolutionary transitions between aposematic and cryptic coloration. Based on phylogenetic results, we synonymize Tribelocephalinae with Ectrichodiinae, describe three new tribes (Ectrichodiini, Tribelocodiini, and Abelocephalini) and two new subtribes (Opistoplatyina and Tribelocephalina), and revise Tribelocephalini.

## **Introduction**

Scientists have long sought to understand the evolution and diverse functions of the astounding diversity of color patterns observed across the animal kingdom (Protas & Patel, 2008). Although many functions have been proposed, it is evident that coloration can be involved in environmental (e.g., thermoregulation), intraspecific (e.g., sexual selection), and interspecific interactions (e.g., predator evasion) (Areekul & Quicke, 2006; Protas & Patel, 2008). Widespread cryptic coloration allows individuals to conceal themselves in the environmental background. In contrast, some structurally or chemically defended species have evolved conspicuous colors that function as a warning, or aposematic, signal to deter potential predators (Poulton, 1890; Edmunds, 1987) and to enhance predator learning and memorization (see Ruxton et al., 2004; Exnerová et al., 2006). Such color patterns may also occur in undefended species but still afford protection against predators that might perceive the species as aposematic (Nilsson & Forsman, 2003) due to, e.g., Batesian mimicry (Bates, 1862) or innate or learned predator aversion (e.g., Smith, 1975; Guilford, 1988). Traditionally, aposematic coloration has been characterized by a combination of black with red, orange, yellow, or white (Cott, 1940; Fabricant et al., 2014). Some studies suggest that metallic or iridescent patterns may also function in aposematic signaling alone or by enhancing traditional aposematic colors (e.g., Arrow, 1951; Schultz, 2001; Fabricant et al., 2014).

Studies on aposematic coloration are prevalent but often restricted to a microevolutionary scale (Härilin & Härilin, 2003; Stevens, 2015). Comparably fewer investigations have studied aposematic coloration in a phylogenetic context, e.g., in poison dart frogs (Anura: Dendrobatidae) (e.g., Hagman & Forsman, 2003; Rudh, 2013), mammals (Stankowich et al., 2011), and holometabolous insects (e.g., Vogler & Kelley, 1998; Sagegami-Oba et al., 2007; Bocakova et al., 2016). Some of these microevolutionary and phylogenetic studies have suggested a positive



relationship between the prevalence or conspicuousness of aposematic coloration and body size in poison dart frogs (Hagman & Forsman, 2003; Rudh, 2013), milkweed bugs (Insecta: Hemiptera: Lygaeidae) (Gamberale & Tullberg, 1996a, 1998), and butterflies and moths (Insecta: Lepidoptera) (Tullberg & Hunter, 1996). In insects, experimental studies by Gamberale & Tullberg (1996a, 1998) concluded that predator aversion increases with larger aposematically colored instars and larger aggregations of *Tropidothorax leucopterus* (Goeze), 1778 (Insecta: Hemiptera: Lygaeidae). Other experimental studies have also shown that nymphal aggregations of the aposematically colored *Spilostethus pandurus* (Scopoli), 1763, (Insecta: Hemiptera: Lygaeidae) result in higher predator aversion (Gamberale & Tullberg, 1996b). Only one phylogenetic comparative study based on independent contrasts concluded that larval gregariousness in Lepidoptera is correlated with aposematic coloration (Tullberg & Hunter, 1996). However, Nilsson & Forsman's (2003) comparative study of 578 moth species (Insecta: Lepidoptera) found no significant difference in larval body size between cryptic and aposematically colored species. Instead, they concluded that an association between coloration and gregarious lifestyles may have confounded their results since transitions to gregariousness were correlated with decreasing larval body length. A study on nudibranchs found a different trend: the prevalence of aposematic coloration decreases as body size increases and may be due to marine habitats (i.e., light environment and visual system of predators), diet, or habitat homogeneity (Cheney et al., 2014).

Insects are excellent models for studying aposematic coloration in an evolutionary context due to their morphological, behavioral, and ecological diversity. However, the two studies that have used phylogenetic comparative methods to explore the relationship between aposematic coloration and body size were restricted to Lepidoptera that exhibit larval gregariousness (Tullberg & Hunter, 1996; Nilsson & Forsman, 2003). These studies have employed phylogenetic

independent contrasts (Felsenstein, 1985) to analyze correlations between characters, a method that is unable to adequately model categorical trait evolution without treating it as continuous “dummy” variables (Ives & Garland, 2010). Phylogenetic comparative studies on aposematic coloration and body size in insects have failed to embrace more recently developed approaches that resolves this issue, e.g., phylogenetic logistic regression (PLR; Ives & Garland, 2010).

The predominantly circumtropical millipede assassin bugs (Heteroptera: Reduviidae: Ectrichodiinae) are likely the largest group of specialized millipede predators with 736 species in 121 genera (Maldonado, 1990; Dougherty, 1995; Carpintero & Maldonado, 1996; Forthman & Weirauch, 2012; Gil-Santana, 2015; Forthman et al., in press). Although adult millipede assassin bugs are typically solitary, communal predatory behaviors occur in some species; out of 32 reports of adult predatory behavior, eight have documented an adult engaging in communal predation with conspecific nymphs and two with conspecific adults (see Forthman & Weirauch, 2012; unpublished data). Aside from communal predatory strategies, aggregations have not been observed, although immatures may be in close proximity to an adult female (pers. obs.). Furthermore, these insects show a remarkable diversity of color patterns: cryptic coloration, traditional aposematic coloration, uniform metallic colors, or a combination of aposematic and metallic patterns. The conspicuous color patterns observed in many millipede assassin bugs may serve an aposematic function. A study on a *Ectrichodia* sp. (Peschke et al., 2002) and our personal observations with *Centraspis ducalis* Distant, 1902, confirm that these brightly colored species emit chemical defenses from the metathoracic glands. In addition, Dougherty (1995) suggested that some species may belong to a Müllerian mimicry complex with species in other genera. Furthermore, Forthman & Weirauch (2012), as well as additional unpublished data, show that at least 20 species of Ectrichodiinae prey on chemically defended millipedes and suggest future investigations into the ability of this predatory group to sequester secondary metabolites

from prey. However, no experimental study has been performed to confirm the hypothesis that conspicuous colors in millipede assassin bugs are an aposematic signal. Regardless, we have referred to these conspicuous patterns as aposematic coloration (i.e., red, yellow, orange, white, black, and/or metallic features).

Some of the smallest and largest species across Ectrichodiinae are either cryptically colored (e.g., *Ectrichodiella* Fracker & Bruner, 1924 [3–5 mm]; *Gibbosella* Chłond, 2010 [5–10 mm]; *Distirogaster* Horváth, 1914 [16–30 mm]; *Xenorhyncocoris* Miller, 1938 [32–37 mm]) or aposematic (e.g., *Antiopuloides* Miller, 1952 [10 mm]; *Schuhella* Dougherty, 1995 [6 mm]; *Centraspis* Schaum, 1862 [25–40 mm]; *Philodoxus* Horváth, 1914 [32–35 mm]). This preliminary observation suggests that coloration and body size may not be correlated in this clade, but this has not been tested in a phylogenetic comparative framework. Given the size of the group and diversity of color forms across a range of body sizes, millipede assassin bugs present an opportunity to explore, for the first time, the evolution of coloration and its association with body length in a non-holometabolous group of specialized predatory insects that are typically solitary as adults.

In addition to the diversity of color patterns, millipede assassin bugs exhibit sexual dimorphism (Fig. 4.1) that ranges from slight morphological differences between males and females in, e.g., body size, wing development, and eye and ocellar size, to extreme cases in which wing reduction is associated with drastic modifications to the female head, thoracic, and/or leg morphology. For species where data was available for both sexes from the literature and/or examined specimens (~200 spp.), females are always larger in size and have reduced antennal setation compared to males. Furthermore, approximately half of these species exhibit some wing reduction in the females when compared to males, which is variable within some species (i.e., pterygopolymorphism; e.g., some *Brontostoma* spp.) (Dougherty, 1995; pers. obs.).

Pterygopolymorphism may also be observed in both sexes in some Old World species (e.g., some *Glymmatophora* spp.), which can make associations between different morphs difficult. Other differences may be observed on the female's head, including reduced depressions and striations, a more pronounced gula, and a shorter scape, among other features. Thoracic morphology in females can include a shorter posterior pronotal lobe and/or a longer and wider anterior lobe, a reduction of sutures and depressions, or a weakly-developed scutellum. Females may further possess modified legs, e.g., stouter forefemora, a larger fossula spongiosa, and tubercles and papillae that may be absent in conspecific males. While any of these differences may occur in species that exhibit limited or extreme sexual dimorphism, a combination of the overall abundance and severity of morphological differences characterizes how extreme sexual dimorphism may appear. Approximately 80% of Ectrichodiinae species for which both sexes are known are, here, assigned to a limited condition, in which a few minor morphological differences are restricted to some structures or body regions (Fig. 4.1). The remainder of species are, here, considered to display extreme sexual dimorphism (e.g., *Racelda* spp., *Distirogaster* spp., *Rhysostethus glabellus* Hsiao, 1973) due to drastic morphological differences in almost every major body region (Fig. 4.1). Such extreme differences often pose significant difficulties in associating sexes when using morphology alone (Zhang & Weirauch, 2011; Forthman et al., in press) and males and females have sometimes been described as different species and even different genera (e.g., Rédei et al., 2012).

Extreme sexual dimorphism is also known to occur in a few other Reduviidae subfamilies: Harpactorinae (some *Zelus* spp.; Hart, 1972), Peiratinae (*Bekilya* genus group; Zhang & Weirauch, 2011), Emesinae (e.g., some *Bergemesa* spp., *Collartida oculata* Villiers, 1949; Wygodzinsky, 1966), and Stenopodainae (e.g., *Pseudobaebius* spp.; Villiers, 1948; Giacchi, 1987). With the possible exception of Emesinae, extreme sexual dimorphism in these subfamilies

does not appear to be as prevalent as in Ectrichodiinae, and its origins have not been investigated using ancestral state reconstruction methods (ASRs). Extreme sexual dimorphism in other groups of insects (e.g., Coleoptera: Elateriformia; Bocakova et al., 2007; Kundera & Bocak, 2011) has been shown to have evolved multiple times from a limited condition, with few or no reversals. Thus, extreme sexual dimorphism may result in convergent female morphology between unrelated taxa (Bocakova et al., 2007; Kundera & Bocak, 2011). As such, evolutionary transitions between limited and extreme sexual dimorphism should be explored in Ectrichodiinae to determine if the extreme condition evolved more than once.

Comprehensive phylogenetic hypotheses using molecular and morphological data are currently unavailable for Ectrichodiinae, impeding evolutionary investigations of coloration and sexual dimorphism. Few Ectrichodiinae have been incorporated in recent phylogenetic analyses of Reduviidae (Weirauch, 2008; Weirauch & Munro, 2009; Weirauch, 2010; Hwang & Weirauch, 2012; Zhang et al., 2016), with the exception of a larger analysis on Madagascan Ectrichodiinae (Forthman & Weirauch, in press). These studies support a clade formed by Ectrichodiinae and Tribelocephalinae, a small subfamily (~130 species, 16 genera; Maldonado, 1990; Weirauch, 2010) of mostly cryptically colored species that often display little sexual dimorphism. Although some of these studies support the monophyly of Ectrichodiinae, others found Ectrichodiinae to be paraphyletic with respect to Tribelocephalinae (Weirauch & Munro, 2009; Weirauch, 2010, Forthman & Weirauch, in press). Thus, relationships between Ectrichodiinae and Tribelocephalinae have remained unresolved. To investigate the evolution of coloration and sexual dimorphism, a resolved, well-supported phylogeny based on molecular and morphological data for a larger sample of Ectrichodiinae and Tribelocephalinae taxa is crucial.

Here, we reconstruct the most comprehensive phylogeny of Ectrichodiinae and Tribelocephalinae to date, using separate and combined morphological and molecular analyses to

test the monophyly of both subfamilies. The resulting phylogenetic framework is used to investigate the evolution of coloration and sexual dimorphism with ASR methods. We predict that 1) the last common ancestor of Ectrichodiinae and Tribelocephalinae was cryptic and exhibited limited sexual dimorphism, 2) aposematic coloration evolved from cryptic coloration multiple times as seen in other groups of animals (e.g., Santos et al., 2003; Bocakova et al., 2016), and 3) extreme sexual dimorphism has evolved multiple times with very few reversals to a limited condition. A phylogenetic comparative analysis is also conducted to test our prediction that there is no significant relationship between the prevalence of aposematic coloration and male body length. Based on our phylogenetic results, we revise the Ectrichodiinae+Tribelocephalinae higher-level classification to reflect morphologically diagnosable clades recovered in the phylogeny.

## **Material and methods**

### *Taxon sampling*

To investigate the evolution of coloration and sexual dimorphism, taxa were selected to include all major color patterns and sexually dimorphic conditions across differently-sized species of Ectrichodiinae and Tribelocephalinae. In total, 156 terminal taxa were examined and comprise 138 Ectrichodiinae, 14 Tribelocephalinae, and four outgroup species belonging to other Reduviidae subfamilies. The morphological character matrix is based on male specimens, where possible, to minimize the effect of sexually dimorphic character states on phylogenetic reconstruction. However, 28 taxa are represented by females because: 1) some represent mono- or ditypic female-based genera (e.g., *Borgmeierina* Wygodzinsky, 1949, *Antiopuloides*) that may be females of male-based species classified in other genera (e.g., *Daraxa* Stål, 1859, *Caloundranus* Miller, 1957), 2) inclusion of female-based genera (e.g., *Schuhella*) permits assignment of

undescribed males into existing genera for future taxonomic description, and 3) inclusion also permits assignment of undescribed females into existing genera (e.g., *Tribelocodia* Weirauch, 2010, *Neoscadra* Miller, 1941, *Katanga* Schouteden, 1903) for future taxonomic work. Of these 28 female-based terminal taxa, male morphology for 12 female-based terminals were coded based on taxonomic descriptions, digital images, and/or examination of specimens at the British Natural History Museum, London, United Kingdom and/or the American Museum of Natural History, New York, USA (see table in Supplementary Material [SM] 4.1). A total of 16 terminals lack information on male morphology because the males remain unknown or the species could not be determined. Despite the possibility for extreme sexual dimorphism in these 16 terminals, we follow Forthman & Weirauch (in press) and infer male morphological characters that are likely static among males and females (e.g., antennal segmentation, relative lengths of labial segments, scutellar apical processes, etc.) based on known patterns of sexual dimorphism among congeners across Ectrichodiinae and Tribelocephalinae.

Specimens examined for this study are deposited in the following institutions: **AM**, Australian Museum, Sydney, Australia; **AMNH**, American Museum of Natural History, New York, USA; **ANIC**, Australian National Insect Collection, Canberra, Australia; **BMNH**, Natural History Museum, London, United Kingdom; **BPBM**, Bernice P. Bishop Museum, Hawaii, USA; **CAS**, California Academy of Sciences, California, USA; **CNC**, Canadian National Collection of Insects, Ontario, Canada; **DEI**, Deutsches Entomologisches Institute, Müncheberg, Germany; **FCAP**, Universidade Federal do Pará, Pará, Brazil; **MNHN**, Muséum National d'Histoire Naturelle, Paris, France; **MRAC**, Musée Royal de l'Afrique Centrale, Tervuren, Belgium; **MTEC**, Montana State University, Montana, USA; **QM**, Queensland Museum, Brisbane, Australia; **SAMC**, South Africa Museum, Cape Town, South Africa; **SU**, Department of Zoology, University of Silesia, Poland; **UCR**, University of California, Riverside Entomological

Research Museum, California, USA; USNM, National Museum of Natural History, Washington D.C., USA.

#### *Morphological methods and data*

External morphology was examined using Nikon SMZ1000 and SMZ1500 dissecting microscopes. Mandibular and maxillary stylets and genitalia were dissected from a subset of taxa and examined under a Zeiss Axioscope compound light microscope and Nikon SMZ1000 and SMZ 1500 microscopes, respectively. For phylogenetic analyses, 168 morphological characters were coded in the Descriptive Language for Taxonomy program (Dallwitz, 1980; Dallwitz et al., 1999) using morphological matrices modified from Weirauch (2008, 2010) and Forthman & Weirauch (in press). Additionally, 30 female characters were coded for species in which specimens of both sexes were available or described in taxonomic publications to investigate patterns of sexually dimorphic characters within terminal taxa. One sexual dimorphism and two coloration characters were also coded for ASR and male body lengths for phylogenetic comparative analyses. All morphological characters and character states used in this study are given in Table 4.1. The morphological matrix is provided in SM 4.2. Morphological terminology follows Dougherty (1995), Weirauch (2008), Forero & Weirauch (2012), Hill (2014), and Forthman & Weirauch (in press).

#### *Molecular methods and data*

Six gene regions (16S rDNA, 18S rDNA, 28S D2 rDNA, 28S D3–D5, rDNA, COI, Wingless [Wg]) were sampled for 59 taxa (47 Ectrichodiinae, nine Tribelocephalinae, three outgroups) (Table 4.2). For some taxa, sequence data were acquired from GenBank, which were contributed by Weirauch & Munro (2009), Hwang & Weirauch (2012), and Forthman &



Weirauch (in press). Molecular vouchers were associated with an ethanol specimen collection number (RCW prefix), as well as a unique specimen identifier (USI) label to link it with specimen information in the Planetary Biodiversity Inventory Arthropod Easy Capture database (<http://www.research.amnh.org/pbi/locality/index.php>) and Heteroptera Species Pages (<http://research.amnh.org/pbi/heteropteraspeciespage/>). All USI labels include an AMNH\_PBI or UCR\_ENT prefix followed by an 8-digit number.

DNA extraction was performed by removing a hind leg and following QIAGEN DNeasy Blood and Tissue Kit protocols. Extracted legs were card mounted and associated with their respective specimens. For new sequences, PCR was performed using either GE Healthcare Life Sciences PuReTaq-Ready-To-Go-PCR-Beads or EmeraldAmp GT PCR Master Mix and Fisher Scientific or BioRad T100 Thermal Cyclers with the protocols listed in SM 4.3. Primer sequences for 16S (16Sa, 16Sb), 18S (18SF, 18SR), and 28S D3–D5 (D3Fa, D5Ra) were taken from Weirauch & Munro (2009); 28S D2 (D2Fa, D2Ra) from Forero et al. (2013); COI (C1-J-2183F) from Simon et al. (1994) and (C1-N-2609R) from Damgaard & Sperling (2000); and Wg (Wg 1A) from Cryan et al. (2001) and (Wg DelR1) from Urban & Cryan (2007). Amplification was assessed using gel electrophoresis with SyberSafe gel stain and a UV illuminator. PCR products were cleaned using Bioline SureClean and sequenced at Macrogen USA (Sanger sequencing) or on an Applied Biosystems 3730x1 DNA Sequencer at UCR's Institute for Integrative Genome Biology. Sequences were assembled and edited in Sequencher v4.8 and are available on GenBank (Table 4.2).

Each gene region was independently aligned in MAFFT (Katoh & Standley, 2013) using the G-INS-i algorithm. Gene regions were concatenated in SequenceMatrix v1.7.8. (Vaidya et al., 2011) and exported with external gaps coded as question marks (matrix provided in SM 4.4). The best partition scheme and models of sequence evolution were determined using the greedy

algorithm and Bayesian information criterion for model selection in PartitionFinder v1.1.1 (Lanfear et al., 2012). Each ribosomal gene was treated as a separate subset in the data blocks, while COI and Wg were subdivided into separate codon positions. The resulting partition scheme and models are: *Partition 1*: 16S, 28S D2, and 28S D3–D5 (GTR+ $\Gamma$ +I); *Partition 2*: 18S, COI codon positions 1 and 2, Wg (GTR+ $\Gamma$ +I); and *Partition 3*: COI codon position 3 (HKY+ $\Gamma$ +I).

### *Phylogenetic analyses*

Morphological and molecular datasets were concatenated using Mesquite v3.04 (Maddison & Maddison, 2015). Separate and combined morphological and molecular phylogenetic analyses were performed using parsimony (TNT v1.1; Goloboff et al., 2008), maximum likelihood (ML) (GARLI v2.01; Zwickl, 2006), and Bayesian inference (MrBayes v3.2.6; Ronquist et al., 2012) optimality criteria. For the morphological and combined phylogenetic analyses, the first 168 morphological characters in Table 4.1 were used. Unless otherwise stated, parameter settings for a given program were used for all three datasets.

All datasets were subjected to equal weights (EW) parsimony analysis using New Technology search, but the combined dataset was also analyzed using implied weights (IW) due to poor phylogenetic resolution from the EW analysis. Internal gaps in the molecular data were treated as fifth state and external gaps as missing data. Prior to analysis, 2,661 uninformative molecular characters were inactivated. Although there is no clear criterion for choosing a particular  $k$ -value and the optimal  $k$ -value is probably matrix dependent, we used three concavity constant ( $k$ ) values for IW analysis (3.000000 [default], 4.266835, and 18.980060) with the latter two based on calculations for average character fit at 58% and 86%. We selected these average character fit values because they encompass a range of  $k$ -values shown to produce results with the highest similarity to the preferred tree based on various measures of performance (see Reemer &

Ståhls, 2013). Default settings for sectorial search, drift, and tree fusing were used, the initial driven search level was set at 100 and checked every 3 hits, the initial addition sequences set to 14, find minimum length set to 100 times, and the random seed set to 4,325. Standard bootstrap resampling with absolute frequencies was conducted with 500 replicates and New Technology search (initial driven search = 38, check level every 3 hits, initial addition sequences = 7, find minimum length 10 times).

The ML analyses were conducted using the best partition scheme and models of molecular evolution determined by PartitionFinder and the Mkv model of morphological evolution (Lewis, 2001). A total of 50 search replicates were performed using random starting trees for each dataset. Non-parametric BS analyses were performed for 500 iterations with the termination condition reduced from 20,000 to 10,000 as recommended in the GARLI manual (Zwickl, 2008).

Bayesian analyses used the same partition scheme and models of molecular and morphological evolution as in our ML analyses. Two simultaneous runs of 20 million generations were carried out with four Markov Chain Monte Carlo (MCMC) chains each. Parameters and trees were sampled every 20,000 generations with the first 25% discarded as burn-in. Stationarity was assessed by the average standard deviation of split frequencies (<0.01 for molecular and morphology only datasets; <0.03 for combined dataset) and the potential scale reduction factor (~1.0 for all datasets). Each chain was also assessed for convergence in Tracer v1.6 (Rambaut et al., 2014) and possessed combined ESS values >200 in for each parameter after burn-in. The majority-rule consensus tree was generated from the sample of post-burn-in trees.

#### *Ancestral state reconstructions: aposematic coloration and extreme sexual dimorphism*

For ASR analyses, the combined ML phylogram was used for the following reasons: 1) a phylogeny with branch lengths is needed for the statistical approaches used, 2) is more resolved

than the Bayesian majority rule consensus tree, and 3) all three phylogenetic approaches recovered similar higher-level relationships. Traditionally, ultrametric trees have been preferred for ASRs as phenotypic change is expected to be related to the amount of time elapsed rather than molecular rates of change along a branch (Bromham et al., 2002; Litsios & Salamin, 2012; Cusimano & Renner, 2014). However, some studies have shown correlations between molecular branch lengths and phenotypic evolution (e.g., Omland, 1997; Smith & Donoghue, 2008); if this is the case, then inferring ASRs on time-calibrated phylogenies could produce inaccurate results as it may not appropriately represent the evolution of the species traits (Litsios & Salamin, 2012). Recently, Litsios & Salamin (2012) and Cusimano & Renner (2014) have studied the effects of time-calibrated versus molecular branch lengths on ASRs of continuous and discrete characters, respectively. Based on these studies, there is no consensus regarding whether ultrametric trees or phylograms produce greater accuracy in ASRs.

Given the potential for branch-length effects on ASRs, we estimated ancestral states on our combined ML phylogram and ultrametric trees. We followed a similar approach to the one implemented by Litsios & Salamin (2012) to transform the phylogram into three ultrametric trees using penalized likelihood in r8s v1.70 (Sanderson, 2002, 2003) under the Powell algorithm and different penalties: parametric estimation (penalty = 0), penalty on rate change (penalty = 10), and global molecular clock (penalty = 10,000). Ultrametric trees had a fixed length of 13.53 to match the length of the phylogram. Outgroup taxa were subsequently pruned from all trees in Mesquite. Although the ML phylogram is completely resolved, the r8s program collapsed eight of the very small, non-zero branch lengths in the generated ultrametric trees. Despite this, we used these trees as generated for ASR since 1) most were near the terminals, 2) all but one were within clades in which terminals had the same state, and 3) we did not expect major differences with regards to the reconstructions of these nodes on the phylogram.

Coloration was treated as categorical data based on the rankings made by five observers independent of the project: cryptic, traditional aposematic, uniform metallic, and aposematic and metallic. Each observer was presented with the same insect specimens, habitus images, and when available, images of individuals in natural habitat. In four cases where a majority opinion was not achieved, ties were broken by the senior author's assignments made prior to surveying the observers. In our first ASR dataset, coloration was coded as (0) cryptic or (1) aposematic, uniform metallic, or aposematic and metallic (character 198). Since some studies suggest uniform metallic appearances may serve a cryptic function (e.g., Parker, 1998; Thomas et al., 2007), we alternatively coded coloration as (0) cryptic or uniform metallic or (1) aposematic or aposematic and metallic (character 199). Sexual dimorphism (character 200) was also investigated, and was coded as either (0) limited or (1) extreme based on the severity and number of differences ( $\geq 6$  for extreme condition) observed between sexes. Some taxa, e.g., *Glymmatophora aeniceps* Horváth, 1914, and *G. dejoncki* Schouteden, 1919, exhibit different morphs within males and/or females; when paired with similar morphs, this species will appear to have little sexual dimorphism, but will appear extremely sexually dimorphic when paired with different morphs. As such, we code these terminals as polymorphic for both conditions.

Mesquite was used for parsimony ASR by tracing characters on the combined ML phylogram, while BayesTraits v2 (Pagel et al., 2004) MultiState was used with ML and MCMC approaches on the phylogram and ultrametric trees. For the ML approach, 1,000 iterations were performed with rate parameters free or equal, followed by likelihood ratio tests (LTR) to compare the two models; we report ASR results based on restricted rates ( $p > 0.09$ ,  $df = 1$ ). Under the Bayesian framework, a hyperprior on an exponential distribution drawn from a uniform distribution (interval between 0 and 10) was used. Twenty million generations were conducted with sampling every 2,000 generations and the first 25% of generations discarded as burn-in

under a free rates and equal rates model. Acceptance rates were between 0.2–0.4. Bayes Factors (BF) were calculated to compare results under the free and equal rates models; we report ASR results based on restricted rates (BF < 2). Tracer v1.6 was used to obtain the mean state value for each node, with ESS values >200 for all nodes reconstructed.

*Phylogenetic comparative approaches: aposematic coloration and male body length*

To examine the relationship between a binary dependent trait (i.e., coloration) and a continuous independent trait (i.e., body length) in a phylogenetic context, PLR with Firth correction was implemented using PLogReg.m (Ives & Garland, 2010) in MATLAB R2015b. This approach uses a two-state Markov process to model the evolution of the dependent trait along a phylogeny and estimate the strength of phylogenetic signal. Subsequently, the independent trait influences the dependent trait, with the evolutionary rate no longer dependent on phylogenetic signal but on the independent trait's regression coefficient (Ives & Garland, 2010).

The combined ML phylogram and ultrametric trees discussed in the previous section were pruned to exclude 23 terminals lacking male body length data and converted to phylogenetic variance-covariance matrices using the PDAP package in Mesquite (Midford et al., 2009). The two coloration datasets for ASR analyses were used to test if increases in male body length were associated with increased prevalence of aposematic coloration. When possible, average male body lengths were measured from the clypeal apex to the posterior margin of the abdomen for up to five specimens per species following Forthman & Weirauch (in press). When male specimens were not available, male body length was determined from taxonomic descriptions; if a range of lengths was provided, the average of the minimum and maximum lengths was used. Prior to analysis, log-transformed male body length was standardized to have a mean equal to zero and standard deviation equal to one following the recommendation of Ives & Garland (2010). A BS

procedure was used for 2,500 replicates to generate confidence intervals and test for statistical significance of the regressions. In all but one case, some replicates failed to converge. Removal of non-converged replicates resulted in slightly different parameter and confidence interval values, which are reported with estimates based on converged and non-converged replicates together.

## Results

### *Ectrichodiinae+Tribelocephalinae phylogenetics*

Results of the separate molecular and morphological analyses are given in SM 4.5–4.7. We report the ML best tree for our combined dataset in Fig. 4.2, with parsimony and Bayesian results in SM 4.8–4.10. Tribelocephalinae monophyly is not supported in all of our phylogenetic analyses, with the phylogenetic position of *Xenocaucus* China & Usinger, 1949, among the recovered Tribelocephalinae lineages being uncertain. Ectrichodiinae monophyly is also not supported, except in the molecular ML and Bayesian analyses (BS <50 and 92, SM 4.5B and C, respectively).

Molecular analyses (SM 4.5) recovered at least three Tribelocephalinae lineages with low to high support: *Tribelocephala+Opistoplatys*, *Afrodecius+Abelocephala*, and *Tribelocodia*. In our molecular parsimony analysis (SM 4.5A), *Tribelocephala+Opistoplatys* is the sister group to an Old World+New World clade (“Green clade”) of Ectrichodiinae with moderate support, while molecular ML and Bayesian analyses (SM 4.5B and C, respectively) recover *Tribelocephala+Opistoplatys* as sister to Ectrichodiinae with weak to high support. In the latter analyses, two major Ectrichodiinae clades are recovered with moderate to high support: the Green clade and an Old World clade (“Orange clade”).

Our morphological parsimony result (SM 4.6) is congruent with our molecular results with respect to the Tribelocephalinae lineages recovered: *Tribelocephala+Centrogastocoris+*

*Opistoplatys*, *Megapocaucus*+*Apocaucus*+*Afrodecius* +*Abelocephala*, and *Tribelocodia*.

However, ML and Bayesian analyses (SM 4.7A and B, respectively) support a paraphyletic Tribelocephalinae with respect to the Ectrichodiinae genus *Ectrichodiella*. *Ectrichodiella* is recovered as the sister group to *Tribelocodia* in all morphological analyses with moderate to high support. The Green and Orange clades are not supported in our morphological analyses, with the exception of a monophyletic Green clade that is weakly supported in our ML phylogeny; our Bayesian analysis recovered a large polytomy near the base of the remaining Ectrichodiinae.

Higher-level relationships between our combined EW and IW ( $k = 18.980060$ ) parsimony (SM 4.8A and 4.9B, respectively), ML (Fig. 4.2), and Bayesian (SM 4.10) phylogenetic results are largely congruent. The same Tribelocephalinae lineages from the morphological parsimony and molecular results are recovered, with *Ectrichodiella* as the sister group to *Tribelocodia*. The *Tribelocephala*+*Centrogastocoris*+*Opistoplatys* clade is weakly supported as the sister group to the remaining Ectrichodiinae, with the exception of our Bayesian phylogeny (unresolved polytomy at the root). In our other IW parsimony analyses (SM 4.8B, 4.9A), *Tribelocodia* is not monophyletic; *Tribelocodia* sp. is sister to *Xenocaucus*, while *T. ashei* Weirauch, 2010, is sister to *Ectrichodiella*. In the IW ( $k = 3.000000$ ) parsimony tree (SM 4.8B), *Tribelocephala*+*Centrogastocoris*+*Opistoplatys* is sister to *Megapocaucus*+*Apocaucus*+*Abelocephala* with poor support, while *Afrodecius* is weakly supported as the sister group to Ectrichodiinae+*Tribelocodia ashei*. In contrast, *Afrodecius*+*Abelocephala* is poorly supported as the sister group to *Tribelocephala*+*Centrogastocoris*+*Opistoplatys* in the IW ( $k = 4.266835$ ) analysis, which together are sister to *Megapocaucus*+*Apocaucus*. This larger Tribelocephalinae clade is weakly supported as the sister group to Ectrichodiinae+*Tribelocodia ashei*. With the exclusion of *Ectrichodiella*, the remaining Ectrichodiinae form a monophyletic group with weak to high support in all combined phylogenies. The Orange clade was recovered in all phylogenetic



analyses with low to high support, while the Green clade was recovered in all but the IW phylogenies (paraphyletic with respect to the Orange clade).

*Evolution of aposematic coloration: ASR and phylogenetic comparative analyses*

We report the result of our ML ASR based on the phylogram when treating cryptic coloration as a separate character from all other color patterns (color character 198; Fig. 4.3). The Ectrichodiinae+Tribelocephalinae ancestor is reconstructed as cryptically colored, with aposematic coloration evolving early in the phylogeny (clade including *Tribelocephala* sp. and *Glymmatophora aeniceps*). There are at least 15 reversals to cryptic coloration, as well as seven additional gains of aposematic coloration (eight total independent origins). In general, parsimony reconstructions (SM 4.11) are highly congruent with the highest probability reconstructions on the phylogram near the tips but are more ambiguous at deeper nodes (at least five gains of aposematic coloration, at least eight reversals to cryptic coloration). Our ML reconstructions on ultrametric trees (SM 4.12) are similar to those based on our phylogram with several exceptions. First, while there is no change with respect to the highest probable state for many nodes, some show significantly more or less probability in their reconstructions than reported in our phylogram (clades indicated by pair of taxa contained within): *Katanga–Cimbus*, *Synavecoris–Cimbus*, *Microstemmatoides–Cimbus*, *Tanindrazanus varicolor–Tanindrazanus tenebricus*, *Tanindrazanus–Marojejycoris*, *Distirogaster–Glymmatophora*, *Cryptonannus–Schottus*, and *Afrodecius–Glymmatophora*. At three nodes, the highest probable state is opposite of that in ML phylogram reconstruction: *Toliarus–Marojejycoris*, *Adrania gracilis–Adrania cylindrica*, and *Scadra–Neoscadra*. Five additional nodes have ambiguous reconstructions in our ultrametric trees compared to the phylogram: *Distirogaster* n. sp.–*Distirogaster tarsalis*, nr. *Neolibavius–Bannania*, *Pseudopothea–Jorgcoris*, *Daraxa ambrosettii–Daraxa* sp., and *Neoscadra ornata–*

*Neoscadra annulicornis*. The MCMC reconstructions on the phylogram and ultrametric trees are highly similar to ML reconstructions (SM 4.12). Results of our PLRs on ultrametric trees indicate that prevalence of aposematic coloration is associated with larger male body length ( $p < 0.05$ ; Table 4.3). However, while the regression coefficient is significantly different from zero for PLR on the phylogram ( $p = 0.0005$ ), the bootstrap mean is not ( $p = 0.4456$ ).

Results of ASRs when treating cryptic coloration and uniform metallic coloration as one character, with all other color patterns as a second state (color character 199), are given in SM 4.13–4.15. All analyses reconstruct a cryptically colored ancestor and support the early evolution of aposematic coloration. Parsimony reconstructions (SM 4.13) are also congruent with ML reconstructions near the tips of the phylogram (SM 4.14) but are more ambiguous at deeper nodes; at least five gains of aposematic coloration and six reversals to cryptic coloration are reconstructed. Based on ML reconstructions on the phylogram, there are at least seven independent origins of aposematic coloration and 18 reversals to cryptic coloration. ML reconstructions on ultrametric trees (SM 4.15) are largely similar to those of the phylogram. We observed similar patterns for some, but not all, of the nodes in which differences are observed between the phylogram and ultrametric trees in our ASR analysis on color character 198. Additionally, one node is ambiguous (*Haematorrhophus linnaei*–*Haematorrhophus pedestris*), and three have the same highest probability state but show significant changes in probability (*Centraspis*–*Philodoxus*, *Centraspis*–*Glymmatophora*, *Microstemmatoides*–*Glymmatophora*). The MCMC reconstructions are similar to ML reconstructions (SM 4.15). The PLRs on the phylogram and ultrametric trees for this color character resulted in the same trends observed in with the other color character (SM 4.16).

### *Evolution of extreme sexual dimorphism*

Despite the amount of missing data, ML ASR reconstructs limited sexual dimorphism as the ancestral state of Ectrichodiinae+ Tribelocephalinae based on the phylogram (Fig. 4.4). At least seven independent transitions of extreme sexual dimorphism from a limited condition is supported. Only one reversal to limited sexual dimorphism is reconstructed at the node containing *Xenocaucus* and *Ectrichodiella*. These results are supported by parsimony-based ASR (with exception of no reversals; SM 4.17), ML reconstructions on ultrametric trees (SM 4.18), and MCMC reconstructions (SM 4.18).

Plotting sexually dimorphic characters at the terminals reveals that taxa exhibiting only limited sexual dimorphism are generally characterized by one to three minor changes. These morphological differences often include a reduction in the female's eye, wing development (brachypterous, micropterous, or apterous), and associated changes in pronotal lobe lengths. Other taxa may exhibit minor differences in other pronotal features, as well as leg and abdominal features. However, in all cases, the presence or absence of the ocellar lens, postclypeal surface, presence or absence of the antennal shield, anterior pronotal width, scapus length, and presence or absence of paramedian abdominal sternal carinae do not differ between sexes. One species, *Haematorrhophus linnaei* (Stål), 1859, was coded as having a limited condition despite the fact that five morphological differences were observed. Females of this species can be easily associated to males despite aptery and an enlarged anterior pronotal lobe. Color patterns are not reduced between sexes in *Haematorrhophus* species, and head and abdominal structures in *H. linnaei* are similar in males and females. Males of this species also possess a relatively large anterior pronotal lobe that is nearly subequal to the posterior lobe, and the pronotal depression and sutures are retained in the females.

Taxa exhibiting only extreme sexual dimorphism tend to have at least six drastic differences between sexes. In these species, the extreme condition is often characterized by apterous females, with drastic changes in pronotal dimensions, or more often, the dimensions of the pronotal lobes. Eyes are generally reduced or completely lacking (e.g., *Xenocaucus*). If ocelli are present in males, these are drastically reduced or absent in the females. In a number of taxa, males have a depressed postclypeus and well-developed paramedian scutellar processes (e.g., *Distirogaster*), but females exhibit a flat postclypeus and weakly-developed scutellar processes. Females may also have increased leg armature (tubercles and/or papillae) and reduced abdominal features.

Two of the four *Glymmatophora* species we included have data on dimorphic males and/or dimorphic females. Based on a survey of the literature for *Glymmatophora* species exhibiting dimorphism within a sex, macropterous morphs have ocelli, small eyes, a flat postclypeus, an antennal shield, a scapus surpassing the apex of the head, an anterior pronotal lobe that is slightly shorter than or subequal to the posterior lobe, a wider posterior pronotal lobe relative to the anterior lobe, thoracic depressions and sutures, well-developed scutellar processes, no small tubercles on the legs but may have papillae, and no paramedian carinae on the abdominal sternites. Micropterous morphs have no or small ocelli, small eyes, a flat postclypeus, an antennal shield, a scapus surpassing the apex of the head, an anterior pronotal lobe that is longer than and as wide as the posterior lobe, reduced thoracic depressions and sutures, weakly-developed scutellar processes, no small tubercles on the legs but often with papillae, and no paramedian carinae on the abdominal sternites. Despite these morphological patterns, we did not code for all of these structures for *G. dejoncki* and *G. aeniceps*, as one representative for each species was available and the taxonomic descriptions sparse in detail (particularly for *G. aeniceps*). However, we still coded both taxa as polymorphic for limited and extreme conditions based on known patterns observed in *Glymmatophora*: males and females of the same morph are very similar

patterns observed in *Glymmatophora*: males and females of the same morph are very similar morphologically, but drastic differences between morphs can make associations between sexes of a species difficult.

## **Discussion**

### *Evolution of aposematic coloration and its association with body length*

Based on our results, the ancestor of Ectrichodiinae+Tribelocephalinae exhibited cryptic coloration, with aposematic coloration evolving in the ancestor of *Tribelocephala*+*Centrogastocoris*+*Opistoplatys*+Ectrichodiinae (excluding *Ectrichodiella*). Our results also support multiple subsequent origins of aposematic coloration from cryptically colored ancestors within this clade, which is an evolutionary pattern observed in other insect groups (e.g., Lepidoptera, Tullberg & Hunter, 1996; Cleroidea, Bocakova et al. 2016) and vertebrates (e.g., Dendrobatidae, Santos et al., 2003; Vences et al., 2003). Our PLR analyses using ultrametric trees for both coloration datasets support the hypothesis that increasing prevalence of aposematic coloration is associated with increasing body size, which is congruent with other phylogenetic comparative studies in poison dart frogs (Hagman & Forsman, 2003; Rudh, 2013). Only one phylogenetic comparative study concluded that larval gregariousness in Lepidoptera is correlated with aposematic coloration (Tullberg & Hunter, 1996). Thus, these studies have suggested an interplay between body size or size of the signal (i.e., aggregation size) and aposematic coloration. Based on these and other experimental studies, it has been hypothesized that species that recently acquired aposematic coloration are under selection for a larger body size (Forsman & Merilaita, 1999; Hagman & Forsman, 2003). Alternatively, a large body size may impair the effectiveness of cryptic coloration, and, thus, aposematic coloration may evolve more easily in

already large species (Hagman & Forsman, 2003). While we did not test for either hypothesis, our positive coloration-body size relationship is congruent with both of them.

Although PLR results based on our phylogram estimated regression coefficients significantly different from zero, parametric bootstrapping resulted in non-significance. For this approach, the parametric bootstrap procedure is currently the best approach to assess the quality of the parameter estimates (Ives & Garland, 2010, 2014), and studies employing this method base their conclusions from the significance of the bootstrap mean. Thus, in our case, there is no association between aposematic coloration and male body length when PLRs are performed on our phylogram. This result may be an artifact of the methodological approach of PLR on phylogenies with non-contemporaneous tips. The PLC approach of Ives & Garland (2010) standardizes tip-to-tip distances for non-contemporaneous trees. However, even if two tips are phylogenetically related (e.g., species A and species B), the standardization procedure may result in one tip being nearest to another less closely related tip (species A and species C) (Ives & Garland, 2010). Only one other phylogenetic comparative study on Lepidoptera larvae has failed to find support for a coloration-body size relationship (Nilsson & Forsman, 2003), which may have been confounded by the association between gregariousness and smaller larval body size and adult wing span.

Other phylogenetic comparative studies in poison dart frogs have shown associations between the prevalence or conspicuousness of aposematic coloration and other traits. Santos et al. (2003) found support for an apparent association with diet specialization (ants, termites, and mites) despite a lack of dietary information for most species. Summers & Clough (2001) found that a positive relationship with toxicity, which is supported by some microevolutionary studies at the species scale (e.g., Maan & Cummings, 2012). Such associations may also occur in millipede assassin bugs given their apparent predatory specialization and chemical defensiveness. However, a lack of dietary and toxicity data for a significant number of species currently hinders any testing

of these hypotheses, although many of the identified millipede-feeding species and genera in Forthman & Weirauch (2012), as well as unpublished images, display aposematic coloration.

*Multiple transitions from limited to extreme sexual dimorphism*

Our study confirms that extreme sexual dimorphism has evolved multiple times from a limited condition throughout the Tribelocephalinae+Ectrichodiinae phylogeny. Based on our results and observations of other species, extreme sexual dimorphism is apparent in taxa that have macropterous males and apterous or some micropterous females (e.g., *Xenocaucus*, *Toxopus*, *Distirogaster*). Compared to males, females generally show drastic reductions in a number of head, thoracic, and abdominal features, while anterior pronotal lobe enlargement and additional leg armature is observed. Given the convergence of these characters in distantly related taxa, we caution the use of female characters in phylogenetic reconstruction and taxonomic decisions, especially for known or potentially extremely dimorphic species. Other phylogenetic studies on cantharoid and elateroid beetles (Bocakova et al., 2007; Bocak et al., 2008; Kunderata & Bocak, 2011) have reached the same conclusions; previous higher-level classifications in these groups were primarily based on characters that are shown to be highly homoplasious among lineages with neotenic females (Bocakova et al., 2007; Kunderata & Bocak, 2011).

Although there is missing data for many species, we are aware of genera in which described or putative macropterous male-based and micropterous or apterous female-based species are observed. For example, both described species of *Apocaucus* Distant, 1909, are known from macropterous males. We have examined apterous females of this genus that exhibit morphological reductions in the head and thorax, which suggests the possibility of extreme sexual dimorphism in these species. These females may be individuals belonging to one of the described species, but we are unable to assign them based on morphology alone. Thus, other sources of

data, e.g., molecular and distributional data, would be needed to confirm conspecificity. Similar situations have been personally observed in, e.g., *Tribelocodia*, *Lynamna* Breddin, 1900, and many other species of *Gibbosella*. Previous taxonomic works have also demonstrated the difficulty of extreme sexual dimorphism in associating males and females; e.g., *Parascadra breuningi* Kerzhner & Günther, 2004, was described from a micropterous female specimen that was later shown to be a female of *Rhysostehus glabellus* males (Rédei et al., 2012). We suspect that similar taxonomic issues are more widespread in Ectrichodiinae given that over half of the genera are monotypic and based on one or few individuals of a single sex; this may even extend to many species within genera.

Wing reduction and/or extreme sexual dimorphism in females has been observed in many other insect orders (e.g., Diptera, Disney, 1996; Phasmatodea, Whiting et al., 2003; Strepsiptera, Kathirithamby, 1989), and hypotheses have been proposed to explain the evolution of this phenomenon. One hypothesis posits that a reduction in wings and other morphological features permits reallocation of resources to increased fecundity in females and/or reduce the age of first reproduction (Darwin, 1859; Roff, 1986; Denno et al., 1989; Roff & Fairbairn, 1991). A second, compatible hypothesis is that these species are under less selection pressure to maintain dispersive flight capabilities in stable environments (Slater, 1977; see Roff, 1990). Non-macropterous morphs of millipede assassin bugs are commonly observed in stable leaf litter microhabitats compared to macropterous morphs that fly in open vegetation (Dougherty, 1995; pers. obs.). However, experimental evidence on allocation trade-offs between flight and reproduction are lacking in this group. Regardless, trade-offs between the production of morphological features for dispersive flight and increased reproduction and the role of stable habitats are plausible hypotheses to explain the evolution of wing reduction and extreme sexual dimorphism in Ectrichodiinae and Tribelocephalinae. Another hypothesis states that flight



capabilities are reduced in taxa that occupy isolated habitats (e.g., oceanic islands, caves, mountaintops), since dispersers are likely to experience higher risks of mortality (Darwin, 1859; Wagner & Leibherr, 1992). Under this hypothesis, we would expect both sexes in a species to exhibit the same wing and associated morphological reductions (i.e., limited sexual dimorphism), which is observed in some Ectrichodiinae species (e.g., *Maraenaspis* Karsch, 1892, *Haematorrhophus* Stål, 1874). However, for species in which only brachypterous to apterous forms are known, it is possible that macropterous forms exist (e.g., *Glymmatophora*) but remain to be discovered. With respect to other species, the isolation hypothesis would not explain why there is sex-specific morphological reductions.

#### *Phylogenetics and reclassification of Ectrichodiinae and Tribelocephalinae*

Previous Reduviidae systematic investigations support a close relationship of Ectrichodiinae with Tribelocephalinae. Although these two subfamilies have drastically distinct habitus, Weirauch (2008, 2010) recovered several synapomorphies supporting this clade: e.g., numerous transverse ridges on the mandibular stylets, lamellate processes on the right maxillary stylet, subdivision of the antennal flagellomeres, and extremely short female external genitalia, among others. However, the monophyly of neither subfamily has been firmly established from previous studies. Weirauch & Munro's (2009) molecular cladistic analyses recovered a monophyletic Ectrichodiinae, with the exception of one analysis supporting paraphyly with respect to the single included Tribelocephalinae representative. Weirauch's (2010) morphological parsimony analysis supported Ectrichodiinae monophyly in half of the parsimonious trees, but *Ectrichodiella minima* (Valdés), 1910 was recovered as the sister group to the Tribelocephalinae in the remaining trees.

Our separate and combined morphological and molecular phylogenetic results based on a larger sample of Ectrichodiinae and Tribelocephalinae taxa supports the non-monophyly of both

analyses, with some discordance with our morphology only results. As such, we use the results of our combined morphological and molecular ML phylogenetic results to reclassify the Ectrichodiinae+Tribelocephalinae higher-level relationships into diagnosable monophyletic groups (Fig. 4.5). We refrain from revising the tribe Xenocaucini or re-assigning *Xenocaucus* to another tribe due to the uncertain phylogenetic position of this genus in on our analyses. Unambiguous character optimizations were examined on the combined ML topology in WinClada v1.00.08 (Nixon, 2002) to inform diagnoses (Fig. 4.5). Diagnoses and descriptions are primarily based on male characters.

### **Taxonomy**

Ectrichodiinae Amyot & Serville, 1843

Fig. 4.5A–T

Ectrichodiinae Amyot & Serville, 1843, 1: 342.

Tribelocephalinae Stål, 1865, 3: 44. New synonymy.

Type genus: *Ectrichodia* Lepeletier & Serville, 1825

REVISED DIAGNOSIS: This subfamily is recognized by the subdivided antennal basi- and/or distiflagellomeres making the antennae appear 5–8-segmented (Fig. 4.5D) (4-segmented in *Schottus* Distant, 1902, *Schuhella* Dougherty, 1995, *Tribelocodia*, *Vilius* Stål, 1863, and *Zirta* Stål, 1859), longer and denser vestiture on the scapus and pedicellus in males (Fig. 4.5D), more than 35 external transverse ridges on the mandibular stylet (47–1; Fig. 4.5F), lamellate ventral processes on the right maxillary stylet interiorly (49–0; Fig. 4.5G) (toothlike in *Vilius*), a medially depressed scutellar disc (83–1; Fig. 4.5K) (flat in some genera, e.g., *Abelocephala* Maldonado, 1996, *Glymmatophora*, and *Haematorrhophus*), and a globular forecoxa (94–0; Fig. 4.5L). Females have short, platelike external genitalia.

REDESCRIPTION: **MALE:** STRUCTURE: HEAD: Short or elongate, variably shaped; scapus and pedicellus with long, dense, erect to semi-erect setae; antennal flagellomeres 5- to 8-segmented with the exception of 4-segmented antenna in *Schottus*, *Schuhella*, *Tribelocodia*, *Vilius*, and *Zirta*; labium with three visible segments (segments II–IV; segment I reduced); mandibular stylet with more than 35 transverse ridges on external surface; right maxillary stylet with interior lamellate ventral processes (toothlike in *Vilius*). THORAX: Scutellar disc medially depressed; globular coxae; tarsi 3-segmented (2-segmented in *Xenocaucus*; 1-segmented and fused to tibia in *Tribelocodia*); hemelytron with two to three membranal cells (one in *Quinssyana funeralis* Distant, 1913). ABDOMEN: Dorsal laterotergites separated from mediotergites by membrane; ventral laterotergites separated from mediosternites by membrane (except segment II in Ectrichodiini, new tribe); mediosternites fused.

DISCUSSION: A taxonomic history of Ectrichodiinae prior to our study is provided by Dougherty (1995). No tribal classification has been previously proposed for this subfamily. Prior to our phylogenetic results and reclassification, Ectrichodiinae was diagnosed by the following features: subdivided basi- and/or distiflagellomeres, making the antennae appear more than 4-segmented; presence of ventral and dorsal processes on the left maxillary stylet; bifurcated scutellum; hemelytral membrane with two or three cells; fossula spongiosa on foretibia, as well as midtibia in many species; absence of the ventral connexival suture on abdominal sternite II; Brindley's gland and associated evaporatorium present, and three dorsal abdominal glands (DAGs) with DAG III ostioles larger than those of DAGs I and II.

The Tribelocephalinae was first recognized as a subfamily of Reduviidae by Stål (1865) and was subsequently divided by Villiers (1943) into two tribes based on venation patterns of the hemelytron: Opistplatyini and Tribelocephalini. A third tribe, Xenocaucini, was later erected by Maldonado (1996) for the female-based genus *Xenocaucus* characterized by a number of unique

features among Tribelocephalinae. Species of Tribelocephalinae have been characterized by less globular eyes, a long first visible labial segment, the strongly bent scapo-pedicellar articulation, the very small corium compared to the large membranous area of the hemelytron, and the presence of dense vestiture that included trichomes with a bulbous base and thin, curved apex. All species of the subfamily lack ocelli, with the exception of *Tribelocodia*.

As mentioned, previous phylogenetic results have provided consistent support for the Ectrichodiinae+Tribelocephalinae clade, with some of the morphological synapomorphies from these studies (Weirauch 2008, 2010) supported by our results. However, the monophyly of both subfamilies remained doubtful, and our phylogenetic results confirm their non-monophyly. As such, we synonymize Tribelocephalinae with Ectrichodiinae, erect three new tribes and two subtribes, and revise the tribe Tribelocephalini. We also include Xenocaucini as a tribe in Ectrichodiinae. In addition to the characters listed in the diagnosis, our character optimizations recovered additional synapomorphies: very dense vestiture with bulbous hairs present on the integument (absent in the Ectrichodiini), eye more than one-half of the head height (reduced in many apterous species or in females), pedicellus ventrally inserted on scapus so that pedicellus and flagellomeres point posteriorly (absent in the Ectrichodiini), pronotum wider than long (exceptions observed in a number of taxa), anterior pronotal lobe shorter than the posterior lobe (exceptions observed in a number of taxa, particularly in individuals with reduced wings), and lateral depression present on the posterior pronotal lobe (absent in some apterous individuals).

Tribelocodiini, new tribe

Fig. 4.5I, N, Q, S

Type genus: *Tribelocodia* Weirauch, 2010

DIAGNOSIS: This tribe is diagnosed by the narrow postocular shape in dorsal view (14–0); presence of ocelli (22–1; Fig. 4.5B); pedicellus ventrally inserted on scapus so that pedicellus and flagellomeres point posteriorly (38–1; Fig. 4.5C); pronotal transverse furrow laterally divided by elevated cuticle (69–1; Fig. 4.5I), which is distinctly ridgelike in some *Ectrichodiella* species; posteriolateral margin of pronotum surpassing medial pronotal margin (71–1; Fig. 4.5I); scutellum with paramedian scutellar processes and a well-developed medial process (75–1, 80–1; Fig. 4.5I); small tubercles uniformly covering the trochanters, femora, and tibiae (97–1, 101–1, 108–1, 113–1, 117–1, 124–1, 129–1, 131–1, 138–1; Fig. 4.5Q); tibiae lacking fossula spongiosa; and proximal parts of M and Cu of forewing forming one vein (143–1; Fig. 4.5N).

DESCRIPTION: **MALE:** STRUCTURE: HEAD: Clypeus without anterior process; postocular narrow in dorsal view; eye reniform with concave posterior margin; ocelli present; pedicellus ventrally inserted on scapus so that pedicellus and flagellomeres point posteriorly; pedicellus moderately to strongly curved; basiflagellomere 1-segmented; constriction between postocular region and neck; labrum sclerotized. THORAX: Pronotum with transverse furrow subdivided by laterally elevated cuticle, ridgelike in *Ectrichodiella*; posteriolateral pronotal margin surpassing medial pronotal margin; metepisternal supracoal lobe of mesocoal cavity present; scutellum with well-developed paramedian processes and a well-developed medial process; trochanters, femora, and tibiae with small, uniformly distributed setigerous tubercles; tibiae lacking fossula spongiosa; proximal parts of M and Cu of forewing forming one vein. ABDOMEN: DAGs I–III ostioles present in adults (except I absent in *Tribelocodia*) and of similar sizes.

DISCUSSION: This tribe includes the Neotropical genera *Ectrichodiella* and *Tribelocodia*.

Abelocephalini, new tribe

Fig. 4.5E

Type genus: *Abelocephala* Maldonado, 1996

DIAGNOSIS: Abelocephalini may be diagnosed by the following characters: bulbous trichomes present (2–1; Fig. 4.5A), eye reniform with concave posterior margin, ocelli absent, antennal shield present (32–1; Fig. 4.5B, E), scapus with ventroposterior apical lobe (37–1; Fig. 4.5E) (absent in *Abelocephala*), pedicellus ventrally inserted on scapus so that pedicellus and flagellomeres point posteriorly (38–1; Fig. 4.5C), posteriolateral margin of pronotum surpassing medial pronotal margin (71–1; Fig. 4.5I), scutellum with well-developed medial process (80–1; Fig. 4.5I, J) (paramedian process also present in *Afrodecius* Jeannel, 1919), tibiae lacking fossula spongiosa, tarsi 2-segmented, hemelytron greatly surpassing abdominal apex, proximal part of M and Cu of hemelytron forming one vein (143–1; Fig. 4.5N, O).

DESCRIPTION: **MALE:** VESTITURE: Dense vestiture on parts of the head, anterior to lateral margins of the pronotum, and near the thoracic-abdominal articulation; bulbous trichomes present. STRUCTURE: HEAD: Clypeus without anterior process; ocelli absent; antennal shield present; scapus with ventroposterior apical lobe (except in *Abelocephala*), pedicellus ventrally inserted on scapus so that pedicellus and flagellomeres point posteriorly; pedicellus strongly curved; distinct constriction between postocular region and neck; labrum completely sclerotized. THORAX: Posteriolateral margin of pronotum surpassing medial pronotal margin; scutellum with well-developed medial process (paramedian process also present in *Afrodecius* Jeannel, 1919); tibiae lacking fossula spongiosa; tarsi 2-segmented; hemelytron greatly surpassing abdominal apex; proximal part of M and Cu of hemelytron forming one vein. ABDOMEN: DAGs II and III ostioles only present and of similar size in adults.

DISCUSSION: *Abelocephala*, *Afrodecius*, *Apocaucus*, and *Megapocaucus* Miller, 1954 are included within this tribe based on our phylogenetic results. Based on our diagnosis, we also include *Enigmocephala* Rédei, 2007, *Gastrogyrus* Bergroth, 1921, *Homognetus* Bergroth, 1923, and *Mantangocoris* Miller, 1940. Rédei (2007) highlight the morphological similarities between *Enigmocephala* with *Gastrogyrus*, *Mantangocoris*, and *Afrodecius*, particularly with head morphology (e.g., the anteriorly declivent head, pubescent postocular and gular regions, robust labium, and/or apical processes on ultimate labial segment). *Homognetus* is also similar to these genera in the pubescent postocular and gular regions. It is worth noting that an undescribed male specimen of *Xenocaucus* was observed for our analysis and also exhibits many of the same diagnostic features of this tribe (e.g., 2-segmented tarsus; hemelytron greatly surpassing abdominal apex), as well as other morphological features (fringe of hairs on postocular and gular areas; DAGs II and III ostioles of similar size). Thus, it is possible that this genus may be a member of Abelocephalini and was recovered as sister to or nested within this tribe in some of our phylogenetic results; however, further phylogenetic testing is needed with more representatives of this tribe for both molecular and morphological data.

Tribelocephalini Villiers, 1943

Fig. 4.5C, J, O, P

Tribelocephalini Villiers, 1943, 10: 9.

Type genus: *Tribelocephala* Stål, 1853

REVISED DIAGNOSIS: Species of this tribe are distinguished by the dense, tomentose vestiture throughout the body (especially the entire head) (Fig. 4.5C), head longer than pronotum (except little shorter than pronotum in *Tomolus* Stål, 1874), clypeus with anterior process (20–1; Fig. 4.5C) (absent in *Opistoplatys* Westwood, 1834 and *Tomolus*), eye adpressed closely to the

head surface and with concave anterior and posterior eye margin (26–0, 27–1; Fig. 4.5C) (concavity slight in *Tomolus*), ventral margin of second visible labial segment (appears dorsal when retracted under the head) is straight or concave (45–0), apex of short prosternal stridulatory process acute (86–0), and DAG I–III ostioles present in adults and of similar sizes (164–1; Fig. 4.5S).

REDESCRIPTION: **MALE:** VESTITURE: Dense tomentose vestiture throughout the entire body, bulbous trichomes present. STRUCTURE: HEAD: Head longer than pronotum (little shorter than pronotum in *Tomolus*); clypeus with anterior process (absent in *Opistoplatys* and *Tomolus*); eye adpressed closely to the head surface and with concave anterior and posterior eye margin; ocelli absent; antennal shield absent; pedicellus slightly to moderately curved; pedicellus ventrally inserted on scapus so that pedicellus and flagellomeres point posteriorly; pedicellus strongly curved; labrum completely sclerotized; ventral margin of second visible labial segment straight or concave. THORAX: Posteriolateral margin of pronotum surpassing or slightly surpassing medial pronotal margin; scutellum with well-developed medial process only; apex of short prosternal stridulatory process acute; tibiae lacking fossula spongiosa; tarsi 3-segmented, hemelytron reaching or extending a little beyond abdominal apex. ABDOMEN: DAG I–III ostioles present in adults and of similar sizes.

DISCUSSION: Villiers (1943) erected the tribe as a member of the Tribelocephalinae and included all of the genera in our new tribe Abelocephalini, as well as *Tomolus* and *Tribelocephala* Stål, 1853. The tribe was once diagnosed by the absence of the proximal m-cu cross vein in the hemelytra. *Acanthorhinocoris* Miller, 1940, was initially assigned to the Tribelocephalinae, but Rédei (2007) transferred the genus to the Opistoplatyini based on the presence of the m-cu cross vein. Based on our results, we revise the Tribelocephalini and establish two subtribes, the Opistoplatyina and Tribelocephalina. Based on our diagnostic features and other morphological



characters, we include *Acanthorhinocoris*, *Centrogastocoris* Miller, 1958, *Distantus* Villiers, 1943, *Opistoplatys*, *Plectrophorocoris* Miller, 1958, *Tribecephala*, and *Tomolus*.

Opistoplatyina, new subtribe

Fig. 4.5J, P

Type genus: *Opistoplatys* Westwood, 1834

DIAGNOSIS: In addition to the tribal diagnosis, this subtribe is diagnosed by the eyes reaching or nearly reaching the dorsal and ventral surfaces of the head (29–1, 30–1), antennal insertion laterally concealing suture between maxillary and mandibular plates (34–1), scapus without apical lobes, a prominent postscutum (84–1; Fig. 4.5J), and the proximal parts of M and Cu forming separate veins in the hemelytron (143–0; Fig. 4.5P).

DESCRIPTION: **MALE:** as in diagnosis.

DISCUSSION: This subtribe retains the same generic composition as the previously recognized Opistoplatyini: *Acanthorhinocoris*, *Centrogastocoris*, *Distantus*, *Opistoplatys*, and *Plectrophorocoris*.

Tribecephalina, new subtribe

Fig. 4.5C, O

Type genus: *Tribecephala* Stål, 1853

DIAGNOSIS: Species of this subtribe are characterized by the antennal insertion ventral to the suture between the maxillary and mandibular plates (34–2; Fig. 4.5C), scapus with ventroanterior and ventroposterior apical lobes (36–1, 37–1; Fig. 4.5E), a reduced or absent postscutum (84–0), and the proximal parts of M and Cu forming a fused vein in the hemelytron (143–1; Fig. 4.5O).

DESCRIPTION: **MALE:** as in diagnosis.

DISCUSSION: We include *Tribelocephala* and *Tomolus* in this subtribe based on the diagnosis.

Ectrichodiini, new tribe

Fig. 4.5A, D, F–H, K–M, R, T

Type genus: *Ectrichodia* Lepeletier & Serville, 1825

DIAGNOSIS: This large tribe is diagnosed by the absence of bulbous trichomes; the distally or slightly ventrally inserted pedicellus (38–0; Fig. 4.5D); basiflagellomere subdivided into two pseudosegments (42–1; Fig. 4.5D) (not subdivided in *Schottus*, *Schuhella*, *Vilius*, and *Zirta*); a subdivided labrum (44–0; Fig. 4.5B) (not subdivided in *Schuhella*); the ventral, internal row of processes on the left maxillary stylet (51–1; Fig. 4.5H); paramedian scutellar processes present (75–1; Fig. 4.5K); a long prosternal stridulatory process that surpasses the posterior margin of the forecoxal cavity (85–1; Fig. 4.5L); the metathoracic gland evaporatorium as modified cuticle (91–1; Fig. 4.5M); and the absence of the ventral connexival suture on abdominal sternites II (154–0; Fig. 4.5T).

DESCRIPTION: **MALE:** VESTITURE: Glabrous to dense, erect to semi-erect vestiture without bulbous trichomes. STRUCTURE: HEAD: Ocellar lens in, at least, macropterous individuals; pedicellus distally or slightly ventrally inserted in scapus; basiflagellomere typically subdivided into two pseudosegments (one-segmented in *Schottus*, *Schuhella*, *Vilius*, and *Zirta*); labrum subdivided by transverse membrane (completely sclerotized in *Schuhella*); left maxillary stylet with ventral, internal row of processes. THORAX: Macropterous forms, at least, with a distinct longitudinal depression on the posterior pronotal lobe; posteriolateral margin of pronotum not or barely surpassing medial pronotal margin; paramedian scutellar processes present; prosternal stridulatory process long, surpassing posterior margin of forecoxal cavity; metathoracic

gland evaporatorium present as modified cuticle; tarsi 3-segmented. ABDOMEN: ventral connexival suture on abdominal sternites II absent; DAGs II and III ostioles present in adults (DAG II ostioles absent in some species; see discussion), with DAG III larger than DAG II.

DISCUSSION: All genera recognized in the previous classification of Ectrichodiinae, with the exception of *Ectrichodiella*, are included in the Ectrichodiini. Several other synapomorphies of the nominal tribe were recovered in our optimizations, but show reversals or future modifications in some of the included taxa. The absence of a single medial scutellar process (79–0) is optimized on our phylogeny, however some genera (e.g., *Ectrychotes* Burmeister, 1835) have very small medial processes. Many genera and species are characterized by more incrassate forefemora compared to the mid femora (95–1), but a number of other genera and species have slender legs (e.g., *Tanindrazanus* Forthman, Chlond, & Weirauch, [in press]; *Pothea* Amyot & Serville, 1843). A fossula spongiosa is commonly observed on the fore- and mid tibiae (111–1, 127–1), but there are several instances of reversals on one or both legs. Although DAGs II and III ostioles are present in many of the species we have observed (Fig. 4.5R), as well as by Weirauch (2006), some species in *Scadra* Stål, 1859, *Pseudopothea* Wygodzinsky, 1951, *Microsanta* Breddin, 1903, and *Adrania* Stål, 1863, lack DAG II ostioles.

## Conclusion

The evolution of aposematic coloration in animals has been largely explored outside a phylogenetic context. This is especially true for insects other than Lepidoptera and Coleoptera. While a few vertebrate and invertebrate studies have shown a positive association of body size and prevalence or conspicuousness of aposematic coloration, others have provided evidence of negative or no associations. However, results indicating no coloration-body size associations in Lepidoptera are likely confounded by the gregarious lifestyles of many species. The typically

solitary millipede assassin bugs presented an opportunity to contribute to a limited body of research that has sought to understand the evolution of coloration and its association with body size. We also had the opportunity to explore the evolution of extreme sexual dimorphism in this group, which is a phenomenon that has not been explored in a phylogenetic framework for other Reduviidae subfamilies. Lastly, we were able to test conflicting phylogenetic hypotheses regarding relationships between Ectrichodiinae and Tribelocephalinae. Phylogenetic hypotheses were generated from separate and combined morphological and molecular cladistic analyses under different optimality criteria, all of which confirmed non-monophyly of both subfamilies. Furthermore, most phylogenetic results recovered similar higher-level clades, which we have recognized in our new classification of Ectrichodiinae.

Based on our phylogenetic hypothesis, aposematic coloration is shown to have evolved from a cryptic ancestor early in the evolutionary history of millipede assassin bugs. Subsequent reversals to cryptic coloration and regains of aposematic coloration are reconstructed. Our PLRs support a positive association between aposematic coloration and body length when performed on ultrametric trees. However, this association is not supported when estimated on phylograms, which may be an artifact of the method used. Due to a lack of data, we were unable to investigate other hypotheses regarding the prevalence of aposematic coloration and toxicity or diet specialization. Thus, characterizing the defense compounds and collecting more natural history observations are critically needed. Future studies should also investigate the role of aposematic coloration and metallic features in aposematic signaling within Ectrichodiinae. Extreme sexual dimorphism is also shown to evolve multiple times from a more limited condition despite the lack of data on both sexes for many species. This gap in knowledge may be, in part, the result of taxonomic decisions based on one sex. Thus, careful consideration must be given to the use of

female morphology in phylogenetic investigations and taxonomic decisions, especially in cases where extreme sexual dimorphism may be apparent based on patterns recovered from our study.

## References

- Areekul, B., Quicke, D.L.J. 2006. The use of colour characters in phylogenetic reconstruction. *Biological Journal of the Linnean Society* 88: 193–202.
- Arrow, G.J. 1951. *Horned Beetles: A Study of the Fantastic in Nature*. W. Junk Publishers, the Hague.
- Bates, H.W. 1862. XXXII. Contributions to an insect fauna of the Amazon Valley. Lepidoptera: Heliconidae. *Transactions of the Linnean Society of London* 23: 495–566.
- Bocak, L., Bocakova, M., Hunt, T., Vogler, A.P. 2008. Multiple ancient origins of neoteny in Lycidae (Coleoptera): consequences for ecology and macroevolution. *Proceedings of the Royal Society B* 275: 2015–2023.
- Bocakova, M., Bocak, L., Hunt, T., Teraväinen, M., Vogler, A.P. 2007. Molecular phylogenetics of Elateriformia (Coleoptera): evolution of bioluminescence and neoteny. *Cladistics* 23: 477–496.
- Bocakova, M., Bocak, L., Gimmel, M.L., Motyka, M., Vogler, A.P. 2016. Aposematism and mimicry in soft-bodied beetles of the superfamily Cleroidea (Insecta). *Zoologica Scripta* 45: 9–21.
- Bromham, L., Woolfit, M., Lee, M.S.Y., Rambaut, A. 2002. Testing the relationship between morphological and molecular rates of change along phylogenies. *Evolution* 56: 1921–1930.
- Cachan, P. 1952. Etude de la prédation chez les Réduvidés de la région Éthiopienne. 1. La prédation en groupe chez *Ectrichodia gigas* H-Sch. *Physiologia Comparata et Oecologia* 2: 378–385.
- Carpintero, D.J., Maldonado, J. 1996. Diagnostic characters and key to the genera of American Ectrichodiinae (Heteroptera, Reduviidae). *Caribbean Journal of Science* 32: 125–141.
- Cheney, K.L., Cortesi, F., How., M.J., Wilson, N.G., Blomberg, S.P., Winters, A.E., Umanzör, S., Marshall, N.J. 2014. Conspicuous visual signals do not coevolve with increased body size in marine sea slugs. *Journal of Evolutionary Biology* 27: 676–687.
- Cott, H.B. 1940. *Adaptive Coloration in Animals*. Methuen & Co., Ltd., London, UK.
- Cryan, J.R., Liebherr, J.K., Fetzner, J.W., Jr., Whiting, M.F. 2001. Evaluation of relationships within the endemic Hawaiian Platynini (Coleoptera: Carabidae) based on molecular and morphological evidence. *Molecular Phylogenetics and Evolution* 21: 72–85.
- Cusimano, N., Renner, S.S. 2014. Ultrametric trees or phylograms for ancestral state reconstruction: does it matter? *Taxon* 63: 721–726.
- Dallwitz, M.J. 1980. A general system for coding taxonomic descriptions. *Taxon* 29: 41–46.

- Dallwitz, M.J., Paine, T.A., Zurcher, E.J. 1999. User's guide to the DELTA Editor. Accessible at <http://delta-intkey.com>.
- Damgaard, J., Andersen, N.M., Cheng, L., Sperling, F.A.H. 2000. Phylogeny of sea skaters, *Halobates* Eschscholtz (Hemiptera, Gerridae), based on mtDNA sequence and morphology. *Zoological Journal of the Linnean Society* 130: 511–526.
- Darwin, C. 1859. *The Origin of Species*. John Murray, London, UK.
- Denno, R.F., Olmstead, K.L., McCloud, E.S. 1989. Reproductive cost of flight capability: a comparison of life history traits in wing dimorphic planthoppers. *Ecological Entomology* 14: 31–44.
- Disney, R.H.L. 1996. A new genus of scuttle fly (Diptera; Phoridae) whose legless, wingless, females mimic ant larvae (Hymenoptera; Formicidae). *Sociobiology* 27: 95–118.
- Distant, W.L. 1904. *Fauna of British India. Rhynchota, Vol. II*. Taylor & Francis, London, UK.
- Dougherty, V. 1995. A review of the New World Ectrichodiinae genera (Hemiptera: Reduviidae). *Transactions of the American Entomological Society* 121: 173–225.
- Edmunds, M. 1987. Color in opisthobranchs. *American Malacological Bulletin* 5: 185–196.
- Exnerová, A., Svádová, K., Štys, P., Barcalová, S., Landová, E., Prokopová, M., Fuchs, R., Socha, R. 2006. Importance of colour in the reaction of passerine predators to aposematic prey: experiments with mutants of *Pyrrhocoris apterus* (Heteroptera). *Biological Journal of the Linnean Society* 88: 143–153.
- Fabricant, S.A., Exnerová, A., Ježová, D., Štys, P. 2014. Scared by shiny? The value of iridescence in aposematic signalling of the hibiscus harlequin bug. *Animal Behaviour* 90: 315–325.
- Felsenstein, J. 1985. Phylogenetics and the comparative method. *American Naturalist* 125: 1–15.
- Forero, D., Berniker, L., Weirauch, C., 2013. Phylogeny and character evolution in the bee-assassins (Insecta, Heteroptera: Reduviidae). *Molecular Phylogenetics and Evolution* 66: 283–302.
- Forsman, A., Merilaita, S. 1999. Fearful symmetry: pattern size and asymmetry affects aposematic signal efficacy. *Evolutionary Ecology* 13: 131–140.
- Forthman, M., Weirauch, C. 2012. Toxic associations: a review of the predatory behaviors of millipede assassin bugs (Hemiptera: Reduviidae: Ectrichodiinae). *European Journal of Entomology* 109: 147–153.
- Forthman, M., Chłond, D., Weirauch, C. (in press). Taxonomic monograph of the endemic millipede assassin bug fauna of Madagascar (Hemiptera: Reduviidae: Ectrichodiinae). *Bulletin of the American Museum of Natural History*.

- Forthman, M., Weirauch, C. (in press) Phylogenetics and biogeography of the endemic Madagascan millipede assassin bugs (Heteroptera: Reduviidae: Ectrichodiinae). *Molecular Phylogenetics and Evolution*.
- Gamberale, G., Tullberg, B.S. 1996a. Evidence for a peak-shift in predator generalization among aposematic prey. *Proceedings of the Royal Society of London B* 263: 1329–1334.
- Gamberale, G., Tullberg, B.S. 1996b. Evidence for a more effective signal in aggregated aposematic prey. *Animal Behaviour* 52: 597–601.
- Gamberale, G., Tullberg, B.S. 1998. Aposematism and gregariousness: the combined effect of group size and coloration on signal repellence. *Proceedings of the Royal Society of London B* 265: 889–894.
- Giacchi, J.C. 1987. Contributions to the morphology and taxonomy of the American Stenopodainae (Heteroptera, Reduviidae). II. General considerations about the subfamily. *PHYSIS (Buenos Aires), Seccion C* 45: 19–30.
- Gil-Santana, H.R. 2015. First record of the genus *Pseudopothea* from South America, with description of a new species from Brazil (Hemiptera: Heteroptera: Reduviidae: Ectrichodiinae). *Zootaxa* 3904: 541–552.
- Goloboff, P.A., Farris, J.S., Nixon, K.C. 2008. TNT, a free program for phylogenetic analysis. *Cladistics* 24: 774–786.
- Guilford, T. 1988. The evolution of conspicuous coloration. *American Naturalist* 131: S7–S21.
- Hagman, M., Forsman, A. 2003. Correlated evolution of conspicuous coloration and body size in poison frogs (Dendrobatidae). *Evolution* 57: 2904–2910.
- Härilin, C., Härilin, M. 2003. Towards a historization of aposematism. *Evolutionary Ecology* 17: 197–212.
- Hart, E.R. 1972. *A Systematic Revision of the Genus Zelus Fabricius (Hemiptera: Reduviidae)*. Ph.D. Dissertation. Texas A&M University.
- Hill, L. 2014. Revision of *Silhouettanus* with description of nine new species (Hemiptera: Heteroptera: Schizopteridae). *Zootaxa* 3815: 353–385.
- Hwang, W.S., Weirauch, C. 2010. Revision of the Malagasy *Durevius* Villiers with descriptions of two new species (Hemiptera: Reduviidae: Reduviinae). *Insect Systematics and Evolution* 41: 123–141.
- Ives, A.R., Garland, T., Jr. 2010. Phylogenetic logistic regression for binary dependent variables. *Systematic Biology* 59: 9–26.



- Ives, A.R., Garland, T., Jr. 2014. Phylogenetic regression for binary dependent variables. In Garamszegi, L.Z. (ed.): *Modern Phylogenetic Comparative Methods and Their Application in Evolutionary Biology: Concepts and Practices*. Springer, Heidelberg, Germany, pp. 231–262.
- Kathirithamby, J. 1989. Review of the order Strepsiptera. *Systematic Entomology* 14: 41–92.
- Katoh, K., Standley, D.M. 2013. MAFFT multiple sequence alignment software version 7: improvements in performance and usability. *Molecular Biology and Evolution* 30: 772–780.
- Kundrata, R., Bocak, L. 2011. The phylogeny and limits of Elateridae (Insecta: Coleoptera): is there a common tendency of click beetles to soft-bodiedness and neoteny? *Zoologica Scripta* 40: 364–378.
- Lanfear, R., Calcott, B., Ho, S.Y.W., Guindon, S. 2012. PartitionFinder: combined selection of partitioning schemes and substitution models for phylogenetic analyses. *Molecular Biology and Evolution* 29: 1695–1701.
- Lawrence, R.F. 1984. *The Centipedes and Millipedes of Southern Africa: a Guide*. A.A. Balkema, Cape Town.
- Lepelletier, A.L.M., Serville, J.G.A. 1825. *Encyclopedie Methodique. Histoire Naturelle, Entomologie, ou Histoire Naturelle des Crustaces, des Arachnides et des Insectes. Tome Dixième*. Agasse, Paris.
- Lewis, P.O. 2001. A likelihood approach to estimating phylogeny from discrete morphological character data. *Systematic Biology* 50: 913–925.
- Litsios, G., Salamin, N. 2012. Effects of phylogenetic signal on ancestral state reconstruction. *Systematics Biology* 61: 533–538.
- Maan, M.E., Cummings, M.E. 2012. Poison frog colors are honest signals of toxicity, particularly for bird predators. *American Naturalist* 179: E1–E14.
- Maddison, W.P., Maddison, D.R. 2015. Mesquite: a modular system for evolutionary analysis. Version 3.02. Available: <http://mesquiteproject.org>.
- Maldonado, J. 1990. Systematic catalogue of the Reduviidae of the world (Insecta: Heteroptera). *Caribbean Journal of Science*, Special Edition: 1–694.
- Maldonado, J. 1996. New taxa and key to the tribes and genera in Tribelocephalinae Stål 1866 (Heteroptera: Reduviidae). *Proceedings of the Entomological Society of Washington* 98: 138–144.
- Midford, P.E., Garland, T., Jr. Maddison, W.P. 2009. PDAP package of Mesquite. Version 1.16. Available: [http://mesquiteproject.org/pdap\\_mesquite/](http://mesquiteproject.org/pdap_mesquite/).

- Miller, N.C.E. 1941. New genera and species of Malaysian Reduviidae (continued). Part II. *Journal of the Federated Malay States Museum* 18: 601–773.
- Miller, N.C.E. 1955. New genera and species of Reduviidae (Hemiptera Heteroptera). *Bollettino del Laboratorio di Zoologia Generale e Agraria della Facolta Agraria in Portici Silvestri* 33: 427–445.
- Nilsson M., Forsman, A. 2003. Evolution of conspicuous colouration, body size and gregariousness: a comparative analysis of lepidopteran larvae. *Evolutionary Ecology* 17: 51–66.
- Nixon, K.C. 2002. WinClada ver. 1.00.08. Published by the author, Ithaca, NY.
- Omland, K.E. 1997. Correlated rates of molecular and morphological evolution. *Evolution* 51: 1381–1393.
- Pagel, M., Meade, A., Barker, D. Bayesian estimation of ancestral character states on phylogenies. *Systematic Biology* 53: 673–684.
- Parker, A. 1998. The diversity and implications of animal structural colours. *Journal of Experimental Biology* 201: 2343–2347.
- Peschke, K. Gack, C., Mahsberg, D. 2002. *o*-Nitrobenzaldehyde sprayed as a sneezing-powder by an assassin bug (*Ectrichodia* spec.) [abstract]. 19<sup>th</sup> Annual Meeting of the International Society of Chemical Ecology, University of Hamburg, Germany.
- Poulton, E.B. 1890. *The Colors of Animals*. Trübner & Co., Ltd., London, UK.
- Protas, M.E., Patel, N.H. 2008. Evolution of coloration patterns. *Annual Review of Cell and Developmental Biology* 24: 425–446.
- Rambaut, A., Suchard, M.A., Xie, W., Drummond, A.J. 2014. Tracer. Version 1.61. Available: <http://tree.bio.ed.ac.uk/software/tracer/>.
- Rédei, D., Ren, S., Bu, W. 2012. A new synonymy in the genus *Rhysostethus* (Hemiptera: Heteroptera: Reduviidae). *Acta Entomologica Musei Nationalis Pragae* 52: 341–348.
- Reemer, M., Ståhls, G. 2012. *Unravelling a Hotchpotch: Phylogeny and Classification of the Microdontinae (Diptera: Syrphidae)*. Ph.D. Dissertation. Leiden University.
- Roff, D.A. 1986. The evolution of wing dimorphism in Insecta. *Evolution* 40: 1009–1020.
- Roff, D.A. 1990. The evolution of flightlessness in Insecta. *Ecological Monographs* 60: 389–421.
- Roff, D.A., Fairbairn, D.J. 1991. Wing dimorphisms and the evolution of migratory polymorphisms among the Insecta. *American Zoologist* 31: 243–251.

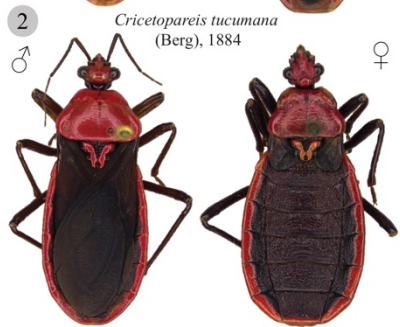
- Ronquist, F., Teslenko, M., van der Mark, P., Ayres, D.L., Darling, A., Höhna, S., Larget, B., Liu, L., Suchard, M.A., Huelsenbeck, J.P. 2012. MrBayes 3.2: efficient Bayesian phylogenetic inference and model choice across a large model space. *Systematic Biology* 61: 539–542.
- Rudh, A. 2013. Loss of conspicuous coloration has co-evolved with decreased body size in populations of poison dart frogs. *Evolutionary Ecology* 27: 755–767.
- Ruxton, G.D., Sherratt, T.N., Speed, M.P. 2004. *Avoiding Attack: The Evolutionary Ecology of Crypsis, Warning Signals, and Mimicry*. Oxford University Press, Oxford, UK.
- Sagegami-Oba, R., Takahashi, N., Oba, Y. 2007. The evolutionary process of bioluminescence and aposematism in cantharoid beetles (Coleoptera: Elateroidea) inferred by the analysis of 18S ribosomal DNA. *Gene* 400: 104–113.
- Sanderson, M.J. 2002. Estimating absolute rates of molecular evolution and divergence times: A penalized likelihood approach. *Molecular Biology and Evolution* 19: 101–109.
- Sanderson, M.J. 2003. r8s: inferring absolute rates of molecular evolution and divergence times in the absence of a molecular clock. *Bioinformatics* 19: 301–302.
- Santos, J.C., Coloma, L.A., Cannatella, D.C. 2003. Multiple, recurring origins of aposematism and diet specialization in poison frogs. *PNAS* 100: 12792–12797.
- Schouteden, H. 1952. Ectrichodiinae novae africanae. *Revue de Zoologie et de Botanique Africaines* 45: 306–312.
- Schultz, T.D. 2001. Tiger beetle defenses revisited: alternative defense strategies and colorations of two Neotropical tiger beetles, *Odontocheila nicaraguensis* Bates and *Pseudoxycheila tarsalis* Bates (Carabidae: Cincindelinae). *Coleopterists Bulletin* 55: 153–163.
- Simon, C., Frati, F., Beckenbach, A., Crespi, B., Liu, H., Flook, P. 1994. Evolution, weighting, and phylogenetic utility of mitochondrial gene sequences and a compilation of conserved polymerase chain reaction primers. *Annals of the Entomological Society of America* 87: 651–701.
- Slater, J.A. 1977. The incidence and evolutionary significance of wing polymorphism in lygaeid bugs with particular reference to those of South Africa. *Biotropica* 9: 217–229.
- Smith, S.M. 1975. Innate recognition of coral snake pattern by a possible avian predator. *Science* 187: 759–760.
- Smith, S.A., Donoghue, M.J. 2008. Rates of molecular evolution are linked to life history in flowering plants. *Science* 322: 86–89.
- Stål, C. 1865. Hemiptera Africana. *Öfversigt af Kongliga Vetenskaps-Akademiens Förhandlingar* 3: 1–200.

- Stankowich, T., Caro, T., Cox, M. 2011. Bold coloration and the evolution of aposematism in terrestrial carnivores. *Evolution* 65: 3090–3099.
- Stevens, M. 2015. Anti-predator coloration and behaviour: a longstanding topic with many outstanding questions. *Current Zoology* 61: 702–707.
- Summers, K., Clough, M.E. 2001. The evolution of coloration and toxicity in the poison frog family (Dendrobatidae). *PNAS* 98: 6227–6232.
- Thomas, D., Seago, A., Robacker, D. 2007. Reflections on golden scarabs. *American Entomologist* 53: 224–230.
- Tullberg, B.S., Hunter, A.F. 1996. Evolution of larval gregariousness in relation to repellent defences and warning coloration in tree-feeding Macrolepidoptera: a phylogenetic analysis based on independent contrasts. *Biological Journal of the Linnean Society* 57: 253–276.
- Urban, J.M., Cryan, J.R. 2007. Evolution of the planthoppers (Insecta: Hemiptera: Fulgoroidea). *Molecular Phylogenetics and Evolution* 42: 556–572.
- Vaidya, G., Lohman, D.J., Meier, R. 2011. SequenceMatrix: concatenation software for the fast assembly of multi-gene datasets with character set and codon information. *Cladistics* 27: 171–180.
- Vences, M., Kosuch, J., Boistel, R., Haddad, C.F.B., la Marca, E., Lötters, S., Veith, M. 2003. Convergent evolution of aposematic coloration in Neotropical poison frogs: a molecular phylogenetic perspective. *Organisms Diversity and Evolution* 3: 215–226.
- Villiers, A. 1943. Morphologie et systématique des Tribelocephalidae africains. *Revue Française d'Entomologie* 10: 1–28.
- Villiers, A. 1948. *Fauna de l'Empire Français. IX. Hémiptères Réduviides de l'Afrique Noire*. Office de la Recherche Scientifique Coloniale, Éditions du Muséum, Paris.
- Villiers, A. 1951. Notes d'entomologie ouest-africaine. II. Observations sur la répartition de quelques Hémiptères Hétéroptères et description de nouvelles espèces. *Bulletin de l'Institut Français d'Afrique Noire* 13: 326–342.
- Villiers, A. 1973. Un nouveau *Microstemma* de l'Ouest africain [Hem. Reduviidae Ectrichodiinae]. *Bulletin de l'Institut Français d'Afrique Noire* 35: 947–950.
- Vogler, A.P., Kelley, K.C. 1998. Covariation of defensive traits in tiger beetles (genus *Cicindela*): a phylogenetic approach using mtDNA. *Evolution* 52: 529–538.
- Wagner, D.L., Liebherr, J.K. 1992. Flightlessness in insects. *Trends in Ecology and Evolution* 7: 216–220.
- Weirauch, C. 2006. Dorsal abdominal glands in adult Reduviidae (Heteroptera, Cimicomorpha). *Deutsche Entomologische Zeitschrift* 53: 91–102.

- Weirauch, C. 2008. Cladistic analysis of Reduviidae (Heteroptera: Cimicomorpha) based on morphological characters. *Systematic Entomology* 33: 229–274.
- Weirauch, C. 2010. *Tribelocodia ashei*, new genus and new species of Reduviidae (Insecta: Hemiptera), has implications on character evolution in Ectrichodiinae and Tribelocephalinae. *Insect Systematics and Evolution* 41: 103–122.
- Weirauch, C., Munro, J.B. 2009. Molecular phylogeny of the assassin bugs (Hemiptera: Reduviidae), based on mitochondrial and nuclear ribosomal genes. *Molecular Phylogenetics and Evolution* 53: 287–299.
- Whiting, M.F., Bradler, S., Maxwell, T. 2003. Loss and recovery of wings in stick insects. *Nature* 421: 264–267.
- Wygodzinsky, P.W. A monograph of the Emesinae (Reduviidae: Hemiptera). *Bulletin of the American Museum of Natural History* 133: 1–613.
- Zhang, G., Weirauch, C. 2011. Matching dimorphic sexes and immature stages with adults: resolving the systematics of the *Bekilya* group of Malagasy assassin bugs (Hemiptera: Reduviidae: Peiratinae). *Systematic Entomology* 36: 115–138.
- Zhang, J., Gordon, E., Forthman, M., Hwang, W.S., Walden, K., Swanson, D., Johnson, K.P., Meier, R., Weirauch, C. 2016. Evolution of the assassin's arms: insights from a phylogeny of combined transcriptomic and ribosomal DNA data (Heteroptera: Reduvioidea). *Scientific Reports* 6: 22177.
- Zwickl, D.J. 2006. Genetic algorithm approaches for the phylogenetic analysis of large biological sequence datasets under the maximum likelihood criterion. Ph.D. Dissertation. University of Texas, Austin.
- Zwickl, D.J. 2008. GARLI manual. Available: <https://code.google.com/archive/p/garli/downloads>.

Figure 4.1. Examples of limited and extreme sexual dimorphism in Ectrichodiinae species. Headers correspond to the color code for the right panel of Fig. 4.3. Numbers in circles correspond to numbers highlighting examples of general morphological differences observed between sexes of the indicated species, but do not reflect how exaggerated the difference may be. Images are not scaled to size.

**Limited Sexual Dimorphism**



**Example Differences:**

<u>1</u>	<u>2</u>	<u>3</u>	<u>4</u>
✓		✓	✓
✓	✓	✓	✓
✓		✓	✓
✓		✓	✓
✓	✓	✓	✓
		✓	✓
✓	✓	✓	✓
		✓	✓

**Extreme Sexual Dimorphism**

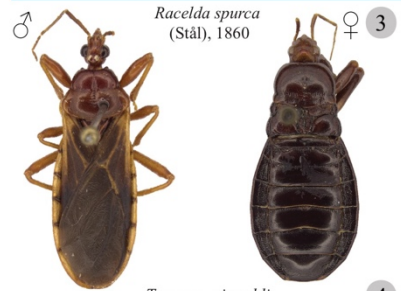


Figure 4.2. ML best tree (final Ln = -44,072.4153) based on 156 taxa (outgroups removed from figure) and a combined morphological and molecular dataset. Molecular data were divided into three partitions and models of sequence evolution: 16S, 28S D2, 28S D3–D5 (GTR+ $\Gamma$ +I); 18S, COI codon positions 1 and 2, Wg (GTR+ $\Gamma$ +I); COI codon position 3 (HKY+ $\Gamma$ +I). Morphological data were analyzed under Lewis' (2001) Mkv model of morphological evolution. Bootstrap values  $\geq 50$  are reported. Branch colors correspond to select clades mentioned in the discussion and serves as a reference to compare with all other molecular and/or morphological phylogenies in the supplementary materials.



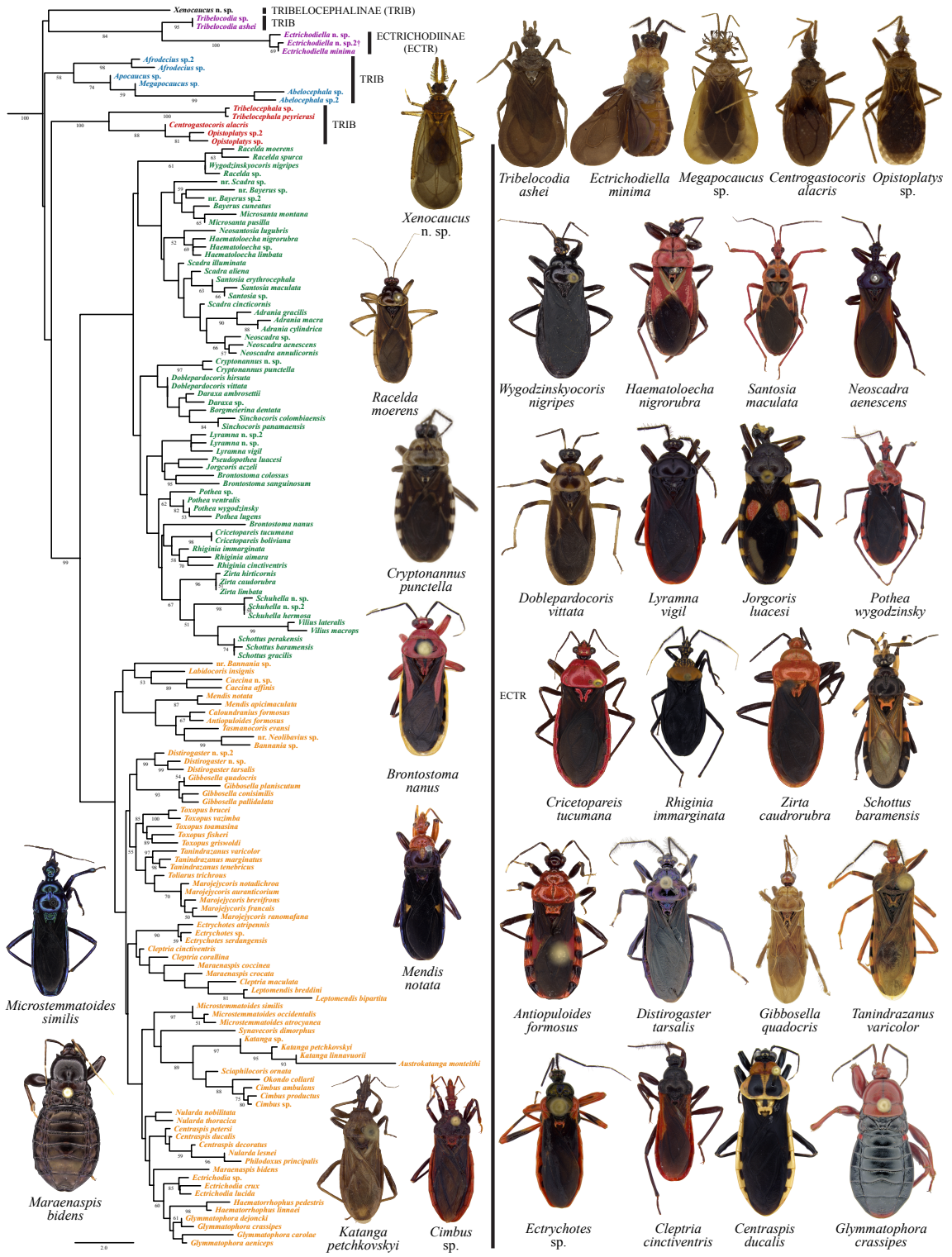


Figure 4.3. BayesTraits Multistate ML ancestral state reconstruction of color character 198 on the ML phylogram (Fig. 4.2) for Tribelocephalinae and Ectrichodiinae taxa (tree converted to cladogram for visual). Terminal taxa coded as cryptic (black) or either aposematic, metallic, or aposematic and metallic (blue). Pie charts at select nodes show probability values from the ML analysis and are used to visualize transitions between states; branches are colored to reflect the highest probability of a color state at that branch.

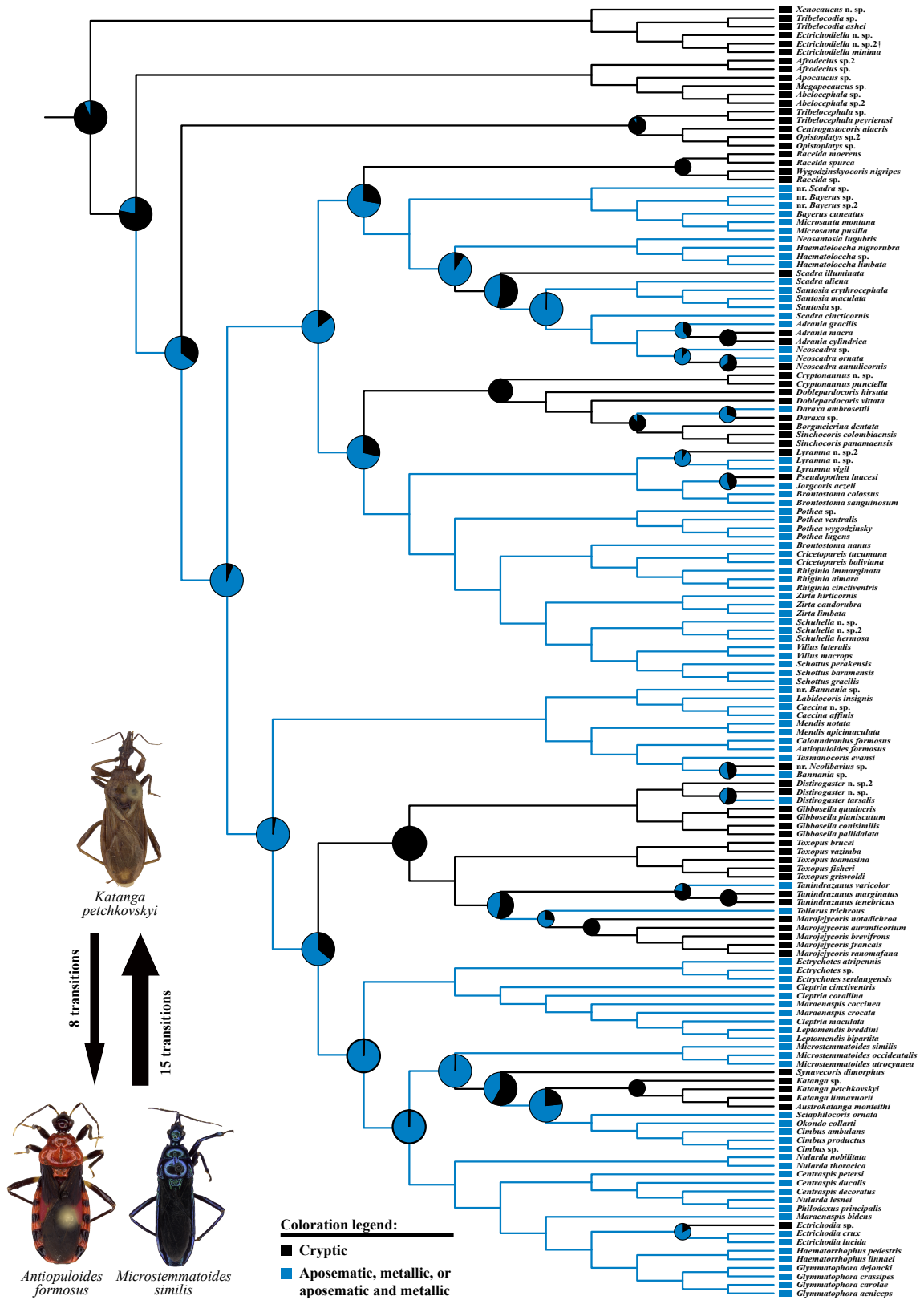


Figure 4.4. BayesTraits MultiState ML reconstruction of ancestral sexual dimorphism states on the ML phylogram (Fig. 4.2) for Tribelocephalinae and Ectrichodiinae taxa (tree converted to cladogram for visual). On the left, sexual dimorphism for terminal taxa is coded as limited (black), extreme (blue), polymorphic (gradient blue and black), or missing data (gray). Pie charts at select nodes show probability values from the ML reconstructions; branches are colored to reflect the highest probability of a color state at that branch; gray internal branches indicate ambiguity in reconstructions. On the right, color codes correspond to Fig. 4.1, with characters that exhibit no differences (black), differences in limited sexual dimorphism (green) or differences in extreme sexual dimorphism (blue) between males and females shown for species in which data on both sexes are known. For taxa that exhibit limited sexual dimorphism, individuals of a sex that exhibit no or minor differences (i.e., polymorphic) are coded gradient black and green. For taxa exhibiting both limited and extreme sexual dimorphism (i.e., males and/or females have different morphs), differences are indicated with gradient green and blue. Missing data indicated in gray.

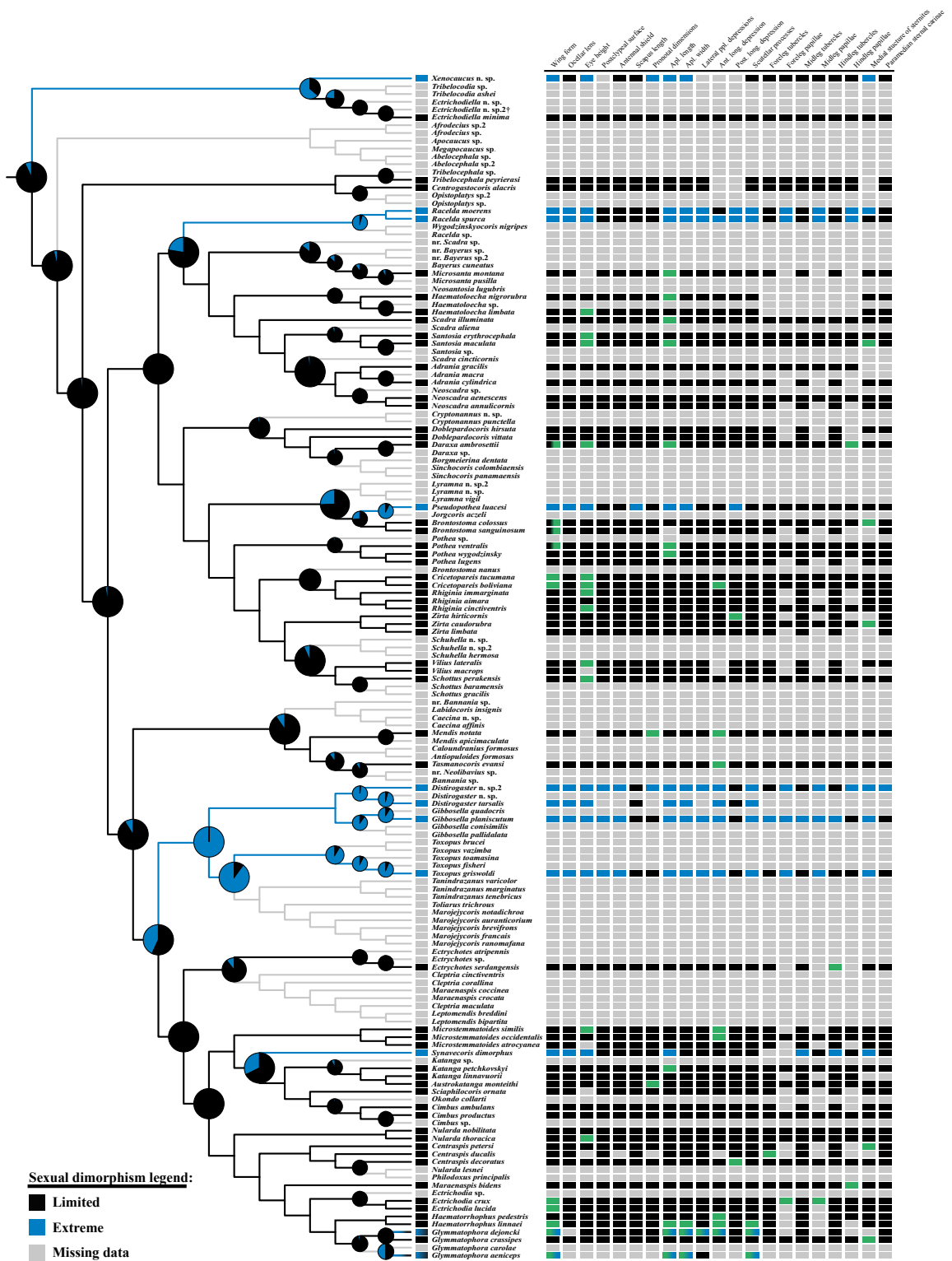


Figure 4.5. Combined morphological and molecular ML phylogram (converted to cladogram) showing new classification of Ectrichodiinae (see Fig. 4.2 for comparison to old classification). Unambiguous character optimizations are shown. A. Bulbous trichome of *Tribelocephala* sp. B. Rostral view of *Distriogaster tarsalis* head. C. Lateral view of *Tribelocephala* sp. head. D. Lateral antennal morphology of *Toxopus insignis* (modified from Forthman et al., in press). E. Dorsolateral-rostral view of *Afrodecius* sp. head and scapus. F. Apex of mandibular stylet in *Rhiginia cinctiventris*. G. Apex of right maxillary stylet in *Centraspis ducalis*. H. Apex of left maxillary stylet in *Rhiginia cinctiventris*. I. Thoracic morphology of *Ectrichodiella* n. sp., in dorsal view. J. Scutellum and postscutum of *Centrogastocoris alacris*. K. Thoracic morphology of nr. *Bannania* sp., in dorsal view. L. Thoracic sternal morphology in *Tanindraanus* sp. (modified from Forthman et al., in press). M. Metathoracic gland evaporatorium of *Tanindrazanus varicolor*, in lateral view. N.–P. Hemelytral morphology of *Ectrichodiella minima* (N.), *Tribelocephala* sp. (O.), and *Centrogastocoris alacris* (P.). Q. Foreleg morphology of *Ectrichodiella minima*. R. Dorsal abdominal glands of *Cricetopareis tucumana* female. S. Dorsal abdominal glands of *Ectrichodiella minima*. T. Ventral connexival suture and ventral abdominal morphology of *Tanindrazanus varicolor*. Character numbers and character state codings are listed in Table 4.1. Abbreviations used in figures: 1A, first anal vein; bf, basiflagellomere; cs, ventral connexival suture; Cu, cubitus; DAG I, dorsal abdominal gland I; DAG II, dorsal abdominal gland II; DAG III, dorsal abdominal gland III; df, distiflagellomere; M, media; mms, suture separating mandibular and maxillary plates; R, radius; s2, abdominal sternite II (fused sternites I and II); s3, abdominal sternite III; vlt, ventral laterotergites III.

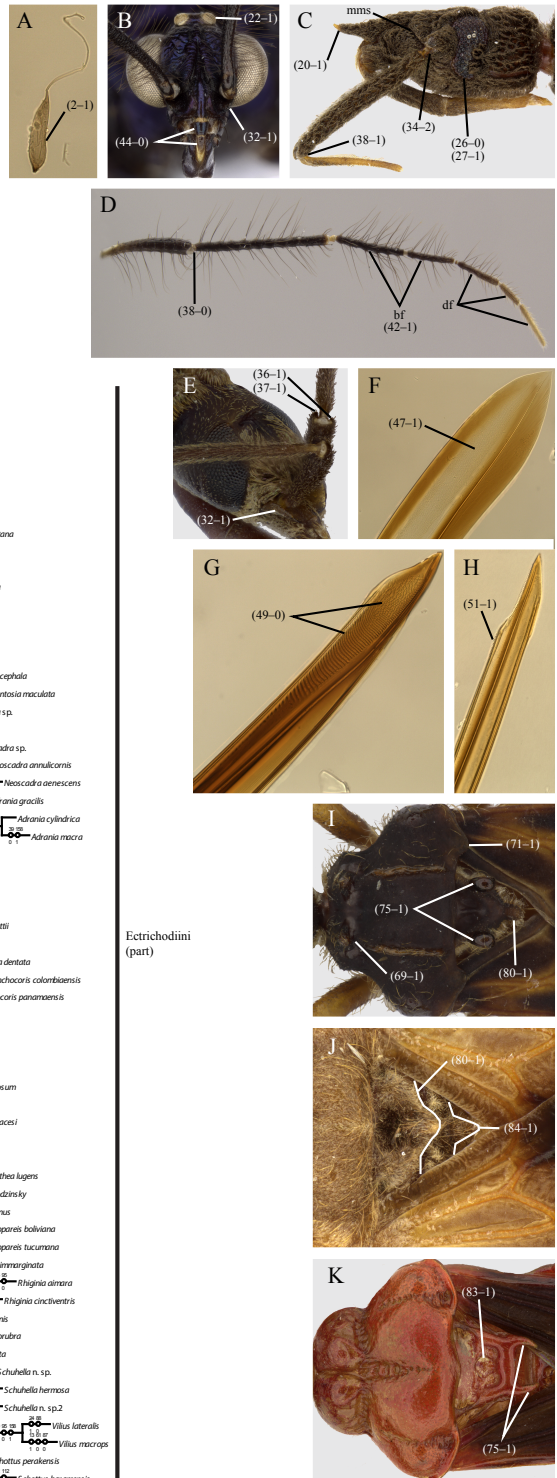
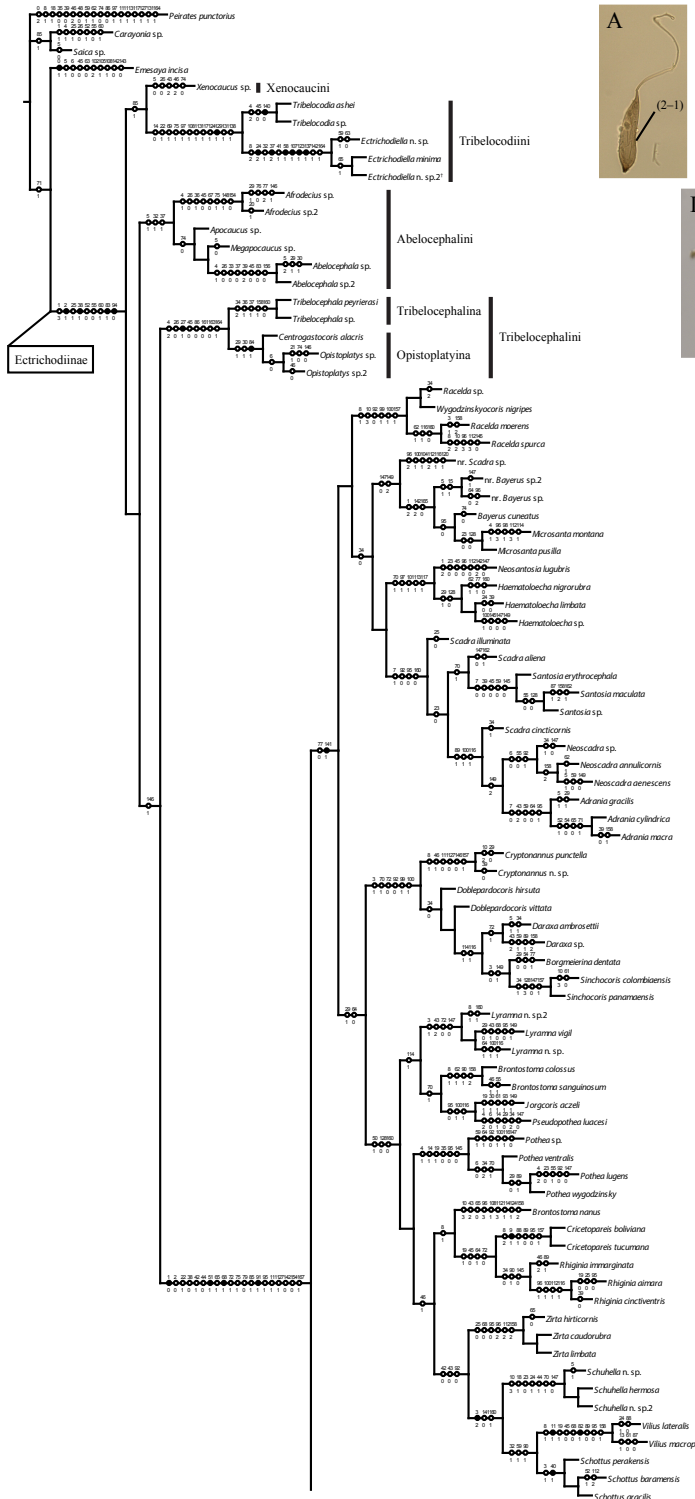
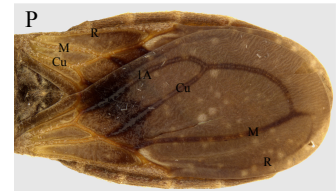
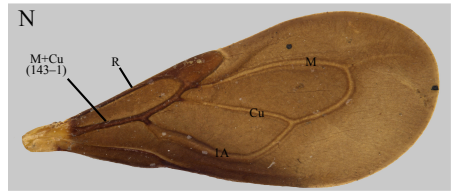
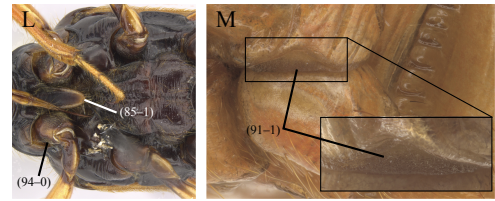
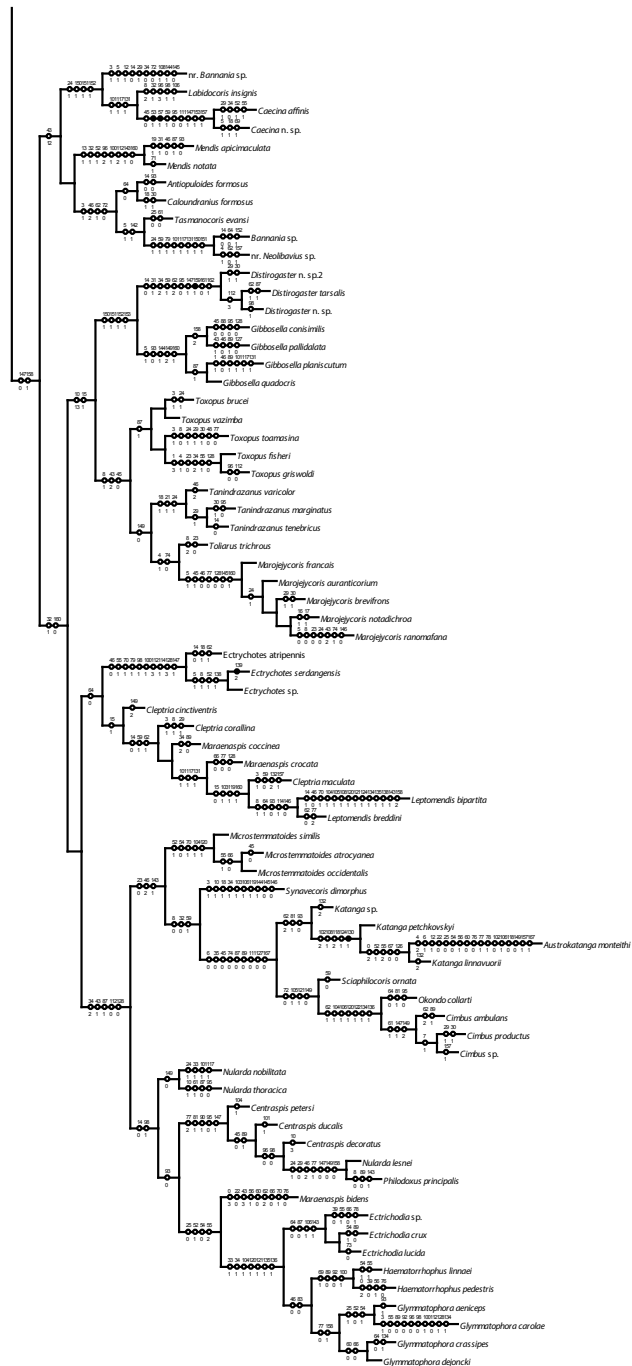


Figure 4.5. (Continued).





Ectrichodini (part)

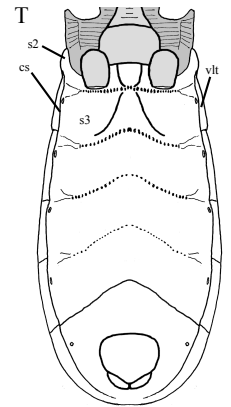
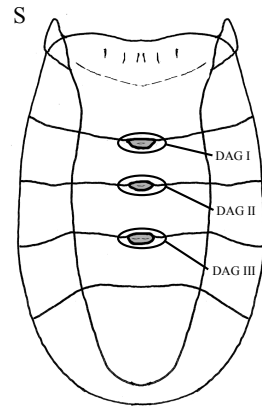
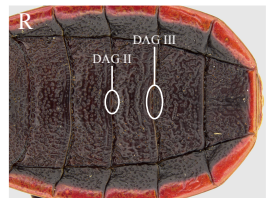


Table 4.1. Morphological characters and character state codings.

Character no.	Character	Character states
<b>GENERAL</b>		
0	Wing form	(0) Macropterous (reaches/surpasses tergite VII); (1) Brachypterous (reaches/surpasses tergite VI but not VII); (2) Micropterous (minute leathery wing pads covering, at most, base of abdomen); (3) Apterous
1	General vestiture, excluding hemelytra and legs	(0) Glabrous to sparse; (1) Moderately dense; (2) Dense; (3) Very dense, conspicuously thick pubescence obscuring integument on all or part of the body
2	Basally bulbous trichomes	(0) Absent; (1) Present
<b>HEAD</b>		
3	Head length (clypeal apex to anterior neck margin) relative to head width in dorsal view	(0) Longer than wide; (1) About as long as wide; (2) Wider than long
4	Head length relative to greatest pronotal length in dorsal view	(0) Shorter than pronotum; (1) About as long as pronotum; (2) Longer than pronotum
5	Anteocular length (clypeal apex to anterior eye margin) relative to postocular length (hind eye margin to anterior neck margin or, if postocular-neck constriction obsolete, halfway to anterior pronotal margin)	(0) Shorter than postocular; (1) About as long as postocular; (2) Longer than postocular
6	Postocular-neck constriction	(0) Absent, gradually transitioning; (1) Present, distinctly constricted
7	Shape of anterolateral neck area	(0) Flat; (1) Bulbous
8	Shape of posterolateral gula area in lateral view	(0) Flat to slightly swollen, conforming to rounded head shape; (1) Moderately swollen; (2) Very swollen
9	Extension of posterolateral gula area relative to ventral head margin	(0) Not or barely surpassing ventral head margin; (1) Distinctly surpassing ventral head margin
10	Ventral medial longitudinal depression	(0) Absent; (1) Restricted to anteocular region; (2) Restricted to postocular region; (3) Present along entire ventral length
11	Ventral paramedial longitudinal depression	(0) Absent; (1) Present
12	Small tubercles on ventromedial region of gula	(0) Absent; (1) Present
13	Ventrolateral tubercle on gula	(0) Absent; (1) Present
14	Postocular shape in dorsal view	(0) Narrow; (1) Broad
15	Postclypeal surface	(0) Flat or convex; at most a thin medial line visible but not distinctly depressed; (1) Distinctly depressed

Table 4.1. (Continued).

Character no.	Character	Character states
16	Extension of postclypeal depression	(0) Posterior clypeal margin to middle of synthlipsis; (1) Posterior clypeal margin to interocular sulcus
17	Width of postclypeal depression	(0) Narrow to slightly wide; (1) Very wide
18	Maxillary plate in lateral view	(0) Not reaching dorsal clypeal margin; (1) Reaching dorsal clypeal margin
19	Dorsal subtriangular elevation of clypeus	(0) Absent, clypeus rounded, sometimes with dorsally swollen area; (1) Present
20	Anterior clypeal process	(0) Absent; (1) Present
21	Clypeal apex relative to labral base	(0) Not elevated; (1) Elevated
22	Ocellar lens	(0) Absent; (1) Present
23	Location of ocelli	(0) On flat to shallowly elevated tubercle; (1) On moderately to extremely elevated cuticle
24	Shape of ocellar tubercle	(0) Convex; (1) Medially concaved, but ocelli appear on one tubercle; (2) Medially concaved, but ocelli appearing on separate tubercles
25	Eye height in lateral view	(0) Less than or half of head height; (1) More than half of head height
26	Eye shape	(0) Adpressed; (1) Globbose
27	Anterior eye margin in lateral view	(0) Convex; (1) Concave
28	Posterior eye margin in lateral view	(0) Concave; (1) Straight
29	Dorsal eye margin in lateral view	(0) Not attaining dorsal head surface; (1) Attaining dorsal head surface
30	Ventral eye margin in lateral view	(0) Not attaining dorsal head surface; (1) Attaining dorsal head surface
31	Antennifer armature	(0) Unarmed; (1) With dorsolateral tubercle
32	Antennal shield	(0) Absent, not expanded; (1) Present, expanded
33	Antennal insertion site in lateral view	(0) Not concealed by antennal shield; (1) Concealed by antennal shield
34	Point of antennal insertion relative to suture between mandibular and maxillary plates in lateral view	(0) Dorsal, ventrally exposing suture between mandibular and maxillary plates; (1) Lateral, largely obscuring suture between mandibular and maxillary plates; (2) Ventral, dorsally exposing suture between mandibular and maxillary plates
35	Scapus length	(0) Not surpassing or barely surpassing clypeal apex; (1) Surpassing clypeal apex
36	Ventroanterior apical lobe of scapus	(0) Absent; (1) Present

Table 4.1. (Continued).

Character no.	Character	Character states
37	Ventroposterior apical armature of scapus	(0) Absent; (1) Lobed; (2) Toothlike
38	Scapo-pedicellar articulation	(0) Pedicellus distally or, at most, slightly ventrally inserted on scape; (1) Pedicellus ventrally inserted on scape, so that pedicellus and flagellomeres point posteriorly
39	Curvature of pedicellus	(0) Straight; (1) Slightly to moderately curved; (2) Very curved
40	Shape of pedicellus	(0) Cylindrical; (1) Laterally flattened
41	Diameter of flagellomeres	(0) Similar to diameter of pedicellus; (1) More slender than diameter of pedicellus
42	Basiflagellomere segmentation	(0) 1-segmented; (1) 2-segmented
43	Distiflagellomere segmentation	(0) 1-segmented; (1) 2-segmented; (2) 3-segmented; (3) 4-segmented; (4) 5-segmented
44	Labrum structural	(0) Subdivided by transverse membrane; (1) Completely sclerotized
45	Ventral margin (appears dorsal when reflexed) of labial segment III (second visible segment)	(0) Straight or concave; (1) Convex
46	Labial segment III length	(0) Shorter than labial segment II; (1) Subequal to labial segment II; (2) Longer than labial segment II
47	Number of transverse ridges on mandibular stylets in external view	(0) Up to 35; (1) More than 35
48	Dorsal row of processes on right maxillary stylet in internal view	(0) Absent; (1) Present
49	Structure of ventral processes of right maxillary stylet in internal view	(0) Lamellate; (1) Hairlike; (2) Tootlike
50	Dorsal row of processes on left maxillary stylet in internal view	(0) Absent; (1) Present
51	Ventral row of processes on left maxillary stylet in internal view	(0) Absent; (1) Present
<b>THORAX</b>		
52	Greatest length of pronotum relative to greatest width	(0) Wider than long; (1) About as long as wide; (2) Longer than wide
53	Pronotal structure	(0) Smooth, striated, rugose, punctate, or any combination of; (1) entirely tuberculate/spinulose
54	Pronotal collar	(0) Absent; (1) Present
55	Anterior pronotal lobe greatest length relative to posterior pronotal lobe	(0) Shorter than posterior lobe; (1) About as long as posterior lobe; (2) Longer than posterior lobe

Table 4.1. (Continued).

Character no.	Character	Character states
56	Anterior pronotal lobe greatest width relative to posterior pronotal lobe	(0) Less than width of posterior lobe; (1) About as wide as posterior lobe
57	Medial expansion of anterior pronotal lobe	(0) Not medially expanded; (1) Medially expanded anteriorly
58	Medial discal tubercles on anterior paramedial pronotal lobes	(0) Absent; (1) Present
59	Lateral carinae on anterior pronotal lobe	(0) Absent; (1) Present
60	Lateral depressions on posterior pronotal lobe	(0) Absent; (1) Present
61	Structure of lateral depressions on posterior pronotal lobe	(0) Not foveate; (1) Foveate
62	Anterolateral armature of pronotum	(0) Absent; (1) Small tubercle; (2) Strong projecting process
63	Presence of pronotal longitudinal depression on anterior lobe	(0) Absent; (1) Present
64	Extension of pronotal longitudinal depression on anterior lobe	(0) Not reaching anterior margin; (1) Reaching or nearly reaching anterior margin
65	Presence of pronotal longitudinal depression on posterior lobe	(0) Absent; (1) Present
66	Structure of pronotal longitudinal depression on posterior lobe	(0) Not foveate; (1) Foveate
67	Pronotal transverse suture development	(0) Absent or weakly developed, anterior and posterior pronotal lobes at most weakly delimited; (1) Well-developed, anterior and posterior pronotal lobes distinctly delimited
68	Paramedial division of pronotal transverse suture	(0) Absent; (1) Present
69	Lateral division of pronotal transverse suture	(0) Absent; (1) Present
70	Pronotal transverse suture structure	(0) Not foveate; (1) Foveate
71	Posteriolateral margin of pronotum	(0) Not or slightly surpassing medial posterior margin of pronotum; (1) Distinctly surpassing medial posterior margin of pronotum
72	Expansion of mesopleuron	(0) Not expanded anterodorsally onto lateral side of pronotum; (1) Expanded anterodorsally onto lateral side of pronotum, overlapping pronotum
73	Shape of mesopleural expansion	(0) Knoblike; (1) Broad, sinuate
74	Metepisternal supracoxal lobe of mesocoxal cavity	(0) Absent; (1) Present
75	Paramedian scutellar processes	(0) Absent; (1) Present

Table 4.1. (Continued).

Character no.	Character	Character states
76	Development of paramedian scutellar processes	(0) Weakly developed; (1) Well developed
77	Distance between paramedian scutellar processes	(0) Extremely narrow, apices nearly contiguous, less than one-tenth of posterior pronotal lobe width; (1) Moderately broad, between one-tenth to one-fourth posterior pronotal lobe width; (2) Very broad, more than one-fourth posterior pronotal lobe width
78	Orientation of paramedian scutellar processes	(0) Horizontally directed; (1) Dorsally directed
79	Presence of single medial scutellar process	(0) Absent; (1) Present
80	Development of medial scutellar process	(0) Weakly developed; (1) Well developed
81	Presence of two medial scutellar processes	(0) Absent; (1) Present
82	Presence of lateral scutellar processes	(0) Absent; (1) Present
83	Scutellar disc surface	(0) Flat; (1) Medially depressed
84	Postscutum development	(0) Absent or reduced, not prominently projecting posteriorly; (1) Present, prominently projecting posteriorly
85	Prosternal stridulatory process length	(0) Short, not reaching or surpassing posterior margin of procoxal cavity; (1) Long, surpassing posterior margin of procoxal cavity
86	Apical shape of prosternal stridulatory process	(0) Acute; (1) Rounded, blunt
87	Transverse suture between meso- and metasterna	(0) Complete; (1) Incomplete, at least partially
88	Medial surface of mesosternum	(0) Flat or convex; (1) Anteriorly depressed; (2) Depressed along entire length
89	Medial surface of metasternum	(0) Flat or convex; (1) Posteriorly depressed; (2) Depressed along entire length
90	Metathoracic gland callus in lateral view	(0) Absent; (1) Present
91	Metathoracic gland evaporatorium	(0) Absent, cuticle along metaxocal rim not modified into ridges or meshlike structures; (1) Present, cuticle along metaxocal rim modified into ridges or meshlike structures
92	Cuticular structure of metathoracic gland evaporatorium	(0) Faint to distinct ridges without meshlike structure; (1) Meshlike structure
93	Metathoracic gland evaporatorium size	(0) Small, not visible in lateral view; (1) Large, visible in lateral view
<b>LEG</b>		
94	Shape of forecoxa	(0) Globular; (1) Elongate

Table 4.1. (Continued).

Character no.	Character	Character states
95	Width of forefemur relative to mid femur	(0) About as wide as mid femur; (1) Wider than mid femur
96	Ventral vestiture of foretrochanter	(0) Glabrous; (1) Long, slender setae; (2) Short, thick setae; (3) Round to rodlike papillae
97	Presence of small tubercles on foretrochanter	(0) Absent; (1) Present
98	Presence of ventral papillae on forefemur	(0) Absent; (1) Present
99	Distribution of papillae on forefemur	(0) Basally; (1) Longitudinally along at least half the length of the femur
100	Presence of ventral longitudinal carinae on forefemur	(0) Absent; (1) Present
101	Presence of small tubercles on forefemur	(0) Absent; (1) Present
102	Distribution of small tubercles on forefemur	(0) Entirely throughout femur; (1) Throughout ventral femoral surface; (2) Restricted to ventroapical half of femur
103	Vestiture of small tubercles on forefemur	(0) Glabrous; (1) With setae
104	Medial projection of forefemur	(0) Absent; (1) Present
105	Anterior subapical projection on forefemur	(0) Absent; (1) Present
106	Posterior subapical projection on forefemur	(0) Absent; (1) Present
107	Paired ventroapical projections on forefemur	(0) Absent; (1) Present
108	Presence of small tubercles on foretibia	(0) Absent; (1) Present
109	Distribution of small tubercles on foretibia	(0) Entirely throughout tibia; (1) Throughout ventral tibial surface; (2) Restricted to ventroapical half of tibia
110	Vestiture of small tubercles on foretibia	(0) Glabrous; (1) With setae
111	Presence of fossula spongiosa on foretibia	(0) Absent or reduced; (1) Present
112	Ventral vestiture of mid trochanter	(0) Glabrous; (1) Long, slender setae; (2) Short, thick setae; (3) Round to rodlike papillae
113	Presence of small tubercles on mid trochanter	(0) Absent; (1) Present
114	Presence of ventral papillae on mid femur	(0) Absent; (1) Present
115	Distribution of papillae on mid femur	(0) Basally; (1) Longitudinally along at least half the length of the femur
116	Presence of ventral longitudinal carinae on mid femur	(0) Absent; (1) Present
117	Presence of small tubercles on mid femur	(0) Absent; (1) Present

Table 4.1. (Continued).

Character no.	Character	Character states
118	Distribution of small tubercles on mid femur	(0) Entirely throughout femur; (1) Throughout ventral femoral surface; (2) Restricted to ventroapical half of femur
119	Vestiture of small tubercles on mid femur	(0) Glabrous; (1) With setae
120	Medial projection of mid femur	(0) Absent; (1) Present
121	Anterior subapical projection on mid femur	(0) Absent; (1) Present
122	Posterior subapical projection on mid femur	(0) Absent; (1) Present
123	Paired ventroapical projections on mid femur	(0) Absent; (1) Present
124	Presence of small tubercles on mid tibia	(0) Absent; (1) Present
125	Distribution of small tubercles on mid tibia	(0) Entirely throughout tibia; (1) Throughout ventral tibial surface; (2) Restricted to ventroapical half of tibia
126	Vestiture of small tubercles on mid tibia	(0) Glabrous; (1) With setae
127	Presence of fossula spongiosa on mid tibia	(0) Absent or reduced; (1) Present
128	Ventral vestiture of hind trochanter	(0) Glabrous; (1) Long, slender setae; (2) Short, thick setae; (3) Round to rodlike papillae
129	Presence of small tubercles on hind trochanter	(0) Absent; (1) Present
130	Basal curvature of hind femur	(0) Not curved or slightly curved posteriorad; (1) Extremely curved posteriorad, almost from 45 to 90 degree angle
131	Presence of small tubercles on hind femur	(0) Absent; (1) Present
132	Distribution of small tubercles on hind femur	(0) Entirely throughout femur; (1) Throughout ventral femoral surface; (2) Restricted to ventroapical half of femur
133	Vestiture of small tubercles on hind femur	(0) Glabrous; (1) With setae
134	Medial projection of hind femur	(0) Absent; (1) Present
135	Anterior subapical projection on hind femur	(0) Absent; (1) Present
136	Posterior subapical projection on hind femur	(0) Absent; (1) Present
137	Paired ventroapical projections on hind femur	(0) Absent; (1) Present
138	Presence of small tubercles on hind tibia	(0) Absent; (1) Present
139	Distribution of small tubercles on hind tibia	(0) Entirely throughout tibia; (1) Throughout ventral tibial surface; (2) Restricted to ventroapical half of tibia
140	Tarsal segmentation	(0) 1-segmented; (1) 2-segmented; (2) 3-segmented



Table 4.1. (Continued).

Character no.	Character	Character states
<b>FOREWING</b>		
141	Extension of corium	(0) Restricted to areas adjacent to basal wing veins and with pterostigmalike appearance; (1) Well-developed corium
142	Vestiture of corium	(0) Long, simple setae absent; (1) Long, simple setae sparse; (2) Long, simple setae dense
143	Proximal part of M and Cu	(0) Separate veins; (1) Forming one vein, at least partially
144	Distal part of M and Cu	(0) Separate veins; (1) Fused basally
145	Extension of distal part of M	(0) Absent; (1) Present
146	Distal part of R	(0) Absent; (1) Present
147	Development of distal R cell	(0) Incomplete; (1) Complete, although may appear faint apically
<b>ABDOMEN</b>		
148	Division of dorsal synlaterotergite I+II	(0) Not divided; (1) Divided
149	Expansion of dorsal synlaterotergite I+II	(0) Not expanded; (1) Laterally expanded; (2) Posterior tubercle
150	Dorsal laterotergite III armature	(0) Unarmed; (1) Posterior tubercle
151	Dorsal laterotergite IV armature	(0) Unarmed; (1) Posterior tubercle
152	Dorsal laterotergite V armature	(0) Unarmed; (1) Posterior tubercle
153	Dorsal laterotergite VI armature	(0) Unarmed; (1) Posterior tubercle
154	Ventral connexival suture on sternite II	(0) Absent; (1) Present
155	Carination of intersegmental suture between sternites II and III	(0) Not carinate; (1) Carinate
156	Development of intersegmental sutures between III and VI	(0) Absent or reduced; (1) Well-developed
157	Sternal surface	(0) Smooth or slightly wrinkled; (1) Punctate, at least laterally
158	Medial structure of sternites	(0) Flat; (1) Medially depressed, at least partially; (2) Convex or keeled, at least partially
159	Paramedian sternal carinae	(0) Absent; (1) Present
160	Spiracle shape	(0) Ovoid or elliptical; (1) Circular
<b>MALE GENITALIA</b>		
161	Anteroposterior thickness of median pygophore process	(0) Flattened; (1) Thickened, at least partially

Table 4.1. (Continued).

Character no.	Character	Character states
162	Lateral thickness of median pygophore process	(0) Flattened; (1) Thickened, at least partially
<b>GLANDS</b>		
163	Presence of meshlike Brindley's gland evaporatorium	(0) Absent; (1) Present
164	Dorsal abdominal scent gland (DAG) I ostioles in adults	(0) Absent; (1) Present
165	DAG II ostioles in adults	(0) Absent; (1) Present
166	DAG III ostioles in adults	(0) Absent; (1) Present
167	Size of DAG III ostioles relative to DAGs I and II ostioles in adults	(0) Similar size; (1) Larger
<b>FEMALE</b>		
168	Wing form	(0) Macropterous (reaches/surpasses tergite VII); (1) Brachypterous (reaches/surpasses tergite VI but not VII); (2) Micropterous (minute leathery wing pads covering, at most, base of abdomen); (3) Apterous
169	Ocellar lens	(0) Absent; (1) Present
170	Eye height in lateral view	(0) Less than or half of head height; (1) More than half of head height; (2) Absent
171	Postclypeal surface	(0) Flat or convex; at most a thin medial line visible but not distinctly depressed; (1) Distinctly depressed
172	Antennal shield	(0) Absent, not expanded; (1) Present, expanded
173	Scapus length	(0) Not surpassing or barely surpassing clypeal apex; (1) Surpassing clypeal apex
174	Greatest length of pronotum relative to greatest width	(0) Wider than long; (1) About as long as wide; (2) Longer than wide
175	Anterior pronotal lobe greatest length relative to posterior pronotal lobe	(0) Shorter than posterior lobe; (1) About as long as posterior lobe; (2) Longer than posterior lobe
176	Anterior pronotal lobe greatest width relative to posterior pronotal lobe	(0) Less than width of posterior lobe; (1) About as wide as posterior lobe
177	Lateral depressions on posterior pronotal lobe	(0) Absent; (1) Present
178	Extension of pronotal longitudinal depression on anterior lobe	(0) Not reaching anterior margin; (1) Reaching or nearly reaching anterior margin; (2) Absent or reduce
179	Presence of pronotal longitudinal depression on posterior lobe	(0) Absent; (1) Present

Table 4.1. (Continued).

Character no.	Character	Character states
180	Development of paramedian scutellar processes	(0) Weakly developed; (1) Well developed; (2) Absent
181	Ventral vestiture of foretrochanter	(0) Glabrous; (1) Long, slender setae; (2) Short, thick setae; (3) Round to rodlike papillae
182	Presence of small tubercles on forefemur	(0) Absent; (1) Present
183	Presence of ventral papillae on forefemur	(0) Absent; (1) Present
184	Presence of small tubercles on foretibia	(0) Absent; (1) Present
185	Presence of ventral papillae on fore tibia	(0) Absent; (1) Present
186	Ventral vestiture of mid trochanter	(0) Glabrous; (1) Long, slender setae; (2) Short, thick setae; (3) Round to rodlike papillae
187	Presence of small tubercles on mid femur	(0) Absent; (1) Present
188	Presence of ventral papillae on mid femur	(0) Absent; (1) Present
189	Presence of small tubercles on mid tibia	(0) Absent; (1) Present
190	Presence of ventral papillae on mid tibia	(0) Absent; (1) Present
191	Ventral vestiture of hind trochanter	(0) Glabrous; (1) Long, slender setae; (2) Short, thick setae; (3) Round to rodlike papillae
192	Presence of small tubercles on hind femur	(0) Absent; (1) Present
193	Presence of ventral papillae on hind femur	(0) Absent; (1) Present
194	Presence of small tubercles on hind tibia	(0) Absent; (1) Present
195	Presence of ventral papillae on hind tibia	(0) Absent; (1) Present
196	Medial structure of sternites	(0) Flat; (1) Medially depressed, at least partially; (2) Convex or keeled, at least partially
197	Paramedian sternal carinae	(0) Absent; (1) Present
<b>COLORATION</b>		
198	Coloration coding 1	(0) Cryptic; (1) Aposematic, uniform metallic, or aposematic and metallic
199	Coloration coding 2	(0) Cryptic and uniform metallic; (1) Aposematic or aposematic and metallic
<b>SEXUAL DIMORPHISM</b>		
200	Sexual dimorphism	(0) Limited; (1) Extreme
<b>MALE BODY LENGTH</b>		
201	Male body length from clypeal apex to posterior abdominal margin	Continuous values

Table 4.2. Taxon sampling, USI codes, ethanol molecular specimen collection numbers (RCW), depositories (Dep.), and GenBank accession numbers for sequenced specimens. Asterisks indicate sequences retrieved from GenBank.

Subfamily	Taxon	USI	RCW	Dep.	GenBank accession no.						
					16S	18S	28S D2	28S D3–D5	COI	Wg	
Ectrichodimae	<i>Bannania</i> sp.	UCR_ENT 00119027	3064	UCR	KU764537	KT221910*	KT221939*	KT221968*	KT221890*		
	nr <i>Bannania</i> sp.	UCR_ENT 00004465	609	UCR		KT221937*	KT221966*	KT221996*	KT221908*		
	nr <i>Bayerus</i> sp.	UCR_ENT 00004096	2001	UCR			KU764586	KU764601	KU764610		
	nr <i>Bayerus</i> sp.2	UCR_ENT 00004102	1969	UCR			KU764585	KU764600			
	<i>Brontostoma colossus</i>	UCR_ENT 00119017	3049	UCR	KU764538	KT221911*	KT221940*	KT221969*	KT221891*	KU764611	
	<i>Brontostoma nanus</i>	UCR_ENT 00123508	4030	UCR	KU764539		KU764574	KU764589	KU764603	KU764612	
	<i>Brontostoma sanguinosum</i>	UCR_ENT 00002617	1243	UCR	KU764540	KT221912*		KT221970*	KT221892*	KU764613	
	<i>Caecina</i> affinis	UCR_ENT 00004440	2636	UCR	KU764541	KT221913*	KT221941*	KT221971*	KT221893*		
	<i>Caecina</i> n. sp.	UCR_ENT 00123509	3959	UCR	KU764542		KU764575	KU764590			
	<i>Centraspis ducalis</i>	UCR_ENT 00119021	3014	UCR	KU764543	KT221914*	KT221942*	KT221972*	KT221894*	KU764614	
	<i>Cimbus</i> sp.	UCR_ENT 00123510	4722	UCR		KU764569	KU764576	KU764591	KU764604	KU764615	
	<i>Cleptria corallina</i>	AMNH_PBI 00218770	14	UCR	FJ230388*	FJ230462*	FJ230543*	FJ230621*, FJ230700*		JQ897871*	
	<i>Cryptonannus</i> n. sp.	UCR_ENT 00002757	1433	UCR	KU764544	KT221915*		KT221973*	KT221895*	KU764616	
	<i>Daraxa</i> sp.	UCR_ENT 00119016	3076	UCR	KU764564	KT221938*	KT221967*	KT221997*	KT221909*		

Table 4.2. (Continued).

Subfamily	Taxon	USI	RCW	Dep.	16S	18S	GenBank accession no.			Wg
							28S D2	28S D3-D5	COI	
	<i>Distrogaster tarsalis</i>	UCR_ENT 00006366	2898	CAS		KT221918*	KT221945*	KT221976*	KR606396*	
	<i>Distrogaster</i> n. sp.	UCR_ENT 00088090	3018	UCR		KT221917*	KT221944*	KT221975*	KR606395*	
	<i>Distrogaster</i> n. sp.2	UCR_ENT 00007158	2881	UCR		KT221916*	KT221943*	KT221974*	KR606393*	
	<i>Ectrichodia crux</i>	UCR_ENT 00119022	3026	UCR		KT221919*	KT221946*	KT221977*	KT221896*	KU764617
	<i>Ectrichodia lucida</i>	AMNH_PBI 00218769	13	UCR	FJ230387*	FJ230461*	FJ230542*	FJ230620*, FJ230699*		JQ897878*
	<i>Ectrichodia</i> sp.	UCR_ENT 00123511	3715	UCR	KU764545		KU764577	KU764592	KU764605	KU764618
	<i>Ectrychotes atripennis</i>	AMNH_PBI 00218932	188	UCR	FJ230424*	FJ230503*	FJ230584*	FJ230661*, FJ230740*	JQ942322*	JQ897879*
	<i>Ectrychotes serdangensis</i>	AMNH_PBI 00218830	76	UCR		FJ230479*	FJ230560*	FJ230638*, FJ230717*		
	<i>Ectrychotes</i> sp.	UCR_ENT 00119028	571	UCR	KU764546	KT221920*	KT221947*	KT221978*	KT221897*	KU764619
	<i>Glymmatophora crassipes</i>	UCR_ENT 00088087	3020	UCR	KU764547	KT221921*	KT221948*	KT221979*	KR606415*	KU764620
	<i>Haematoloecha</i> sp.	UCR_ENT 00123512	3956	UCR	KU764548		KU764578	KU764593	KU764606	KU764621
	<i>Katanga</i> sp.	UCR_ENT 00123513	4029	UCR	KU764549		KU764579	KU764594	KU764607	KU764622
	<i>Maraenaspis coccinea</i>	AMNH_PBI 00218772	16	UCR	FJ230389*	FJ230463*	FJ230544*			JQ897889*
	<i>Marojejycoris brevifrons</i>	UCR_ENT 00006480	2923	CAS	KU764550	KT221922*	KU764580	KT221980*	KR606417*	KU764623

Table 4.2. (Continued).

Subfamily	Taxon	USI	RCW	Dep.	GenBank accession no.							Wg
					16S	18S	28S D2	28S D3-D5	COI			
	<i>Mendis apicimaculata</i>	UCR_ENT 00119026	2647	UCR	KU764551	KT221923*	KT221949*	KT221981*	KT221898*	KU764624		
	<i>Microstemmatoides atrocyanea</i>	UCR_ENT 00119029	3029	UCR	KU764552	KT221924*	KT221950*	KT221982*	KT221899*			
	nr <i>Neolibavius</i> sp.	UCR_ENT 00003965	1830	UCR	KU764565	KT221925*	KT221951*	KT221983*	KT221900*			
	<i>Neoscadra</i> sp.	UCR_ENT 00004990	2139	UCR	KU764553		KU764581	KU764595	KU764608	KU764625		
	<i>Pothea lugens</i>	UCR_ENT 00119018	3304	UCR	KU764554	KT221927*	KT221953*		KT221902*	KU764626		
	<i>Pothea</i> sp.	UCR_ENT 00123514	4028	UCR	KU764555		KU764582	KU764596	KU764609			
	<i>Racelda</i> sp.	AMNH_PBI 00218801	41	UCR	FJ230398*	FJ230472*	FJ230553*	FJ230631*, FJ230710*		JQ897915*		
	<i>Rhiginia aimara</i>	UCR_ENT 00119019	3077	UCR	KU764556	KT221928*	KT221954*		KT221903*	KU764627		
	<i>Rhiginia cinciventris</i>	AMNH_PBI 00218891	139	UCR	FJ230410*	FJ230490*	FJ230571*	FJ230648*, FJ230727*	JQ942333*	JQ897917*		
	nr <i>Scadra</i> sp.	UCR_ENT 00052213	699	UCR	JQ897793*		JQ897634*	JQ897710*		JQ897880*		
	<i>Santosia</i> sp.	UCR_ENT 00004138	2046	UCR	KU764557	KT221930*	KT221956*	KT221986*	KT221905*	KU764628		
	<i>Tanindrazanus marginatus</i>	UCR_ENT 00006887	2902	CAS	KU764558	KT221931*	KT221957*	KT221987*	KR606423*			
	<i>Tanindrazanus tenebricus</i>	UCR_ENT 00006723	2931	CAS	KU764559	KT221932*	KT221958*	KT221988*	KR606430*			
	<i>Tanindrazanus varicolor</i>	UCR_ENT 00006482	2949	UCR		KT221933*	KT221959*	KT221989*	KR606433*			

Table 4.2. (Continued).

Subfamily	Taxon	USI	RCW	Dep.	16S	18S	GenBank accession no.			Wg
							28S D2	28S D3–D5	COI	
	<i>Toxopneustes brucei</i>	UCR_ENT 00045338	2901	UCR			KT221960*	KT221990*	KR606438*	
	<i>Toxopneustes fisheri</i>	UCR_ENT 00045431	2910	CAS	KU764560		KT221961*	KT221991*	KR606446*	
	<i>Toxopneustes griswoldi</i>	UCR_ENT 00006435	2916	CAS		KT221934*	KT221962*	KT221992*	KR606448*	
	<i>Toxopneustes toamasina</i>	UCR_ENT 00007056	2951	SU			KT221963*	KU764597	KR606457*	
	<i>Toxopneustes vazimba</i>	UCR_ENT 00006472	2953	CAS	KU764561	KT221935*	KT221964*	KT221993*	KR606462*	
Emesinae	<i>Emesaya incisa</i>	AMNH_PBI 00219017	282	UCR	FJ230436*	FJ230515*	FJ230598*	FJ230672*, FJ230751*	JQ942323*	
Peiratinae	<i>Peirates punctatus</i>	AMNH_PBI 00218960	216	UCR	FJ230430*	FJ230508*	FJ230590*	FJ230666*, FJ230745*		
Saicinae	<i>Saica</i> sp.	AMNH_PBI 00218796	42	UCR	FJ230399*	FJ230473*	FJ230554*	FJ230632*, FJ230711*		
Tribelocephalinae	<i>Abelocephala</i> sp.	UCR_ENT 00106432	3308	UCR	KU764536	KU764567	KU764572	KU764588	KU764602	
	<i>Abelocephala</i> sp.2	UCR_ENT 00107032	3963	UCR		KU764566	KU764571	KU764587		
	<i>Afrodecia</i> sp.	UCR_ENT 00092725	4711	SAMC		KU764568	KU764573			
	<i>Opisthoplatys</i> sp.	UCR_ENT 00119024	3056	UCR		KT221926*		KT221984*		
	<i>Opisthoplatys</i> sp.2	UCR_ENT 00052187	1592	UCR	JQ897851*	JQ897612*	JQ897682*	JQ897767*		
	<i>Tribelocephala peyeriasi</i>	AMNH_PBI 00219033	287	CAS	FJ230440*	FJ230521*	FJ230601*			

Table 4.2. (Continued).

Subfamily	Taxon	USI	RCW	Dep.	16S	18S	GenBank accession no.			
							28S D2	28S D3-D5	COI	Wg
	<i>Tribeloccephala</i> sp.	UCR_ENT 00119023	3048	UCR	KU764562		KT221965*	KT221994*	KT221906*	
	<i>Tribelocodia</i> sp.	UCR_ENT 00122389	4721	UCR		KU764570	KU764583	KU764598		
	<i>Xenocaucus</i> n. sp.	UCR_ENT 00020656	4707	CNC	KU764563		KU764584	KU764599		



Table 4.3. Phylogenetic logistic regression parameter estimates for the effects of log-transformed male body length on coloration in 133 Ectrichodiinae and Tribelocephalinae species. Aposematic, uniform metallic, and aposematic and metallic color patterns are treated as one state with cryptic coloration as a separate state. Parameters are phylogenetic signal ( $a$ ), intercept ( $b_0$ ), and the regression coefficient ( $b_1$ ), and p-values  $\leq 0.05$  are indicated in bold. Abbreviations: Phylo, phylogram; Ultra0, ultrametric tree generated with penalty set to 0; Ultra10, ultrametric tree generated with penalty set to 10; Ultra10000, ultrametric tree generated with penalty set to 10000; Par, parameter; SE, standard error; BS, bootstrap; C, converged; NC, non-converged.

Tree/dataset	Par	Estimate	SE	p-value	BS mean (C and NC) <sup>a</sup>	BS confidence interval (C and NC) <sup>a</sup>	BS p-value (C and NC) <sup>a</sup>	BS mean (C only)	BS confidence interval (C only)
Phylo TA+UM									
	$a$	-1.1535			-1.5274	(-2.8656, -0.3581)	<b>0.0088</b>	-1.4888	(-2.8745, -0.2869)
	$b_0$	0.3058	0.4420	0.4902	0.1196	(-2.2624, 2.3698)	0.8072	0.0866	(-2.2315, 2.0630)
	$b_1$	0.8857	0.2498	<b>0.0005</b>	0.4773	(-2.5222, 2.7345)	0.4456	0.4658	(-2.3864, 2.3871)
Ultra0 TA+UM									
	$a$	-0.6208			-0.0933	(-3.4576, -0.4597)	<b>0.0180</b>	-0.9365	(-3.4482, -0.4129)
	$b_0$	-0.2458	0.5559	0.6592	-0.2911	(-1.5741, 1.0498)	0.6160	-0.2907	(-1.5610, 1.0496)
	$b_1$	0.7983	0.2325	<b>0.0008</b>	0.8945	(-0.3999, 1.6021)	<b>0.0008</b>	0.8944	(-0.4014, 1.6000)
Ultra10 TA+UM									
	$a$	-0.0238			-0.5099	(-3.9681, 1.0646)	<b>0.0240</b>	-0.5153	(-3.8989, 1.0510)
	$b_0$	-0.1197	0.6771	0.8598	-0.1841	(-1.5967, 1.3802)	0.7960	-0.1840	(-1.5876, 1.3802)
	$b_1$	0.7557	0.2309	<b>0.0014</b>	0.8776	(0.3657, 1.5349)	< <b>0.0001</b>	0.8774	(0.3666, 1.5349)
Ultra10000 TA+UM									
	$a$	-0.0233			-0.4881	(-2.9396, 0.9724)	<b>0.0168</b>	-0.4949	(-2.9080, 0.9357)
	$b_0$	-0.1195	0.6772	0.8598	-0.1738	(-1.5637, 1.3268)	0.7976	-0.1741	(-1.5596, 1.3268)
	$b_1$	0.7557	0.2308	<b>0.0014</b>	0.8779	(0.3791, 1.5382)	<b>0.0008</b>	0.8786	(0.3798, 1.5382)

<sup>a</sup>Parametric bootstrapping was performed to obtain means and confidence intervals, as well as test the null hypotheses that there is no phylogenetic signal in the residuals, ( $H_0: a = -4$ , 1-tailed test) and that the regression coefficient equals 0 ( $H_0: b_1 = 0$ , 2-tailed test).

Supplementary material 4.1. Twenty-eight female-based terminal taxa, with sources of male morphological data indicated. Abbreviations: I, digital images; T, taxonomic descriptions; M, direct examination of specimens at the British Natural History Museum, London, United Kingdom and/or the American Museum of Natural History, New York, USA.

<b>Female-based terminal taxa</b>	<b>Male morphological source</b>	<b>Taxonomic reference</b>
<i>Adrania gracilis</i>	I, T, M	Miller (1955)
<i>Afrodecius</i> sp.2		
<i>Antiopuloides formosus</i>		
<i>Apocaucus</i> sp.		
nr <i>Bayerus</i> sp.		
nr <i>Bayerus</i> sp.2		
<i>Borgmeierina dentata</i>		
<i>Centraspis decoratus</i>	I, T, M	Villiers (1951)
<i>Centraspis petersi</i>	M	
<i>Cimbus productus</i>	M	
<i>Ectrichodia lucida</i>	I, T, M	Lepeletier & Serville (1825); Stål (1865); Villiers (1948)
<i>Ectrichodiella minima</i>	T	Gil-Santana et al. (2013)
<i>Glymmatophora dejoncki</i>	T	Villiers (1948)
<i>Haematorrhophus pedestris</i>	I, T, M	Distant (1904)
<i>Katanga</i> sp.		
<i>Maraenaspis coccinea</i>		
<i>Mendis notata</i>	M	
<i>Microstemmatoides occidentalis</i>	T, M	Villiers (1973)
<i>Microstemmatoides similis</i>	T	Schouteden (1952); Villiers (1973)
<i>Neoscadra</i> sp.		
<i>Okondo collarti</i>		
<i>Opistoplatys</i> sp.2		
<i>Pothea</i> sp.		
<i>Racelda</i> sp.		
nr <i>Scadra</i> sp.		
<i>Schottus perakensis</i>	I, T	Miller (1941)
<i>Schuhella hermosa</i>		
<i>Tribelocodia</i> sp.		

Supplementary material 4.2. Morphological character matrix in nexus format for cladistic analysis.

#NEXUS

BEGIN TAXA;

TITLE Taxa;

DIMENSIONS NTAX=156;

TAXLABELS

Adrania\_cylindrica Adrania\_gracilis Adrania\_macra Antiopuloides\_formosus  
Austrokatanga\_monteithi Bannania\_sp nr\_Bannania\_sp Bayerus\_cuneatus Borgmeierina\_dentata  
Brontostoma\_colossus Brontostoma\_nanus Brontostoma\_sanguinosum Caecina\_affinis  
Caecina\_n\_sp Caloundranus\_formosus Centraspis\_decoratus Centraspis\_ducalis  
Centraspis\_petersi Cimbus\_ambulans Cimbus\_productus Cimbus\_sp Cleptria\_cinctiventris  
Cleptria\_corallina Cleptria\_maculata Cricetopareis\_boliviana Cricetopareis\_tucumana  
Cryptonannus\_punctella Cryptonannus\_n\_sp Daraxa\_ambrosettii Daraxa\_sp  
Distirogaster\_tarsalis Distirogaster\_n\_sp Distirogaster\_n\_sp2 Doblepardocoris\_hirsuta  
Doblepardocoris\_vittata Ectrichodia\_crux Ectrichodia\_lucida Ectrichodia\_sp  
Ectrichodiella\_minima Ectrichodiella\_n\_sp Ectrichodiella\_n\_sp2 nr\_Bayerus\_sp2 nr\_Bayerus\_sp  
Ectrychotes\_serdangensis Ectrychotes\_sp Ectrychotes\_atripennis Gibbosella\_conisimilis  
Gibbosella\_pallidalata Gibbosella\_planiscutum Gibbosella\_quadocris Glymmatophora\_aeniceps  
Glymmatophora\_carolae Glymmatophora\_crassipes Glymmatophora\_dejoncki  
Haematoloecha\_limbata Haematoloecha\_nigrorubra Haematoloecha\_sp  
Haematorrhophus\_linnaei Haematorrhophus\_pedestris Jorgcoris\_aczeli Katanga\_linnavuorii  
Katanga\_petchkovskiy Katanga\_sp Labidocoris\_insignis Leptomendis\_bipartita  
Leptomendis\_breddini Lyramna\_vigil Lyramna\_n\_sp Lyramna\_n\_sp2 Maraenaspis\_bidens  
Maraenaspis\_coccinea Maraenaspis\_crocata Marojejycoris\_auranticorium  
Marojejycoris\_brevifrons Marojejycoris\_francais Marojejycoris\_notadichroa  
Marojejycoris\_ranomafana Mendis\_apicimaculata Mendis\_notata Microsanta\_montana  
Microsanta\_pusilla Microstemmatoides\_atrocyanea Microstemmatoides\_occidentalis  
Microstemmatoides\_similis nr\_Neolibavius\_sp Neosantasia\_lugubris Neoscadra\_annulicornis  
Neoscadra\_ornata Neoscadra\_sp Nularda\_lesnei Nularda\_nobilitata Nularda\_thoracica  
Okondo\_collarti Philodoxus\_principalis Pothea\_lugens Pothea\_ventralis Pothea\_wygodzinsky  
Pothea\_sp Pseudopothea\_luacesi Racelda\_moerens Racelda\_spurca Racelda\_sp Rhiginia\_aimara  
Rhiginia\_cinctiventris Rhiginia\_immarginata Santasia\_erythrocephala Santasia\_maculata  
Santasia\_sp Scadra\_aliena Scadra\_cincticornis Scadra\_illuminata nr\_Scadra\_sp  
Schottus\_baramensis Schottus\_gracilis Schottus\_perakensis Schuhella\_hermosa Schuhella\_n\_sp  
Schuhella\_n\_sp2 Sciaphilocoris\_ornata Sinchocoris\_colombiana Sinchocoris\_panamaensis  
Synavecoris\_dimorphus Tanindrazanus\_marginatus Tanindrazanus\_tenebricus  
Tanindrazanus\_varicolor Tasmanocoris\_evansi Toliarius\_trichrous Toxopus\_brucei  
Toxopus\_fisherii Toxopus\_griswoldi Toxopus\_toamasina Toxopus\_vazimba Vilius\_lateralis  
Vilius\_macrops Wygodzinskyocoris\_nigripes Zirta\_caudorubrum Zirta\_hirticornis Zirta\_limbata  
Abelocephala\_sp Abelocephala\_sp2 Afrodecius\_sp Afrodecius\_sp2 Apocaucus\_sp  
Centrogastocoris\_alacris Megapocaucus\_sp Opisthoplatys\_sp Opisthoplatys\_sp2  
Tribelocephala\_peyrierasi Tribelocephala\_sp Tribelocodia\_ashei Tribelocodia\_sp Xenocaucus\_sp  
Emesaya\_incisa Carayonia\_sp Saica\_sp Peirates\_punctorius

END;

```

BEGIN CHARACTERS;
  TITLE Character_Matrix;
  DIMENSIONS NCHAR=201;
  FORMAT DATATYPE = STANDARD GAP = - MISSING = ? SYMBOLS = " 1 2 3 4
5";
  MATRIX
    Adrania_cylindrica 111113211-111121--111121112111111-
12111211??121????21121111221211-221122122211-1121221?2121112211-21--11111--2211-
21--11111--22111--11111-
321?12221311111221111212????1211122212112?1?1??1?1??1?1?11111
    Adrania_gracilis 111112211-111121--111121112112111-
1211121123121????112211112212122212121222211-112122132121112211-21--11111--
2211-21--11111--21111--11111-321?12221311111221111??2112-
12111212122211112111111111??221
    Adrania_macra 111113211-111121--111121112111111-
1211111123121????21121111221211-2212221222211-1121221?2121112211-21--11111--
2211-21--11111--22111--11111-
321?12221311111221211??2????????????????????????????????????11?
    Antiopuloides_formosus 1112132131112111--11112??22111111-
3211121122123????112111112222122221111-122311-112122132122112211-11--11111--
2211-11--11111--22111--11111-
311112211311111221212??2????????????????????????????????????22?
    Austrokatanga_monteithi 311133211-112121--11111--12111111-
3111121224113????211321121-3221-1211122121221-212122131122112111-
121111212211111-12111111221111(1 2)22111111-3-----
111111222122121222311111332121112121121211211111111
    Bannania_sp 121212211-112111--111122222111111-
221112112212322112112111122222122221211-1222121112122133122211211-123111111--
2211-123111111--221123111111-
3121122113222112212122121222????????????????????????????????22?
    nr_Bannania_sp 121212211-112111--111122222112111-
1211121122122221121121111221222221211-122211-112122133122212211-11--
11112312211-11--11111--22111--11111-
3111212113222112212122121222????????????????????????????????22?
    Bayerus_cuneatus 131113211-111121--111122222111111-
1211121124111????11221112221222221?121122111-1121221?3122111211-11--11111--
2211-11--11111--22111--11111-
323??1??1311111221112??2????????????????????????????????????22?
    Borgmeierina_dentata 111113211-111111--11112??22111111-
12111211??121????111?11112212122221211-222211-112122?3?1??12?1??2??11111--
2?1??21--11111--22111--11111-
321?112212111112?1?2??2????????????????????????????????????11?
    Brontostoma_colossus 1111132121111111--111122122112111-
22111211241212212211211111221222212121222111-112122133222112412111--11111--
2412111--11111--21111--11111-321112221111112213112121222(1
2)2211211121224121141211111111221

```

*Brontostoma nanus* 1112132121411111--111122122112111-  
221112112312221221121111211211-2211121222111-112122133122112411-11--  
11112312412111--111123121111--11111-  
321112221111112213112121222????????????????????????????????22?  
*Brontostoma sanguinosum* 1111132121111111--111122122112111-  
22111211241222122112211121221222121222111-112122133222112412111--11111--  
2412111--11111--21111--11111-321112221111112213112121222(1 2  
3)221121?12122?1?1??1?1??1?1??1221  
*Caecina affinis* 131113211-111121--11112222112111-  
1211121122112????22?212122232222211122122311-112122133122211211-122111111--  
1211-122111111--121122111111-  
31311222132221222122121222????????????????????????????????22?  
*Caecina n\_sp* 131112211-111121--21112222111111-  
2211121123112????12?112122232222221122122311-112122133122211211-122111111--  
1211-122111111--121122111111-  
31311222132221222122121222????????????????????????????????22?  
*Caloundranus formosus* 1112132121112121--211122122111211-  
3211121122123????1121111122212221111-122311-11212213?122211211-11--11111--  
2211-11--11111--22111--11111-  
311112211311111221212?12????????????????????????????????22?  
*Centraspis decoratus* 11121321?411111--  
11112212211211213211121122112????11211111211222?2211122222311-  
21212223222111111-11--11111--2111-11--11111--21111--11111-  
31111222121111221211?2????12212211122121111111111111121221  
*Centraspis ducalis* 1112132121111111--  
111122122112112132111211221122112111112112222211121222311-  
212122232221114121123111111--2412111--11111--21111--11111-  
31111222121111221211122122212?1?2?112?22?1?1??1?1??1?1??1221  
*Centraspis petersi* 11121321??111111--  
11112212211211213211121122122????112111112112222211121222311-  
21212223222111?1??11--21111--2?1??11--11111--21111--11111-  
31111222121111221211?2????122122111222241211?1?11111111211  
*Cimbus ambulans* 111133111-111121--111122122111111-  
3111121124113????11211112223222221111-122311-112122132122212111-11--22211--  
1111-11--22211--11111--22211-  
311212221311111221212?12????122111112222?1?1??1?1??1?1??1221  
*Cimbus productus* 111133121-111121--111122122112211-  
3111121124113????11211112222222221111-122311-1121221?1122212111-11--22211--  
1111-11--22211--11111--22211-  
311212221311111221212?2????122111111222111111111111111121221  
*Cimbus sp* 111133121-111121--111122122111111-  
3111121124113????11211112222222221111-122311-112122111122212111-11--22211--  
1111-11--22211--11111--22211-  
31121222131111122212??21221????????????????????????????????22?  
*Cleptria cinctiventris*  
111113213111112211?111221221111121221121124122????112111112112122221112222221  
1-112122?3?122212211-11--11111--2211-11--11111--22111--11111-  
311112211311111221?11112????????????????????????????????22?

*Cleptria\_corallina*  
 111213212111112111112212211211212211121124122221??1121112212212222111222222  
 11-112122132122212211-11--11111--2211-11--11111--22111--11111-  
 311112211211112212111121222????????????????????????????????22?

*Cleptria\_maculata* 1112132131111111--  
 1111221221111121221112112?122?????1121111121221222211122222211-  
 1121221321222124121122211111--2211-122211111--221123211111-  
 311112211311111222212??2????????????????????????????????22?

*Cricetopareis\_boliviana* 1112132132111111--12112212211??11-  
 221112112?112?????11211111221222221111-222111-112122221122111111-11--11111--  
 2111-11--11111--21111--11111-3???????121111222111??21222(2  
 3)211121112122111111111111111221

*Cricetopareis\_tucumana* 1112132132111111--121122122111111-  
 2211121124112?????11211111221222221111-222111-112122221122111111-11--11111--  
 2111-11--11111--21111--11111-32111222121111222111??21222(2  
 3)211121112222111111111111111221

*Cryptonannus\_punctella* 1112132121311111--111122122111111-  
 2211121124122?????112111112212122221211-222111-112122233121112212221--11111--  
 1211-11--11111--12111--11111-321?121-  
 111111222112??2????????????????????????????????????11?

*Cryptonannus\_n\_sp* 1112132121111111--111122122112111-  
 221111112412222112112111112212122221211--22111-112122233121112212221--11111--  
 1211-11--11111--12111--11111-321?121-  
 121111222112212????????????????????????????????????11?

*Daraxa\_ambrosettii* 111212211-111111--111122122112111-  
 22111211241212?1?21121111221212222121222111-112122233121112412221--11111--  
 2412221--11111--22111--11111-32111122111111221122121222(1  
 2)2111213121224121141211411111221

*Daraxa\_sp* 111213211-111111--111122122112111-  
 12111211231212?1?21121111221212222121222111-112122232121112412221--11111--  
 2412221--11111--22111--11111-  
 321?12221111112213122121222????????????????????????????????11?

*Distirogaster\_tarsalis*  
 1111132131411112111111221221112213211121124122221221121111221222222111222222  
 11-112122233122211411-21--11111--2411-11--11111--22111--11111-  
 3121122212222212212211221222411??2?32?121????????????????????212

*Distirogaster\_n\_sp*  
 1111132131411112111111221221112213211121124122221221121111221322222111222222  
 11-112122133122211412111--11111--2411-11--11111--22111--11111-  
 3121122212222212212211221222????????????????????????????????11?

*Distirogaster\_n\_sp2*  
 1111132131411112111111221221122213211121124122221221121111221322222111222222  
 11-112122133122211111-21--11111--2211-11--11111--22111--11111-  
 312112221222221221221122122241112332112141211412114121111112

*Doblepardocoris\_hirsuta* 111213211-111111--111122122112111-  
 1211121124121?????112111112212122221211-222111-1121221?3121112412221--11111--  
 2211-11--11111--22111--11111-  
 32111?22111111221112212?????12211211121222?1?1?1?1?1?1?11111

*Doblepardocoris\_vittata* 111213211-111111--111122122112111-  
1211121124121????1121111221212221211-222111-112122???1???12412221--11111--2??1-  
11--11111--2??11--11111-  
32111122111111221??2212????1221121112122?1?1??1?1??1?1??111  
*Ectrichodia\_crux* 1111132131111111--  
1111221121111222211121122122221222123111121121222211122222211-  
112122131122112412111--22211--2412111--22211--21111--12211-  
31121221121111221111??21222221122231212241212412211111111221  
*Ectrichodia\_lucida* 1111132131111111--  
1111221121111222211121122122221222113111121121222211121222211-  
112122133122112?1??11--22211--2?1??11--22211--21111--12211-  
31121221121111221111??21222221122231212241212412211111111221  
*Ectrichodia\_sp* 1111132131111111--  
1111221121121122221111112?12222122211211112112121221112222221-  
112122133122112412111--22211--2412111--22211--21111--12211-  
31121221121111221111121222????????????????????????????????11?  
*Ectrichodiella\_minima* 14?2132131????11--  
11112232211211222213221214221????11111121211222?212121-222222112122??11--  
11?21-121211122121221-1212112212122121211122121221?1-2-11111?11112--  
2222112212211122222121221212212111111  
*Ectrichodiella\_n\_sp* 1422132131????11--  
111122322112112222132312??221????111111222111-1-212121--2222221121????1??11?1?-  
121211122121????121211122121??121211122121221?1---  
1111??????2?22221????????????????????????????????11?  
*Ectrichodiella\_n\_sp2* 14?21321????11--  
111?223221121122?213221214??1????1111112?211222?21212??2?22221121?2??1??11??  
??212??2212??2??212??2212??2121211122121221?1-2-  
1111?1131????2221????????????????????????????????11?  
*nr\_Bayerus\_sp2* 131112211-1111221111112??211??11-  
121111112411221?2112?11122212222211121222211-11212213312?112211-11--11111--  
2211-11--11111--22111--11111-32311222131111?221112??2112-  
????????????????????????????????22?  
*nr\_Bayerus\_sp* 131112211-1111221111112??2211??11-  
121111112412122112112?111222121222211121222211-112122133122112311-11--11111--  
2211-11--11111--22111--11111-32311221131111?221112??2112-  
????????????????????????????????22?  
*Ectrychotes\_serdangensis* 1111122121111121--  
11112212211111212211121124121221122122111121121222212122222121112122133122212  
412121--11111--2412111--11111--24111--  
1111233111122212111122111121212221221222212122?1?1??1?1??1?1?11221  
*Ectrychotes\_sp* 1111122121111121--  
11112212211111212211121124121221122122111121121222212122222121112122133122212  
412121--11111--2412121--11111--24111--  
111122311112221211112213112121222????????????????????????????????22?  
*Ectrychotes\_atripennis* 1111132131111111--  
21112212211111212211121124121221121122111121221222212122222121112122133122212  
412121--11111--2412121--11111--24111--11111-  
311112221211112212112121222????????????????????????????????22?

*Gibbosella\_conisimilis* 111112211-  
 211122111112212211111212211121124122?1??112111112112222221112222221-  
 112122113122111211-11--11111--2211-11--11111--21111--11111-  
 3121222113222212213122121222????????????????????????????????11?

*Gibbosella\_pallidalata* 111112211-  
 2111221111122122111112122111211221212?1??112111112112222221112222221-  
 112122132122112211-11--11111--2211-11--11111--12111--11111-  
 3121222113222212213122121222????????????????????????????????11?

*Gibbosella\_planiscutum* 121112211-  
 211122111112212211111212211121124121????1121111121122222211122222211-  
 112122232122112211-121211111--2211-121211111--221121211111-  
 3121222113222212212122121222411112132112142221421212212111112

*Gibbosella\_quadocris* 111112211-  
 2111221111122122111112122111211241222?1??112111112112222221112222221-  
 112122233122112211-11--11111--2211-11--11111--22111--11111-  
 3121222113222212212122121222????????????????????????????????11?

*Glymmatophora\_aeniceps* 1111132131111111--  
 1111221221121122221112112?1??????1123111121122222211122222111-  
 111122233122212412111--22111--2412111----11--24111--12211-  
 311?1221121111221211??2????(1 4)????(2 3)(1 2)2??(1 2)????????????????22(1 2)

*Glymmatophora\_carolae* 1112132131111111--  
 111122122111112222111211221212212211111211222(1 2)2211122222111-  
 111122231121112111-21--22111--2111-11--22111--22111--22211-  
 311112211211112212111221222????????????????????????????????22?

*Glymmatophora\_crassipes* 4111132131111111--11111--  
 121111122221112112212122122211321111-121212211122221111-111122233122112412111--  
 22111--2411-11--22111--24111--22211-3-----  
 121111221211122122241112223211214121141211411111221

*Glymmatophora\_dejoncki* (1 4)1111321311111111--  
 1111??1211211222211121122121????(2 3)1?3(1 2)111(1 2)-12(1 2)2122111222(1 2)111-  
 11112223312?112?1??11--22111--2?1??11--22111--2?111--12211-311-1---  
 1211111221?11??212224111222321121412114121141111122(1 2)

*Haematoloecha\_limbata* 111113211-111121--111122122112111-  
 1211111124121????112211122212222212121222111-112122?3?122112221-122111111--  
 2221-122111111--21111--11111-  
 32111222121111221112??2????121112121222????????????????11221

*Haematoloecha\_nigrorubra* 111113211-111121--111122222112111-  
 1211121124121????112211122222222212121222211-112122?3?122112221-122111111--  
 2221-122111111--21111--11111-  
 32111222121111221111212????122112131222????????????????11221

*Haematoloecha\_sp* 111113211-111121--111122222112111-  
 1211121124121?211211221122212222212121222111-112122233122112221-2221111111--  
 2221-122111111--21111--11111-  
 321111211111112211122121222????????????????????????????????22?

*Haematorrhophus\_linnaei* 1111132131111111--  
 11112221211211222211121122121????212211112112222221122222211-  
 111122231121112412121--21111--2412111--22211--23111--12211-  
 311?122?121111122111122????(3 4)211222322121?1?1?1?1?1?1?1?1?1?1221



Haematorrhophus pedestris 3111132131111111--  
 11112??121121122221111122121????2113211211222222112221211-  
 11112223112112?1??21--21111--2?1??11--22211--23111--12211-3-----  
 1211111221111??21222321122232212141211412114111111211  
 Jorgcoris\_aczeli 111113211-111111--121122122112211-  
 2?????????121?????1121111221212222121222111-112122?3?122211412121--11111--  
 2412121--11111--21111--  
 1111?32111222121111221111??2????????????????????????????????22?  
 Katanga\_linnavuorii 311123111-111121--111121122111111-  
 3111121??113?????21231112213211-1211122122311-2121?????122112??1-  
 123111112211??1-1231111122111122311111223-----  
 1211111221??2??212213221112312112?21211212112121??111  
 Katanga\_petchkovskiy 111123111-111121--111121122111111-  
 311112112?113?????11211112213211-2211122122311-2121221?1122112111-  
 123111112221111-  
 12311111222111222111112231121222121111221112??212211221111212112121212121  
 2111111  
 Katanga\_sp 111123111-111121--11112??22111111-  
 311112112411322?11112111122132??2211122122311-212122131122112111-122111111--  
 1111-122111111--111123111111-  
 31121221121111221112??21221????????????????????????????????11?  
 Labidocoris\_insignis 1311132131??121--  
 1111222221111121221121123122?????112111112222222211122222311-  
 1121?????1222124121122111211--2?????122111111-----122111111-  
 312?1221132221?221?22?2????????????????????????????????????22?  
 Leptomendis\_bipartita 1111132121111121--  
 111122122111112122111211??121?????21211112212222221212222211-  
 11212213?1221124121122222112222412112222112222?11222211223112121-  
 1211111221312??2????????????????????????????????????22?  
 Leptomendis\_breddini 1111132121111111--  
 11112212211111212211121124122?????212111122112222211122222311-  
 112122?3?1221124121122211111--24121122211111--241122211111-3111121-  
 1211111221212?12????????????????????????????????????22?  
 Lyramna\_vigil 111213211-111111--111122122111111-  
 2211121122121?????112111112112122211111-222111-1121222?2122111412111--11111--  
 2412111--11111--21111--11111-  
 321112211211111221111212????????????????????????????????????22?  
 Lyramna\_n\_sp 111213211-111111--111122122112111-  
 2211121123121?????11211111211222221111-222111-11212222122112412121--11111--  
 2412121--11111--21111--11111-  
 32111221111111221111?121222????????????????????????????????22?  
 Lyramna\_n\_sp2 1112132121111111--111122122112111-  
 2211121123121?????112111112112122221111-222111-112122211122112412111--11111--  
 2412111--11111--21111--11111-  
 32111221111111?221112??21222????????????????????????????????11?

*Maraenaspis bidens* 4111132131111111--11111--  
121111121321112112412222122211321111-322212212122221211-112122233122112412111--  
11111--2412111--11111--24111--11111-3-----  
12111112211111221222411122232122141211412111111111211  
*Maraenaspis coccinea*  
?1111321??11111??1111??211??1??321112112412222111?1??1122?2??221??22??11-  
112122131122212?1??11--11111--2?1??11--11111--2?111--11111-  
3??????111111221?11??21222????????????????????????????????22?  
*Maraenaspis crocata*  
111113213111111211111221221111121221112112?122????212111122122121221112222211  
1-1121221331222124121122111111--2411-122111111--21122111111-  
31111221111111221211??2????????????????????????????????22?  
*Marojejycoris auranticorium*  
1211222121211122111111222221111121221112112212122112111111211222222111221221  
11-112122133122212211-11--11111--2211-11--11111--21111--11111-  
313111211111112212122121222????????????????????????????????11?  
*Marojejycoris brevifrons*  
12112221212111221111112222211221212211121122121221111121111211222222111221221  
11-112122133122112211-11--11111--2211-11--11111--11111--11111-  
313111211111112212122121222????????????????????????????????11?  
*Marojejycoris francais*  
12112221212111221211112212211111212211121123121????112111112112222221112212211  
1-112122133122212211-11--11111--2211-11--11111--11111--11111-  
313111211111112212122121222????????????????????????????????11?  
*Marojejycoris notadichroa*  
1211222121211122221111222221111212211121122121????112111112112222221112212211  
1-112122133122212211-11--11111--2211-11--11111--21111--11111-  
313111211111112212122121222????????????????????????????????11?  
*Marojejycoris ranomafana* 121121211-  
2111221111112112211111212211121123121????1121111121122222211122222111-  
112122133122112211-11--11111--2211-11--11111--11111--11111-3131111-  
1111112212122121222????????????????????????????????11?  
*Mendis apicimaculata* 111113211-111221--  
1211221221111221221112112312122122212111122122222212122222311-  
112122233122112311-21--11111--2311-11--11111--22111--11111-  
31121221121111221211121222????????????????????????????????22?  
*Mendis notata* 111113211-111221--  
11112??211??12132111211??122????2121111221222221222222311-  
112122133122212311-21--11111--2311-11--11111--22111--11111-  
31121221121111221211??2????122122111212231111311113111121221  
*Microsanta montana* 131123211-111121--111121122111111-  
1211111??111??112211122212222211121222111-112122??3122111412111--11111--  
2412111--11111--21111--11111-  
323?1221131111221112212????12?1121112222?1?1??1?1??1?11221  
*Microsanta pusilla* 131113211-111121--111121122111111-  
121111112?111????112111122212222211121222111-112122133122111211-11--11111--  
2211-11--11111--21111--11111-323?11211311111221112??2112-----22?



*Philodoxus\_principalis* 111213211-111111--  
11112222211111213211121??113????1121111121122212211121222211-  
212122231222111111-11--11111--2111-11--11111--21111--11111-  
31121221111111221111?221222????????????????????????????????22?  
*Pothea\_lugens* 111133111-111121--121121122111111-  
311112112412122122112211221212222121222111-11212213212111111-11--11111--  
2111-11--11111--21111--11111-  
3211112111111122111121212221221111212122?1?1??1?1??1?1?11221  
*Pothea\_ventralis* 111123111-111121--121122122112111-  
3111121124121????112111112212122212221222111-112122133122111111-11--11111--  
2111-11--11111--21111--11111-321111221211111221111??2????(1  
2)2211112121221111111111111111221  
*Pothea\_wygodzinsky* 111123111-111121--121122122111111-  
311112112?121????1121111122121222212221222111-112122132122111111-11--11111--  
2111-11--11111--21111--11111-  
321111221211111221111?12????122111121212211111111111111111221  
*Pothea\_sp* 111123211-111121--121122122112?11-  
2111121124121????112111122212222211121222111-112122133121111?1??21--11111--  
2?1??21--11111--21111--11111-  
321111211311111221111??21222????????????????????????????????22?  
*Pseudopothea\_luacesi* 111133111-111121--111122122111111-  
3211121124121????11211111211212222121222111-112122133122111412121--11111--  
2412121--11111--21111--11111-32111?21111111221111??2112-  
311111132211241211412111111111112  
*Racelda\_moerens* 1112132121411111--111122222111111-  
2211121124121????112111112122222211121222111-112122233121112212221--11111--  
2211-21--11111--22111--11111-  
321112221211111222311??21222411112132121141212412124121211112  
*Racelda\_spurca* 1111132131311111--111122222111111-  
221112112?121????112111112122222211121222111-112122231121112411-21--11111--  
2411-21--11111--22111--11111-  
321111221211111222111??21222411112132111141212412124121211112  
*Racelda\_sp*  
?1111321???1111???1111???211?1??3211121124121?????1??111??1??????????22??11-  
112122231121112?1??21--11111--2?1??11--11111--  
2??1?????????3??????1?1111?222?12??1222????????????????????????????????11?  
*Rhiginia\_aimara* 1111132121111111--111122112111111-  
12111211241122212211211111211222221111-222111-112122233222111211-21--11111--  
2211-21--11111--21111--11111-  
32111122121111122111121212221211121112222?1?1??1?1??1?1?11221  
*Rhiginia\_cinctiventris* 1111132121111111--121122122111111-  
12111111241122212211211111211222221111-222111-112122133222112211-21--11111--  
2211-21--11111--21111--11111-  
321111221211111221111212122212111211122221112111111111111221  
*Rhiginia\_immarginata* 1111132121111111--121122122112111-  
12111211241132212211211111211222221111-222111-11212213222112111-11--11111--  
2111-11--11111--21111--11111-  
32111122121111122111121212221211121112222?1?1??1?1??1?1?11221

*Santosia erythrocephala* 111113211-111121--111121122111111-  
121111112?111????112211112212222212121222111-112122133121111211-11--11111--  
2211-11--11111--22111--11111-  
3211112212111122111112????121112121222221111211112111111221  
*Santosia maculata* 111113211-111121--111121122111111-  
1211111124111????112111112212222212121222111-1121222?3121111211-11--11111--  
2211-11--11111--21111--11111-  
32111122121111?221311?22????12111212122222111121111111111221  
*Santosia sp* 111113211-111121--111121122111111-  
12111111241112????112111112212222212121222111-112122133121111211-11--11111--  
2211-11--11111--21111--11111-  
321111221211112211112121222????????????????????????????22?  
*Scadra aliena* 111113221-111121--111121122111111-  
1211121??121????112211122212222212121222111-112122??1?1??11211-11--11111--2211-  
11--11111--22111--11111-  
32111221121111221111122????????????????????????????????22?  
*Scadra cincticornis* 111113221-111121--111121122111111-  
221112112?121????112211122212222211121222111-112122??2121111211-21--11111--  
2211-21--11111--2??11--11111-32111222121111221111??2112-----22?  
*Scadra illuminata* 111113221-111121--111121121111111-  
1211121124121????112111122212222211121222111-112122?3?121111211-11--11111--  
2211-11--11111--22111--11111-32111222121111221111??2112-  
1211121212222211112111121111111111  
*nr Scadra sp* 11111321??111121--11112??2211??11-  
121112112412122112112?1111221222221?121222211-1121222331221123?1-  
2??2111??23?1-  
2??2111??22?1??1111??32111221131111221?12??21222????????????????????????2?  
2?  
*Schottus baramensis* 111213211-111111--  
11112212211211212211122111121????2121112221212222112222211-  
112122?33221112311-11--11111--2311-11--11111--21111--11111-  
31111222121111221112?12????????????????????????????????22?  
*Schottus gracilis* 111213211-111111--  
11112212211211212211122111121????1121112221212222112222211-  
112122?33221112311-11--11111--2111-11--11111--21111--11111-  
311112221211112211122121222????????????????????????????????22?  
*Schottus perakensis* 111213211-111111--  
11112212211211212211122111121????1121112221212222112222211-  
11212223322112?11-11--11111--2?11-11--11111--21111--11111-  
31111222121111221112?2????12112211121223111131111111111221  
*Schuhella hermosa* 11131321??411111--21112??2211??11-  
2211121111222????112?1111221222221211-222111-11212223312?112111-11--11111--2111-  
11--11111--21111--11111-31111221121111222112??2????????????????????????????22?  
*Schuhella n\_sp* 111312211-411111--211121222111111-  
2211121111223????1121111221222221211-222111-11212223312?112111-11--11111--  
2111-11--11111--21111--11111-  
311112211211112221122121222????????????????????????????????22?

Schuhella\_n\_sp2 111313211----111--211121222111111-  
2211121111222?????11211111221222221211-222111-1121?????1???12111-11--11111--2111-  
11--11111--2??11--11111-31111221121111?222??2212????????????????????????????????????22?  
Sciaphilocoris\_ornata 111132111----121--111121122111111-  
3111121122113?????21211111211222221111-122311-1121221??122212111-11--12111--1111-  
11--12111--1?111--1?111-  
31?21?2?1111111221112212?????12?1112112222?1?1?1?1?1?1?1?11221  
Sinchocoris\_colombianaensis 111113211-411111--111121122112111-  
221112112?121?????112211112112122221211-222111-112122233121112412221--11111--  
2412221--11111--24111--11111-  
321112211211111222112??2????????????????????????????????????11?  
Sinchocoris\_panamaensis 111113211-111111--111121122112111-  
2211121124121?????112311112212122221211-222111-112122133121112412221--11111--  
2412221--11111--24111--11111-  
321112211211111222112??2????????????????????????????????????11?  
Synavecoris\_dimorphus 111212211-211121--211121122111111-  
2211121122123?????1121111221122222111?222211-112122233122212111-122211211--  
2111-122211111--211122211111-3112211-  
1211111221212??212223111121212222?2??21212112111112  
Tanindrazanus\_marginatus  
1311132121211122122112222211221212211121123112221121121111121122222111222222  
11-11212213312221211-11--11111--2211-11--11111--22111--11111-  
313112211111112212112121222????????????????????????????????????11?  
Tanindrazanus\_tenebricus  
13111321212111121221122222112112211121123112?21121121111121122222111222222  
11-112122133122212211-11--11111--2211-11--11111--22111--11111-  
313112211111112212112121222????????????????????????????????????11?  
Tanindrazanus\_varicolor  
131113212121112212211222221111212211121123113?21121121111121122222111222222  
11-112122133122212211-11--11111--2211-11--11111--22111--11111-  
313112211111112212112121222????????????????????????????????????22?  
Tasmanocoris\_evansi 1312122131111121--111121112111111-  
3211121122123?????11211111212222221211-122211-112122133122211211-11--11111--  
2211-11--11111--22111--11111-  
31211221121111122121212?????121112111212221111211112111121221  
Toliarus\_trichrous  
131123213121112211111121122111112122111211221122?1?1121111121122222111221222  
11-112122133122212211-11--11111--2211-11--11111--22111--11111-  
313112211111112212112121222????????????????????????????????????22?  
Toxopus\_brucei  
1212132121411112221111222221111212211121123112221121121111121122222111222222  
11-112122233122212211-11--11111--2211-11--11111--22111--11111-  
3131122112111112212112121222????????????????????????????????????11?  
Toxopus\_fisheri  
14112321212111221111112112211111213211121123112221??1122111121222222111222222  
11-112122233122212211-11--11111--2211-11--11111--21111--11111-  
3131122112111112212112121222????????????????????????????????????11?

*Toxopus griswoldi*  
 1411232121211122111111211221111121321121123112221??11221111211222222111222222  
 11-112122233122212111-11--11111--2111-11--11111--21111--11111-  
 313112211211112212112121222411112132112131111311111111111112  
*Toxopus toamasina* 121213211-  
 411112221111222221122121221112112311221112112111121222222211122222111-  
 112122233122212211-11--11111--2211-11--11111--22111--11111-  
 313112211211112212112121222????????????????????????????????????11?  
*Toxopus vazimba*  
 121113212141111222111122122111112122111211231122211211211111211222222111222222  
 11-112122233122212211-11--11111--2211-11--11111--22111--11111-  
 313112211211112212112121222????????????????????????????????????11?  
*Vilius lateralis* 1113132121121111--  
 12112222211211212211121111113?????112111122212222211111-222311-  
 122122211221111111-11--11111--2111-11--11111--21111--11111-  
 311112221211111222212?12????1211221112222?1?1??1?1??1?1?21221  
*Vilius macrops* 1113132121121211--  
 12112212211111212211121111132231211211122112222211111-222311-  
 122122131221111111-11--11111--2111-11--11111--21111--11111-  
 3111122212111112222122122212?1221112222?1?1??1?1??1?1??221  
*Wygodzinskyocoris nigripes* 1111132121411111--111122222111111-  
 221112112?121????1121111122122222211121222111-11212223112?112212221--11111--  
 2211-11--11111--22111--11111-  
 3211122?1211111222112??2????????????????????????????????????11?  
*Zirta caudorubrum* 111113211-111111--111122112111111-  
 2211121111122????1121111121121222111121222111-112122233121111311-11--11111--  
 2311-11--11111--21111--11111-  
 32111222111111221311??2????12111211121223111311111111111221  
*Zirta hirticornis* 111113211-111111--111122112111111-  
 2211121111122????11211111211211-2111121222111-112122233121111311-11--11111--  
 2311-11--11111--21111--11111-  
 32111222111111221311??2????12111211121222?1?1??1?1??1?1??1221  
*Zirta limbata* 111113211-111111--111122112111111-  
 2211121111122221221121111121121222111121222111-112122233121111311-11--11111--  
 2311-11--11111--21111--11111-  
 32111222111111221311??2122212111211121222?1?1??1?1??1?1??1221  
*Abelocephala\_sp* 142123211----121--11111--  
 2111221211211231224211????11?1111?211??1-2??21-11---22111112??1??11211-11--  
 11111--1211-11--11111--12111--11111-2132111-1111112?1111221?1221?????????????-  
 ?????????????????11?  
*Abelocephala\_sp2* 142122211----121---1111--  
 2111111211211231??211????11211111211??1-2??121-11---22111112??1??11211-11--  
 11111--1211-11--11111--12111--1111?21?2111-1111112?11112??11221?????????????-  
 ?????????????????11?  
*Afrodecius\_sp* 142122211----12??11111--  
 2111211221222221224212????11?11111211??1-112121-2213222112112??1??11211-11--  
 11111--1211-11--11111--12111--11111-213212212-  
 11111?2111221221221????????????????????????????????????11?

Afrodecius\_sp2 142122211----2???11111--  
 21111122122221???213????11?11111211??1-112121-22222211?112???1???11211-11--  
 11111--1211-11--11111--12111--11111-2132121-2-  
 11111??1??2??21221????????????????????????????????11?  
 Apocaucus\_sp 142?1221?????????????1?1--  
 221??1????21222114?21????11?11111211????????12??11---2211??12????????11211-11--  
 11111--1211-11--11111--12111--11111-21?22??111111????12???1221?????????????-  
 ?????????????????11?  
 Centrogastocoris\_alacris 142133211---121--11211--21212211-  
 2211221???212????11?11111211????2??11-21---22112211????????11211-11--11111--1211-  
 11--11111--12111--11111-31311222??11112?21112??22211121121112??-  
 211112111121111?1111  
 Megapocaucus\_sp 142?11211----1?????1?1--  
 2211?11221212221214?21????11?1111?211??1-2??121-11---22112112??11--11211-11--  
 11111--1211-11--11111--12111--11111-2132211-1111112?21112??11221????????????-  
 ?????????????????11?  
 Opistoplatys\_sp 142133111---121--11121--21212211-  
 2211221???212221111111111211221-211111-11---22112211??11--11211-11--11111--1211-  
 11--11111--12111--11111-3131121-1111112?211121212221????????????-????????????????11?  
 Opistoplatys\_sp2 142133111---121--11111--21212211-  
 2211221???211????11111111211221-211111-21---22112211??11--11211-11--11111--1211-  
 11--11111--12111--11111-31311221111112?21112??12221????????????-  
 ?????????????????11?  
 Tribelocephala\_peyrierasi 142133211---1121--11211--21211111-  
 3222221115212????11?11111211221-211121-21---22112111????1--11211-11--11111--1211-  
 11--11111--12111--11111-31321??1111112?2121111?22211121121112??-  
 211112111121111?1111  
 Tribelocephala\_sp 142133211---1121--11211--21211111-  
 32222211152122211111111111211221-211121-21---22112111??11--11211-11--11111--1211-  
 11--11111--12111--11111-31321?221111112?21211112221????????????-  
 ?????????????????11?  
 Tribelocodia\_ashei 14?133211-41?111---11?22122111211-  
 2211231111111????1111111211??1-112121-222111-1121221111??11221-  
 12121111212121-  
 121211112121221211112111321222111111??21212??1221????????????????????????????  
 11?  
 Tribelocodia\_sp ???13?211-  
 4121????11????11??1??2211231111?11????????11????????????1-  
 2????????221111??11221-1??1111??12?1-  
 1??1111??12?1??1111??1????11111??21?12??1221????????????????????????????11  
 ?  
 Xenocaucus\_sp 142211211---121---1111--21112211-  
 1211221113?23????11?1111??12??211121-11---2211?122?1111--11211-11--11111--1211-  
 11--11111--12111--11111-2132121-1111112?2131????1221413?12232??-  
 2111121111211111112  
 Emesaya\_incisa 231112111-111121--11111--12121111-  
 2211121111211????311321111-11-1-211121-21---22111?1221111--22211-123112112211211-  
 11--11111--1?111--11111-31111?1-1111112121112??2111-----



```

Carayonia_sp      121123211-111121--11111--21121111-
121112111122112211212111112112121211111-21---22111?22???11--21211-11--11111--1211-
11--11111--12111--11111-31?21?1-1111112?21112???111-----
Saica_sp          131111211-111121--11111--12121111-
1211121111221????311311111-1211-211111-21---22111?2221111--21211-11--11111--1211-
11--11111--12111--11111-31321???1111112121112???111-----
Peirates_punctorius 3311132121111121--21111--12111111-
11111111122311211312321121-22121211111-11---2211111121111--22221-122111111--2221-
122111111--221122111111-3-----1111112121112???22221-----
;
END;

```

BEGIN CHARACTERS;

```

TITLE Character_Matrix2;
DIMENSIONS NCHAR=1;
FORMAT DATATYPE = CONTINUOUS GAP = - MISSING = ?;
MATRIX
Adrania_cylindrica      12.62
Adrania_gracilis       11.00
Adrania_macra          12.95
Antiopuloides_formosus ?
Austrokatanga_monteithi 10.80
Bannania_sp            5.92
nr_Bannania_sp         6.64
Bayerus_cuneatus       8.68
Borgmeierina_dentata   ?
Brontostoma_colossus   25.11
Brontostoma_nanus      12.48
Brontostoma_sanguinosum 25.87
Caecina_affinis        8.71
Caecina_n_sp           10.89
Caloudranius_formosus  9.03
Centraspis_decoratus   36.00
Centraspis_ducalis     24.56
Centraspis_petersi     ?
Cimbus_ambulans        21.00
Cimbus_productus       21.00
Cimbus_sp              19.12
Cleptria_cinctiventris 11.72
Cleptria_corallina     13.97
Cleptria_maculata      7.87
Cricetopareis_boliviana 15.75
Cricetopareis_tucumana 19.65
Cryptonannus_punctella 5.75
Cryptonannus_n_sp      6.44
Daraxa_ambrosettii     8.41
Daraxa_sp              7.97
Distirogaster_tarsalis 15.93

```

Distirogaster_n_sp	21.48
Distirogaster_n_sp2	29.20
Doblepardocoris_hirsuta	9.56
Doblepardocoris_vittata	9.38
Ectrichodia_crux	?
Ectrichodia_lucida	23.00
Ectrichodia_sp	21.83
Ectrichodiella_minima	3.79
Ectrichodiella_n_sp	4.82
Ectrichodiella_n_sp2	3.30
nr_Bayerus_sp2	?
nr_Bayerus_sp	?
Ectrychotes_serdangensis	10.23
Ectrychotes_sp	12.51
Ectrychotes_atripennis	10.91
Gibbosella_conisimilis	8.07
Gibbosella_pallidalata	5.18
Gibbosella_planiscutum	5.32
Gibbosella_quadocris	5.57
Glymmatophora_aeniceps	17.93
Glymmatophora_carolae	15.77
Glymmatophora_crassipes	17.94
Glymmatophora_dejoncki	21.00
Haematoloecha_limbata	10.23
Haematoloecha_nigrorubra	14.98
Haematoloecha_sp	11.13
Haematorrhophus_linnaei	23.61
Haematorrhophus_pedestris	22.00
Jorgcoris_aczeli	16.75
Katanga_linnavuorii	12.07
Katanga_petchkovskyi	11.64
Katanga_sp	?
Labidocoris_insignis	10.86
Leptomendis_bipartita	8.98
Leptomendis_breddini	8.09
Lynamna_vigil	11.58
Lynamna_n_sp	13.69
Lynamna_n_sp2	15.60
Maraenaspis_bidens	21.85
Maraenaspis_coccinea	?
Maraenaspis_crocata	17.67
Marojejycoris_auranticorium	8.96
Marojejycoris_brevifrons	6.58
Marojejycoris_francais	6.50
Marojejycoris_notadichroa	7.94
Marojejycoris_ranomafana	5.46
Mendis_apicimaculata	14.91
Mendis_notata	?

Microsanta_montana	11.32
Microsanta_pusilla	8.61
Microstematoides_atrocyanea	13.03
Microstematoides_occidentalis	12.25
Microstematoides_similis	17.50
nr_Neolibavius_sp	4.51
Neosantasia_lugubris	12.25
Neoscadra_annulicornis	19.16
Neoscadra_ornata	16.92
Neoscadra_sp	?
Nularda_lesnei	31.30
Nularda_nobilitata	26.58
Nularda_thoracica	16.66
Okondo_collarti	?
Philodoxus_principalis	34.35
Pothea_lugens	13.93
Pothea_ventralis	19.85
Pothea_wygodzinsky	16.95
Pothea_sp	?
Pseudopothea_luacesi	11.37
Racelda_moerens	13.51
Racelda_spurca	16.16
Racelda_sp	?
Rhiginia_aimara	15.59
Rhiginia_cinctiventris	17.84
Rhiginia_immarginata	17.37
Santosia_erythrocephala	15.49
Santosia_maculata	13.27
Santosia_sp	13.49
Scadra_aliena	11.33
Scadra_cincticornis	12.88
Scadra_illuminata	11.27
nr_Scadra_sp	?
Schottus_baramensis	11.52
Schottus_gracilis	9.50
Schottus_perakensis	11.50
Schuhella_hermosa	?
Schuhella_n_sp	5.92
Schuhella_n_sp2	6.05
Sciaphilocoris_ornata	7.33
Sinchocoris_colombiaensis	7.65
Sinchocoris_panamaensis	9.85
Synavecoris_dimorphus	5.76
Tanindrazanus_marginatus	15.33
Tanindrazanus_tenebricus	16.12
Tanindrazanus_varicolor	16.00
Tasmanocoris_evansi	7.68
Toliarus_trichrous	8.22

Toxopus_brucei	11.84
Toxopus_fisheri	10.29
Toxopus_griswoldi	9.39
Toxopus_toamasina	12.02
Toxopus_vazimba	11.99
Vilius_lateralis	17.45
Vilius_macrops	20.16
Wygodzinskyocoris_nigripes	16.38
Zirta_caudorubrum	14.21
Zirta_hirticornis	14.29
Zirta_limbata	17.95
Abelocephala_sp	3.25
Abelocephala_sp2	2.53
Afrodecius_sp	7.98
Afrodecius_sp2	?
Apocaucus_sp	?
Centrogastocoris_alacris	6.59
Megapocaucus_sp	4.24
Opistoplatys_sp	7.54
Opistoplatys_sp2	?
Tribelocephala_peyrrierasi	?
Tribelocephala_sp	12.50
Tribelocodia_ashei	2.27
Tribelocodia_sp	?
Xenocaucus_sp	3.40
Emesaya_incisa	-
Carayonia_sp	-
Saica_sp	-
Peirates_punctorius	-

;
  
END;

Supplementary material 4.3. PCR protocols for each targeted gene region. Denaturation, annealing, and extension were conducted for 35 cycles for all gene regions sampled.

<b>Gene region</b>	<b>PCR step</b>	<b>Temperature (°C)</b>	<b>Time</b>
16S, 18S rDNAs	Initial denaturation	94	5 min
	Denaturation	94	30 sec
	Annealing	48	30 sec
	Extension	72	45 sec
	Final extension	72	7 min
28S D2, 28S D3–D5 rDNAs	Initial denaturation	94	5 min
	Denaturation	94	30 sec
	Annealing	50	30 sec
	Extension	72	45 sec
	Final extension	72	7 min
COI mtDNA	Initial denaturation	94	10 min
	Denaturation	94	45 sec
	Annealing	43	45 sec
	Extension	72	1 min
	Final extension	72	7 min
Wg npcDNA	Initial denaturation	94	5 min
	Denaturation	94	30 sec
	Annealing	54	1 min
	Extension	72	1 min
	Final extension	72	7 min

Supplementary material 4.4. Molecular character matrix in nexus format for cladistic analysis.

```
#NEXUS
BEGIN TAXA;
  TITLE Taxa;
  DIMENSIONS NTAX=59;
  TAXLABELS
    Abelocephala_sp Abelocephala_sp2 Afrodecius_sp Bannania_sp
nr_Bannania_sp nr_Bayerus_sp nr_Bayerus_sp2 Brontostoma_colossus Brontostoma_nanus
Brontostoma_sanguinosum Caecina_affinis Caecina_n_sp Centraspis_ducalis Cimbus_sp
Cleptria_corallina Cryptonannus_n_sp Daraxa_sp Distirogaster_n_sp Distirogaster_n_sp2
Distirogaster_tarsalis Ectrichodia_crux Ectrichodia_lucida Ectrichodia_sp Ectrychotes_atripennis
Ectrychotes_serdangensis Ectrychotes_sp Emesaya_incisa Glymmatophora_crassipes
Haematoloecha_sp Katanga_sp Maraenaspis_coccinea Marojejycoris_brevifrons
Mendis_apicimaculata Microstemmatoides_atrocyanea Neoscadra_sp nr_Neolibavius_sp
Opisthoplatys_sp Opisthoplatys_sp2 Peirates_punctorius Pothea_lugens Pothea_sp Racelda_sp
Rhiginia_aimara Rhiginia_cinctiventris Saica_sp Santosia_sp nr_Scadra_sp
Tanindrazanus_marginatus Tanindrazanus_tenebricus Tanindrazanus_varicolor Toxopus_brucei
Toxopus_fisheri Toxopus_griswoldi Toxopus_toamasina Toxopus_vazimba
Tribelocephala_peyrierasi Tribelocephala_sp Tribelocodia_sp Xenocaucus_n_sp
;
END;

BEGIN CHARACTERS;
  TITLE Character_Matrix;
  DIMENSIONS NCHAR=4055;
  FORMAT DATATYPE = DNA GAP = - MISSING = ?;
  MATRIX
Abelocephala_sp      ??????????????????ATCTTTTTTTG-----TTATTATATA-
AAATTTGGCCTGCCCTATGA---
AAATGATTAAATGGCCGCAGTATTTTAACTGCGCTAAGGTAGCATAATAAATTGTCT
TTTAATTAAGGGCTTGTATGAATGGTTGGACGAAATAAAAACCTATTTTG--TTTTAT--
TTTTTGAATTTAATTTTTTTAGTTAAAAGGCTAA-
GATTTTATTAAGGGACGAGAAGACCCTATAAAGCTTTATACGTGA-GGTA--
GGCACCCAATTTTAG-ATTTTTAAT-
AAATTGGGTGCTTTATTTGTGGCGTATTA AATTGGGGTGATTGGGGAATTC-
TTTTAACTTCTGA--ATA-CTGATTTCGTTAATAT--TCGAT--TTAATTGATCCAGT---
TTTATTGATTATAAGATTAAGTTACTTTAGGGATAACA-
GCGTAATTTTCTCAGAGAGTTCAAATTGATGAGATAGTTTGCACCTCGATGTTGGAT
TAAAA-TTAGCC-GTTGGTGCAGCAATTA-ATTGG-
CTTGGTCTGTTTCGACCATTGAAAT-
TTTACATGAT????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????CGAGGATCCA-
TTGGAGGGCAAGCCT--
GGTGCCAGCAGCTGCGGTAATTCCAGCTCCAATAGCGTATATTA AAGTTGTTGCGGT
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGTA
ACTGGCATGTCGCGGCATGTCTGTGCGGTGGTGAACCAAATTC-----
-----TTGGTAT---AAACTAAGCT---TAAATGCTGAAT-T-TGTAT-CCA-----TA-G-
```

TAATTTT--GTTTGCCGA---  
TTTCTCTATTTTCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAAAATCTGCCTGAATAGTGGTGCATGGAATGATAGAATAGGACCTCAGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGACGGGGGCATTCGTA  
TTGCGACGTTAGAGGTGAAATTCTTGGACCGTCGCAAGACGCACTAGAGCGAAAGC  
ATTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCTGAAGGCGATC  
AGATAACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCC  
TATAATGACTCGGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAA??  
??  
????????????GGTAAACTCCATCTAAGGCTAAATATAACCACGAGACCGATAGCGAACA  
AGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACGTGA  
AACCGCTCAGGGGTAAACGGAAAAGAACTGAA-  
TAATCCAAAAGGGGAGATTCACGTCTTACCGC---TAGCTTAACTTTAAGATGT-----  
TTTGAGATGGTCATTTTC-GCCCTATCCAG---TGCAA--ACTGTG--TTAG-  
TGGGTTTGATTACTCGCATTTT-ATAA--  
GATGTCAAGGCAGCGGTGGACCGCACTTCTTCTTAGT-  
AGGACGTTGCGACTTGTCACTA-GAAT----ACTTTAAGT--AT-A----GTAA---  
AGTGCCTGTAT-T-TAACAA--TTAGACTTT-----ACTATTTAATTA-----  
TAGTAATTTTCATTTTATTACAGC-TTTAA--CAT--AAA-----ATACCGAGTATAT-----  
----  
ACTTGTGACAATAACGATCGATGGTTGTTGAGCCGCTAAAAATTATAAATAATTTTT  
ATTTATA----G-----TTATAAGCGTT-----TATTGGCCT-TTATACAA-  
GTGGTTGTCAAGTTCAAGTGTTAATT-A-  
TTGTCAATATGTCCTTTTTATGTTTCAGTCATG-TTA-ATCTTACTA-CTGTTGGTA---  
TAAT-TG--TG-CCTT--AACAG--TGCTTA--  
CAACACTGGTCGGCAGCGATCAAG????????????????????????????????????  
??  
??  
??  
ATTATCAA  
GGGAAGATGG--AAAT-----ATTATACAAGTAAC-----  
AAATTTTT-----GTT-AATTTATAAATAAT-  
TTCCCGCATTCCCAGGGCGTTTGT-TTACTCTCA---TG-  
AGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTTTCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGAT-----T----  
TTAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTGC-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAT????????????????????  
??  
??  
????????????????TTGGTCCACCAGAAGTATACATTCTAATTTTACCCGCCTTTGGAATTATT  
CACATATCATTTCAATAAAAACCGGAAAAAATGAAACATTTGGTGCCTTAGGAATAA  
TTTACGCCATATTAGCAATTGGCCTTTTAGGATTTATTGTATGAGCCCACCATATATT  
TACCATCGGAATAGATGTAGACACACGAGCATATTTTACCTCAGCTACAATGATCAT  
TGCAGTACCTACAGGCATTAATACTTTAGATGATTAGCAACCCTTCATGGAAGCAT  
TTTACATCCTAGTCCCAGAACTCTTTGAGCACTAGGGTTTGTTCCTATTTACCATA  
GGGGGTATAACTGGTATTATTCTTGCCAATTCGAGAATTGATATCTCTTCATGATA





GGGGGGAAAAAAG-----ACTCTTCTTTTTTT-----TTT-  
TTTTTTTTAAGGGAAGATGG--AAAT-----GTTAATTAATTAAC-----  
---AACTTTT-----GTT-AATTTCTAATAAT-  
TTCCCGCATTCCCAGGGCGTTTG-TTACTCTCA---AG-  
TGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTTTCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTTCCTCAGGATAGCTTGCCTCGAT-----T-----  
TTAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCTATGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCC??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

Afrodecius\_sp

??  
??  
??  
??  
??  
??  
??  
??  
??AA  
TTACCCACTCCCGGCACGGGGAGGTAGTGACAAAAAATAACGATACGGGACTCTATT  
GAGGCCCGTAATCGGAATGAGTACACTTAAAT-CCTTTAAACAAGGATCCA-  
TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCGCCGCTGTGCGGTGA  
ACTGGCATGTGCGGCATGTCTGTCGGTGGTGAATGGAATCC-----  
-----CTGGTTC---AGACTAGGCT-TTA--CTGCCGAAG-T-TTGTT-CCG-----TG-G-  
TGATACC--GTTTGCCGA---  
TTTCTCTACTTCGGTGCTCTTAAACGAGTGTTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAATAGGACCTCAGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGCGATTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCAGTCTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAAATGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGC?AAGCTGAACT  
TAAAGGAATTGACG??  
??AGT

GCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCACGAGACCGAT  
AGCGAACAAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGA  
GTACGTGAAACCGCTCAGGGGTAAACGGAAAAGACCTGAA--  
TATCCGAAAGGGGAGATTACGT-TTACGAC---TAGCCTAGCTTTAAAGTGC-----  
TTAGAGATGGTTA-TTC-GCCCTATCCAG---TGCA---ACTGTG--GTAG-  
AGGGTTTTGCCGCTCGCATTTT-TTT-----  
AGTGAAGGAAGTGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGACCTGTAAAT--AAT----ATTCTAAGT--AT-A----GTAA--  
AGTGTCTGTAT-TTTAGGTT--ATACTGAAA-----  
GGTTTACCTTTTTTTACAGC-TTTAA--CAAT-----TAC-ATACCGAATA-----  
TTAATTAACAGTATCGATTGGTGG-TGTTGAGCCACTTTTAAAATTTATATA--  
TTTATATATA----GT----TTAAAAGTGTG-----TTTGGTC-GTCTGCAA-GTGGGT---  
-----AGATATTAATT-A-ATGTCGGGCTT-TT-----GTCCGTCTAG-TTA-AATCTCTTA-  
CTGTTGGTA---GAT-AG--TG-CCTCA--GACCG--TGCTTA--  
CAACACCGTTCGGCAGCGATCCAGGTTCAAGGCTCTTATCCGACCCGTCTTGAAACAC?  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
????????

Bannania\_sp                    ???????TATCAAAAACATGTCCTTTATG--TT-----  
TTTAATTTTA-AGGTTTAACCTGCCCAATGA----  
TTTATTTAATGGCCGCAGTATATTAAGTGTGCTAAGGTAGCATAATCATTGTCTTTT  
AATTGTAGGCTAGTATGAAAGGTTGGACGAGGAATTAACCTTTCTTT-ATTATAA--  
TTCTTGAATTTAATTTTTAAGTTAAAAAGCTTA-  
AATTTTTTTATGGGACGAGAAGACCCTATAGATCTTTATATTTAT-ATTT--  
ATTTATTAATTTTAGGATTGA-----  
TGATTTTTTTAAGTGAATTAATTATTTAATTGGGGTGATTGAAGAATTT-  
TGTGAACCTTCTT--TTTGTGTTTTCATAATTT--TATGT--  
ATATTTGATCCA??  
??  
??  
??  
??  
CCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTAAACAAGGATCCA-TTGAGGGCAAGTCT--

GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAATTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTTTCGCACTGTCGGTTCGCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGAATGTCCTGTCGGTGGTGAATAAAAATTC-----  
-----CTGGTGC----TAACCTAGGC-TTTTATAGCTGAAG-T-CGGTG-CCG-----TG-  
GTTGATTTT--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTCTGAAGTAGGCCGACACATTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAAGATCTGCCTAAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGTCAGCCAGCGATCCGCCAATGTTTCTT  
TAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGG??  
??  
GTGCAGCCCTAATTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCACGAGTCCG  
ATAGCGAACAAGTACCGTGAGGAAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAA  
TAGTACGTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA-  
TATTTGAAAGAAGAGATTCACGTCTTAGCAC---TTTGTTTACTTTTAAGTAC----  
CTTTGAGATGGTAA-TTCAGTTTTATCCAG----TGAAT--ACTGTG--GTAA-  
GAGGTTTTGCCGCTCGTACTTT-ATT---  
AAGTAAGCTGTAGTGCTGGACCGCACTTCTCTTTTAGT-  
AGGACGTTGCAACCTGTCAGAT--AAT-----ACTCTAAGT--AT-G-----ATTAA---  
AGTGTCTCTGT-A-AATGTA--TTAAGAC-----TTGTTTT-----  
ATTATATCTACAACAGC-TTTA--TCA--AT---TTT---ACTGGGTATTTA-----  
TTTTTTGACAGTATCAACTGATGG-TGTAGAGCCGTTGATTACTA-----  
---AGCGAGCAGCCG-----TTTTACAA-GTTAGGTG-----TATATAAGTT-A-  
ATGTCTAACAT-CT-----GTTAGTCTAA-CTT-TATCACCTA-CTGTTTGCA----AAA-TG--  
TG-CCTTGC-ATTAG--TGCCTA--  
CAACACCGGTTGGCAGCGATTCAGGTTTCAAGTTCTTATCCGACCCGTCTTGAAACA??  
??  
AAAGTAAAGGATAAAAAGATAAAAG-----AAGTTAAG-----  
CAATTTTCATCTTTGAT-----TCT-TTTATCCTAGGGAAGATGG--ATGT-----  
CTTGTTTTAAAAGC-----TTT-TA-----GT-----TT-  
TCTAAACATAGAT-CTCCCGCATTCCCAGGGCGCTTG--CACTTGCTT--TG-  
CAAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTTCCCTTAGGATAGCTTGCCTCGAT-----T---  
TATAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCTGGCTTTCTTAAAGAAAAAATTTTTTTTTTTTGAAGTCTTGGGATTAGGATCA?  
??  
TTCTAATTCTACCTGGATTTGGACTTATTTACATATTATCGCTATAGAAACCGGGAA  
AAATGAAGCCTTCGGCTCATTGGGGATAATTTATGCTATACTAACAATTGGTTTATTA  
GGATTCATTGTTTGGCCATCATATATTCACAGTAGGGATAGATGTAGACACCCGA  
GCCTACTTTACCTCTGCAACTATAATTATGCTGTACCTACTGGAATTAATTTTCA

GTTGATTAGCTACTCTTCACGGTAGCGTATTGAACCTTCCCTAGATTATTATGAGC  
ATTGGGATTCGTTTTCTTATTCACTATTGGAGGATTAACCGGAGTTATTTTAGCAAAC  
TCAAGAATTGATATTGTACTTCATGATACATACTATGTTGTAGCCCATTTCC?????  
??  
??  
??  
??  
??  
????????????

nr *Bannania* sp

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
GTAATTCAGCTCCAATAGCGT  
ATATTAAAGTTGTTGCGGTTAAAAGCTCGTAGTTGGTCTGCGTGCCGCGCTGTCGG  
TTCGCCGCCTGTCGGTGTACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAA  
TTCC-----CCGGTGC----TACGTAGGCTATTTATTAGCTGAAG-  
T-ATGTG-CCG-----CG-G-TGAGTTC--ATTCGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTCTGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACAAGAGCGAAAGC  
ATTTGCCAAGTATGTCTTAAATTGATCAAGAACGAAAGTTAGAGGTTTCGAAGGCGATC  
AGATAACGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCC  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGG????????????????????????  
??  
??  
??  
AA  
CTCCATCTAAGGCTAAATAGAGCCACGAGTCCGATAGCGAACAAAGTACCGTGAGGG  
AAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGG  
TAAACGGAGAAGACCTGAA--TATTCGAAAGAGGAGATTCACGTCTTACCAC----  
TGCCCGACATTGAAGTAC----ACTGAGATGGTAA-TTT-GCCCTGTCCAG----TTTAT--  
ACTGTT--GCAG-AGGGTTTTGCCGCTCGTATTTT-ATTT--  
AAGTAAGGTGCAGTGGTGGATCGCACTTCTCTTTAGT-  
AGGACGTTGCGACCTGTCAGCA--AAT----ACTCTAAAT--AT-G----ATTAA---  
AGAGCCTGCGT-A-TTGGTA--ATGGAAAAT-----TTTATTATTTTTTA-----  
TGCCCTTATGCAGC-TTTAA--TCAG-AT---TTT---ATTGAGTATTTA-----  
TTTACTGACAGTAACGACTGATGG-TGTCGAGCCGCTTCTGTAAA-----  
----AGGAAGCGTC-----CTTACAAAGCC-GTTTACAA-GTGGGCTAG-----  
GTAATTGGTT-A-ATGTCGAACTT-TT-----GTTCGTCTAA-CTA-GGCCTCCTA-  
CTGTTTGTGA---AAC-TG--TG-CCCTG--GACAG--TGCTTA--  
TACCGCCGGTCCGCGAGCGATTAGGTTTAGGCTC????????????????????  
??ACGCAAGTCATTGGGTGAT-



GGGTAAACGGAGAAGATCTGAA--TTTTCGAAAGGGGAGATTCACGTCTTACCATA--  
TTGTTTAACTTATAAATGTA---TTAAGAGATGTTTA-TAC-GCCCTATTTAG----TGCAA--  
ACTGTA--ATAG-TGGGTTTATTCGCTCGCATTATATT---  
TAGTAAGAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTGTCAGAT--AAT----ACTCTAAGT--AT-G----CATTAA---  
AGTGTCTGTAT-T-TAAATGTGGTTGAAATTGAA--ATTTTTTTTTGTTTTG-----  
CTATATTTTTTTTACAGC-TTAA--TCA--AT---TATAATACTGAGTATTT-----  
TATACTGATAGTAACAAATGATAG-TGTTGAGCCACTTGCT-----T--  
-TTTAAAAGTG-----TCTAGCC-GTTTACAA-GTAGGTATTTAGG----  
CTTAAGCT-A-ATTTTCGGGCTT-GT-----TAGTCCGTATAG-CTG-GGCCACTTA-  
CTGTTGGTA---AACGTGTTTATTCTTA--  
GACAGATTGCTTACAAAACACTGATCAGCAGCGATTGAGG????????????????  
??  
AATGAAAGTAAATAGCTATTTAGTTAACTTAAGAGGAAATTAGAGGGAAAAAATTT  
ATT-TTTTTCTTTATTTTACTTTTT---TGTATAAACTA-  
AATACTTAAAGGAAGAAAG--  
ATTTATATAAAAAAATTTTTTTTATATCTTCATTAAGAGGGAAA-----  
AAAAAAAATTTTT-TT-----TT-----TT-TTTCTTAATGGAA-  
ATCTTGCATTCTAGGGCGCTTT-ATTCTCGCAT--TG-  
CGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTTATT-----A---  
GTTGTGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCTCGGCTT??  
??  
CCAGGATTTGGTATTATCTCCCATGTAATTGCAATAGAAACTGGAAAAAATGAAGCA  
TTCGGGGCTTTGGGTATAATTTATGCTATGTTGGCAATTGGTTTATTAGGGTTCATTG  
TATGGGCACATCACATGTTTACTGTAGGAATAGATGTAGACACACGAGCATATTTCA  
CATCCGCCACAATAATTATTGCAGTCCCAACAGGAATCAAATTTTTTAGATGACTAG  
CCACCTTACATGGTAATAAACTAATTCTCACTCCAAGAAATTTTATGATCCTTGGGATT  
TGTTTTTCTATTTACAATAGGAGGCCTAACAGGAGTAGTACTAGCTAATTCAGAATT  
GATATTATTCTACACGATACATACTATGTAGTAGCCACT????????????????  
??  
??  
??  
??  
??  
??  
??  
??

nr\_Bayerus\_sp2

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??



AATTAATTTTAATGGCCGCAGTATTTAACTGTGCTAAGGTAGCATAATCAATTGTCT  
TTTAATTGAAGGCTTGTATGAAAGGTTGAATGAAACATATACTTTCTTT-ATATTGT-  
AATTTTGAATTTAATTTGTTAGTTAAAAGGCTAA-  
CATTTTTTTATGGGACGAGAAGACCCTATAGATCTTTATATTAAT-AGTT--  
TATATTTAAATTG---TTTTA-----  
TTAATTTAATAAAAAGTTATTAATATTTAATTGGGGTGATTTAGGAATTT-  
TATTAACCTCTAG--TTT--TTATTTCAATTAATA---TATGT--TTAATTGATCCAAT---  
TTTATTGATTTTAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTCTTCGGAGAGTTTCAATTGATGAAGGAGATTGCGACCTCGATGTTGGA  
TTAAA-TAAGTT-TTAGGCGTAGAAGTTG-AATTA-  
CTAGGTCTGTTCGACCTTTAATAT-  
TTTACATGATCTGAGTTCAAACCGG????????????????????????????????????  
??  
CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGGAATCC-----  
-----CCGGTGC---TTCTTGGGCT-TCT-TTAGCCTAAG-T-ATGTG-CCG-----TA-C-  
GGATGCC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGCTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGG??  
??  
AG  
AGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACACTACGAGTCC  
GATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCA  
AGAGTACGTGAAACCGCTCAGGGGTAAACGGAGAAGAACTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATA--TTGTTTAGCTTTTAAATGT----  
TCAAGAGATGTTTA-TTC-GCCCTATTCAG----TGCAA--ACTGTA--ATAG-  
AGGGTTTATTCGCTCGCATTTA-ATA---  
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTGTCAGTT--AAT----ACTCTAAGT--AT-G----GTAA---  
AGTGTCTGTAT-A-AAGAAA--ATTTA-----GTTTCGGCTT-----  
TTTTTTCTTTTTACAGC-TTTAA--TCAT-----TTT-ATACTGAGTATTT-----  
TAACTTATGACAGTATCAATTGATGG-TGTTGAGCCACTTTATAACTT-----  
----TT--TCTAAAGTGTC-----TTTAGCCGTTTACAA-GTAGGTTT-----  
AGTAGTT-A-ATTTCCGGCTT-TT-----CGGTCCGTATAA-CTA-GGCCACTTA-  
CTGTTAGTA----AAC-TG--TG-CCTTA--GACTG--TGCTTA--  
TAACACTGATCAGCAGCGATCCAG????????????????????????????????????  
??  
ACCCAAAGGCATAATAAAAAGTAAA  
GAGTAATTTAATTATAA-----AAG-----TGTTAGTACTTTTTTT-----  
TATTA-AATCTTAAAGGAAGATAG--ATTT-----



AATTTCTTTAATAGAAAAAGAGTGAT-----ATTTTTTATTTCTC-TT-----TT-----  
-----TT-TTTTTTTAGAAAT-ATCTTTCATTCTAGGGCGCTTG-TTTCTCGCAT--TG-  
CGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT-----  
GAAACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCTCGGCT??  
??  
??  
TATATATTTAATCTTACCCGGA  
TTTGGTATTTTACACTCTATCAGTATAGAGACCGGAAAAAATGAAACCTTTGGA  
GCACTAGGAATAATCTATGCTATAATTGCCATTGGATTACTAGGATTTATTGTATGAG  
CACACCATATATTTACTGTAGGAATAGATGTAGATACACGAGCATATTTCACTTCAG  
CAACAATAATTATTGCCGTACCCACAGGAATTAATAATTTTATAGATGACTTGCAACCTT  
ACACGGCAAAAACATCGAGTTCACGCCAAGTATATTATGAACATTAGGATTCGTTTT  
CCTATTCACCATTGGAGGATTAACCTGGTATCATTCTAGCAAATTCAAGAATTGATATT  
ATTCTACATGACACTTATTATGTAGTAGCCATTTCC????????????????????  
TGTCTGGCTTYTGTACAGTTAAAACATGCTGGATGCGTTTACCACCATTTTCGTCTGT  
TGGAGATAATTTAAAAGATCGATTTGATGGCGCATCACGTGTAATGGTTAGTAATGC  
TGGCTCAATACGTGCAA--  
CTGGAGGACGTAGAAATCGTTATAATTTCCAATTRAACCSTACAATCCTGATCATA  
AACCGCCTGGTGTAAAGATCTAGTATACCTTGAACCATCGCCAGTATTCTGTGAAC  
GTAATCCTAGATTAGGTATAACAAGGAACACACGGACGCCAGTGTAACGCTACATCAC  
AGGGTGTTGAAGGATGC????????????????????  
          Brontostoma\_nanus          ????????????????CATGTCCTTTTGT--TT----  
GATTTATTTA-AGGTCGGCCCTGCCCATGA----  
TTCGTTTAAATGGCCGCAGTATTTAACTGTGCTAAGGTAGCATAATCAATTGTCTTTT  
AATTGGGGGCTAGGATGAACGGGTTTATGAAGCACTTCTTTCTTT-TTATTAT-  
TTATATGAATTTAATTTTATAGTTAAAAAGCTAA-  
TATTTTATTAAGGACGAGAAGACCCTATAGAGCTTTATATTTAA-TTAT--  
TTGTTTAAATTTTT---GTTAATAT-----  
TTTTTATACAATTGAGTTAATTATTTGATTGGGGTGATTTTGGTATTT-  
TTATAACTACTTT--TTA--AAGTTTCATTTATG---AATGT--TTTTATGATCCATT---  
ATTTATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTTATTTTCTTGGAGAGTTCATTATTGATAAGGAAGTTTGCACCTCGATGTTGGAT  
TAAAA-TAAGTA-ATGGGCGTAGCAGTTC-ATTTA-  
CTGGGTCTGTTTCGACCTTTAAAGT-  
TTTACA??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

??  
????????????GAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAAC  
TACGAGTCCGATAGCGAACAAGTACCGTGAGGGGAAAGTTGAAAAGAAGCTTTGAAGA  
GAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGGAGAAGAAGTAA--  
TTTTCGAAAGGGGAGATTACGTCTTACCATATTATGTTAACTTTTAAGTGT----  
ACAAGAGATGTTTA-TTC-GCCCTATTCTG----TGCAA--ACTGTA--ATAG-  
AGGGTTTATTTCGCTCGCATTTA-TAT----  
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTATCAAAT--AAT----ATTCTAAGC--AT-G----GTTAA---  
AGTGTCTGTAT-A-AGAAGA--ATAGT-----TTTATTTTAT-----  
AAAGTTATTTTTTTTTTATACAGC-TTTAA--TCA--AT---TTT-ATGCTGAGTATAA-----  
-T--TATATTGATAGTATCAAATGATAG-TGTTGAGCCACATTT-----  
AA--ATAAATGTGTT-----TTAAGCC-GTTTACAA-GTAGGTTT-----  
AGTAGTT-A-ACTTCGGACTT-TT-----CGGTCCGTATAA-CTATCACCCTTA-  
CTGTTAGTA----AAC-TG--TG-CCTTT--AACTG--TGCTTA--  
TAACACTGATCAGCAGCGATCCAGGTTCACTCTTATCC????????????????????????  
??AGGCATAATGAAAGT  
AAAGGGCAATTTAAATAAGT-----AGCAATAC-----TTT-----  
TTTTA-AATTCCTAAAGGAAGAGAG--ATAT-----TTTTTTCCTTTGTATTTTAA---  
-----TA-----TT-AAAGGAAAAAATT-  
ATCTTGCATTCTAGGGCGCTTG-TAACTCGCAT--  
TGTCGAGTACATGCGCACCCAAAGCGTACACGATGATACCCGAAAGATGGTGAAC  
ATGCCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTG  
ACGTGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATC  
TAGTAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTTATT-----  
GATGCGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAT-  
CCTCGGCTT??  
??  
CCAGGATTCGGCATTATCTCACATACAATTGCCATAGAAGCAGGAAAAAATGAAGCT  
TTTGGAGCATTAGGAATAATTTATGCAATACTAGCTATTGGTTTACTTGGATTCTTG  
TCTGAGCACACCATATATTTACTATCGGGATGGACGTTGATACACGTGCATACTTTAC  
CTCAGCCACCATAATTATTGCAGTACCCACAGGTATTAATAATTTTTAGATGACTAGCC  
ACATTTTACGGAAGAAATCTTCCCCTTACGCCAAGAACATTATGATCATTGGGGTTC  
GTGTTTCTATTCACTATTGGGGGATTAACAGGGGTAATCTTGGCTAATTCTAGAATTG  
ATATTATTCTTCATGATACCTATTATGTAGT????????????????????????????????  
ATGTCTGGCTCATGCACAGTGAAAACATGCTGGATGAGATTACCACCATTTTCGTGTT  
GTCGGCGATAATCTGAAAGATCGCTTTGACGGTGCATCCCGCGTGATGGTCAGCAAT  
GCTGGCTCAATACGCGCAA---  
CTGGCGGTCTGAGAAATCGATATAACTTTCAATTGAAACCATAACAATCCTGATCATA  
AACCGCCAGGCGTTAAAGATCTAGTATACCTTGAACCGTCGCCAGTCTTTTGCGAAC  
GTAATCCTAGATTAGGTATAAAGGCACGCACGGGCGGCAATGTAATGCAACGTCAC  
AGGGCGTTGAAGGATGCGATCTGATGTGTTGTGG  
Brontostoma\_sanguinosum ??????????AAAAACATGTCTTCTTGT--T-----  
TATTTATTG-AAGTTTAACTGCTCAATGG--  
GTTTTCTAAATGGCCGAGTATCCTAACTGTGCTAAGGTAGCATAATAAATTGTCTT  
TTAATTGAAGGCTTGATGAAAGGTTGGATGAAATATAAGCTTTCTTT-ATATTAA-  
AATCATGAATTTAATCTTTTGGTTAAAAAGCCAA-

TATCTTTTTAAGGGACAAGAAGACCTATAGATCTTAATATCTGA-GTAG--  
TTATACTTAACTT---AGTTAATT-----  
TGTAAGTTATAAATTCTTAAATATTTAATTGGGGTGATTTTTAGTATTT-  
TAATAACTTCTAA--CT---TTCTTTCATTGATT---TATGT--TTATTTGATCCAGT---  
TATATTGATTATAAGACTAAGTTACCTTAGGGATAACA-  
GCGTTATTTCTTCGGAGAGTTCATATCGATGAAGGAGTTTGCACCTCGATGTTGGAT  
TAAA-TAAGTC-  
TTGGATGTAGTG??  
??GTAATCGGA  
ATGAGTACACTTTAAAT-CCTTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCTGTCGGTGGTGAATGGAATCC-----  
-----CCGGTGC---TGCCTGGGCT-TCT-TTAGCCCAAG-T-ATGTG-CCG-----TA-C-  
GGACGCC--ATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCGGGGGAAGTATGGTTGCAAAG????????????  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
TACGCAAGTCATTGGGTAA  
T--TAAAACCAAAGGCGCAATAAAAAGTAAAGAGCAATTTAAATAAGT-----  
ATTAGTAC-----TTTTT-----TTTTA-AATTCTTAAAGGAAGACAG--  
ATTT-----TTAACTGCTTTAAT-----AAGAAAATTC-----  
-----TT-TTTTAAAGGAGAA-ATCTTGCATTCTAGGGCGCTTG-TTTCTCGCAT--TG-  
CGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGCG  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT-----  
AAAACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCTCGGCTTTCTTAAA-----  
TTGAAGCCGTAGGATTTGGATCAGAGTGC????????????????????????????????????

??TATATTCTAATTCTACCTGGATTTCGGAATTATTTCTC  
ACACTATTGCAATAGAAACCGGTA AAAAATGAAGCATTGGAGCACTAGGTATAATTT  
ATGCAATAATTGCTATTGGATTATTAGGATTCATTGTATGAGCCATCATATATTTAC  
AGTAGGAATAGATGTAGATACACGTGCATACTTCACCTCAGCAACTATAATCATTGC  
TGTTCTACAGGAATTA AAAATCTTCAGATGACTAGCCACATTACACGGAAGTAATAT  
TAACCTTACCCCAAGTATAATGTGAACTTTAGGGTTGTATTCTTATTACCATTGGA  
GGATTA ACTGGAGTCATTTTAGCAAATTCAAGTATTGACATTATTTTACATGATACCT  
ACTATGTAGTAGCTCACTTCC????????????????????????????????????ATGTCTGGCTCATGTAC  
AGTTAAAACATGCTGGATGCGTTTACCACCGTTTCGTGTTGTTGGAGATAATTTAAAA  
GATCGTTTCGATGGTGCATCACGTGTAATGGTTAGTAATGCTGGCTCAATACGTGCA  
A---  
CTGGCGGACGTAGGAATCGTTATAACTTTCAATTA AAAACCCTATAATCCTGATCATA  
AACCTCCTGGTGTTAAAGATCTAGTATATCTTGAACCATCGCCAGTATTCTGTGAACG  
TAATCCAAGATTAGGTATACAAGGAACACATGGACGTCAGTGAATGCTACGTACA  
GGGTGTTGAAGGATGCGATCTGATGTGTTGTGG  
Caecina\_affinis ???  
TGTTTATTTA-  
AGGTCTGGCCTGCCAATGAATATTAATTTTGAATGGCCGCAGTATCTTAACTGTGCT  
AAGGTAGCGTAATCATTTGTCTTTTAAATTGTTGGCTAGTATGAAGGGTTGGACAAAGT  
GTAATCTTTCTTT-ACTTTAA--TTAATGAATTTAATTTTAAAGTTAAAATGCTTA-  
AATTTTATTAAGGGACGATAAGACCCCGTAGAGTTTAAATTTTAT-TTAA--  
TTAATTTTAAAGTATTTGATTA-----  
TGTATAAATTAATTTATTTGAAATATTGGATTGGGGTGATTATTGAATTT-  
GTTTAACTTCTAT--TTATTTAGTTTCATTTATT---TTTGT--TTTTTTGATCCATC---  
AGTGATGATTA AAAAGATTAAATTACCTCGGGGATAACA-  
GCGTAATTTTTTCGAAAGTTCATATTGATGAAAGAGTTTGCACCTCGATGTTGGAT  
TAAAA-TTAGTT-TTTGATGCASTAGTTA-ATTAA-CTAGGTCTGTTTCGACCTTTAAATT-  
TTTA??  
??  
??  
TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCGGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGCCGGTTCGCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAATC-----  
-----CTGGTAC----TGCGTAGGCT-TTAAATAGCTGAAG-T-TTGTG-CCG-----TG-G-  
TAATTTCT--ATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACATTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATCTCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
ATAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCC????????????????????????????????????  
??  
??  
??  
??  
??  
AAATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGA  
AAAGA ACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGA

GAAGATCTGAA--TGTTGCGAAAGAGGAGATTCACGTCTTACCAG---  
TTGCTCTACTTTGAAATAC----TTTTGAGATGGTAA-TTC-GCCCTATCCAG----TGTAT--  
ACTGTG--GTGG-TGGGTTTTGCCGCTCGTATTC-ATT----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTATCAGAC--AAT----ACTCTAAGT-TTT-A----ATTAA---  
AGCGCCTGTGT-A-TTGGTT--GTATATT-----TATTTATA-----  
TAATCTTTATACAGC-TTTAA--TAA--AT----TTT---ACTGGGTATTTA-----  
TTTTCTGATAGTAACGACTGATGG-TGTAGAGCCACTTTAC-----TATATA----  
ATG----TAAAAGTGTCC-----GATTTTTTGGGC-GTTTACAA-GTGAGTTTAG----  
GCTTTTTAGTT-A-ATGTCGAACCTT-TT-----GTCGTCTAA-CTA-TGCCGCTTA-  
CTGTTTGTA---AAC-TG--CG-CCTG--GACAG--TGCTTA--  
CAACACCGGTTCGGCAGCGATTACAGG????????????????????????????????  
??  
??  
TTTTTTTTTAATCTTTGAT-----TCT-  
TTGATTCAAGGGAAGATGG--GTTA-----TCTTAATATTTTTTTTCT-----  
-----TT-TAATTAAG-----GA-AAAAAAGTTTTGA-  
TTCTCGCATTCCCAGGGCGCTTGTTTTCTCGCT--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGA????????????????  
??  
??  
AGTTTATATTTTAATTTTACCCGATTGGATTAATTTCCCAT  
ATTATTTCTATAGAATCCGGAATAAATGAATCATTGGTTCATTAGGAATAATTTACG  
CGATATTAACCTATTGGATTATTGGGATTATTGTATGAGCCATCATATATTTACAGT  
AGGAATAGATGTAGATACTCGAGCTTACTTCACATCAGCAACAATAATTATTGCTGT  
TCCTACAGGGATTAATAATTTTATAGATGAATTGCAACACTTCACGGATCAAACCTAAC  
CCTATCTCCTAGAATATTATGAGCATTAGGATTCGTGTTTCTTTACTATTGGAGGA  
TTAACGGGAGTAATACTTGCAAATTCAGAATCGATATTACCCTCCATGATACTTATT  
ACGTAGTTGCACACTTCCATTAT????????????????????????????????  
??  
??  
??  
??  
??  
??  
Caecina\_n\_sp ??????????????ACATGTCCTTAAGT-----  
TTTAATTTTT-AGGTTTGACCTGCCAATGA--  
ATAATATTTGAATGGCTGCAGTATTTAACTGTACTAAGGTAGCATAATCATTGTCT  
TTTAATTGTTGGCTGGAATGAATGGTTGGACAAAATATAAACTTTCTTT-TTTTTAA--  
TTAATTAATTTAATTTTAAAGTTAAAAAGCTTA-  
AATTTTTTTAAGGGACGATAAGACCCCGTAGATTTTTATATCTAT-TTTT--  
ATTGAATTATTG---  
GTTATAATTTTATAATTTGTTTTATTTAATTAGGTATTAAGTTGGGGTGACTATTGAAT  
TT-TTTAACTTCAAT--TTTATATATTTCAATTTATTTAGTTGTT--TTTTTTGATCCAAT--  
TTTTTTGATTAGAAGATTAAATTACCTCGGGGATAACA-  
GCGTAATTTCTTTGGAAAGTTCAATTTGATAAAGAAGTTTGCACCTCGATGTTGGAT  
TAAATTTATTT-TTTGGTGCAGTAATTA-



??  
??  
??  
??

Centraspis\_ducalis      ???????TATCAAAAACATGTCCTTTAGT--TG----  
TTTATATTTA-AGGTCTAACCTGCCCAATGA---  
TTTTTATTTAATGGCCGCAGTATTTTAACTGTGCTAAGGTAGCATAATCATTGTCTTT  
TAATTGAGGGCTGGAATGAATGGTTGGACGAGAACTTTCTTCTTT-TATTTAA--  
TTAATGAATTTAACTTTTTAGTTAAAAGGCTTA-  
ACTTTTTTTATGGGACGAGAAGACCCTATAGATCTTTATATTTTAAATAT-  
TATATATTTGTTA--GTTATT-----  
GATTAATTTTATTATTTGATTTATTTTGTGGGGTGATGCAGGAATTT-  
AATGAACTTCTG--TTTTATTTATTCATTTATT---TATGT--TTTTTTGATCCATT----  
TTTATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTTCGGAGAGTTCATATCGATGAAGAAGTTTGCACCTCGATGTTGGAT  
TAAA-TAAGTT-TTGGCGTAGTAGTTC-AATTA-CTAGGTCTGTTCGACCTTTAATTT-  
TTTACATGATCTGAGTT??  
??  
??  
??  
CCTTTAACAAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTGCGGTGT  
ACTGGCATGTCGCGCATGTCCTGTCGGTGGTGAATGAAATCC-----  
-----CCGGTGC---TACGTAGGCT-TTTATAAGCTGAAG-T-ATGTG-CCG-----TG-G-  
TGACTTC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCC-  
ACCCGGGAAACCAAAGCTTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAA??  
??  
??  
??  
??  
AGTTGAAAAG  
AACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGAGAAG  
ATCTGAA--TATCCGAAAGAGGAGATTACGTCTTACCAC---  
TTGCTCGACTTTAAAGTGC---TTTTGAGATGGTAA-TTC-GCCCTATCCAG----TGTAT--  
ACTGTG--GTAG-AGGGTTTTATTGCTCGCATTTT-ATTT--  
AAGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAAAT--AAT----ACTCTAAGT--AT-G----GTTAT--  
AGTGCCTGTCT-A-AAGGT--ATAAAAAT-----TTTTTTA-----  
TCACTTTTTTACAGC-TTTAA--TCA--AA---TTT---ACTGGGTATTTG-----  
TTTTTTGACAGTAACGATTGATGG-TGTAGAGCCACTTAAACATTTTT-----  
ATG----TTAAAGTGTT-----AATTTATTTGCC-GTTTACAA-GTGGGTTA-----  
GGTATTAGTT-A-ATGTCGAACTT-TT-----GTTCGTCTAA-CTA-AGCCGCCTA-  
CTGTTTGTA---AAC-TG--TGCCCTTG--AACAG--TGCTTA--

CAACACCGGTCGGCAGCGATC??  
??AGTAAAGAGTAGA  
AAGATAAAAG-----ATAAATTTT-----CTAATTTTCATCTTTGAT-----  
TCTCT-CTCTCTCAAGGGAAGATGG--AAAT-----CTTATTTTGAATAA-TATTTAG----  
-----TAT---TT-ATATTAAG-TAA-----GTA-TTTAAAATATAGA-  
TTCCCGCATTCCCAGGGCGCTTGTTTTCTCGCTT--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTC??  
??  
TTTCCCATATTATTGCAATAGAAACAGGTAAAATGAAGCCTTTGGGTCAATTAGGTA  
TAATTTATGCTATATTAACAATTGGATTATTAGGATTTATTGTATGAGCTCATCATAT  
ATTCACAGTAGGAATAGATGTTGACACTCGAGCTTACTTTACCTCAGCCACTATAATT  
ATTGCTGTTCCCTACAGGAATTAATAATTTTATGTTGATTAGCTACTTTACATGGAAGAA  
TTTTATTATTACCCCTAGACTTTTATGAGCACTGGGATTCGTATTTCTATTCACTATT  
GGGGGATTAACAGGAGTAATTCTAGCAAATTCAGAATTGATATTATTCTACATGA??  
??  
ATGTCTGGATCATGTACAGTA  
AAAACATGTTGGATGCGTTTACCACCATTTCGTGTAATTGGTGATAATTTAAAAGATC  
GTTTTGATGGAGCATCACGTGTTATGGTAAGCAATTCTGGATCTGTACGCGGTA---  
CTGGTGGACGTAAGAGTCGTTATAATTTCCAATTGAAACCCTATAATCCAGATCATA  
AACCACCAGGTGTTAAAGATCTAGTATATTTAGAACCATCACCATTATTTGTGAAC  
GTAATCCAAGACTAGGWATACAGGGTACGCATGGACGACAGTGTAATGCAACATCG  
CAAGGTGTTGAAGGTTGTGATCTAATGTGTTGTGG

Cimbus\_sp

??  
??  
??  
??  
??  
??  
??  
??AGCAGGCAC  
GCAAATTACCCACTTCCGGCACGGAGAGGTAGTGACAAAAAATAACGATACGGGAC  
TCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCTGTCGGTGGTGAATGTGATCC-----  
-----CCGGTGC---TACCTAGGCT-TGAATTAGCTGAAG-T-ACGTATCCG-----TG-G-  
TGATGGC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATTTCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA



TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTAATGACTCGGCGGGGAGCTTCCGACCCGGGAAACCAAAGCTTTTGGGTTCCGGGG  
GAAGTATGGTTGCAAAGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGA  
GTGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAACCTCACCAGGCCAGGACATT  
GGAAGGATTGACAGATTGATAGCTCTTTCTTGATTGAGTGGGTAGTGGTG?????????  
??  
AGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGA  
GTACGTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTACCAA---TAGCTCTACTTTAAAGTGT----  
ATTTGAGATGATAA-TTC-GCCCTATCCAG----TGTATACACTGTG--GTAG-  
AGGGTTTTATTGCTCGCACTTTTATTT--  
AAGTAAGGTGCAATGGTGGACCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAA--ATTA---ACTCTAAGTAAAT-A----ATTAA---  
AGTGTCTGTTT-T-AAAATT--TTAGTAA-----ATTTTTTA-----  
TTAACTTTTTATACAGC-TTTAA--TAA--TTAATGTTT---ACCGAGTTTTTTTTTG--  
CTTTATTTCTGACAGTAACGACAGAAAG-TGTAGAGCCGCTACTTAACA-----  
ATGTGTGT-----TAAAAAGCGTT-----CATTGTTTGCCT-ATTTACAA-  
GTAGGTTTTA----GGTAATTAGTT-A-ATGTCGAACTT-TA-----GTTTCGTCTAA-CTA-  
AGCCGCCTA-TTGTTTGTA---AAC-TG--TG-CCTTG--AACAG--TGCTCA--  
CAACACTGGTCGGCAGCGATCCAGGTTTACTCTTATCCGACCCGCTTGAACAC  
GG????????????????????GACCAAGGAGTTTATTGTGTACGCGAGTCATTGGGCGAT-  
AAAAACCTAAAGGCGTAATGAAAGTAAAGAGTTGAAAGATAAGAA-----  
AATAAATT-----TTCCTTTTATCTTTGAT-----TCT-  
TTCTCTCAAGGGAAGATGG--AAA-----CTCTAAATATTTTT-----  
-----TAGAA-TTCCCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTTCGCG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGA-----TG--  
AAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCTGATCAGAGTGCAAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGC-----  
GTCAACATTTATTTTGGATTTTTTGGACATCCAGAAGTATATATTTTAATTTTACCAGG  
ATTTGGATTAATTTACATATTTTCTATAGAAACAGGAAAAAATGAAGCATTG  
CTCATTAGGAATAATTTATGCTATAATTACTATTGGACTTTTAGGATTCATTGTATGA  
GCTCATCATATTTACTGTAGGAATAGACGTAGACACACGAGCATACTTCACATCA  
GCAACAATAATTATGCAAGTGCCTACAGGTATTAATAATTTTATAGATGATTAGCTACT  
TACATGGCAGAAATTTAATTTTTCCCAAGACTATTATGATCATTAGGATTTGTATT  
TTATTTACTATTGGAGGGTTAACAGGAGTTATTTTAGCTAATTCGAAGATCGATATT  
GTACTCCATGACACTTATTATGTAGTAGCCATTTCCATTATGTTTTATCAATAGGAG  
CAGTATTCGA????TGGATCATGCACAGTAAAAACATGTTGGATGCGTTTACCACCGT  
TTCGTTTAATTGGCGATAATTTAAAAGATCGTTTCGACGGTGCATCACGCGTTATGGT  
AAGCAATTCTGGTTCGATACGCAATA---  
CTGGTGGACGTAAAAGTCGTTATAATTTCCAATTGAAACCGTATAATCCGGATCATA

AACCACCTGGAGTTAAAGATCTAGTATATCTAGATCCGTCGCCTTTATTCTGTGAACG  
TAATCC??  
?????

Cleptria\_corallina CGCCTGTTTATCAAAAACATGTCCTTTTGT--  
TTAATTTAATGTATTTA-AGGTCTGGTCTGCCCAATGAT-  
TTAATTATTTAATGGCCGCAGTATTTAACTGTGCTAAGGTAGCATAATCATTGTCT  
TTTAATTGAGGGCTTGTATGAATGGCTTGACGAATCATTATCTTTCTTT-TGATTAA--  
TATATGAATTTAATTTTTTTGTTAAAAAGCTAAAGATTTTTTTATGGGACGAGAAGAC  
CCTATAGAGTTTAATATTTAT-ATTT--ATTAATTTTTTTA---GTTA-----  
TTTTTTATTTAATTAATTGTGTATTTTCGTTGGGGTGTATGTTAAATTT-  
TAATAACTTTTAT--TTTTATGTTTTTCATTTTTT---TATGT--TTTTTTGATCCATT---  
GTTTATGATTATAAGATTAAATTACCTTAGGGATAACA-  
GCGTAATTTCTTTGGAGAGTTCTTATTGATAAGGAAGTTTGCGACCTCGATGTTGGAT  
TAAA-TAAGTT-TTGGGTGTAGTAGTTAATTTA-  
CTAGGTCTGTTTCGACCTTTAATTT-  
TTTACATGATCTGAGTTCAAACCGGAGGCAGCAGGCACGCAAATTACCCACTCCCGG  
CACGGAGAGGTAGTGACAAAAATAACGATACGGGACTCTATTGAGGCCCGTAAT  
CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAATCC-----  
-----CCGGTGC---TATGTAGGCT-TTTATTAGCTGAAG-T-ACGTG-CCG-----TT-G-  
TGACTTC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACCACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTGACAGATTGATAGCT  
CTTTCTTGATTGAGTGGGTAGTGGTGCATGGCCGTTTC?CGTGTGCTTGATAGTGCAG  
CCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCACGAGTCCGATAGCG  
ACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGAGTAC  
GTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCAC---TTGCTCGACTTTGAAGTGC----  
TTTTGAGATGGTAA-TTT-GCCCTATCCAG----TTTTT--ACTGTG--GTAG-  
AGGGTTTTACCGCTCGCATTTT-ATA----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT-----ACTCTAAGT--AT-G-----GTTAA---  
AGTGCCTGTCT-T-TAGGTT--ATAAAAT-----TTATTTA-----  
TAATCTATTTACAGC-TTTAA--TCA--AT---TTT---ACTGGGTATTTA-----  
TTTTCTGACAGTAACGATTGATGG-TGTAGAGCCACTTTT-----  
ATAAAGTGTCTCT-----TTTTTTTTGCC-GTTTACAACGTGGGTTTA-----  
GGTATTAGTT-A-ATGTCGAACTT-TT-----GTTTCGTCTAA-CTA-ACCCGCCTA-





TAATCCAAGACTAGGTATACAGGGCACCCACGGACGTCAGTGTAACGCTACATCACA  
GGGTGTGGAAGGTTGTGATTTAATGTGTTGTGG  
Daraxa\_sp ??????????????ACATGTCCTTTAGT--TT----TATTTATTTA-  
AGGTCTGTCCTGCTCTATGA---  
TTAATATTTAATGGCCGCAGTATATTAAGTGTGCTAAGGTAGCATAATCAATTGTCCT  
TTAATTGTGGGCTAGAATGAATGGTTGAATGAGGCATTATCTTTCTTTTATTTAT--  
TTTTTGAATTTAACTTTTAAGTTAAAAGGCTTA-  
TATTTTATTAAGGGACGAGAAGACCCTATAGATTTTAATTAGTTA-TTAT--  
TTTAATTTTATTTATTGTTTAAAT----  
TTAATGGTCTATTTAATAATTAATTTAGTTGGGGTGATTTTAGAATTA-  
TATTAACCTCTAT--TTTTAATTTTTCATTAATT---TATGT--TTATTTGATCCATT---  
ATTATTGATTATAAGATTAAATTACCTTAGGGATAACA-  
GCGTAATTTTTTCGGAGAGTTCTTATTGATGAAAAAGTTTGCGACCTCGATGTTGGAT  
TAAA-TTAGTA-AAGGGTGTAGCAGTTC-ATTAA-  
CTAGGTCTGTTTCGACCTTTAAAAT-  
TTACATG??  
??  
??  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCCTGTCGGTGGTGAATGGAATCC-----  
-----CCAGGAC----TTCTTAGGCT-TC--ATAGCTTATG-T-ATGTT-CTG-----TA-A-  
GGACTCC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGG??  
??  
??  
AG  
AGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATATAACTACGAGTCC  
GATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCA  
AGAGTACGTGAAACCGCTCAGGGGTAAACGGAGAAGAAGTAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATA--TTGTTAACTTTTAAATGT----  
ATATGAGATGTTTA-TTC-GCCCTATTGAG----TGCAA--ACTGTA--ATAG-  
AGGGTTTATTCGCTCGCATTTA-ATT----  
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTGTTAGAT--AAT----ACACTAAGT--AT-G----ATTAA---  
AGTGTCTGTAT-T-GAAGAA--AGGAA-----ATTTTTT-----  
CTTTCTATTTTACAGC-TTAA--TCA--AT----TAT-ATACTGTGTATTT-----T-  
ATTTACTGACAGTATCAATTGATGG-TGTTGAGCCACTTATT-----  
-TTTCAAGTGTC-----TTAGCCTGTTTACAA-GTAGGTTT-----ATTAGTT-A-  
ATTTCCGGGCTT-TT-----TGGTCCGTATAA-CTA-GGCCACTTA-CTGTTAGTT----AAC-  
TG--TG-CCTTA--GACTG--TGCTTA--









????????????????????????????????AAGTATACATTTTGTATTTACCCGGATTTGGTTTAATTTCC  
CCACATTATTGCTATAGAAACAGGAAAAAATGAAGTTTTTGGGACATTAGGTATAAT  
TTATGCCATACTAACTATCGGTTTATTAGGATTCATTGTTTGGAGCACATCATATATTT  
ACAGTAGGTATAGATGTAGATACACGAGCCTACTTTACATCAGCAACAATAATCATC  
GCTGTACCTACAGGTATTAATAATTTTCAGATGATTAGCCACCCTACATGGAAGAATT  
TTAATATTTTCACCCAGCTTGTATGAGCACTAGGATTCGTTTTCTATTACAATTGG  
GGGGTTAACGGGAGTAGTTTTAGCAAATTCAGAATTGATATTATCTTACATGATAC  
ATATTATGTAGTAGCACA??  
??  
??  
??  
??  
??  
??

*Distirogaster tarsalis*

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
CCGTAATCGGAATGAGT  
ACACTTTAAAT-CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGGAATCC-----  
-----CCGGTGC---TACGTAGGCT-TTTATTAGCTGAAG-T-TTGTA-CCG-----TG-G-  
TGAATTC--ATTTGCCGA--  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCAGGGGGCATTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAGCTTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACG??  
??  
??  
????????????????GTTGTAATACTCCATCTAAGGCTAAATATGACCACGAGTCCGATAGCGAA  
CAAGTACCGTGAGGGAAAGTTGAAAAGAAGCTTTGAAAGAGAGGTTCAAGAGTACGT  
GAAACCGCTCAGGGGTAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTACCAC---TTGCTCGACTTTAAAGTGC---  
TCTTGAGATGGTAA-TTC-GCCCTATCCAG---TGCAT--ACTGTG--ATAG-  
AGGGTTTTATCGCTCGCATT--ATT---  
AGTAAGGCGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAGT--AT-A----ATTAA---  
AGTGCCTGTCT-A-AAAAGG--TTTTATT-----AATTTATT-----  
ACCTTTTTTTACAGC-TTTAA--TTA--AT---TTT---ACTGGGTATTTA-----



TTGCGACGTTAGAGGTGAAATTCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACG??  
??  
??  
??  
AGTTGA  
AAAGA ACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGA  
GAAGATCTGAA--TATCCGAAAGAGGAGATTCACGTCTTACCAC---  
TTGTTCAACTTTAAAGTGC---TTTTGAGATGGTAA-TTC-GCCCTATCCAG---TGTAT--  
ACTGTG--ATAG-AGGGTTTTATTGCTCGCATT--ATTT--  
AAGTAATGTACAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT-----ACTCTAAGT--AT-G----GTTAA---  
AGTGTCTATTT-A-AAGGTT--ATAAATTA-----TAATTTAT-----  
TACCCTTTTATAGC-TTTAA--TCA--AT---TCT---ACTGAGTGTTT-----  
TTTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTAAGCATT-----TTATA-  
---ATG----TAAAAGTGTCTA-----AATTATATTGCC-GTTTACAA-GTGGGTTTTA-----  
GGTATTAGTTAA-ATGTCCA ACTT-TT-----GTTCTGCTAA-CTA-AGCCGCCTA-  
CTGTTTGTA---AAC-TG--  
TG??  
??  
ACCCAAAGGCGTAATG  
AAAGTAAAGAGTGAGTAGAGAAGAAAGATTAAT-----AAGATAAAAATT-----  
TTATTTTTAACTTTGCTTTATCT--CTCTTTCTCT-CTTACTCAAGGGAAGATAG--AAAT-  
-----CTTATTTTAGATAA-TATTTTAAATAATTTTTTTTTTTTTTTTATAAAT---TT-  
ATATTAAG-TA-----AA-TTTAAAATGTAGATTTCTCGCATTCCCAGGGCGCTTG-  
TTTCTCGCTT--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGA ACTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAA ACTCAAATAGATGAGAC-  
CCCCGGCTTTCT??  
??  
AGTATATATTCTTATTTTACCAG  
GATTCGGCTTAATCTCCCATATTATTGCTATAGAGACAGGGAAAAATGAAGCATTG  
GATCACTTGAATAATTTATGCTATATTA ACTATTGGATTATTAGGATTCATCGTTTG  
AGCTCATCATATATTTACAGTAGGAATAGACGTTGATACACGAGCTTATTTTACATCA  
GCTACTATAATTATTGCTGTACCTACAGGAATTTAAATTTTCAGATGATTAGCAACCT  
TACATGGAAGTATTTTAAGATTTTACCAAGATTACTTTGAGCATTAGGCTTCGTGTT  
CCTTTTACAATCGGTGGATTAACCGGAGTAATTTGGCTAATTCAAGAATTGATATT  
ATTTTACATGACACTTATTATGTTGTTGCACACTTCCA????????????????  
??TCTGGATCATGTACAGTAAAGACAKGTTGGATGCGTTTATCACCATTTCTGTATT  
GGTGATAATTTAAAAGATCGTTTTGATGGCGCATCACGTGTTATGGTAAGCAATTCT  
GGATCAATACGAGGGA---  
CTGGTGGACGAAAGAGTCGTTATAATTTCCAATTGAAACCCTATAATCCAGATCATA  
AACCACCAGGAGTTAAAGATTTAGTATATTTAGAACCATCGCCATTATTCTGTGAAC

GTAATCCAAGACTAGGTATACAGGGTACGCATGGACGACAGTGTAACGCAAC??????  
????????????????????????????????

*Ectrichodia lucida* CGCCTGTTTATCAAAAACATGTCCTTTAGT-----  
TTTAAATTTA-AGGTCTGGCCTGCCAATGA---  
TTTTTATTTAATGGCCGCAGTATTCTAACTGTGCTAAGGTAGCATAATCAATTGTCTT  
TTAATTGAGGGCTGGAATGAAAGGTTGGACGAGGAATTTACTTTCTTT-TAAATAA--  
TTAATGAACTTGACTIONTTAGGTTAAAAGGCTTA-  
AATATTTTTGTGGGACGAGAAGACCCTATAGATCTTTATATTTTT-ATAT-  
TTTATATTTTTTTT---GTTA-----  
TAAAAAATATAATTTATTATGTATTTTCATTGGGGTGATGCAGGAATTC-  
AATTAACCTTCTG--TTATTTTATTTCAATTAATT---TATGT--TTTTTTGATCCATT----  
ATTATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTTCGGAGAGTTCATATCGATGAAGGAGTTTGGCAGCTCGATGTTGGAT  
TAAAA-TTAGTT-TTGGGCGTAGAAGTTC-AATTA-CTGGGTCTGTTCCGACCTTTAATTT-  
TTTACATGATCTGAGTTCAAACCGGAGGCAGCAGGCACGCAAATTACCCACTCCCGG  
CACGGAGAGGTAGTGACAAAAAATAACGATACGGGACTCTATTGAGGCCCGTAAT  
CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAATCC-----  
-----CCGGTGC---TACGTAGGCT-TTTATTAGCTGAAG-T-ATGTG-CCG-----TG-G-  
TGATTTT--ATTTACCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACATTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACCACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTGACAGATTGATAGCT  
CTTTCTTGATTGAGTGGGTAGTGGTGCATGGCCGTTTC?CGTGTGCTTGATAGTGCAG  
CCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCACGAGTCCGATAGCG  
ACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTAC  
GTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTACCAC---TTGTTCAACTTTAAAGTGC----  
TTTTGAGATGGTAA-TTC-GCCCTATCCAG----TGTAT--ACTGTG--ATAG-  
AGGGTTTTATTGCTCGCATTTT-ATTT--  
AAGTAAGGTACAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAGT--AT-G----GTTAA---  
AGTGCCTATTT-A-AATGTT--ATAAATT-----TTA-TTTA-----  
TGACATTTTTATAGC-TTTAA--TCA--AT---TCT---ACTGGTTTTT-----  
TTTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTATTACAT-----TTTATA-  
---ATGT--TAAAAGTGTC-----TAAATTGCC-GTTTACAA-GTGGGTTA-----  
GGTATTAGTTAA-ATGTCGAACTT-AT-----GTTTCGTCTAA-CTA-  
AGCCGCCTAGCTGTTTGTA---AAC-TG--TG-CCTTA--GACAG--TGCTTA--

CAACACCGGTCAGCAGCGATCCAGGTTCAAGGCTCTTATCCGGTCCGT????????????  
????CTTGAAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTGAT-  
ACAAAACCAAAGGCGTAATGAAAGTAAAGAGTGAGTAGAGAAAGATTTTAAAA---  
---GGATAAGAATTT-----TTATTTTAACTTTGATT-----TCTTTATCT-  
CTCTCTCAAGGGAAGATGG--AAAT-----CTTATTTTAGATTAATATTTAGTA-----  
-----TATTTT---TT-ATATTAAG-TA-----AA-  
TTTAAATGTAGATTTCTCGCATTCCCAGGGCGCTTG-TTCTCGCTT--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACACTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTTGGGATTCTGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGCGCTGTGG??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

*Ectrichodia* sp                                 ????????????????????A AAAACATGTCCTTTAGT-----

TTTAAATTTA-AGGTCTAACCTGCCAATGA--  
TTATTTATTAATGGCCGAGTATTTAACTGTGCTAAGGTAGCATAATCAGTTGTCT  
TTTAAATGAGGGCTGGAATGAAAGGTTGGACGAGGAATTTACTTTCTTT-TAAATAA--  
TTAATGAATTTGACTTTTAAAGTTAAAGGCTTA-  
AATTATTTGTGGGACGAGAAGACCCTATAGATCTTTATATTTTT-  
ATTTAATTATGTTTTTTTT---GTTA-----  
TTAAAAATTTAATTTATTAAGTATTTTCATTGGGGTGATGTAGGAATTT-  
ATTTAACTTCTTG--TTTTTAAATTTCATTTATT---TATGT--TTTTTTGATCCATT-----  
TTATGATTA AAAAGATTAAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTTCGGAGAGTTCATATCGATGAAGGAGTTTGCACCTCGATGTTGGAT  
TAAA-TTAGTT-TTGGGCGTAGTAGTTC-ATTTT-CTAGGTCTGTTTCGACCTTTAATTT-  
TTTACATGATCTGAGTTC????????????????????????????????????  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

???  
 ???  
 ???  
 ???  
 ???AGTCCGATAGC  
 GAACAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTA  
 CGTCAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
 TATCCGAAAGAGGAGATTCACGTCTTACCAC---TTGCTCATATTTAAAATGC----  
 TTTTGAGATGGTAA-TTC-GCCCTATCCAG---TGTAT--ACTGTG--ATAG-  
 AGGGTTTTATTGCTCGCATTTT-ATTT--  
 AAATAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
 AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAGT--AT-G----GTTAA--  
 AGTGCCTATTT-A-AATGTT--ATAAATT-----CTA-TTTA-----  
 TTACCTTTTATAGC-TTTAA--TCA--AT---TCT---ACTGGGTGTTTA-AA--  
 ATTTTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTAAGCATT-----  
 TTTATA-----ATG---TAAAAGTGTCTA-----AATTAATTGCC-GTTTACAA-  
 GTGGGTTA-----GGTATTAGTTAA-ATGTGCAACTT-TT-----GTTGCTCTAA-CTA-  
 AGCCGCCTA-CTGTTTGTGTA---AAC-TG--TG-CCTTA--GACAG--TGCTTA--  
 CAACACCGGTCAGCAGCGA??  
 ???AAGTAAAGAGTGAG  
 TAGAGAAAGA---TAAA-----GGATAAAAATTT-----TTATTTTAACTTTGATT-----  
 TCTTCTCT-CTCTCTCAAGGGAAGATGG--AAAT-----CTTATTTTAGATAA-  
 TATTTAG-----TATTT-TT-ATATTAAG-TA-----GA-  
 TTTAAAATGTAGATTTCTCGCATTCCCAGGGCGCTTG-TTCTCGCTT--TG-  
 CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
 CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
 GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
 AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
 GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
 AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAC-  
 CCCCAGGCTT??  
 ???TTGGACATCCAGAAGTATATATTCTTATTTTA  
 CCAGGATTTGGTTAAATTTCCCATATTTGCAATAGAGACAGGAAAAAATGAAGCA  
 TTTGGATCTTTAGGAATAATTTATGCTATATTAACCTATTGGACTTTTGGATTCATTGT  
 ATGAGCCCATCATATATTTACAGTAGGTATAGATGTTGATACACGAGCTTACTTCAC  
 ATCCGCTACCATAATTATTGCTGTACTACAGGTATTAATAATTTTCAGATGATTAGCA  
 ACATTACATGGAAGAATTTAAGATTTTCACCTAGATTATTATGAGCCTTAGGTTTTG  
 TATTTCTTTTACAATTGGTGGACTAACTGGAGTAATTTTAGCAAACCTCAAGAATTGA  
 TATTATTTTACATGATACTTATTATGTTGTTGCACATTTTCATTACGTTTTATCAATAG  
 G????????????????TGGATCATGTACAGTAAAACATGTTGGATGCGTTTTATCACCATT  
 TCGTGTTATTGGTGATAATTTAAAAGATCGTTTTGATGGCGCATCACGTGTTATGGTA  
 AGCAATTCTGGATCAATACGCGGCA---  
 CTGGTGGACGAAAGAGTCGTTATAATTTCCAATTGAAACCCTATAATCCAGATCATA  
 AACCACCAGGCGTTAAAGATCTAGTATATTTAGAACCGTCGCCATTATTCTGTGAAC  
 GTAATCCAAGACTAGGTATACAGGGTACGCATGGACGACAGTGTAACGCAACATCG  
 CAAGGCGTTGAAGGTTGTGATCTAATGTGTTGTGG

Ectrychotes atripennis CGCTGTTTATCAAAAACATGTCCTTATGG--TTA---  
 TTATTTATTTA-AGGTTAACCTGCCCAATGA----  
 TTAATTTAATGGCCGCAGTATTTAACTGTGCTAAGGTAGCATAATCAATTGTCTTT

TAATTGAGGGCTTGAATGAATGGTTGGACAAAATATCTACTTTCTTT-TAATTA--  
ATAATGAATTTAATTTTTAGGTGAAAAAGCTTA-  
AATTTATTTAAGGGACGAGAAGACCCTATAGAGCTTTATATTTAT-ATTT--  
ATTTATTTAATTT---GTTA-----  
TTTTTGAATAGATTTTTTGTGTATTATATTGGGGTGATAGTAAAATAT-  
TATTAACTTTTAT--TTTTTATTTACATTAATT---TATGT--TATTTTGATCCATT---  
ATTAATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTATTCGGAGAGTTCATATTGATGAATAAGTTTTCGACCTCGATGTTGGAT  
TAAA-TTAGGT-TTAAGGCGTAGAAGTT-ATTAT-  
CTAGGTCTGTTCGACCTTTAAAAT-  
TTTACATGATCTGAGTTCAAACCGGAGGCAGCAGGCACGCAAATTACCCACTCCCGG  
CACGGAGAGGTAGTGACAAAAATAACGATACGGGACTCTATTGAGGCCCGTAAT  
CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCCAGTCT--  
GGTGCCAGCAGCCGCGTAATTCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAGTCC-----  
-----CCGGTAC---TGCGTAGGCT-TTTATTAGCTGAAG-TTGTTTA-TCG-----TG-G-  
TGACTTC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACCACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTGACAGATTGATAGCT  
CTTTCTTGATTGAGTGGTGTGATGGCCGTTTC?CGTGTTGCTTGATAGTGCAG  
CCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCACGAGTCCGATAGCG  
ACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGAGTAC  
GTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TATTCGAAAGAGGAGATTCACGTCTTACCAC---TTGCTCGACTTTAAATTGC----  
TTTTGAGATGGTAA-TTC-GCCCTATTCAG----TATTT--ACTGTG--GTAG-  
AGGGTTTTACCGCTCGCATTTT-ATA----  
TGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAAT--AT-G----ATTAA---  
AGTGACTGTCT-A-GAGGTT--ATAGAAT-----TTATTTTA-----  
TAATCTTTTTACAGC-TTTAA--TTA--AT---TTT---ATTGAGTATTTA-----  
TTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTTA-----ATT---  
--TTAAAGTGTTT-----ATTTGTATTGCC-GTTTACAA-GTGGGTTTT-----  
GATATTGGTT-A-ATGTTGAACTT-TT-----GTTGCTCTAA-CTA-AGTCGCCTA-  
CTGTTTGTA----AAC-TG--TG-CCTTG--AACAG--TGCTTA--  
CAACACCGGTCAGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT?????????????  
?????CTTGAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTGAT-  
ACAAAACCTAAAGGCGAAATGAAAGTAAAGAGTAGTAAGATAGAGG-----  
AAGAAATT-----TATTTTTTCATCCTTGAT-----TCT-

TGCTCTCAAGGGAAGATGG--TTAT-----CTTATTTTAAATTA-----  
----TAT---T---TAATTAT-AT-----TA-TTTAAGATGTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-TTTCTCGCTT--TG-  
CGAGTTTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTTCTCTCAGGATAGCTTGCCTCGAT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGCGCTGTGG????????????????????????????????????CATTTTAATCCTTCCGGGATT  
TGGTTAATTTCCCATATTATTCAATAGAAACAGGAAAAAGAGAAGCATTGGTCC  
TTAGGTATAATTTATGCAATAATTACTATCGGACTTTTAGGATTTATTGTATGGGCA  
CACCATATATTCAGTGTAGGTATAGATGTTGATACACGAGCATATTTACCTCAGCTA  
CTATAATTATTGCAGTTCCTACAGGAATTAATAATTTTAGATGATTAGCAACTTTACA  
TGGAAGAATATTAACCTTCTCTCCTAGACTTTTATGAGCATTAGGCTTCGTATTCCTT  
TTACTATTGGAGGATTAACAGGAGTAATTTAGCAAATCAAGTATTGATATTGTACT  
ACACGATACTTACTATGTAGTAGCACATTT????????????????????????????????ATGTCT  
GGATCATGTACAGTAAAACATGTTGGATGCGTTTACCACCGTTTYGTGTGTCGTCGGA  
GATAATTTAAAAGATCGTTTCGACGGCGCATCACGCGTTATGGTAAGCAATTCTGGT  
TCATTACGCGGCA---  
ATGGCGGACGTAAGAGTCGTTATAATTTACAATTGAAACCATAACAATCCAGATCATA  
AGCCACCAGGCGTTAAAGATCTAGTATATTTAGAACCATCGCCATTATTCTGCGAAC  
GTAATCCAAAGCTTGGCATAACAGGGTACACATGGA????????????????????????????  
????????????????????????

*Ectrychotes serdangensis*

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
GCAGCAGGCAC  
GCAAATTACCCACTCCCGGCACGGAGAGGTAGTGACAAAAAATAACGATACGGGAC  
TCTATTGAGGCCCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCCAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAGTCC-----  
-----CCGGTAC---TGCGTAGGCT-TTTATTAGCTGAAG-TTGTTTA-TCG-----TG-G-  
TGACTTC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA



GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTGACAGATTGATAGCT  
CTTCTTGATTAGTGGGTAGTGGTGCATGGCCGTTTC?CGTGTGCTTGTAGTAGTGCAG  
CCCTAAGTAGGTGGTAACTCCATCTAAGGCTAAATACAACCACGAGTCCGATAGCG  
ACAAGTACCGTGAGGAAAGTTGAAAAGAAGTTGAAGAGAGAGATTCAAGAGTAC  
GTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TATTCGAAAGAGGAGATTCACGTCTTACCAC---TTGCTCGACTTTAAAGTGC----  
TCTTGAGATGGTAA-TTC-GCCCTATTCAG---TATTT--ACTGTG--GTAG-  
AGGGTTTTACCGCTCGCATT---ATA----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAAT--AT-G----ATTAA---  
AGTGACTGTCT-A-GAGGTT--ATAGAAT-----TTATTTTA-----  
TAATCTTTTTACAGC-TTTAA--TTA--AT---TTT---ATTGAGTATTTA-----  
TTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTTA-----ATT---  
--TTAAAGTGTTC-----ATTTGTATTGCC-GTTTACAA-GTGGGTTTT-----  
GATATTAGTT-A-ATGTTGAACTT-TT-----GTTCGTCTAACCTA-AGTCGCCTA-  
CTGTTTGTGTA---AAC-TG--TG-CCTTG--AACAG--TGCTTA--  
CAACACCGGTCAGCAGCGATCCAGGTTTCAAGCTCTTATCCGGTCCGT?????????????  
?????CTTGAAACACGGACCAAGGAGTTTATTGTGTACGCAAGTCATTGGGTGAT-  
ACAAAACCTAAAGGGCAAATGAAAGTAAAGAGTAGTAAGATAGAGG-----  
AAGAAATT-----TATTTTTCATCCTTGAT-----TCT-  
TGCTCTCAAGGGAAGATGG--TTAT-----CTTATTTTA-----  
-----AATTAT-AT-----TA-TTTAAGATGTAGA-  
TTCCCGCATTCCAGGGCGCTTG-TTTCTCGCTT--TG-  
CGAGTTTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGGCAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTTCTCTCAGGATAGCTTGCGCTCGAT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGCGCTGTGG??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

Ectrychotes\_sp ??????????AACATGTCCTTATGG--T-----

TATATATTTA-AGGTTTAACTGCCCAATGA---  
AAATTTTTTAAATGGCCGCAGTATTTAAGTGTGCTAAGGTAGCATAATCAATTGTCTT

TTAATTAAGGGCTTGAATGAATGGTTGGACAAAATATTAACCTTCTTT-TTATTAA--  
TTAATGAATTTAATTTTTAAGTGAAAAAGCTTA-  
AATATATTTAAGGGACGAGAAGACCCTATAGAGCTTTATATTTAT-ATTT--  
ATTTAATGAATTT---GTTA-----  
TGTATTTTATAATATTTTGTGTATTTTGTGGGGTGACAGTAAAATTT-  
TATTAACCTTTTAT--TTT-TTATTATCATTTATT---TATGT--TACTTTGATCCATT---  
ATTAATGATTAGAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTATTCGAAGAGTTCATATTGATGAATAAGTTTGCACCTCGATGTTGGAT  
TAAA-TTAGAT-TTAGGCGCAGAAGTTT-ATTAT-CTAGGTCTGTTCGACCTTTAATTT-  
TTTACATGATCTGA??  
??  
??  
CCTTAAACAAGGATCCA-TTGGAGGGCCAGTCT--  
GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCCGTTCCGCCCTGTCCGGTGT  
ACTGGCATGTCCGCGCATGTCTGTCCGGTGGTGAATAAAGTCC-----  
-----CCGGTAC---TGCGTAGGCT-ATTATTAGCTGAAGTT-GTTA-TCG-----TG-G-  
TGACTTT--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGG??  
??  
??  
??  
AAACTCCATCTAAGGCTAAATACAACCACGAGTCCGATAGCGAA  
CAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACGT  
GAAACCGCTCAGGGGTAAACGGAGAAGAACTGAA--  
TATTCGAAAGAGGAGATTCACGTCTTACCAT---TTGCTCGACTTTGAAGTGC---  
TTTTGAGATGGTAA-TTC-GCCCTATTCAG---TATTT--ACTGTG--GTAG-  
AGGGTTTTACCGCTCGCATTTT-ATA---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAAT--AT-G----ATTAA---  
AGTGACTGTCT-A-GAGGTT--ATAGAAT-----TTATTTTA-----  
TAATCTTTTTACAGC-TTTAA--TTA--AT---TTT--ATTGAGTATTTA-----  
TTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTTA-----ATT---  
--TTAAAGTGTTT-----ATTTGTATTGCC-GTTTACAA-GTGGGTTTA-----  
GATATTAGTT-A-ATGTTGAACTT-TT-----GTTCGTCTAA-CTA-AGTCGCCTA-  
CTGTTTGTAA---AAC-TG--TG-CCTTG--AACAG--TGCTTA--  
CAACACCGGTCAGCAGCGATCCAGGTTTCACTCTC????????????????????????  
??  
ACAAAACCTAAAGGCGAAATGAAAGTAAAGAGTAGTAAGATAGAGG-----  
AAGAAATT-----TATTTTTTCATCCTTGAT-----TCT-  
TACTCTCAAGGGAAGATGG--TTAT-----CTTATTTTAAATTA-----  
----TAT---T---TAATTAT-AT-----TA-TTAAAGATGTAGA-

TTCCCGCATTCCCAGGGCGCTTG-TTTCTCGCTT--TG-  
 CGAGTTTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
 CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
 GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
 AGCTGGTTCCCTCTGAAGTTTCTCTCAGGATAGCTTGCCTCGAT-----T---  
 GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
 AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
 CCCC GGCTTTCTTAAAA-----  
 TTGAAGCCGTGGGATTCGGATCAGAGTGCCAA-----  
 G????????????????????????????????????AAGTGTATATTTAATTCTTCCCGGATTTGGATTAA  
 TTTCTCATATTATTTCTATAGAAACAGGAAAGAGAGAAGCTTTTGGACCTTTAGGTAT  
 AATTTATGCAATAATTACTATTGGTCTTTTAGGGTTTATTGTATGAGCACATCATATA  
 TTTACTGTAGGTATAGATGTTGATACACGGGCTTATTTACCTCTGCTACTATAATTA  
 TTGCAGTACCTACTGGTATTAAGATTTTTAGATGATTAGCTACATTACATGGAAGAAT  
 TTTAACCTTTTCTCCTAGACTTTTATGGGCCCTAGGTTTTGTGTTTCTTTTCACTATTG  
 GAGGATTAACCTGGAGTAATTTAGCTAATCAAGAATTGATATTGTTTTACACGATAC  
 TACTATGTAGTAGCACATTTTCA????????????????????????????????????ATGTCTGGATCATG  
 TACAGTAAAACATGTTGGATGCGTTTACCACCGTTTCGTGTCGTTGGCGATAATTTA  
 AAAGATCGTTTCGACGGCGCATCACGCGTTATGGTAAGCAATTCTGGTTCATTACGC  
 GGCA---  
 ATGGAGGACGTAAGAGTCGTTATAATTTACAATTGAAACCATAACAATCCAGATCATA  
 AACCACCAGGTGTTAAAGATCTAGTATATTTAGAGCCATCGCCATTATTCTGCGAAC  
 GTAATCCAAAGCTTGGTATACAGGGTACACATGGACGTCAATGTAATGCTACATCGC  
 AGGGTGTGTAAGGTTGTGATTTAATGTGTTGTGG  
                   Emesaya\_incisa                  CGCCTGTTTATCAAAAACATGTCCTTTTGA--TT----  
 TTTAAAATTTT-AGGTCTAACCTGCTCTATGA----  
 TTTATTAATGGCCGCGGTATTCTAACTGTGCTAAGGTAGCATAATTATTTGTCTTTT  
 AATTGAAGGCTGGAATGAATGGTTGGACAAATAACTTTCTTTCTTT-ATTTTAA-  
 TTTCTTGAATTTTATTTTTTAGTCAAAAAGCTAA-  
 AATTTTTTTGTAGGACGAGAAGACCCTATAGATCTTTATATAAGC-TTTA--  
 ATAATAATATTTT---GGTAAGTG----  
 TAATTTTTATTATTATGGTTTGTATTTTGTGGGGTGATGTCAGAATCT-  
 AACTAACTTCTGT--TAC-ATTATTACATTAATT---TATGT--TTATATGATCCATT---  
 TTTAATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
 GCGTAATTTTCTTGGAGAGTTCTTATCGATAAGAAAGTTTGCACCTCGATGTTGGAT  
 TAAAA-TTAGTA-TTTGGTGCAGTCAT---AAAAA-CTAGGTCTGTTCGACCTTTAAAAT-  
 TTTACATGATCTGAGTTCAAACCGGAGGCAGCAGGCACGCAAATTACCCACTCCCGG  
 CACGGGGAGGTAGTGACAAAAAATAACGATACGGGACTCTATTGAGGCCCGTAAT  
 CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
 GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
 TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCCGCCCTGTTCGGTGA  
 ACTGGCATGTCGCGGCATGTCCTGTTCGGTGGTGAACGGGATCC-----  
 -----CCGGTGC---TAACTGAGAC-TTC--TTGTCAAAG-TAATGTG-CCG-----TGTG-  
 TGATGTC--GTTTGCCGA---  
 TCTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
 AACAAATTAGAGTGCTTAAAGCAGGCT-  
 AAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAAATAGGACCTCGGTTCTATT  
 TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA

TTGCGACG-  
TAGAGGTGAAATTCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCATTGCTTGGCCAAG  
TATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCAGATACCGCC  
CTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCTATAATGACT  
CGGCGGGGAGCTTCT-  
ACTCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAAGGCCACCACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTGACAGATTGATAGCT  
CTTTCTTGATTAGTGGGTAGTGGTGCATGGCCGTT?CGTGTTGCTTGATAGTGCAG  
CCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATACAACCACAAGACCGATAGCG  
ACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTGAAAGAGAGAGTTCAAGAGTAC  
GTGAAACCGTTACAGGGTAAACGGAAAAGATCTGAA--  
ATCCGAAAGGGGAGATTCACGTCTTACCAT---TTACTTATCTTTTTTATGTT---  
TTAAGAGATGGCAA-GTC-GCCCTATCCAG----TGCAA--ACTGTG--GTAG-  
AGGGTCTGGTTCGCTCGCATATT----A--  
AAGTTTTGGTAGTGGTGGACCGCACTTCTCCCTCAGT-  
AGGACGTTGCGACCTGTCATTT--TAC----ATTCTAAGT--GC-GTTTGTGTAA---  
AGTGCCTGTAT-T-TAGATA--ATGTCGCAA-----  
GATATTATCTTTTACAGC-TTTAA--CAA--TA----TGC---ACCGAGTGTTAA-----  
TTTTTATGACAGTAACAAATGATGG-TGTTGAGCCACTTAAAATTGTTGTCAT----  
TATTGATG----ATAA---TGGAGAGTGTG-----TTAGGCCGGTTTGCAA-  
GTGGGCTG-----TTTTATAGTT-A-TATTCGGACTT-TT-----GTCCGTCTTG-CTA-  
TTGCTCTCA-CTGTTGGTA---GAC-TG--TG-CCTCT--GACAG--TGCTTA--  
CAACATCAGTCAGCAGCGATCAAGGTTTCAAGTTTATCCTCGGTCCGT?????????????  
?????CTTGAAACACGGACCAAGGAGTTTATCGTGTACGCGAGTCATTGGGTAAA--  
GAAAACCCAAAGGCGAAATGAAAGTAAATAGTGGGAAATTGATT-----  
-----TATTCTATATTATCAG-----TTT-CCCCTATAAGGGAAAGATGG--AAAA-  
-----CTATATCAGTAATG-----GTGTGATAATATCA-TA-----TT-----  
-----GT-TGTTGATACAAGT-TTCCCGCACTCCCGGGGCGCTTG-TTTCTCGCTT--TG-  
CGAGTCACTGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGC ACTCGGT-----T---  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAACCCCCGGCTTTCTTAA  
AA-----  
ATGAAGCCCGTGGGATTCGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAAC  
TGGCGCTGTGG???  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

Glymmatophora\_crassipes ?????????????AACATGTCCTTTAGT-----  
TTTAAATTTA-AGGTTTGACCTGCCCAATGA--  
TGAATAATTTAATGGCCGCAGTATTTTAAGGTAGCATAATCAGTTGTCT  
TTTAATTGAGGGCTGGAATGAAGGGTTCGGACGAGGAATAGTCTTTCTTT-TTAGTAA--  
TTAATAAATTTGACTTTTTAGTTAAAAGGCTAA-  
AATTTTTTTGTGGGACGAGAAGACCTATAGATCTTTATATTTAT-ATAT-  
AATTTAAGTTTTTA---GTACA-----  
TGAATAATTGAATTTATTAGGTATTTTGTGGGGTGTATACGAGAATTT-  
TTTTAACTTCTTG--TTATTTTAATTCATTTATT---TATGT--TTTTATGATCCATT---  
TTTATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTTCGGAGAGTTCATATCGATGAAGAAGTTTGCACCTCGATGTTGGAT  
TAAA-AAAGTT-TTGGGTGTAGTAGTTC-AATTA-CTAGGTCTGTTTCGACCTTTAATTT-  
TTTACATGATCT??  
??  
??  
CCTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAATTC-----  
-----CCGGTGC---TACGTAGGCT-TTTATTAGCTGAAG-T-ATGTG-CCG-----TG-G-  
TGATTTT--ATTTGCCGA--  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCCGCCAGCGATCCGCCGATGTTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
??  
??  
??  
??  
GCGAACAAGTACCGTGAGGGAAAGT  
TGAAAAGAAGTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAC  
GGAGAAGATCTGAA--TATCCGAAAGAGGAGATTCACGTCTTACCAC---  
TTGCTCGACTTTAAAGTGC---TTTTGAGATGGTAA-TTC-GCCCTATCCAG---TGTAT--  
ACTGTG--GTAG-AGGGTTTTATTGCTCGCATT--ATT--  
AAGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAGT--AT-G----GTTAA---  
AGTGTCTATCT-A-AAGGT--ATAAATTTA-----TCTAAGTTTA-----  
TTACCTTTTTATAGC-TTTAA--TCA--AT---TAT-ACACTGGGTGTTTG-----  
TTTTTCTGACAGTAACGATTGATGG-TGTAGAGCCACTTGAACATTATGATAA--  
TTTTATTATA-----ATG----TAAAAGTGTG-----CATGTGTTGCC-GTTTACAA-  
GTGGGTTTA-----GGTATTAGTT-A-ATGTCGAACTT-TT-----GTTTCGTCTAA-CTA-  
AGCCGCCTA-CTGTTTGTAA---AAC-TG--TG-----  
-----  
C??GTAATGAAAGTAA  
AGAGAGTGTAGAGGAGGGGAAAGAAA-----AGGATAAAAGTTTT-----  
TATACTTTAACTTTGATTTCTCTCTCTCTCTCTCTCTCTCAAGGGAAGATGG--

AAAT-----CTTATTTTAAGTGA-TGT-----TA-----  
 ----AA-TTTAAAATGTTGA-TTCCCGCATTCCCAGGGCGCTTG-TTTCTCGCTT--TG-  
 CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
 CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
 GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
 AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGCTCGGT-----T---  
 GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
 AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
 CCCC GGCTTTCTTAAAA-----

TTGAAGCCGTGGGATTCTGATCAGAGTGCCAAGTGGGCCAATTTTGGTAA?????????  
 ???  
 AATTTACATATTATCGCAATAGAAACAGGAAAAACGAGGCATTTGGATCACTAGG  
 AATAATTTACGCTATACTAATTTGGTCTCTTAGGGTTTATTGTATGAGCACACCAT  
 ATATTTACAGTAGGAATAGATGTAGATACTCGAGCTTACTTTACATCAGCCACTATA  
 ATTATTGCTGTACTACTGGAATTAATAATCTTTAGATGATTAGCCACTTTACATGGAA  
 GAATTTAATATTACACCTAGACTATTATGAGCATTAGGATTTGTATTTTTATTTAC  
 AATTGGAGGATTA ACTGGAATTATTCTTGCTAATTCAAGAATTGATATTATTCTTCAT  
 GACTTACTATGTAGTAGCCACTTCCA?????????????????????????????????  
 ATGTCTGGA  
 TCATGTACAGTAAAAACATGTTGGATGCGTTTATCACCATTTCTGTAATTGGTGATA  
 ATTTAAAAGATCGTTTTGATGGGGCAGCAGTGTATGGTAAGCAATTCTGGATCTAT  
 ACGCGGTA---

CTGGTGGACGTA AAAAGTCGTTATAATTTCCAATTA AAAACCATATAATCCAGATCATA  
 AACCACCAGGCGTTAAAGATCTAGTATAATTTAGAACCATCACCATTATTCTGTGAAC  
 GTAATCCAAGGCTAGGTATACAGGGTACGCATGGACGACAGTGTAACGCAACAT????  
 ???

Haematoloecha\_sp          ???

GTATTTATATA-AGGTCTAGCCTGCCCAATGA---  
 TATTTATTTAATGGCCGAGTATATTA ACTGTGCTAAGGTAGCATAATCAATTGTCTT  
 TTAATTGAAGGCTGGTATGAAGGGTTGGATGAAATATTGACTTTTTTC-TTATTAA--  
 TTCTTGAATTTAATTTATCGATTAAAAAGTTGA-  
 TATAAAATTAAGGGACGAGAAGACCCTATAGATCTTTATAATTA-  
 ATTTTATTATTTATTTT---GTTTATTT-----  
 TTTAAAGATAATGATTTTATTTATTTAATTGGGGTGATTTTAGAATTT-  
 TAATAACTTCTAT--TAT--AATATTCATTAATC---AATGT--TTAATTGATCCTAT---ATT-  
 TAGATTA AAAAATTAAGTTACCTTAGGGATAACA-  
 GCGTAATTTTTTCAGAGAGTT CATATTAATGAAAAAGTTTGCACCTCGATGTTGGAT  
 TAAA-TAAGTT-TTGGGCGCAAAGTTC-ATTAA-  
 CTAGGTCTGTTCGACCTTTAAAAT-

TTTACATGATCTG??  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???

??  
??  
????????????????GAGAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATAT  
AACTACGAGTCCGATAGCGAACCAAGTACCGTGAGGGGAAAGTTGAAAAGAAGCTTGA  
AGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TTTTCGAAAGGGGAGATTACGTCTTACCATA--TTGTTTAACTTATAAATGTA---  
TTAAGAGATGTTTA-TTC-GCCCTATTCAG----TGCAA--ACTGTA--ATAG-  
AGGGTTTATTCGCTCGCATTATTTT----  
TAGTAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTGTCAGAT--AAT----ACTCTAAGT--AT-G----ATTAA---  
AGTGTCTGTAT-T-AATGTA--TTATAAA-----TTTA-----  
TTTTGTATTTTACAGC-TTTAA--TCA--AT---TATAATACCGAGTATTT-----  
TATACTGATAGTAACAATTGATAG-TGTTGAGCCACTTTAA-----A--  
-ACAATAAAGTG-----TCTAGCC-GTTTACAA-GTAGGTTTTTAGG---  
CTTTTTAGCT-A-ATATCGGGCTT-TT-----TGGTCCGTATAG-CTA-GGCCACTTA-  
CTGTTGGTA---GGCGTG--CT-CCTTA--GACAGATTGCTTA--  
TAACACTGATCAGCAGCGATCCAGGTTCAGTCTCTTATCCGACCCGTCTTGAA???????  
??TTATTGTGTACGCAAGTCATTGGGTAAT--  
TAAACCCAAAGGCGTAATAAAAGTAAAGAGCAATTTAATTGAAT-----  
AGCAATAT-----TTT-----AATTA-AATTCTTAAAGGAAGATAG--  
ATAT-----TATTTTCTTTAAA-----  
GTT-TTCTTTTAAATGAA-ATCTTGCATTCCTAGGGCGCTTG-TCTCTCGCAT--TG-  
CGAGTATATGCGCACCTAAAGCGTACACGATGATACCCGAAAGATGGTGAAGCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCATT---TTT---T---  
TAGTTGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCTCGGCTTTCTTAAA-----  
TGAAGCCGTAGGATTTGGATCAGAGTGCAAAGTGGGCCAATTTTGGTAAGC?????????  
????????????????????GATTTTTTGGACATCCCGAAGTTTATATTTTAAATTTTACCAGGAT  
TTGGAATTATTTCTCACGTAATTGCTATAGAAACAGGAAAAAATGAAGCCTTCGGTA  
CTTTAGGGATAATTTATGCTATACTTGTATTGGACTTTTAGGATTTATTGTATGAGC  
ACATCACATATTTACAGTAGGAATAGATGTAGATACACGAGCCTACTTTACTTCAGC  
AACTATAATCATTGCAGTCCAACAGGAATTAATTTTTCAGTTGATTAGCAACACTT  
CATGGAAGAAAAATTAATATACTCCCAGATTACTGTGAGCTCTCGGATTCGTATTC  
TTATTTACCATTGGGGGATTAACGGGAGTTATTTTAGCCAACTCAAGTATTGATATTA  
TTTTACATGATACTTATTATGTAGTAGCACACTTCCATTATGTATTATCAATA?????????  
????????TGGATCATGTACAGTGAAAACCTGTTGGATGAGGTTACCACCATTTTCGTAT  
TGTTGGTGATAATTTAAAAGATCGTTTTTGACGGTGCATCAAGAGTAATGGTTAGCAA  
TGCAGGTTCAATACGTGCAA---  
CTGGTGGGCGTAGAAATCGTTATAACTTTCAATTA AAA ACCATA CAAT CCTGATCATA  
AACCGCCAGGTGTTAAAGATCTTGTTTATCTTGAACCGTCGCCAGTATTTGTGAGCG  
TAATCCAAGATTAGGTATACAGGGTACGCATGGACGCCAGTGTAATGCAACATCGCA  
GGGTGTTGAAGGTTGTGACTTAATGTGTTGTGG  
Katanga\_sp ?????????????AAAAACATGTCTTTAAGT-----  
TTTTATTTA-AAGTTTAACTGCCCTATGA--  
TTAACTATTAATGGCCGCAGTACTTTAACTGTGCTAAGGTAGCATAATAATTTGTCT









CGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGTTA  
AAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCCTGTCGGTGTAC  
TGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAATCC-----  
---CCGGTGC---TACGTAGGCT-TTTATTAGCTGAAG-T-TTGTG-CCG-----TG-G-  
TGACTION--ATTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAACGATGCCAGCCAGCGATCCGCCGATGTTCTT  
TTAATGACTCGGC??  
??  
??  
??  
????????????????????CTAAGGCTAAATACAACCACGAGTCCGATAGCGAACAAGTACCG  
TGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCT  
CAGGGGTAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCAG---TTGCTCAACTTTGAAGTGC---  
TTTTGAGATGGTAA-TTC-GCCCTATCCAG---TGCAT--ACTGTG--GTAG-  
AGGGTTTTATCGCTCGCATT--ATT---  
AGTATGTTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAAAT--AAT----ACTCTAAGT--AT-A----GTAA---  
AGTGCCTGTCT-A-AAGGTT--ATAAAAT-----TTATTTA-----  
TAACCATTTTACAGC-TTTAA--TTA--AT---TTT---ACTGGGTATTTA-----  
TTTTTTGACAGTAACGACTGATGG-TGTAGAGCCACTTAAATATT-----ATTTATA--  
---ATA----TAAAAGTGTGTA---ATTGTTTTTTGTTGCC-GTTTACAA-GTGGGTTTTTA---  
AGGTTTTAGTT-A-ATGTCGAACCT-TT-----GTTTGTCTAA-CTA-AGCCGCCTA-  
CTGTTTTGTA----AAC-TG--TG-CCTTA--GACAG--TGCTTA--  
CAACACCGGTCGGCAGCGATCCAGGTTCCAGGC????????????????????????  
????????????????????????TACGCAAGTCATTGGGTGAT-  
ATAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGAAGGATAAAAG-----  
AAGGAAAT-----TAATTTTCATCTTTGAT-----TCT-  
TTCTCTCAAGGGAAGATGG--ATAT-----CTTTATTTTAAAT-----  
-----TA-----TA-TTTAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-GTTCTCGCTA--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTTCCCTCAGGATAGCTTGCGCTCGAT-----T---  
TTAAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTAGGATCAGAGTGCCAAGT????????????????????????  
????????????????????????????????TTTTACATTTAATTTTACCCGGATTTCGGAATTTT  
CACATATCATTGCCAGAGAAACGGGAAAAACGAAGCGTTTGGGGCTATTGGAATA  
ATTTACGCAATAATACTATTGGATTATTAGGATTTATTGTATGAGCCCATCATATAT  
TCACTGTAGGTATAGATGTTGATACACGAGCATACTTTACTTCTGCAACTATAATTAT

TGCCGTTCCCACAGGAATCAAATCTTTAGTTGACTAGCAACATTACACGGAAGAGT  
CATAGAATTCTCCCCTAACATACTCTGAGCCCTAGGATTTGTATTTTTATTCACTATC  
GGAGGCCTTACTGGAGTAATACTCGCTAATTCTAGAATTGATATTGTACTTCACGAC  
ACATACTATGTAGTAGCCATTTC????????????????????????????????TCATGTA  
CAGTAAAACATGCTGGATGCGTTTACCACCATTTCGTATTGTAGGTGATAATTTAA  
AAGATCGTTTTGATGGTGCATCACGTGTTATGGTAAGCAATGTTGGATCATTAAAGAG  
GTA---  
CTGGTGGACGTAGAAAGTCGTTATAATTTCCAATTA AAAACCATATAATCCAGATCATA  
AACCACCAGGTGTTAAAGATCTAGTATATTTAGAACCATCACCACCATTTTGTGAAC  
GTAATCCAAGACTTGGTATACAGGGTACGCACGGACGTCAATGTAACGCAACATCAC  
AGGGTGTGAAAGTGTGATTTAATGTGTTGTGG  
Mendis\_apicimaculata ??????????AAAACATGTCCTTTAGTAATT-----  
GTTGTATTTA-AGGTTTAACTGCTCAATGA---  
TTTATTTAAATAGCCGCAGTATTTTAACTGTGCTAAGGTAGCATAATCACTTGTCTTT  
TAATTGTTGGCTTGTATGAATGGTTGGACGAGGAACCTTACTTTCTTT-  
TTTTTAGTTTGTGTTGAATTTAACTTTTAAATTA AAAAGGTTTA-  
AATTTTATTAAGGGACGAGAAGACCCTATAGATTTTTTATAATTA A-GTTA--  
TTTGTTATTTTTT---GTTA-----  
TGTTTTTTATAATTAATTTAATTATTTGGTTGGGGTGACTAAAGAATTT-  
CAAGAACTTCTTT--TTT--TAATTTCAATTTATT---TATGT--TTTTATGATCCATA--  
TTTTGTTGATTATAAGATTAAATTACCTTAGGGATAACA-  
GCGTAATTTTTTCGGAGAGTTCTTATTGATGATAAAGTTTGCACCTCGATGTTGGAT  
TAAA-TAAGTT-TTAGGTGTAGTTGTTT-AATGA-CTAGGTCTGTTTCGACCTTTAAATT-  
TTACATGATCTGA??  
??  
??AT-CCTTTAACAAGGATCCA-  
TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCCCCGCCTGTGCGGTGT  
ACTGGCATGTCGCGGCATGTCTGTGCGGTGGTAAATAAAAATTC-----  
-----TCCGCGGTA CTTACCTAGGCT-TTTATTAGCTGAAG-T-TTGTG-TTG-----TG-  
G-TGACTTT--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCAAGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAAGATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGA?????  
??  
??  
????????????????CCATCTAAGGCTAAATACAACCACGAGTCCGATAGCGAACAAGTACCGT  
GAGGGAAAGTTGAAAAGA ACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTC  
AGGGGTAAACGGAGAAGATCTGGAATATCTCGAAAGAGGAGATTACGTCTTACCA  
T---TTGCTCTACGTTGAAGTAC---CTTTGAGATGGTAA-TTC-GCCCTATCCTG---  
TGCAT--ACTGTG--ATAG-TGGGTTTCACCGCTCGTATTTT-ATT---  
AGTAAGCTGCAGTGGTGGACCGCACTTCTCCTTTAGT-

AGGACGTTGCGACCTGTCAGAT--AAT----ACCCTAAGT--AT-A----ATTAA---  
AGTGCCTGTGT-T-AAATAT-AATAAAAA-----TTGTTTTA-----  
TTCAATTTATAACAGC-TTTAA--TAA--AT----TTT---ACTGGGTATTTA-----  
TTTTCTGACAGTATCGACTGATGG-TGTAGAGCCACTTGTTTTTAA-----  
CATTAAATA-----ATGT--TAAAAAGTGTT-----TTTCAGCC-ATTTACAA-  
GTTTGGTGA-----TTATTTAGTT-A-ATGTCAAACCTT-TT-----GTTCGTCTAA-CTA-  
GATCACCTA-TTGTTTGTA---AAT-TG--TG-TCTTG--AACAG--TGCTTA--  
TATCACCGGTCGGCAGCGATTTCAGGTTTCAGA????????????????????????  
????????????????????GTGTACGCGAGTCATTGGGTAAA--  
TAAAACCTAAAGGCGCAATGAAAGTAAAGGGTGGAAAAGATATAGG-----  
AAGGAAAG-----TAATTTTCATCCTTAAT-----  
TCTATACACCCGAGGGAAGATGG--TTTA-----CCTTGTTTTCAAAT-----  
-----TT-TTTTTAAA-----ATT-TTTAAACGTAGGT-  
TTCTCGCATTCCCAGGGCGCTTG-TTCTTGCTT--TG-  
CAAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT-----T---  
GATTGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTGC-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCTGGCTTTCTTAAAT-----  
TTTTTGAAGTCTTGGGATTAGGATCAGAGTGCCAAG????????????????????  
????????????????????AGTATATATTTAATCCTACCAGGATTCGGATTAA  
TCTCACATATTATTTCAATAGAAACAGGAAAAATAGAAGCATTTTGGATCAATAGGAA  
TAATTTATGCTATATTAACAATTGGACTATTAGGATTTATTGTTGAGCACACCATAT  
GTTACAGTAGGAATAGATGTTGACACTCGAGCTTATTTACTTCAGCAACAATAATT  
ATTGCTGTCCCTACAGGGATTAATAATTTTCAGATGATTAGCCACATTACATGGAAGA  
GTATTAATCTATCCCCTAGATTGATATGAGCCTTAGGATTCGTATTTTTATTACCA  
TTGGAGGATTAACCGGAGTAATTCTAGCTAATTCCAGTATTGATATTATTTACATGA  
TACTTATTATGTAGTGGCCATTTT????????????????????ATGTCTGGATCA  
TGCACAGTAAAACCTGTTGGATGCGTTTACCGCCATTTTCGTATTATTGGCGACCATT  
TAAAGGATCGTTTTGATGGCGCTTCTCGTGTAATGGTTAGCAATGCTGGCTCATTACG  
TGGCAGTGGTGGTGGACGTAAAAGTCGCTATAGTTTCCAATTGAAACCATATAATCC  
AGAGCATAAACCACCTGGTGTAAAGATCTTGTATATCTAGAGCCATCACCGTTATT  
CTGCGAACGTAATCCAAGACTTGGTATACAAGGTACGCATGGAAGACAATGTAATGC  
A????????????????????????????????

Microstemmatoides atrocyanea ??????????AAAAACATGTTCTTTAGT-----

TTTAAATTTA-AGGTTTGACCTGCTCTATGA--  
CTTTGGGTTAAATGGCCGCAGTATTCTAACTGTGCTAAGGTAGCATAATCATTGTCT  
TTTAAATTGGAGGATTGAATGAAAGGTTGGACGAGAAACAACTTTCTTT-TATATAG--  
TTTATGAATTTAATTTTTTAGTTAAAAAGCTAA-  
AATGTATTTAAGGGACGAGAAGACCCTATAGATCTTTATATTTGT-GAA---  
ATATTTTTTTTTG---GTTAATG-----  
AAAAGATTTTATTGAACCAAGTATTTTATTGGGGTGATTCAAAAATTA-  
ATTTAACTTTTTG--TTTATTATTTTCATTAATT---TATGT--TTTATTGATCCATT---  
GTTATGATTAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTTTGGAGAGTTCTTATCGATAAAAAAGTTTGGCAGCTCGATGTTGGAT  
TAAA-TAAGTT-TTTGGCGTAGCAGTTT-TAATA-CTAGGTCTGTTCGACCTTTAATTT-

TTTACATGATCTGAGTT??  
??  
??  
CCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAATCC-----  
-----CCGGTAC----TACGTAGGCT-TTTATTAGCTGAAG-T-ATGTG-CCG-----TG-G-  
TGATTC--ATTTACCGA--  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTC-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAACT  
TAAAGGAATTGACGGAAGG??  
??  
??  
??  
??  
??  
??  
??  
??  
CCACGAGTCCGATAGCGAACAAGTACC  
GTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGC  
TCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGAGGAGATTCACGTCTTGCCAC---ATGCTCGACTTTAGAGTGCT---  
TTTTGAGATGATAA-TTC-GCCCTATCCAG----TGTAT--ACTGTGTATAGG-  
AGTGTTTTATTGCTCGCACTTT-  
CTTTTAAAGTAGGATGCTGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAAT--AT-G----GTTTT---  
AGTGCCTGTGC-T-AAGGCT--ATTGGAAAAAAGTGTCTACATTTTTTTTTA-----  
ATATAACCTTTTTACAGC-TTTAA--CCA--AT---TCT-A-TTTGGGTATTT-----  
GTTTTCTTGACAGTAACGACTGATGG-TGTAGAGCCACTTTTAAGCATT-----  
TTCAAAGAA----ATG---TAAAAGTGTTTT-----TTTTTTATTTGCC-GTTTACAA-  
GTGGGCTTCA----AGGTATTAGTT-A-ATGTGCAACTT-TT-----GTTCTGCTAA-CTA-  
AGCCGCTA-CTGTTTGTGTA----AAC-TG--TGCACTTA--AACAG--TTCTTA--  
CAACACCGGTGCGCGGCATCCAG??  
??  
??  
ACCAAAGGCGTAATGAAAGTAAA  
GAGTAGAAAAGAGAGAAGATGAAAA-----GGTAAATTATAT-----  
ATTTTTTTGTCTTTGATATTC----TCTCTCTCT-TTCTCTCTAGGGAAGATGG--AAAT----  
-----CTTATTTTAAATATATAC-----AATATCTTG-TA-----TT-----  
---AT-ATTAATAATTTAGA-TTCCCGCATTCCAGGGCGCTTG-TTTCTCGCTT--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCCCGGCTTTCT??  
??  
??  
AAGTATATATTTAATTTTACCA  
GGATTTGGGTTAATCTCACATATTATTTCAATAGAAACTGGCAAAGAGAAGCCTTT

GGCTCCTTAGGAATAATTTATGCAATGTTAACTATTGGACTACTAGGATTCATCGTTT  
GAGCTCACCATATATTTACAGTAGGAATAGATGTTGATACACGAGCTTATTTTACTTC  
TGCAACAATAATTATTGCAGTACCTACAGGGATTA AAATCTTTAGATGATTAGCCAC  
CTTACATGGCAGACCTTTATCTTCTCCCTAGTCTTTTATGAAGATTAGGATTCGTAT  
TTTTATTTACAATTGGAGGATTA AACTGGAATTATTTTAGCTAATTCAAGAATTGATAT  
TATTCTTCATGATACTTATTATGTTGTTGCCCA????????????????????????  
??  
??  
??  
??  
??

Neoscadra\_sp                     ????????????AAAACATGTCCTCTTTG--T-----

TATTTATATG-AGGTCTGACCTGCCCGATGA-----  
TGTTTTCAATGGCCGCAGTATACTGACTGTGCTAAGGTAGCATAATCATTTGTCTTTT  
AATTGAAGGCTGGAATGAAGGGTTGGACGAGATATTAGCTTTCTTT-TTATTAC--  
ATTCTAAATTTAACTTTTAGGTTAAAAGGCCTA-  
TATTATATTAAGGGACGAGAAGACCCTATAGATCTTTATATTTAT-  
TATTATTTATCATCCTTTA---GTTAGATA-----  
ATTGATGGTATATTTAATATATATTTAATTGGGGTGATTACAGAATTA-  
CTATAACTTCTGT--TTT--AGCTTGCAATTGATT---TATGT--TTTAGTGATCCATT---  
ATTATTGATTGTAAGATTAAGTTACCTTAGGGATAACA-  
GCGTTATTCTTTCAGAAAGTTCTTATTAATGAAAGAGTTTGCGACCTCGATGTTGGAT  
TAAAA-TAAGTT-TTGGGCGCAAAGTCT-AATTA-  
CTTGGTCTGTTTCGACCATTA AATT-

TTTACATGATCTGA??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

ATATACTACGAGTCCGAT  
AGCGAACAAAGTACCGTGAGGGAAAGTTGAAAAGA AACTTTGAAGAGAGAGTTCAAGA  
GTACGTGAAACCGCTCAGGGGTA AACGGAGAAGATCTGAA--  
TTTTCGAAAGGGGAGATTACGCTTACCATA--TTGTTTAACTTATAAATGTA---  
TAAAGAGATGTTTA-TTC-GCCCTATTCAG----TGCAA--ACTGTA--ATAG-  
AGGGTTTATTCGCTCGCATTATTTT---  
TAGTAAAAACAATGGTGACC GCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTATCAATT--AAT-----ACTCTAAGT--AT-G-----TTTAA---  
AGTGACTGTAT-T-AATGTA--ATT-----TTTTTTTA-----  
AATTATATTTTACAGC-TTTAA--GCA--AT---TATAACTGAGTATTT-----T--  
AATATTGATAGTAACAATTGATAG-TGTTGAGCCACTTTAA-----  
TTATAAAAAGTG-----TCTAGCC-GTTTTACAA-TTAAGTAATAGGC--  
TTATTTTAGCT-A-ATTTCTGGGCTT-TA-----TGGTCCGTATAG-TTA-TGCCACTTA-

ATGTTTGTA----AACGTG--CT-CCTTA--GACAGATTGCTAA--  
TAACACTGATCAGCAGCGATCCAGGTTGAGTCTCTTATCCGACCCGTCTTGA????????  
??GTA  
AAGAGAATTAATTTAAGT-----AGAAATATTTTT-----  
AATTA-AATTCTTGAAGGAAGAAAG--ATTA-----  
ATTTATTTTATTAAAAA-----TTTT-TT-----TT-----  
TT-TTTTATTAATGAA-GTCTTGCATTCTAGGGCGCTTG-TCTCTCGCAT--TG-  
CGAGTATATGCGCACCTAAAGCGTACACGATGATACCCGAAAGATGGTGAACATATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTTCTCTCAGGATAGCTTGCCTCATT---TTTTATAAAT---  
TATTTGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTGC-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCTCGGCTT??  
??  
CGGGATTTGGGATCATTTCCTATGTTATCGCCATAGAAACAGGTA AAAATGAAGCCT  
TTGGGACACTAGGTATAATTTATGCAATACTAGCAATTGGATTATTAGGATTTATCGT  
ATGAGCCCATCATATATTCACAGTCGGAATAGACGTTGATACTCGAGCCTATTTTACC  
TCCGCTACCATAATTATTGCTGTTTCTACTGGTATTA AAAATCTTCAGATGACTAGCAA  
CCTTACATGGAAGTAAGATCACATTCACGCCAAGAATAATATGAGCATTAGGATTTG  
TATTTTATTACCATTGGAGGATTAACCGGAGTGATTCTAGCTAATTCAAGAATTGA  
CATTGTATTACATGATACTTATTATGTAGTCGCACA????????????????????????????  
??ATGTCTGGCTCATGTACAGTTAAAACCTGCTGGATGCGTTTACCACCGTTTCGTATT  
GTCGGTGATAATTTAAAAGATAGATTTGATGGTGCCTCACGTGTAATGGTTAGCAAT  
GCAGGCTCAATACGTGCAA---  
CAGGCGGACGTAAGAATCGTTATAATTTACAATTA AAAACCATTTAATCCAGATCATA  
AACCGCCGGGTGTA AAAAGATCTTGTATTCTTGAACCATCGCCAGTATTTTGTGAACG  
TAATCCAAGATTAGGTATAAAGGTACACACGGACGTCAGTGTA????????????????????  
????????????????????????????  
nr\_Neolibavus\_sp ?????????????????ACATGTCCTTTATG--T-----  
GTAAATTTA-AGGTTTGACCTGCCAATGA----  
TTAATTTAATGGCCGAGTATATTA ACTGTGCTAAGGTAGCATAATCATTTGTCTTTT  
AATTGTAGGCTAGTATGAAAGGTTGGACGAGAAATTGACTTTCTTT-TATATAA--  
TAATTGAATTTAATTTTAAAGTTAAAGAGCTTA-  
AATTTTTTATGGGACGAGAAGACCCTATAGATCTTTATAATTAT-ATTT--  
TTTTATTTATTTAGAGTTGA-----  
TTATTTTTTTTAGTGAATTAATTATTTGATTGGGGTGATTGAAGAATTT-  
AAATAACTTCTTT--TGT-TTTTTTTCATAATTT---TATGT--ATTTTTGATCTTTT---  
ATTAAGGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTTTGGAGAGTTCTTATCGATAAAGAAGTTTGGCAGCTCGATGTTGGAT  
TAAAA-TAAGTT-TCGGGTGCAGTAATTT-GATTA-CTAGGTCTGTTCGACCTTTAAATT-  
TTTACATGATCT??  
??  
??  
ATCGGAATGAGTACACTTTAAAT-  
CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCAGCTCCAATAGCGTATATTA AAAATTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTTCGCACTGTCGGTTCACCGCCTGTGCGGTGT  
ACTGGCATGTCGCGGAATGTCTGTGCGGTGGTGAATAAAGTTC-----  
-----CTGGTGC---TACCTAGGCT-TTTTATAGCTGAAG-T-GTGTG-CCG-----TG-G-



TGACTTT--ATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAAGTAGGCCGACACGTTTACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAAGATCTGCCTAAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAAGACGAGGTAATGATCAATGTGGACAGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTTCT  
TTAATGACTCGGCGGGGAGCTTC-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAAT??  
??  
?????????TGGTAAACTCCATCTAAGGCTAAATATAACCACGAGTCCGATAGTAAACA  
AGTACCGTGAGGAAAAGTTGAAAAGAAGCTTTGAAGAGAGAGTTCAATAGTACGTGA  
AACCGCTCAGGGGTAAACGGAGAAGATCTGAA-  
TATCTCGAAAGAAGAGATTCAAGTCTTAGCAT---TTTGTTTACTTTTAAGTAC----  
CTTTGAGATGGTAA-TTCAGTATTATCCAG---TAAAT--ACTGTG--GTAA-  
GAAGTTTTGCCGCTCGTACTTT-ATTTT-  
AAGTAAGCTGTAATGCTGGACCGCACTTCTCTTTTAGTAAGGACGTTGCAACCTGTC  
AGAT--AAT----ACTCTAAGT--AT-G----ATTAA---AGTGTCTCTGT-A-AATGTA--  
ATTTAAAAC-----TTGTTTTA-----ATATTATATCTTGCAGTAGC-TTTTA--  
TCA--AT---TTTACTACTGGGTGTTTA-----TATTCTGACAGTATCAACT-ATGG-  
TGTAGAGCCACTAAAATTA-----AGTAAGCATCCG-----  
-TTTTACAA-GTTTGGTA-----ATTATAAGTT-A-ACGTCTAACTT-TT-----  
GTTAGTCTAA-CTT-TATTACCTA-CTGTTTGTA---AAA-TG--TA-ACTTGCTTTTAG--  
TGCTCA--  
TCACACCGGTTGGCAGCGATTTCAGGTTTCAGATTCTTATCCG????????????????  
??AGGCGTAATGAAAGT  
AAAGGATAAAAAGATAGAAG-----AAGTTAAG-----  
CAATTTTCATCTTTGAT-----TCT-TTTATCCTAGGGAAGATGG--ATAT-----  
CTTGTTTTAAAAA-----AAATTTT-TT-----TT-----TT-  
TTTAAACATAGAT-CTCCCGCATTCCCAGGGCGCTTG--GACTTGCTT--TG-  
CAAGTCTATGCGCACCTAGAGGGTACACGATGACACCCGAAAGATGGTGAAGTACTATG  
CCTGGCCAGGCCGAAGCCAGGGGAAACCCTGGGGGAGGTCCGCAGCGATTCTGACG  
TGCAAATCGATCGTTAGAGCTGGGTATAGGGGGGAAAGACCAATCGAACCATCTAG  
TAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGAT-----T---  
TTAAAGAGTTTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCAGGCTTTCTTAAAGAGAGAAAAAAGAAAATTTTAA????????????????????  
??TTTGGGCACCCAGAGGTCTAT  
ATTCTAATTCTACCGGGGTTTGGCCTAATTCACATATTATCGCCATAGAGACAGGGA  
AAAATGAAGCCTTTGGTTCCTTGGGAATAATTTATGCTATGCTAACCATTGGCCTTTT  
AGGATTTATTGTATGAGCACATCATATATTTACTGTAGGAATAGATGTAGATACACG  
AGCCTACTTTACTTCAGCAACAATAATATTGCTGTACCCACCGGAATTAATAATTTT  
AGATGGCTAGCAACGCTACATGGAAGAGTTCTAAGATTATCCCTAGATTATTGTGA  
GCCCTGGGATTCGTTTTCTTATTCACTATTGGAGGGTTGACTGGAGTTATTTTAGCAA  
ATTCAAGAATTGATATTGTTCTCCATGATACATACTATGTAGTTGCCCATTTT??????  
??



AGCTGGTTCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCCCGGCTTTCT??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
Opisthoplatys\_sp2 ??????????CAAAAACATGTCTTTTTGT-----  
AATTTATTTA-ATGTTGGTCTGCTCAATGA--  
TTTTATATTTAATAGCCGCAGTATATTAAGTGTGCTAAGGTAGCATAATCAATTGTCT  
TTTAATTAAGGCTGGAATGAAAGGTTTGACGAGGTAATAGCTTTCTTT-AATTTAA-  
TTAATTGAATTTAATTTTTAGGTTAAAAAGCTTA-  
AATTAATTAAGGGACGAGAAGACCCTATAGATTTTAATATTTTG-TTTA--  
ATTTTATTGATTTA--GTTA-----  
TTATTAATAAAATTGATTTATTTATTTAATTGGGGCGATTTAGAAATTT-  
ATTGAACCTTTTTA--TTT--ATTTTTCATTAATT---TATGT--TTTATTGATCCGTT--  
ATTAGTGATTATAAGATTAAATTACCTTAGGGATAACA-  
GCGTAATTTTTCCAGAGAGTTCCTTATTGATGGATAA????????????????  
??  
??  
??  
CGTAATCGGAAT  
GAGTACACTTTAAAT-CCTTTAACAAGGATCTA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTTGCCGCTGTCGGTGA  
ACTGGCATGTCGCGGTATGTCCTGTCGGTGGTGAATGTAATGG-----  
-----CAACGG----TGCCAGGGTA-TGCTTTATCAAAG-T-ATACG-  
CGGGTTAACTTGTTGTT-CAATTAC--ATTTGCCGT---  
TTTTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAGAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTGCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
??  
??  
??  
??  
ACCACGAGACCGATAGCGAACAAGTACCGTGAGGG  
AAAGTTGAAAAGAAGCTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGTTCAGGGG  
TAAACGGAAAAGACCTGAA--TATTCGAAAAGGGGAGATTCACGTCTTACCGC---  
CTGTTTAACTATTGAATGTTATTCAATTGAGATGTTTC-TTC-GCCCTGTCCAG--

TTTATAT--ACTGTG--ATGG-TGGGTTGAAACGCTCGCATTTA-ATT-----  
 AGTAAAAACAGCGGTGGACCGCACTTCTCCCTCAGT-  
 AGGACGTTGCGATTTGTCAAAA-AAAT-----ATTCTAAGC--ATCC-----GTAA---  
 AGTGTCTGTAT-T-TGTGAA--AGGTGTAGGGGA-----ATTTTTTCTTTTA-----  
 CATTTTTTTTCCCTATACAGC-TTTAA--TAA--ATAA-----ATGCTGAATA-----  
 TATTTTAACAGTAACAAATGATGG-TGTTGAGCCACTTTA-----  
 CAAAAAGTGTC-----TTGGCCA-ATTTACAA-GTGGGGTATG-----AACATTAGTT-  
 A-ATGTCGGTCTTAAT-----GTCCGTCTAA-CTG-GTTCTCCTA-CTGTTGGTA---AAT-  
 TGTATG-CTTTG--GACAG--CGCTAA--  
 CAGCACCGATCGGCAGCGATCAAGGTTTCAAGGCTCTTATCCGACCCGTCTTGAAAC??  
 ???  
 AGTAAAAAGGTGTTTGGATGAT-----GGGGGGGAAAAAT-----  
 TTTTTTYTTCCTTCAATTTAT-----CCT-TAATCCTAAGGGAAGATGA--TAAT-----  
 ---TACTTTTTTTTCC-----CCTGA---TAAAGGGG-----  
 AGAGAAAAGAAAACGAT-TGTCCGCATTCCCAGGGCGCTTG--TTCTTGCTT--TG-  
 CAAGTCTATGCGCACCTAGAGCGTACACGATAATACCCGAAAGATGGTGAACCTATGC  
 CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
 GCAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
 AGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
 GAGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
 AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAA-  
 CCCCAGCTTTCTTAAAA-----  
 TT??  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ???  
 ?

Peirates\_punctorius CGCCTGTTTATCAAAAACATGTCCCTTTTG--TT-----  
 TATTTATTTA-AGGTCTGTCCTGCCCAATGA-----  
 GTGTTTTAATGGCCGCGGTATATTAACCGTGCTAAGGTAGCGTAATCATTGTCTTTT  
 AATTGTGGCCTTGATGAAGGGTTCGAACGAGGCACATTCTTTCTTT-ATCTTGA--  
 TTTTAGAAGTTATCTTTTTAGTCAAAAAGACTAA-  
 AATTTATCTGTGGGACGAGAAGACCCTATAGATCTTTATATTACT-TTTT--  
 ATTGAGCTTGTTT---AGCAAAAT-----  
 TTATGGTTTAATTAATTGTATTATTTGATTGGGGTGATTACAGAGAAT-  
 CAGCAACCTCTGT--TTT-AGTATTCGTTTATG---GGCGT--TTAATGATCCATT---  
 AATGATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
 GCGTAATCTTCCGGAGAGTTCCCTATCGATGGAGGGGTTTGCACCTCGATGTTGGA  
 TAAAAA-TAAGTA-TTGGGTG-AGCGATTT-AATTA-  
 CTGGGTCTGTTCCGACCCTTAATTT-  
 TTTACATGATCTGAGTTCAAACCGGAGGCAGCAGGCACGCAAATTACCCACTCCCGG  
 CACGGGGAGGTAGTGACAAAAAATAACGATACGGGACTCTTTTGAGGCCCGTAAT

CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGA  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAACGGGATCC-----  
-----CCGGTGC---TGTTGAGGCC-GTC--AGGTCAAAG-C-ATGTA-CCG-----TGTG-  
TGTTTCC--GTTTGCCGA---  
TCTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAACAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACTCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACCACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCACCAGGCCAGGACATTGGAAGGATTGACAGATTGATAGCT  
CTTTCTTGATTGAGTGGTGTAGTGGTGCATGGCCGTTTC?CGTGTGCTTGATAGTGCAG  
CCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATATAACCACGAGACCGATAGCA  
AACAAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTGAAAGAGAGAGTTCAAGAGTAC  
GTGAAACCGCTCAGGGGTAACGGAAAAGACCTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCAC---TTGCTCAGCTTTTAAATA-----  
TTATGAGATGGCCA-TTC-GCCCTGTCCAG---TGTA--ACTGTG--ATGG-  
TGGGTTTGTTCGCTCGTATTTA--TT----  
ATGCAAGGGTGGTGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGACCCGTCAAAC--AAT-----GCTCTAAGT--ATTT-----GTTTT---  
AGTGTCTGTAT-A-AGTGTA-----TACCGTTA-----  
GGTATATTTATCTTGTACAGC-TTTAA--CAA--TT---TAT-ATACTGGGTATTTG-----  
TTAGACGGTAACAAACGATGG-TGTCGAGCCACTTTAATTT-----  
TTTAAAGTGTC-----CACC GGCT-GTTTGCAA-GTGAGTTTG-----GTTAATAAAT-  
T-AGTCGGGCTTT-TG-----GTCCGTCTAA-TTA-TTCCACTTA-CTGTTGGCG----GAC-  
AT--AG-CCTCG--GACAG--TGCTTA--  
CAACACCAGTCGGCAGCGATCCAGGTTCCAGGCTCTTATCCGGTCCGT?????????????  
?????CTTGAAACACGGACCAAGGAGTTTATCGTGTACGCAAGTCATTGGGTTAG--  
CTAAACCCAAAGGCGAAATGAAAGTAAATAGTGTTTAATAGTGGT-----  
-----TTTTCCATTT-----TTT-TACTTAAAGGGAAGATAG--AATC-----  
--TATTTGTAGATTTT-----AAATTTTT-TTTATAAAAAATTT-----  
---AG-TTTACAAAATAGT-TTCTCGCACTCCCTGGGCGCTTG-TTCTCGTTTT--TG-  
CGAGTCAATGCGCACCCAGAGCGTACACGCTGATACCCGAAAGATGGTGAACCTATG  
CCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACG  
TGCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAG  
TAGCTGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCGCTCGGA-----  
TAAAGAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCCCGGCTTTCTTAAAA-----  
CTGAAGCCGTGGGACTCGGATCAGAGTGCCAAGTGGGCCAATTTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??

??  
??  
??  
??  
??  
??  
??  
??  
??  
??

Pothea\_lugens ??????????AAAACATGTCCTTTAGT--AT----

TAATTATTTA-AGGTTTGCCTGCCCAATGA----  
TTTATTTAATGGCCGCAGTATATTAAGTGTGCTAAGGTAGCATAATAAATTGTCTTTT  
AATTGAAGGCTGGAATGAAGGGTGAATGAGATATCAGCTTTCTTT-TTATTAA-  
TCTTTTAAATTTAATTTATTAGTCAAAAAGCTAA-  
TATGTTATTAGAGGACGAGAAGACCCTATAGATCTTTATATTTAA-TTTT--  
TAAATTTTATTTA---TATAATTT-----  
TTATATTTTATATTAATTAATTAATTTAATTGGGGTGATTTTAGAATTT-  
TTTTAACTACTAT--CT--ATTTTTATTGATG---GATGT--TTAATTGAGCCAAT---  
TTTATTGATTGGAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTCTTTGGAGAGTTCATATCAATAAGGGAGTTTGCACCTCGATGTTGGAT  
TAAA-TTAGTA-GTGGGTGCAACAATTT-ACTAA-  
CTAGGTCTGTTCGACCTTTAAAGT-  
TTTACATGATCT??  
??  
CCGTAATCGGAATGAGTACACTTTAAAT-  
CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCTGTCGGTGGTGAATGGGATCT-----  
-----CCGGTGC---TGCCTAGGCT-TC--ACAGCTGAAG-T-ACGTG-CCG-----TT-T-  
TGATGCC--ATTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGG??  
??  
??  
??  
??  
GGTGGTAAACTCCATCTAAGGCTAAATACAACTACGAGTCCGATAGCG  
ACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGAGTAC  
GTGAAACCGCTCAGGGTAAACGAGAGAAGAACTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATATTATGTTTAACTTTTAAATGT----  
ACAAGAGATGTTTA-TTC-GCCCTATTCTRG----TGCAA--ACTGTT--ATAG-  
AGGGTTTATTCGCTCGCATTTA-ATT----  
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTATCAGAT--AAT----ACTCTAAGT--AT-G----GTTAA---

AGTGTCTGTAT-A-AAAATA--ATTGG-----TTTCGGCT-----  
AATTTTTTTTTTACAGC-TTTAA--TCA--AT---TAT-ATACTGAGTATAA-----C-  
ATTCTCTGATAGTATCAAATGATGG-TGTTGAGCCACTTGTT-----TTTTTA-----  
-AT---TAAAAAGTGTC-----TTTAGCC-GTTTACAA-GTAGGTTT-----  
AGTAGTT-A-ATTTCTGGGCTT-TT-----CGGTCCGTATAA-CTA-TGCCATTTA-  
CTGTTTGTA----AAC-TG--TG-CCTTT--GACTG--TGCTTA--  
TAACACTGATCAGCAGCGAT???  
???  
???  
???  
???  
???  
???  
???  
???  
???  
AAGTCTACATTCTAATTTTACCAGGATTTGGGCTTATTTCCCATATTATCGCAATAGA  
GACAGGAAAAACGAAGCTTTTGGAGCTCTGGGTATAATTTATGCAATAATCGCCAT  
TGGAATTTTGGGATTTATTGTATGAGCCCATCATATTTACAGTAGGAATAGATGTT  
GACACACGGGCCTACTTTACTTCCGCAACTATAAATTATTGCTGTACCTACAGGAATTA  
AAATTTTTAGATGACTTGCCACACTGCATGGTAGAAGAATCATATTTACCCCTAGAA  
TTTTATGGGCCTTGGGGTTTCGTGTTCTTTTTACTGTAGGGGGCTTAACAGGAGTTAT  
TTTAGCTAATTCAAGAATTGATATTATCTTACATGATACATATTATGTCGTAGCACAC  
TTCC????????????????????????????????????ATGTCTGGCTCATGTACAGTTAAACATGCTGG  
ATGCGATTACCACCATTTCTGTGTTGTTGGTGATAATTTAAAAGATCGTTTTGATGGAG  
CATCACGTGTAATGGTTAGCAATGCTGGCTCAATACGCGCTA---  
CTGGCGGACGGAGAAATCGCTATAACTTCCAATTAAAACCTTACAATCCAGATCATA  
AACCGCCTGGCGTTAAAGATCTAGTATACCTTGAACCATCGCCTGTTTTCTGTGAACG  
CAATCTAAGGCTAGGCATACAAGGAACCTACGGGGCCAGTGTA????????????????????  
????????????????????????????????  
    Pothea\_sp            ????????????????AACATGTCCTTTTGT--TT----AATTTATTTA-  
AGGTTTGCCCTGCCCAATGA-----  
TTTATTTAATGGCCGAGTATATTAACGTGTGCTAAGGTAGCATAATCAATTGCCTTTT  
AATTGGAGGCTGGAATGAAGGGGTGACTGAGGAATTATCTTTCTTT-TTATTAA--  
AACTTGAATTTAATTTGTTAGTTAAAAAGCTAA-  
CATTTTATTAAGGGACGAGAAGACCCTATAGATCTTAATATTTAA-TTCA--  
TTATTTTATTATG---GGTTATA-----  
ATTATAAAATAAAGCATTATGTATTTAATTGGGGTGATTTTAGAATTT-  
TTTGAACACTAT--TTT--AATTATCATTAATT---ATTGT--TTTATTGATCCAAT---  
ATTATTGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTTATTTCTTCGGAGAGTTCATATTGATGAAGAAGTTTGGCACCCTCGATGTTGGAT  
TAAAA-TAAATA-ATGGGTGCAGTAATTC-ATTGA-  
TTAGGTCTGTTTCGACCTTTAAATT-  
TTACATGATCT??  
???  
???  
???  
???  
???  
???





Racelda\_sp CGCCTGTTTATCAAAAACATGTCCTTTTGG-----  
AATTTATATA-AGGTATAACCTGCTCTATGA---  
TATTATTAATGGCCGAGTATATTAAGGTAGCATAATCAATTGTCTTT  
TAATTAAGGGCTAGAATGAATGGTTGGACGAGATGATGACTTTCTTT-TTGTTAT--  
TTTTTTAATTTGACTTGTGAGTAAAAAGGCTGA-  
TATATTTTTATGGGACGAGAAGACCTATAGATCTTTATATTATT-TTGA--  
ATTTGTTATTTT---TTAA-----  
TTTATAAATTAATTTATTGTTTTATTTAATTGGGGTGATTGTGGAATTT-  
CTTTAACTTCTTT--TAT--TATTTTTATAGATT---AGTAT--TTATATGATCTTAT---  
TTTATAGATTTTAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTCCGGAGAGTTTATATTGATGGATTAGTTTTCGACCTCGATGTTGGAT  
TAAAA-TAGGTT-TTGGGTGAAGAAGTCT-AATAA-  
CTAGGCTGTTCGACCTTTATAAT-  
TTTACATGATCTGAGTTCAAACCGGAGGCAGCAGGCACGCAAATTACCCACTCCCGG  
CACGGGGAGGTAGTGACAAAAATAACGATACGGGACTCTATTGAGGCCCGTAAT  
CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCCTGTCGGTGGTGAATAAGATCC-----  
-----CCGGTGC---CTGCGTAGGCT-TCTA-TAGCTGAAG-TATGGTG-CCG-----TA-C-  
GGATGTT--ATTTGCCGA---  
TTTTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGTAATAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGCTCGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACCACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAATCTCACCAGACCAGGACATTGGAAGGATTGACAGATTGATAGCT  
CTTTCTTGATTGAGTGGTGTGATGGCCGTT?CGTGTGCTTGATAGTGCAG  
CCCTAAGTAGGTGGTAAACTCCATCTAAGGCTAAATATAACTACGAGTCCGATAGCG  
ACAAGTACCGTGAGGGAAAGTTGAAAAGAAGTTTGAAGAGAGAGTTCAAGAGTAC  
GTGAAACCGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATA--TTGTTAACTTTTAAATGTA---  
TTCTGAGATGTTTA-TTC-GCCCTATTCAG----TGCAA--ACTGTA--ATAG-  
AGGGTTTATTCGCTCGCATTTT-ATT----  
TAGTAAGAACAGTGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTGTCAGAT--AAT-----ACTCCAAGT--AT-G-----GTTAA---  
AGTGTCTGTAT-T-TAAAAT--AATCTATCTTG-----GGGTTTACTCTAGG----  
GTAGGTAAAATTATTTTTTCTTACAGC-TTTAA--TCA--ATTTATTAA-  
ATACTGAGTATT-----TATATTGATAGTATCAATTGATGG-  
TGTTGAGCCACTTAAA-----CA---AAAAAAGTGTC-----  
TTTAGCC-GTTTACAA-GTAGGTTT-----AATAGTT-GTTTTTCGGGTTT-TT-----  
CGATCCGTATAA-CTA-GAACACTTA-CTGTTTGTA---AAC-TG--TG-CCTTA--GACTG--  
TGCCTTATAAACACTGATCAGCAGCGATCCAGGTTTCAAGGCTCTTATCCGGTCCGT???



AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCT-  
ACCCGGGAAACCAAAGCTTTTGGGTTCGGGGGAAGTATGGTTGC?????????????  
??  
??  
??  
AGAGTGCAGCCCT  
AAGTAGGTGGTAAACTCCATCTAAGGCTAAATACTACGAGTCCGATAGCGAAC  
AAGTACCGTGAGGGAAAGTTGAAAAGAAGTTGAAAGAGAGAGTTCAAGAGTACGTG  
AAACCGCTCAGGGGTAAACGGAGAAGAAGTAACTGAA--  
TTTTCGAAAGGGGAGATTCACGTCTTACCATAATATGTTTAACTTTTAAATGT----  
ACAAGAGATGTTTA-TTC-GCCCTATTCAG----TGCAA--ACTGTA--ATAG-  
AGGGTTTATTCGCTCGCATTTA-ATT----  
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTATCAGAT--AAT----ACTCTAAGC--AT-G----ATTAA---  
AGTGTCTGTAT-T-AGAATT--ATTAG-----TTTTAATTA-----  
ATTTTTTCTTTTACAGC-TTTAG--TCA--AT---TAT-ATGCTGAGTATTTGTTGTTGT-  
TTCTGCTGATAGTATCAAATGATGG-TGTTGAGCCACTTATTTT-----  
TA---TTTAAAGTGTG-----TTTAGCC-GTTTACAA-GTAGGTTT-----AGTAGTT-  
A-ATTTCCGGGCTT-TT-----CGGTCCGTATAA-CTA-GGACACTTA-CTGTTTGTA---AAC-  
TG--TG-CCTTA--GACTG--TGCTTA--  
TAACACTGATCGGCAGCGATTACAG????????????????????????????????  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
TGAAGTTTACATTTTAAATTTTACCAGGATTTGGGCTTATTTACATATTATTGCTATAG  
AAACAGGAAAAAATGAAGCTTTCGGAGCCTTAGGTATAATTTACGCTATATTAGCAA  
TCGGACTTTTAGGATTTATTGTATGAGCCCACCACATATTTACAGTAGGAATAGATGT  
TGACACTCGAGCTTATTTTACATCAGCAACAATAATTATTGCCGTACCTACAGGAATT  
AAAATTTTTAGTTGATTAGCCACTTTACATGGTAGACCCATTACATTTACTCCCAGAA  
TTATATGAGCATTAGGATTTATTTTTCTATTCATATAGGAGGATTAAGTGGTGTAAT  
TTTAGCAAATCCAGAATTGATATTATTTCTTCATGATACATATTATGTAGTAGCACAT  
TTTCATTA??ATGTCTGGTTCATGCACAGTTAAACATGCTG  
GATGAGATTACCACCATTTCGTGTTGTTGGTGATAATTTAAAAGATCGCTTTGATGGT  
GCCTCACGAGTAATGGTCAGCAATGCTGGCTCAATACGCGCTA---  
CTGGCGGACGTAAAAATCGTTATAACTTCCAATTGAATCCCTACAATCCAGATCATA  
AACCACCCGGTGTTAAAGATCTAGTTTATCTTGAACCATCGCCAGTATTCTGTGAACG  
CAATCCAAGACTGGGTATACAAGGAACACACGGGCGTCAGTGTAATGCAACATCGC  
AGGGTGTGAAAGGATGCGATCTCATGTGTTGTGG

Rhiginia\_cinctiventris CGCCTGTTTATCAAAAACATGTCCTTTTGT--TT----  
 GTATTTATTTA-AGGTTTGCCCTGCCCAATGA-----  
 TTAATTTAATGGCCGCAGTATACTAAGGTAGCATAATCAATTGTCTTTT  
 AATTGGAGGCTGGAATGAAGGGGTGGATGAAGTATTATCTTTCTTT-TTACTAA--  
 ATTTTGAACTTAATTTTTTGGTTAAAAAGCTAA-  
 AATTTTATTGTGGGACGAGAAGACCCTATAGATCTTTATATTTAT-TTAT--  
 TTTATGATGATTTA--GTTTAATT-----  
 ATCTATTTTAAAATAATTTTTTATTTAATTGGGGTGATTTTAGAATTT-  
 AAATAACTACTAT--TTA--ATTTATCATTAAATG---AGTGT--TTTATTGATCCTTT---  
 ATTATTGATTTTAAAGATTAAGTTACCTTAGGGATAACA-  
 GCGTAATTTTTTCGGAGAGTTCATATTGATGAAGAAGTTTGGCAGCTCGATGTTGGAT  
 TAAAG-TAAGTT-ATGGGCGCAGTTGCCT-ATAAA-  
 CTGAGTCTGTTCGACTTTTAAATT-  
 TTTACATGATCTGAGTTCAAACCGGAG????????????AAATTACCCACTCCCGGCACGG  
 GGAGGTAGTGACAAAAAATAACGATACGGGACTCTATTGAGGCCCGTAATCGGAA  
 TGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
 GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
 TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGT  
 ACTGGCATGTCGCGGTATGTCCTGTCGGTGGTGAATGGGATCC-----  
 -----CCGGTGC---TGCCTAGGCT-TTT-ATAGTCGAAG-C-ACGTG-CCG-----TA-C-  
 GGATGCC--ATTTGCCGA---  
 TTTCTCTACTCCGGTGCTCTTAAACGAGTGTCGAGGTAGGCCGACACGTTCACTTTG  
 AACAAATTAGAGTGCTTAAAGCAGGCT-  
 GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
 TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTCTGTA  
 TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
 TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
 GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
 ATAATGACTCGGCGGGGAGCTTCT-  
 ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
 TAAAAGGAATGACGGAAGGGCACCACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
 CAACACGGGAAACCTCA??  
 ?????????????????CGTGTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTA  
 AGGCTAAATACAACACTACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAA  
 AAGAACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGAG  
 AAGAACTGAA--  
 TTTTCGAAAGGGGAGATTCACGTCTTACCATGATATGTTTAACTTTTAAATGT----  
 ACAAGAGATGTTTA-TTC-GCCCTATTCAG----TGCAA--ACTGTA--ATAG-  
 AGGGTTTATTCGCTCGCATTTA-ATT----  
 TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
 AGGACGTTGCGATCTATCAGAT--AAT-----ACTCTAAGC--AT-G-----ATTAA---  
 AGTGTCTGTAT-T-AGAATT--ATTAGT-----TTTTTGGCTA-----  
 ATCTTTTCTTTTACAGC-TTTAA--TCA--AT----TAT-ATGCTGAGTATTT-----T-  
 ACTTTTGTAGATATCAAATGATTG-TGTTGAGCCACTTATT-----A-  
 --TTTAAAGTGTC-----TTTAGCC-GTTTACAA-GTAGGTTT-----AGTAGTT-A-  
 ATTTCCGGGCTT-TT-----CGGTCCGTATAA-CTA-GGACACTTA-CTGTTTGTA----AAC-  
 TG--TG-CCTTA--GACTG--TGCTTA--  
 TAACACTGATCAGCAGGATTCAGGTTCACTCTTATCCGGTCCGT????????????????

?????CTTGAAACACGGACCAAGGAGTTTAT?GTGTACGCAAGTCATTGGGTAAT--  
 TAAACCCAAAGGCGTAATAAAAAGTAAAAAGTGATTTAAAAATAT-----  
 -----TTTTA-AATTCTTAAAGGAAGAAAG--ATTT-----  
 TTTAAATTTTTAC-----TA-CGAGTAAA-----AA-  
 AAAATTTTAAAAA-ATCTTGCATTCTAGGGCGCTTG-TCTCTCGCAT--TG-  
 CGAGTATATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
 CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
 GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
 AGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCATT-----  
 GAAGCGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
 AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAA-  
 CCTCGGCTTTCTTAAAA-----  
 TGAAGCCGTAGGATTCGGATCAGAGGCAAGGG-----  
 CCTGGTAAGCAGAAGTGGCGCTGTGG????????????????????????????TGAAGTCTACA  
 TTTAATTCTACCCGGATTTCGGACTAATCTCACACATCATTGCTATAGAAACAGGAA  
 AAAATGAAGCCTTTGGAGCTCTCGGAATAATCTATGCAATATTAGCAATTGGTCTTCT  
 AGGATTCATTGTATGGGCCACCATATATTTACAGTAGGTATAGACGTAGATACTCG  
 AGCCTATTTCACTTCAGCAACAATAATTATTGCAGTACCCACAGGAATTAATTTTC  
 AGTTGATTAGCCACTTTACACGGAAGTCCCATCACTTTACCCCTAGCATTCTTTGAG  
 CACTAGGATTTGTATTCTTATTACAAATTGGGGGATTAAGTGGAGTTATCTTAGCAA  
 TTCTAGAATTGATATTATCCTTACGATACCTACTACGTAGTAGCTCACTTC?????????  
 ?????????????????????ATGTCTGGTTCATGCACTGTAAAACATGCTGGATGAGATTAC  
 CACCATTTCGTGTTGTTGGTGATAATTTAAAAGATCGCTTTGATGGTGCATCACGCGT  
 AATGGTCAGCAATGCTGGYTCAATACGCGCAA---  
 CTGGCGGACGTAAAAATCGTTATAATTTCCAATTGAATCCTTACAATCCTGATCATAA  
 ACCACCAGGCGTTAAAGATCTAGTTTACCTTGAACCATCGCCAGTATTCTGTGAACG  
 CAATCCAAGACTGGGTATACAAGGAACCTCATGGACGTCAATGCAATGCAACATCAC  
 AGGGCGTTGAAGGATGCGATCTCATGTGTTGTGG  
           Saica\_sp          CGCCTGTTTATCAAAAACATGTCCTTTAGT-----  
 TATTTATGTATAGGTTTGGCCTGCTCTATGA----  
 AATTTTAAATGGCCGCGGTATTATAACTGTGCTAAGGTAGCATAATAAATTGTCTTTT  
 AATTGAAGACTTGTATGAAAGGTTGGACGAAGATATTTCTTTCTTT-GATTTAA-  
 TAATTAGAATCTAATTTCTTAGTTAAAAAGCTAA-  
 GATGTTAAAGAGGGACGAGAAGACCCCATAGATTTTATTATTATA-AATA--  
 TTTGATTTTGT----AGTTAA-----  
 TTATTTTTTTTAAATAGTAAAATAAGTTTAATTGGGGTGATTATAGAATTA-  
 TTTTAACTTCTAT--TCT--ATTTTTCATTTATA---TATGT--TTTTTTGATCCATT---  
 TATTATGATTATAAGATTAAAATACCTTGGGGATAACA-  
 GCGTTATTTTTTTTAAAGAGTTCTTATCGATGAAGGAGTTTGGCACCCTCGATGTTGGAT  
 TAAAG-TAAATT-TTAGGTGTAGAAGCTT-AGATT-  
 TTAGGTCTGTTTCGACCTTTTAAAC-  
 TTTACATGATCTGAGTTCAAACCGGAG????????????????AAATTACCCACTCCCGGCACGG  
 GGAGGTAGTGACAAAAAATAACGATACGGGACTCTTTTGAGGCCCCCGTAATCGGAA  
 TGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
 GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
 TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCGCCGCTGTGCGGTGTA  
 ACTGGCATGTCGCGGCATGTCTGTGCGGTGGTGAACGGGGGCA-----  
 -----CCGGTAC---TGCTTTGGCT-TTT-TAAGTCGATT-C-TAGTG-CCG-----GG-C-

TGTTTCCCTGTTTGCCGA--  
TCTCTCTACTTCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATGATAAAATAGGACCTCGGTTCTATT  
TTGTTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATTCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
ACTCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACC  
TTAAAGGAATGACGGAAGGGCACACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAATCTCA??  
????????????????CGTGTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTA  
AGGCTAAATACAACCACGAGACCGATAGCGAACAAAGTACCGTGAGGGGAAAGTTGAA  
AAGAAGTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGAA  
AAGACCTGAA--TTTTCGAAAGGGGAGATTCACGTCTTACCGC---  
TTAAATTTCTTTTTATGTA---TAAAGAGATGGTCA-CTC-GCCCTATTCAG----TGCAA--  
ACTGTA--ATAG-AGGGTTTGGTTCGCTCGCATTTA-----  
AAGATTTGTTTAGTGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGACCTGTCAAATAAAAT----GCTCTAAGT--ATTT----TTTAA---  
ATTGTCTGTAC-A-ATGGTT--ATACTGTAA-----  
AAGTTTTTCCATATACAGT-TTTAA--ATA--AT----TTT-ATACTGAGTA-----  
TATTTTGACAGTAACAAACGATGG-TGTTGAGCCACTTATTATTTTTTTA-----  
-----ATGTAAGTGTC-----TTTGGCC-GTTTGCAA-GTGGGTAA-----  
GGTAATAGTT-T-TAGTCGGACTA-TT-----GTCCGTCTAG-TTA-TTCCTCCTA-  
CTGTTGGTA---AAC-TG--TG-CCTTA--GACTG--TGCTTA--  
CCACACCAGTCGGCAGCGATCCAGGTTTCAGGCTCTTATCCGGTCCGT????????????  
????CTTGAAACACGGACCAAGGAGTTTAGCGTGTACGCAAGTCATTGGGTATT--  
TAAAACCCAAAGGCGAAATGAAAGTAAAGAGTAGTAAATTGTTGC-----  
AGAAATT-----TATTTTTTGTATATAAT-----TTCTCTTTAGGGGAAGATGG--  
AAAG-----TTAATCATAAATTG-----ATTATTGATAATT-  
TTTTATCTATAATT-----GT-TTTTGATATAGCT-  
TTCCCGCATTCCCGGGGCGCTTG-CATCTCGCTT--TG-  
CGAGTCAGTGCACCTAGAGCGTACACGCTGATACCCGAAAGATGGTGAAGTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCGGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAA-----T---  
GTGACGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCAAAAACGAC  
CTCAACCTATTCTCAAACCTCAAATAGATGAGAA-CCCCGGCTTCTTGAAAA-----  
-----

ATGAAGCCGTGGGATTCGGATCAGAGTGCCAAAGTGGGCCAATTTTGGTAAGCAGAA  
CTGGCGCTGTGG??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

??  
??  
??

Santosia\_sp                      ??????????A AAAACATGTCCTTTTGT--T-----

TATTTTTTTA-AGGTCTAGCCTGCCAATGA---  
AGTTTTATTTAATGGCCGAGTATATTAAGTGTGCTAAGGTAGCATAATCAGTTGTCTT  
TTAATTTAAGGCTGGAATGAAGGTTGGACGAGATATTATCTTCTTTATTTTTAA--  
AATTTTAACTTTGAGTTAAAAGGCTCA-  
TTTAAATTTAAGGACGAGAAGACCCTATAGATCTTTATATTTAT-ATTT--  
TTGTTTTTTGTA--GCTAAATA----  
AAAATTTATAAAGCAGTATTTATTTAATTGGGGTGATTTTCAGAATTT-  
TAAAAACTTCTGT--AT---TTTTAACATTGATG---GATGT---TTTGTGATCCAAT---  
ATTATTGATTATTAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATCTTTTGGAGAGTCTTATCGATAAAAAGAGTTTGGCAGCTCGATGTTGGAT  
TAAA-TGAGTT-TGGGTGCAAAAGTTC-AATTA-  
CTGGTCTGTTGACCATTAAGT-  
TTACATGATCTGA??  
??  
CCTTAACCAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCAGCAGCCGCGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGTTCTGCGTGCCGCGCTGTCGGTTCGTCGCCTGTCGGTGT  
ACTGGCATGTCGCGCATGTCCTGTCGGTGGTGGATGGAATCC-----  
-----ACGTTAC---TGCTTTAGTTTACTATAAGCTGAAG-T-ATGTA-TCG-----TT-C-  
GGGCGCC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
GAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGCGATTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAAATTGATCAAGAACGAAAGTTAGAGGTTGAAGGCGATCA  
GATACCGCCTAGTTCTAACATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCA  
ATAATGACTCGGCGGGGAGCTTCC-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAAT??  
??  
??  
??  
??  
GTTGAAA  
AGAACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGATCAGGGGTAAACGGAGA  
AGATCTGAA--TTTTCGAAAGGGGAGATTACGTTTACCATA--  
TTGTTAACTTATAAATGTA---TTAAGAGATGTTTA-TTC-GCCCTATTTAG----TGCAA--  
ACTGTA--ATAG-AGGGTTTATTCGCTCGCATTATTTT----  
TAGTAAAAACAATGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGATCTGTCAGTT--AAT----ACTCGAAGT--AT-G----ATTAA---  
AATGTCTGTAT-T-TATGTA--ATTTT-----TTTTTTTTA-----  
AATTATTTTTTTACAGC-TTAA--TCA--AT---TTAATACTGAGTATTT-----T--  
AATATTGATAGTAAACAATGATAG-TGTTGAGCCCTTTAA-----  
--AAAAAGTG-----TCTAGCC-GTTTACAA-GTATGTTTTTAGG--CTTATTTAGCT-  
A-ATTTCCGGTTTT-TA-----TGATCCGTATAG-CAA-GGCCACATA-CTGTTGGTA---  
GATGTG--CT-CCTA--







TTTAATTAAGGCTTGAATGAACGGTTGGACGAGGTGTTATCTTTCTTT-TAATTAT--  
TTGATTAATTTAATTTTAAAGTTAAAAAGCTTA-  
AATTTTTTTATGGGACGAGAAGACCTATAGATCTTTATATTTAT-AGGG--  
ATTAAATTTTTTG---GTTTGAT-----  
ATTTTATTTGATTTTTTTATGTATTAAATTGGGGTGATTGAAGAATTT-  
TTTTGACTTCTAT--TGTTTATTATTCATTTATT---TATGT--TTTATTGATCCATT----  
TTATTGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTCTTTTGGAGAGTTCTTATCGATAAATGAGTTTTCGACCTCGATGTTGGAT  
TAAAT-TAAGTT-TTAGGTGTAGTGGCTT-ATTA-CTAGGTCTGTTTCGACCTTTAAAT-  
TTTACATGATCTGA??  
??  
??  
CCTTAAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGTAATCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCCGTTCCGCCCTGTCCGGTGT  
ACTGGCATGTCCGCGCATGTCTGTCCGGTGGTGAATGAAATCC-----  
-----CTGGTGC---TTACGTAGGCT-TTTATTAGCTGAAG-T-TTGTG-CCG-----TG-G-  
TGACTTC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCC-  
ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACG??  
??  
??  
??  
CTAAGGCTAAATACAACCACGAGTCCGATAGCGAACAAGT  
ACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAAGAGTACGTGAAAC  
CGCTCAGGGGTAAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTCACGTCTTACCGC---TTGCACGACTTTGAAGTGC----  
TTTTGAGATGGTAA-TTC-GCCCTATCCAG----TGAAT--ACTGTG--GTAG-  
AGGGTTTTACCGCTCGCATTTT-ATT----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAGT--AT-A----GTTAA---  
AGTGCCTGTCA-A-AAGGTT--ATAAAAT-----TTATTTTA-----  
TAACCATTTTACAGC-TTTAA--CTA--AT---TTT---ACTGGGTATTTA-----  
TTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTAAAAATT-----ATTTATA-  
----ATT----TAAAAGTGTC-----TTTTGGTTGCC-GTTTACAA-GTGGGTTTTT----  
AGGTATTAGTT-A-ATGTCGAACTT-TT-----GTTTCGTCTAA-CTA-AGCCGCCTA-  
CTGTTTGTGTA----AAC-TG--TG-CCTTA--GACAG--TGCTTA--  
CAACACCGGTCGCGCAGCGATCCAGG????????????????????????????????????  
??  
GTACGCAAGTCATTGGGTGAT-  
ATAAAACCAAAGGCGTAATGAAAGTAAAGAGTAGAAGGATAGAAG-----  
AAGAAAAT-----TAATTTTCATCTTTGAT-----TCT-  
TTCTCTCAAGGGAAGATGGATATAT-----CTTATTTTAAATAA-TATTTAG-----  
-----TAT--TT-ATATTAAG-TA-----TA-TTTAAAATTTAGA-

TTCCCGCATTCCCAGGGCGCTTG-TATCTCGCTC--TG-  
CGAGATTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTCTTAAAA-----

TTGAAGCCGTGGGATTAGGATCAGAGTGCCAA????????????????????????  
??  
CACACATTATTGCAACAGAACTGGCAAGAGAGAAGCATTGGATCATTGGGTATAA  
TTTACGCTATAATTACTATCGGACTTCTGGATTATTGTATGAGCTCACCATATGTT  
ACTGTTGGGATAGATGTTGACACTCGAGCCTACTTCACCTCAGCAACTATAATTATG  
CCGTACCCACAGGAATTAATACTTTCAGATGACTAGCAACATTACATGGAAGAGTAA  
TAACCTTTTCTCCTAGAATACTATGAGCCTTAGGATTTGTGTTCCCTGTTCACTATTG  
AGGCTTAACAGGAGTCATGCTAGCTAATTCCAGAATTGATATTATCTTACACGATAC  
ATACTATGTAGTAGCACATTTTC????????????????????????????????  
??  
??  
??  
??  
??  
??

*Tanindrazanus\_tenebricus*

??  
??  
GGTTGGACGAGGTATTGACTTTCTTT-TAATTAT--  
TTAATTAATTTAATTTTAAAGTTAAAAAGCTTA-  
ATTTTTTTTATGGGACGAGAAGACCCTATAGATCTTTATATTTAT-AAAG--  
TTTAAATTTTTTA---GTTTTA-----  
TTTTTTGTTGATCTTTTATGTATTTAATTGGGGTGATTGAAGAATTT-  
TTTTAACTTCTAA--TGTTTTTTTTTCATTTATT---TATGT--TTTATTGATCCATA---  
TTAATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTCTTTGGAGAGTTCTTATCGATAAATGAGTTTGCACCTCGATGTTGGAT  
TAAAT-TAAGTT-TTAGGTGTAGTGGCTT-ATTAA-CTGGGTCTGTTTCGACCTTTAAAAT-  
TTTACATGATCTGAGT????????????????????????????????????  
??  
?????GCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTCGCCGCCTGTCGGTGT  
ACTGGCATGTCGCGGCATGTCCTGTCGGTGGTGAATGAAATCC-----  
-----CCGGTGC---TTACGTAGGCT-TTTATTAGCTGAAG-T-TTGTG-CCG-----TG-G-  
TGACTTC--ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGCATTCTGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCTT  
TTAATGACTCGGCGGGGAGCTTCC-

ACCCGGGAAACCAAAGCTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAG??  
??  
??  
??  
AGTCCGATAGCGAACAAAGTACCGTGAGGG  
AAAGTTGAAAAGAAGTCTTGAAGAGAGAGTTCAGAGTACGTGAAACCGCTCAGGGG  
TAAACGGAGAAGATCTGAA--TATCCGAAAGGGGAGATTACGTCTTACCGC---  
TTGCACGACTTTGAAGTGC---TTTTGAGATGGTAA-TTC-GTCCTATCCAG---TGAAT--  
ACTGTG--GTAG-AGGGTTTTACCGCTCGCATTTT-ATT---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT----ACTCTAAGT--AT-A----GTTAA--  
AGTGCCTGTCA-A-AAGGTT--ATAAAAT-----TTATTTA-----  
TAACCATTTTACAGC-TTTAA--CTA--AT---TTT---ACTGGGTATTTA-----  
TTTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTAAAAATT-----ATTTATA-  
---ATT---TAAAAGTGC-----TTTTGGTTGCC-GTTTACAA-GTGGGTTTTT----  
AGGTATTAGTT-A-ATGTCGAACTT-TT-----GTTCTGTCTAA-CTA-AGCCGCCTA-  
CTGTTTGTA----AAC-TG--TG-CCTTA--GACAG--TGCTTA--  
CAACACCGGTCGGCAGCGATCCAGGTTTCAGGCTCTTATCCGACCCGTCTTGAACAC  
GGACCAAGGAGA????????????????????????????????????ACGCAAGTCATTGGGTGAT-  
ATAAAACCAAAGGCGTAATGAAAGTAAAGAGTAGAAGGATAGAAG-----  
AAGAAAAT-----TAATTTTCATCTTTGAT-----TCT-  
TTCTCTCAAGGGAAGATGGATATAT-----CTTATTTTAAATAA-TATTTAG-----  
-----TAT--TT-ATATTAAG-TA-----TA-TTTAAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTT-TATCTCGCTC--TG-  
CGAGATTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAAGTATGCG  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAC-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTAGGATCAGAGTGCCAA????????????????????????????????  
????????????????????????????????????GTATATATTCTGATTTTACCAGGATTTGGACTCATT  
CACATATTATTGCTACAGAAACCGGTAAAAGAGAAGCGTTGGGTCACTAGGAATAA  
TTTATGCAATAATTACTATTGGACTTTTAGGATTCATTGTCTGAGCTCATCATATATC  
ACTGTGGGAATAGATGTTGATACTCGAGCCTATTTCACTCAGCAACTATAATCATTG  
CTGTACCTACAGGAATTAATACTTTCAGATGATTAGCAACATTACATGGAAGAACCA  
TGACATTTTCCCCAAGAATATTATGAGCATTAGGATTTGTATTCCCTATTCACAATTGG  
AGGTTTAAACAGGAGTAATACTAGCCAACCTCAAGAATTGATATTGTTCTTACGATAC  
ATACTATGTAGTAGCACATTTTAC??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

Tanindrazanus\_varicolor

??  
??  
??  
??



TTATTTTCGCACATTATTGCAATAGAAAACCGGTAAAAATGAAGCATTCGGCTCATTGG  
GGATAATCTATGCAATAATCACTATTGGATTATTGGGGTTCATTGTCTGAGCTCATCA  
CATATTTACTGTAGGGATAGATGTAGATACTCGAGCCTACTTCACTTCGGCAACTATA  
ATTATCGCTGTACCAACAGGAATCAAAATCTTTAGATGATTAGCAACATTACAAGGC  
AGTATCATATTATTTTCTCCTAGAATGATATGAGCCCTTGGCTTCGTATTTTATTAC  
AATTGGAGGATTAACAGGTGTCATACTCGCTAATTCAAGAATTGATATTATCCTCCAT  
GATACTTATTACGTAGTAGCACATTTCCA?????????????????????????????  
???  
???  
???  
???  
???

Toxopus brucei

??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
????????????????????????CGAGTCCGATAGCGAACAAAGTACCGTGAGGGAAAGTTGAAAAGAAC  
TTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAACCGGAGAAGATC  
TGAA--TATCCGAAAGGGGAGATTACCGTCTTACCAC---TTGCTCGACTTTGAAGTGC--  
--TCTTGAGATGGTAA-TTC-GCCCTATCCAG----TGCAT--ACTGTG--ATAG-  
AGGGTTTTACCCTCGCATTTT-ATA---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGGA--AAT----ACTCTAAGT--AT-A----GTAA---  
AGTGCCTGTTT-A-AATGTT--ATAAAATTT-----ATATTTTA-----  
TAACTTTATTTTACAGC-TTTAA--TTA--AT----TTT---ACTGGGTATTTT----  
TTTTTGTTCTGACAGTAACGACTGATGG-TGTAGAGCCACTTAAACATT-----  
ATTTTTTATA-----ATG----TAAAAGTGTGCGCT-TATTCTTTTTTTGCT-GTTTACAA-  
GTGGGTTTAA-----GGTATTAGTT-A-ATGTCGAACTT-TT-----GTTTCGTCTAA-CTA-  
AGCCGCTA-CTGTTTGTAA----AAC-TG--TGCCCTTG--GACAG--  
TGC??  
????????????????????????ACGCAAGTCATTGGGTGAT-  
AGAAAACCCAAAGGCGTAATGAAAAGTAAAGAGTATAAAGATAGAAG-----  
AAGAAAAT-----TCATTTTCATCTTTGAT-----TCT-

TTCTCTCAAGGGAAGATGG--ATAT-----CTTATTTTAAATAA-TATTTAG-----  
-----TAT--TT-TTATTAAG-TA-----TT-TTTAAAATTTAGA-  
TTCCCGCATTCCAGGGCGCTTG-TATCTCGCTC--TG-  
CGAGTCTACGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGTC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCAAT-----T---  
GTGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGAT????????????????????????????????????  
??TATATATTCTCATTTTACCAGGATTTGGCTTAATTTACAT  
ATTATCGCAACAGAGACTGGTAAAAGTGAAGCTTTTGGATCATTAGGAATAATTTAT  
GCAATATTAACAATTGGATTATTAGGATTTATTGTTGAGCACACCATATATTTACTG  
TAGGTATAGATGTAGACACCCGAGCATATTTTACCTCTGCAACCATAATTATTGCAGT  
ACCTACAGGAATTAATAATTTTATAGATGACTAGCCACTCTTCATGGAAGATCTACAAA  
ATTCTCACCAAGAATACTATGAGCATTAGGATTCGTATTCCTTTTCACTATTGGTGG  
TTAACGGGAGTAGTACTTGCAAACCTCAAGAATTGATATTATTTTACATGACACTTATT  
ATGTAGTAGCCCACTTCCA??  
??  
??  
??  
??  
??  
??  
??

Toxopus\_fisheri

??  
????????????????CTGTGCTAAGGTAGCATAATCAATTGTCTTTTAATTGTGGGCTAGTATG  
AAAGGTTGCACGAAACATCTACTTTCTTT-TAATTAT--  
TTAATGAATTTAATTTTAAAGTTAAAAGCTTA-  
AATATTTTGGAGGACGAGAAGACCCTATAGATCTTTATATATAA-TAAG--  
ATTGATTTTTTTT---GTTTATAA-----  
TTGTATTATCAATTTTATAATTATTTCATTGGGGTGATAGGAGAATTA-  
TAATAACTTCTCT--TTA-ATTGTTTCATATATG---TATGT--TTTTTTGATCCATT---  
GTGAATGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTAATTTTTTTGGAGAGTTCCTTATTGATAAAGAAGTTTGCGACCTCGATGTTGGAT  
TAAA-TAAGTT-ATAAGTGCAGGAGCTT-ATTTA-  
CTGGGTCTGTTTCGACCTTTAAATT-  
TTTACATGATCT??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

??AAGGCTAAATACAACCACGAGTCCG  
ATAGCGAACAAAGTACCGTGAGGGAAAGTTGAAAAGAACTTTGAAGAGAGAGTTCAA  
GAGTACGTGAAACCGCTCAGGGGTAACGGAGAAGATCTGAA--  
TATCCGAAAGGGGAGATTACGTCTTACCAC---TTGCTCGACTTTGAAGTGC----  
TCTTGAGATGGTAA-TTC-GCCCTATCCAG----TGCAT--ACTGTG--GTAG-  
AGGGTTTTACCGCTCGCATT---ATT----  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAAAT--AAT-----ACTCTAAGT--AT-A----GTTAA---  
AGTGTCTGTCT-A-AAGGT--ATAAAAT-----TTATTTTA-----  
TAACATTTATTAACAGC-TTTAATTTTA--AT---TTT---ACTGGGTATTTA-----  
TTTTTTTTTGACAGTAACAAATGATGG-TGTAGAGCCACTTAAACATT-----  
ATTTATA----ATG----TAAAAGTGTCCA-----TTTTTGCC-GTTTACAA-  
GTTGGTTAAGGT-ATTTTTTTAGTT-ATTAGTCGAACTT-TT-----GTTTCGTCTAA-  
CTA-AGCCGCCTA-CTGTTTGTGTA---AAC-TG--TG-CCTTG--GACAG--TGCTTA--  
CAACACCGTTCGGCAGCGATCCAGG????????????????????????????????????  
????????????AGGAGTTTATTGTGTACGCAAGTCATTGGGTGAT-  
AGAAAACCCAAAGGCGTAATGAAAGTAAAGAGTAGAAAGATAGAAG-----  
AAGAAAAT-----TAATTTTCATCTTTGAT-----TCT-  
TTCTCTCAAGGGAAGATGG--ATAT-----CTTATTTTAAATAA-TATTTAG-----  
-----TAT--TT-ATATTGAG-TA-----TA-TTTAAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-TATCTCGCTC--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCCTCGCTCGGT-----T---  
GAGAAGAGTCTCATCCGCTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCCCGGCTTTCTTAAAA-----  
TGAAGCCGT??  
????????????????????TACATTTTAATTCTACCCGGATTTCGGATTAATTTCCCATATTATC  
GCAATAGAAACAGGAAAAAATGAAGCATTGATCTCTAGGGATAATTTATGCAAT  
ACTTACTATTGGACTATTAGGATTTATTGTTGGGCCCATCATATATTACAGTTGGA  
ATAGATGTAGATACCCGAGCTTATTTCACTTCAGCAACAATAATTATTGCCGTACCTA  
CCGGTATTAGGATTTTATGTTGACTAGCTACAATACATGGAAGTATTATAACTTTCTC  
ACCCAGAATATTATGAGCATTAGGATTCGTATTTTATTACAAATTGGAGGATTAACA  
GGAGTTATTTTGGCAAACCTCTAGAATTGATATTATTTTACATGATACTTATTATGTAG  
TTGCACACTTCCAT??  
??  
??  
??  
??  
??  
??  
??

Toxopus\_griswoldi

??  
??  
??  
??  
??  
??  
??



??  
??  
??TTCCAGCTCCAATAGCGTAT  
ATTAAAGTTGTTGCGGTTAAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTT  
CGCCGCCTGTCGGTGTACTGGCATGTCGCGGCATGTCTGTGCGGTGGTGAATGAAA  
TCC-----CCGGTGC---TACGTAGGCT-TTTATTAGCTGAAG-T-  
TTGTG-CCG-----TG-G-TGACTTC--ATTTGCCGA---  
TTTCTCTACTCCGGTGTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTAATTGATCAAGAACGAAAGTTAGAGGTTGAAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCT  
TTAATGACTCGGCGGGGAGCTTCT-  
A??  
??  
??AAGGCT  
AAATACAACCACGAGTCCGATAGCGAACAAGTACCGTGAGGGAAAGTTGAAAAGAA  
CTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGAGAAGAT  
CTGAA--TATCCGAAAGAGGAGATTCACGTCTTACCAC---  
TTGCTCGACTTTGAAGTGC---TCTTGAGATGGTAA-TTC-GTCCTATCCAG----TGAAT--  
ACTGTG--GTAG-AGGGTTTTACCGCTCGCATTTT-TTT---  
AGTAAGGTGCAGTGGTGGATCGCACTTCTCCTTTAGT-  
AGGACGTTGCGACCTGTCAGAT--AAT-----ACTCTAAGT--AT-A-----GTTAA---  
AGAGTCTGTTT-A-AAGGT--ATAAAAT-----TTATTTTG-----  
TAACTTTTTTTTACAGC-TTTGA--TTA--AT---TTT---ACTGGGTATTT-----  
TTATTCTGACAGTATCGACTGATGG-TGTAGAGCCACTTAAACATT-----  
ATTTATA----ATG---TTTAAAGCGTCCA-----TTGTTGCC-GTTTACAA-  
GTGGGTTAAG-GT--ATTAATTAGTT-A-ATGTCGAACTT-TT-----GTTGCTCTAA-CTA-  
AGCCCCCTA-CTGTTTGTA----AAC-TG--TG-CCTTG--GACAG--TTCTTA--  
CAACACCGGTCAGCAGCGATCCAGG??  
??GCGTAATGAAAGTAAAGA  
GTAGAAAGATAGAAG-----AAGAAAT-----TTATTTTCATCTTTGAT-----  
-----TCT-TTCTCTCAAGGGAAGATGG--ATAT-----C-TAATTTAAATTA-----  
-----TAT--TTATTATTAAGTTA-----TA-TTTAAAATTTAGA-  
TTCCCGCATTCCCAGGGCGCTTG-TATTTGCTT--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCACTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCGGATCAGAGTGCCAAGTGGGCAATTTTGGTAAGCAGAAC  
TGGC-----  
G??  
CACATATTATTTCAATAGAAACTGGGAAAAACGAAGCGTTTGGATCACTTGGGAATAA



TTCCCGCATTCCCAGGGCGCTTG-TATCTCGCTC--TG-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAACCTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACGT  
GCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGCTCGGT-----T---  
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTC??  
??  
????????AGTTTATAATTTAATTTACCAGGATTGGTTAATTTCCCACATTATCGCAA  
CAGAAACCGGAAAAACGAAGCATTTCGGATCACTAGGTATAATTTATGCTATACTTA  
CCATTGGATTATTAGGGTTCATCGTCTGAGCCCATCACATATTTACAGTGGGAATAG  
ATGTAGACACTCGAGCCTATTTACATCAGCAACCATGATTATTGCAGTACCTACAG  
GAATTAATAATTTTAGATGATTAGCAACAATACACGGTAGAATTATAACCTTTTAC  
CCAGAATACTATGAGCCTTGGGATTCGTTTTCCTATTTACTATTGGGGGACTTACAGG  
AGTTATCCTAGCAAATTCAAGAATCGATATTATTCTCCATGATACTTATTATGTAGTT  
GCCCACTTCCATT??  
??  
??  
??  
??  
??  
Toxopus\_vazimba ??????????AAAACATGGTTTTAGT-----  
TAATTATTTA-AGATTGGGCCTGCCAATGA---  
TGTTTATTTAATGGCCGCAGTATTTAACTGTGCTAAGGTAGCATAATCATTTGTCTTT  
TAATTAAGGGCTTGTATGAATGGTCGGACGAGGTATTATCTTTCTTT-TAATTAT--  
CTTGTGAATTTAATTTTGTAGTTAAAAAGCTTA-  
AATTTTTTTGAGGGACGAGAAGACCCTATAGATCTTAATATTTAT-AGAT--  
TTTTGATTTGTTT---GTTTTAT-----  
TGAAGAATTTAATGTTTTAGGTATTTGTTGGGGTGACAGGAGAATTTATAAATAACTT  
CTTT--TAGTTTTTATCATTTATT---TATGT--  
TTTTTTGATCCATTTATTTAAATGATTATAAGTTTAAAGTTACCTTAGGGATAACA-  
GCGTAATTTTCTTGGAAGTTCTTATTGATGGGGAAGTTTGCACCTCGATGTTGGAT  
TATAA-TAAGTTGTAAGGTGTAGTAGCTT-ACTTA-  
CTGGGTCTGTTTCGACCTTTAAATT-  
TTACATGATCTGAGTT??  
??  
??  
??  
TAAAAAGCTCGTAGTTGGT  
TCTGCGTGCCGCGCTGTCGGTTCGCCGCTGTCGGTGTACTGGCATGTCGCGGCATG  
TCCTGTCGGTGGTGAATGAAATCC-----CCGGTGC----  
TACGTAGGCT-TTTATTAGCTGAAG-T-ATGTG-CCG-----TG-G-TGACTTC--  
ATTTGCCGA---  
TTTCTCTACTCCGGTGCTCTTAAACGAGTGTGCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATATCTGCCTGAATAGTGGTGCATGGAATAATAAAATAGGACCTCGGTTCTATT  
TTATTGGTTTTAGGAATATGAGGTAATGATCAATGTGGACTGGCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCTTGGATCGTCGCAAGACGCACTAGAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTTCT  
TTAATGACTCGGCG??



TATGGATTAAATTG---GTGTTAAT-----  
AATTATTTTTATTTAGTTGAATATTTTTGTTGGGGTGATGTTGAAATTT-  
TAATAACTTTTCAT--AT---ATTTTTCATTAAATT---AGTGTA-TTTAATGATCTTTT---  
ATTATTGATTATAAGATTAAGTTACCTTAGGGATAACA-  
GCGTTATTTTTACGGAGAGTTCTTATCGATGTATGAGTTTGCACCTCGATGTTGGAT  
TAAA-TAAATG-ATAGGTGTAGTAGTTT-ATTAA-  
TTAGGTCTGTTGACCTTTAAATT-  
TTTACATGATCTGAGTTCAAACCGGAGGCAGCAGGCACGCAAATTACCCACTCCCGG  
CACGGGGAGGTAGTGACAAAAAATAACGATCGGGGACTCTATTGAGGCCCGTAAT  
CGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCCA-TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGAATAATTCCAGCTCCAATAGCGTATATTAAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTCGGTTTGCCGCTGTCGGTGA  
ACTGGCATGTCTCGGCATGTCTGTCGGTGGTGAATGTAATCTGGCCAATAGGTTTA  
AGGGTTAGTAGTTCGATGTGCGCTAATGCGC----TACACGGGCT-  
TTTCCCCTTCTAACTTGTGTTGGT-TTT-----TT-T-TGATTAT--  
GTTTACCGATTTTTTTCTACTCCGGTGCTCTTAAACGAGTGTCGAAGTAGGCCGAC  
ACGTTCACTTTGAACAAATTAGAGTGCTTAAAGCAGGCTGGAATATCTGCCTGAAT  
AGTGGTGCATGGAATGATAAAATAGGACTCGGTTCTATTTTATTGGTTTTAGGAATA  
TGAGGTAATGATCAATATGGACTGGCGGGGGCATTTCGTATTGCGACGTTAGAGGTGA  
AATTCTTGATCGTCGCAAGACGCACTAGAGCGAAAGCATTGCCAAGTATGTCTTA  
ATTGATCAAAAACGAAAGTTAGAGGTTCCAA-  
GCGATCAGATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGA  
TGTTCCTTTAATGACTCGGCGGGGAGCTTTT-  
ACCCGGGAACCAAAGCTTTTTGGGTTCCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACCCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCA??  
????????????????CGTGTTGCTTGATAGTGCAGCCCTAAGTAGGTGGTAAACTCCATCTA  
AGGCTAAATAGTACCACGAGACCGATAGCGAACAAAGTACCGTGAGGGGAAAGTTGAA  
AAGAACTTTGAAGAGAGAGTTCAAGAGTACGTGAAACCGCTCAGGGGTAAACGGAA  
AAGACCTGAA--TATTCGAAAGGGGAGATTCACGTCTTACCGC---  
ATGTTAACTTTTTATCTT----TTATGAGATGATTT-TTC-GCCCTGTCCAGTTTTTTTAT--  
ACTGTG--ATGGTTGGGTAAATCGCTCGTATATA-ATT-----  
AGTAAAAACAGCGGTGGACCCGCACTTCTCCCTTAGT-  
AGGACGTTGCGACTCGTGGGAC--AATATATTATTCTAAGC-TAT-----ATTAA---  
AGTGTCTTTAT-A-TATTGT--ATGTAGCAT-----TGTTA-----  
TTTATAATATATATAGC-TATAA--TAA--ATAA-----ATGCTGAATA-----  
ATTTCCGACAGTAACGAATGATAG-TGTTGAGCCGCTAATTTTTT-----  
---TCTTTAGCGTT-----TTGACCA-ATTTACAA-GTGGGGTAKG-----  
AATAATTATT-A-ATGTCGAACTT-TT-----GTTCGTCTAA-TAT-GTTCTCCTA-  
CTGTTGGTA----AAT-TGTTTGT-TTCA--AACAA--GGCTTA--  
TAACACTGATCAGCAGCGATCCAGTTTCAGGCTCTTATCCGGTCCCGT????????????  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??



---TCTTTAGCGTT-----TTGACCA-ATTTACAA-GTGGGGTATG-----  
AATAATTATT-A-ATGTCGAACTT-TT-----GTTTCGTCTAA-TAT-GTTCTCCTA-  
CTGTTGGTA---AAT-TGTTTG-TTTCA--AACAA--GGCTTA--  
TAACACTGATCAGCAGCGATCCAGTTTCAGGTACTTATCCGACCCCGTCTTGAAA????  
??TGTACGCAAGTCATTGGGTAAATATTAACCC  
AAAGGCATAATAAAAGTAAATATAATGTAAAAA-----GGAAAAAT----  
---ACTTTTTTAATTTTTT-----TTACATTACAAAGAGAAGATGG--TATA-----  
--TTGTTTTTGTACT-----CTT-  
TGGTATAAAAGAT-TGCCCGCATTCTCAGGGCGCTATTACTCTCGCTT--TG-  
CGAGTGTATGCGCACCCAGAGCGTACACGATAATACCCGAAAGATGGTGAACCTATG  
CCTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCCGCAGCGATTCTGACG  
TGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAG  
TAGCTGGTTCCCTCTGAAGTTTCCCTCAGGATAGCTTGCGCTCGTG-----AT---  
GTGATTAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAA-  
GGCCGGCTTCTTAAAA-----TTGAAGCCG-----  
GTCAGATATATTTGTTTTGTTTTGGATAA????????????????????????????????????  
AGTATATATTCTTATTCTACCGGGATTTCGGACTCATTTACATATTATTTCTATAGAA  
ACAGGAAAAAATGAAGCTTTTGGAGTACTAGGAATAATTTATGCCATATTAGCTATT  
GGACTTCTGGGATTTATTGTTTGGAGCACATCATATGTTTACAATCGGAATGGACGTA  
GATACACGGGCCTATTTTACATCAGCTACAATAATTATTGCTGTGCCAACAGGTATTA  
AAATCTTCAGATGATTAGCCACATTGCATGGAAGAGTAATTATATATAATCCAAGAA  
TTTTATGAGCCTTAGGATTTGTATTTCCTATTACAATCGGGGGATTAACAGGAGTTAT  
TCTCGCTAATTCAAGAATTGACATTGTACTACACGACACCTATTATGTAGTAGCCAC  
TTCCA??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??

Tribelocodia\_sp

??  
??  
??  
??  
??  
??  
??  
??  
??  
TACCCACTCCCGGCACGGGAGGTAGTGACAAAAAATAACGATACGGGACTCTATT  
GAGGCCCGTAATCGGAATGAGTACACTTTAAAT-CCTTTAACAAGGATCTA-  
TTGGAGGGCAAGTCT--  
GGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTAAGTTGTTGCGGT  
TAAAAGCTCGTAGTTGGTTCTGCGTGCCGCGCTGTTCGGTTCCGCCCTGTTCGGTGT  
ACTGGCATGTTCGCGCATGTCCTGTCGGTGGTGAACGAGAATC-----  
-----CTGGTAC---TGCTTAGGCC---TAACGGCCAAAG-T-TTGTA-CCG-----TG-G-  
TGTTTTTC--GTTTGCCGA---  
TTTCTCCTACTCCGGTGCTCTTAAACGAGTGTTCGAGGTAGGCCGACACGTTCACTTTG  
AACAAATTAGAGTGCTTAAAGCAGGCT-  
AAAATCTCTGCCTGAATAGTGGTGCATGGAATGATAAAATAGGACCTCGGTTCTATT

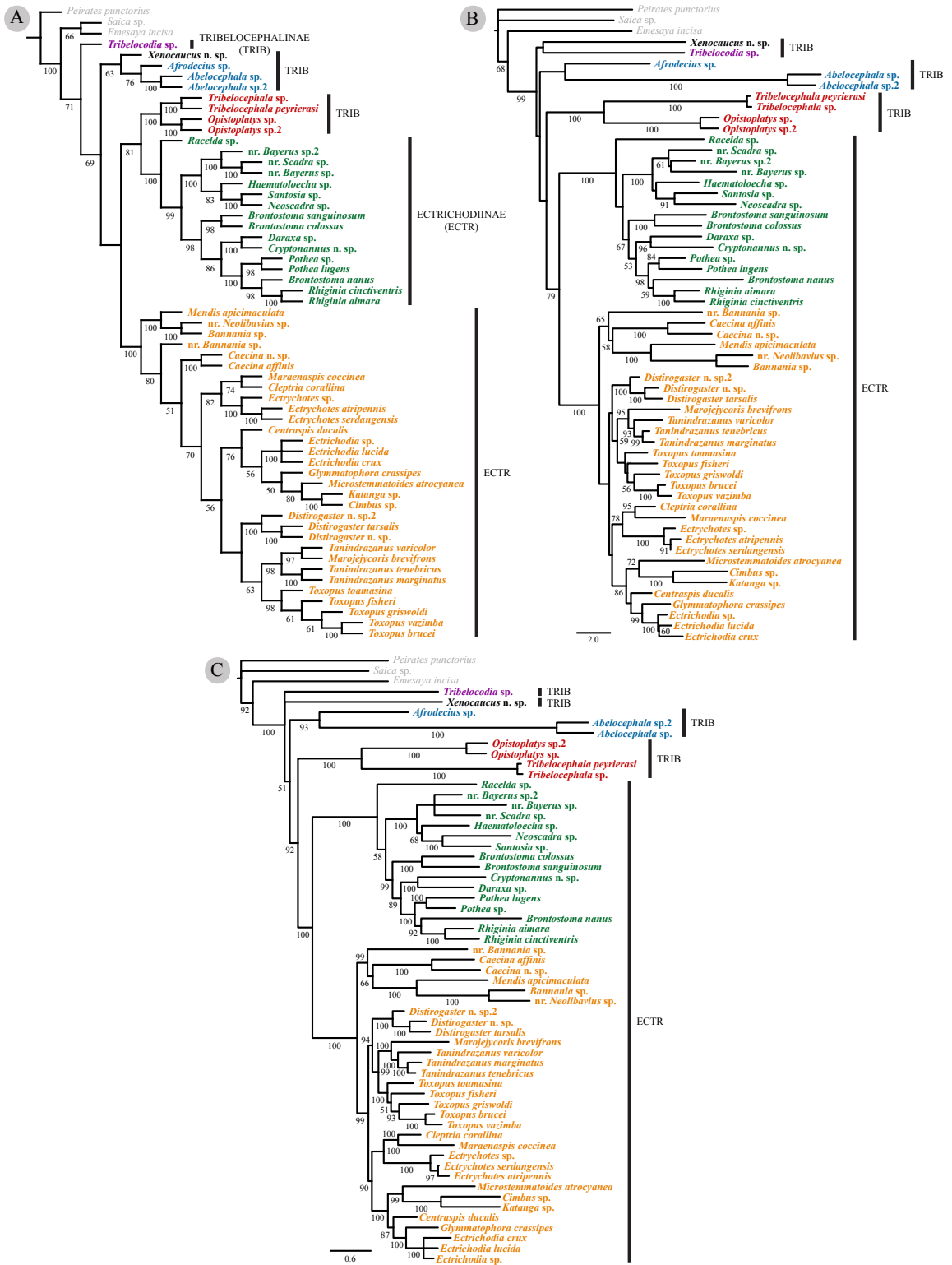
TTATTGGTTTTAGGAATATGAGGTAATGATCAATATGGACAGTCGGGGGCATTTCGTA  
TTGCGACGTTAGAGGTGAAATCCTTGGATCGTCGCAAGACGCACTAAAGCGAAAGCA  
TTTGCCAAGTATGTCTTAATTGATCAAGAACGAAAGTTAGAGGTTCTGAAGGCGATCA  
GATACCGCCCTAGTTCTAACCATAAACGATGCCAGCCAGCGATCCGCCGATGTTCCCT  
TTAATGACTCGGGCGGGAGCTTCC-  
ACCCGGGAAACCAAAGCTTTTGGGTTCGGGGGAAGTATGGTTGCAAAGCTGAAACT  
TAAAGGAATTGACGGAAGGGCACACCAGGAGTGGAGCCTGCGGCTTAATTTGACT  
CAACACGGGAAACCTCACCAGGCCAGGACATTGTAATGATTGACAGATTGATAGCTC  
TTTCTT??  
AGGCTAAATACAACCACGAGACCGATAGTAAACAAGTACCGTGAGGGGAAAGTTGAA  
AAGAAGTTGAAGAGAGAGTTCATAAGTACGTGAAACCGCTCAGGGGTAAACGGAG  
AAGACCTGAA--TATCCGAAAGGGGAGATTACGTCTTACCAT---  
ATGTTCAACTATAAAAATGT-----TTGAGATGGCCA-TTC-GCCCTATCCAA---TGCAA--  
ATTGTG--ATAG-AGGGTTTTGTTACTTCGCATTTTTTTT---  
TAGTAAGAACAGTGGTGGACCGCACTTCTCCCTTAGT-  
AGGACGTTGCGACCTGTCAAAT--AAT----ACTCTAAGT--AA-A----TGTTAA---  
AGTFACTGTAT-T-TATAGG-----  
TTTTTTTAAATTGTAAAAGATTTAAATGCCATTTTTTACAGC-TTTAA--CAATAAA----  
ATT---ACTGGATATT-----TGTATTGACAGTATAGACTGATGT-  
TGTTGAGCCACTTATTATAA-----ACATTTAATAAGTGTC-----  
TTTGGCC-GTTTTGCAA-GTAGGTAGGT-----ATTTTTAGTT-A-ATGTTGGGCTT-TT-----  
GTCCGCTAA-CTA-ATCCTCTTA-CTGTTGGTA----AAC-TG--TG-TCTCT--GACAG--  
TGCTTA??  
???GACCAAGGAGTTTATTGTGTACGCAAGTCATTGGATAAT--  
TAAATCCAAAGGCGAAATGAAAGTAAAGAGTTTTTTTAAGTTTTTAATA-----  
-----ATTTTATTAATCTC-----TTGGA-AAAACTCAAGGGAAGATGG--  
ACA-----CTAATTTAATAAT-----TGATAATA-  
TTTCTTTAATATTATTG-----GTT-ATTTAATATAGG-  
TTCCCGCATTCCAGGGCGCTTG-TTTCTCGTTT--TA-  
CGAGTCTATGCGCACCTAGAGCGTACACGATGATACCCGAAAGATGGTGAAGTACTATGC  
CTGGCCAGGACGAAGCCAGGGGAAACCCTGGTGGAGGTCGCGAGCGATTCTGACGT  
GAAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCGAACCATCTAGT  
AGCTGGTTCCTCTGAAGTTCCCTCAGGATAGCTTGCACTCAAT-----T---  
GAGACGAGTCTCATCAGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-  
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAT-  
CCCCGGCTTTCTTAAAA-----  
TTGAAGCCGTGGGATTCCGATCAGAGTGCCAAGTGGGCAATTTTGGTAAGCAGAA--  
---  
ACTGGCG??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??  
??



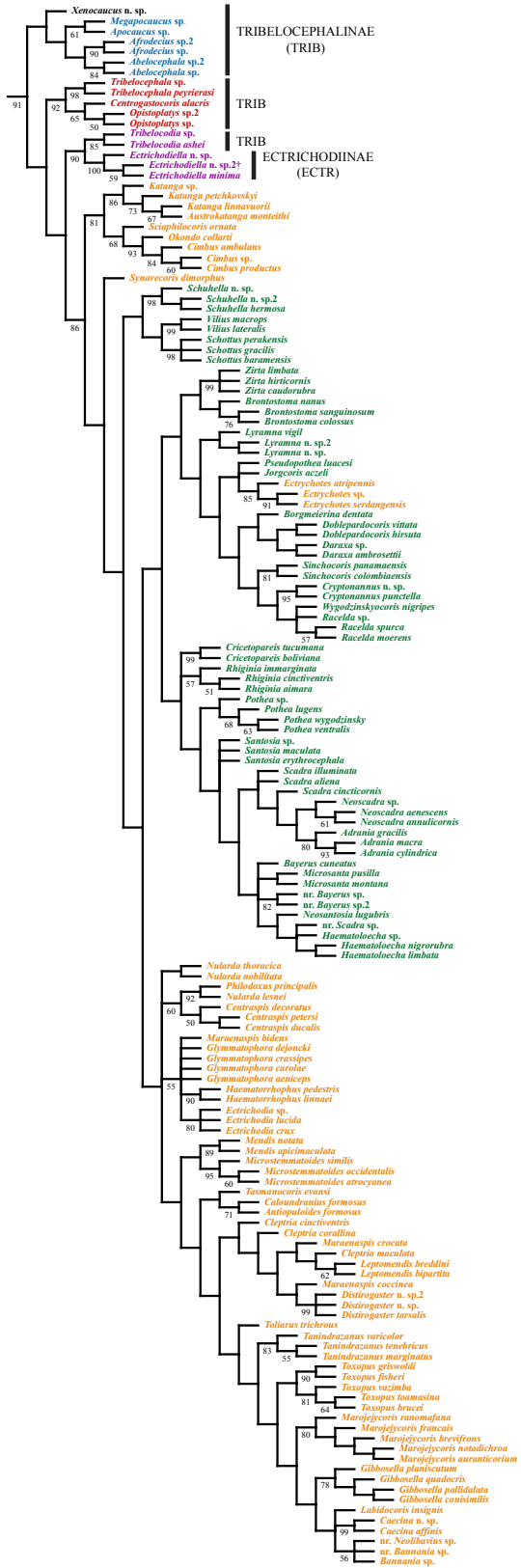


```
TGCAAATCGATCGTCAGAGCTGGGTATAGGGGCGAAAGACCAATCAAACCGTCTAG
TAGCTGGTTCTTCTGAAGTTTCCCTCAGGATAGCTTGCCTCGGT-----T---
GAGAAGAGTCTCATCCGGTAGAGCGAATGATTAGAGGCATTGGGGTCG-
AAACGACCTCAACCTATTCTCAAACCTCAAATAGATGAGAG-
CCCCGGCTTTCTTAAAA-----
TTGAAGCCGTGGGAATAGGATCAGAGTGCCAAGTGGGCAATTTTGGTAAGCAGAA
CTGGC-----
G????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
????????????????????????????????????????????????????????????
;
END;
BEGIN SETS;
    CHARSET 16S = 1-599;
    CHARSET 18S = 600-1624;
    CHARSET 28S_D2 = 1625-2475;
    CHARSET 28S_D3_D5 = 2476-3243;
    CHARSET COI = 3244-3714;
    CHARSET Wg = 3715-4055;
END;
```

Supplementary material 4.5. Molecular phylogenetic analyses of 59 taxa. Terminal taxon names are colored by clades recovered in the combined morphological and molecular ML best tree in Fig. 4.2 for comparison. A. Strict consensus tree from two parsimonious trees (length = 9,181 steps; RI = 0.484; CI = 0.306). B. ML best tree (final  $L_n = -38,481.3284$ ) based on three partitions and models of sequence evolution: 16S, 28S D2, 28S, D3–D5 (GTR+ $\Gamma$ +I); 18S, COI codon positions 1 and 2, Wg (GTR+ $\Gamma$ +I); COI codon position 3 (HKY+ $\Gamma$ +I). C. Bayesian MCMC majority-rule consensus tree based on the same three partitions and models of sequence evolution. Bootstrap values  $\geq 50$  are reported in A. and B. Posterior probabilities  $\geq 50$  are reported in C.

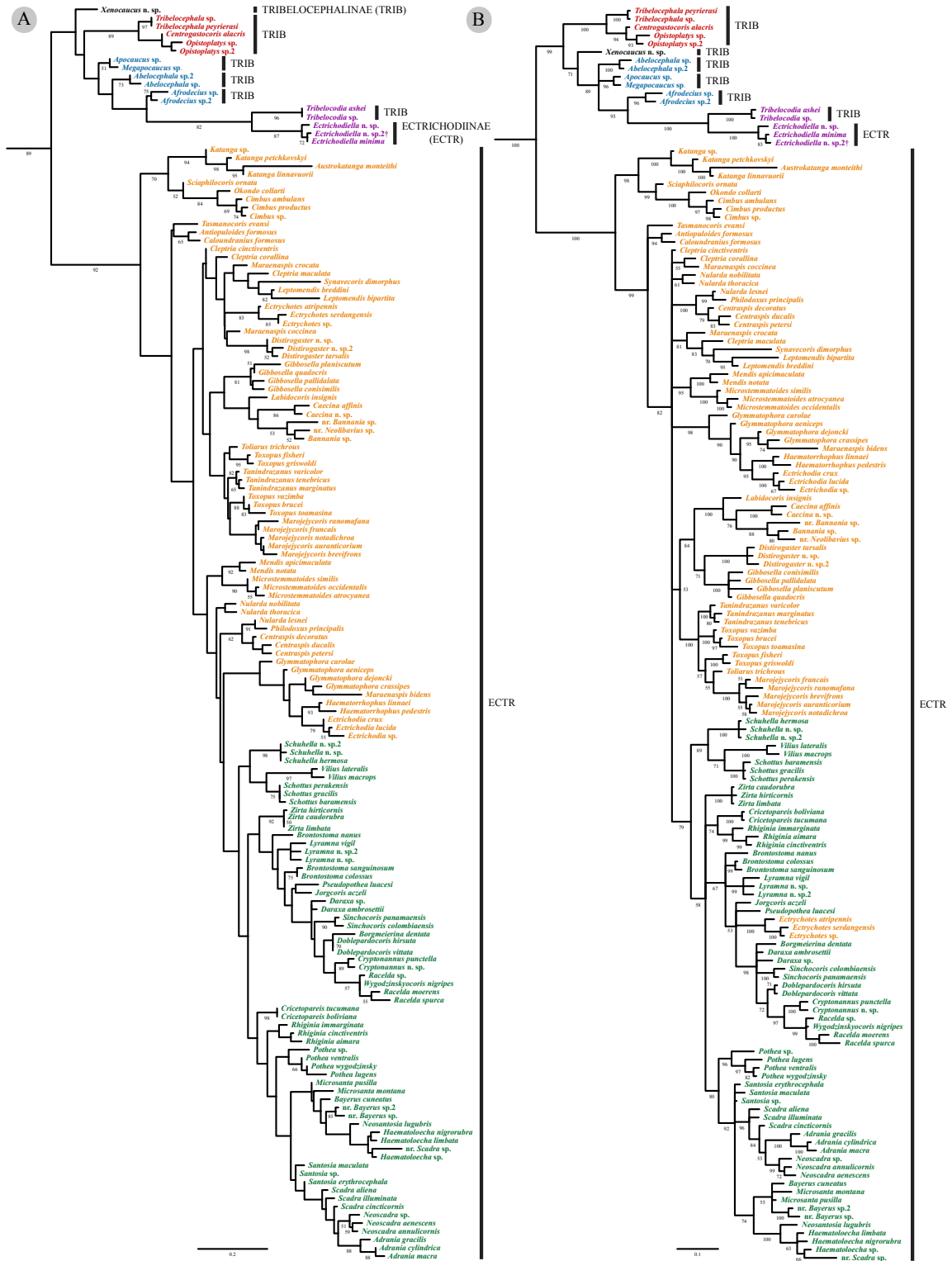


Supplementary material 4.6. Strict consensus tree from 269 parsimonious trees (length = 1,203 steps; RI = 0.705; CI = 0.174) based on TNT analysis of 156 taxa (outgroups removed from figure) and morphological data. Terminal taxon names are colored by clades recovered in the combined morphological and molecular ML best tree in Fig. 4.2 for comparison. Bootstrap values  $\geq 50$  are reported.



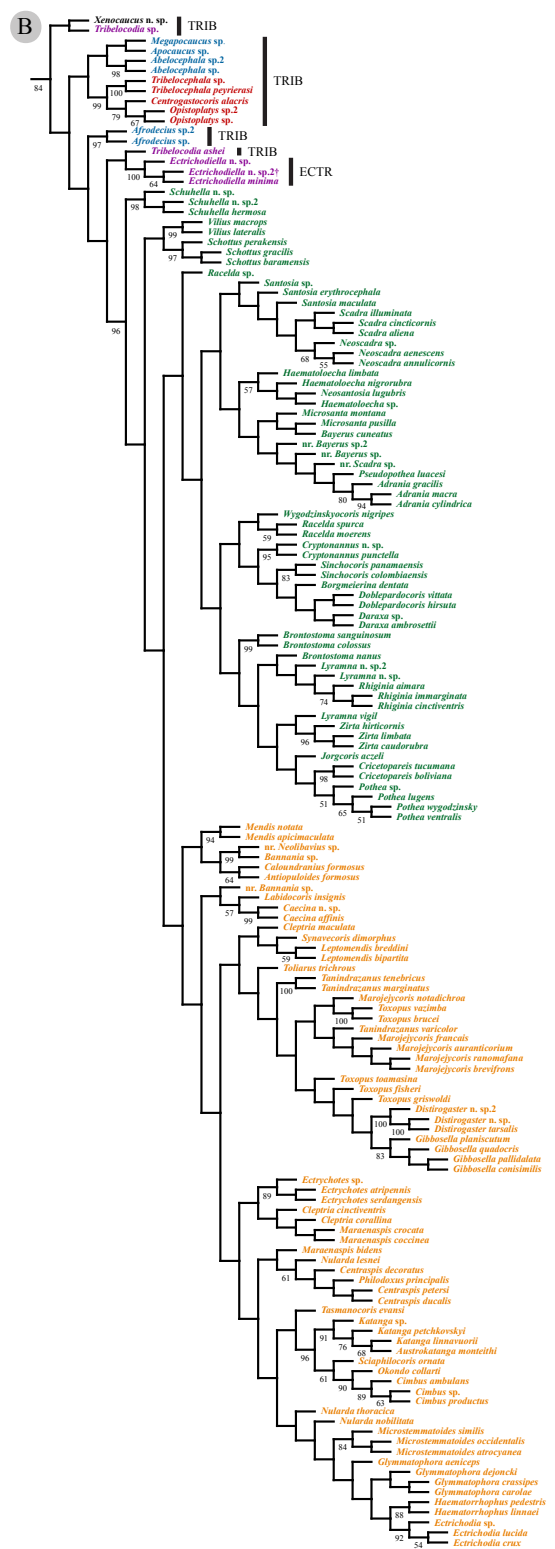
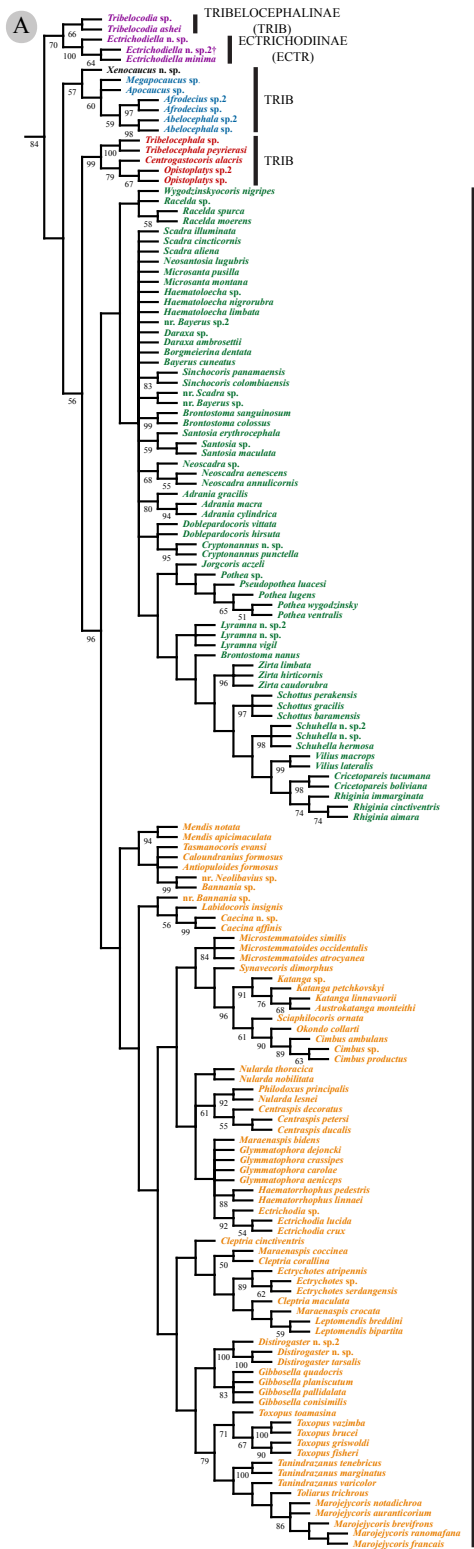
ECTR

Supplementary Material 4.7. Morphological phylogenetic analyses of 156 taxa (outgroups removed from figure). Terminal taxon names are colored by clades recovered in the combined morphological and molecular ML best tree in Fig. 4.2 for comparison. A. ML best tree (final Ln = -5,215.5335) based on Lewis' (2001) Mkv model of morphological evolution. B. Bayesian MCMC majority-rule consensus tree based on the same models of morphological evolution. Bootstrap values  $\geq 50$  are reported in A. Posterior probabilities  $\geq 50$  are reported in B.

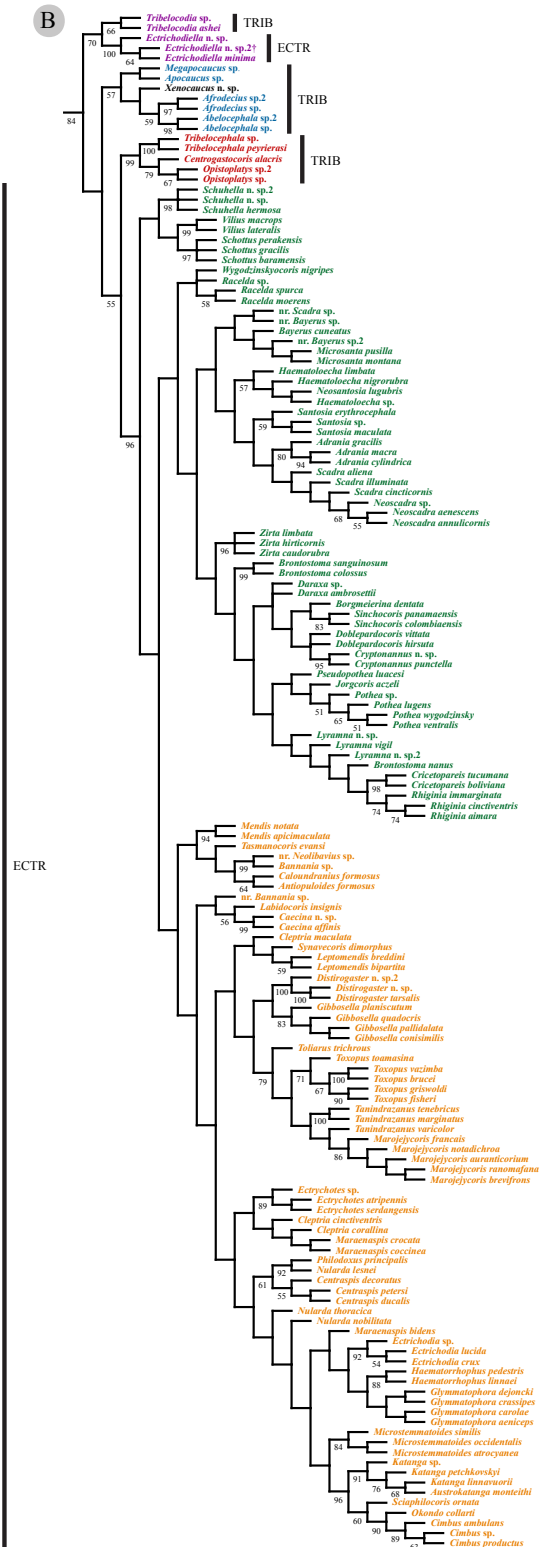
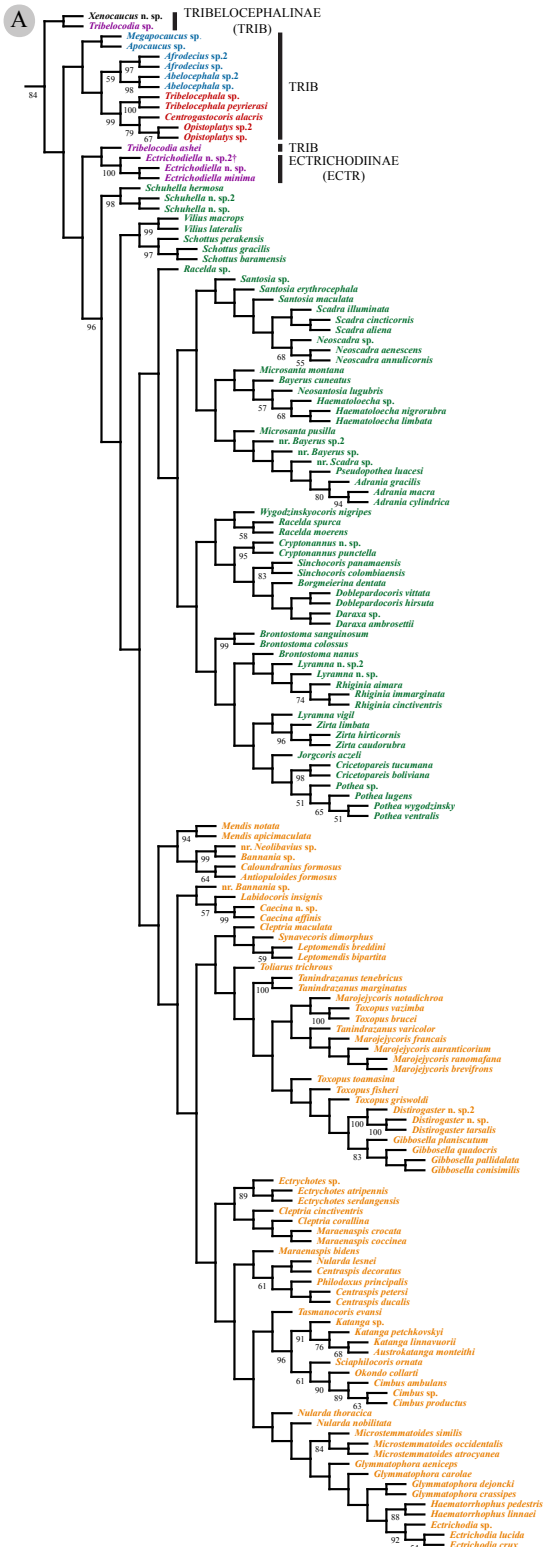




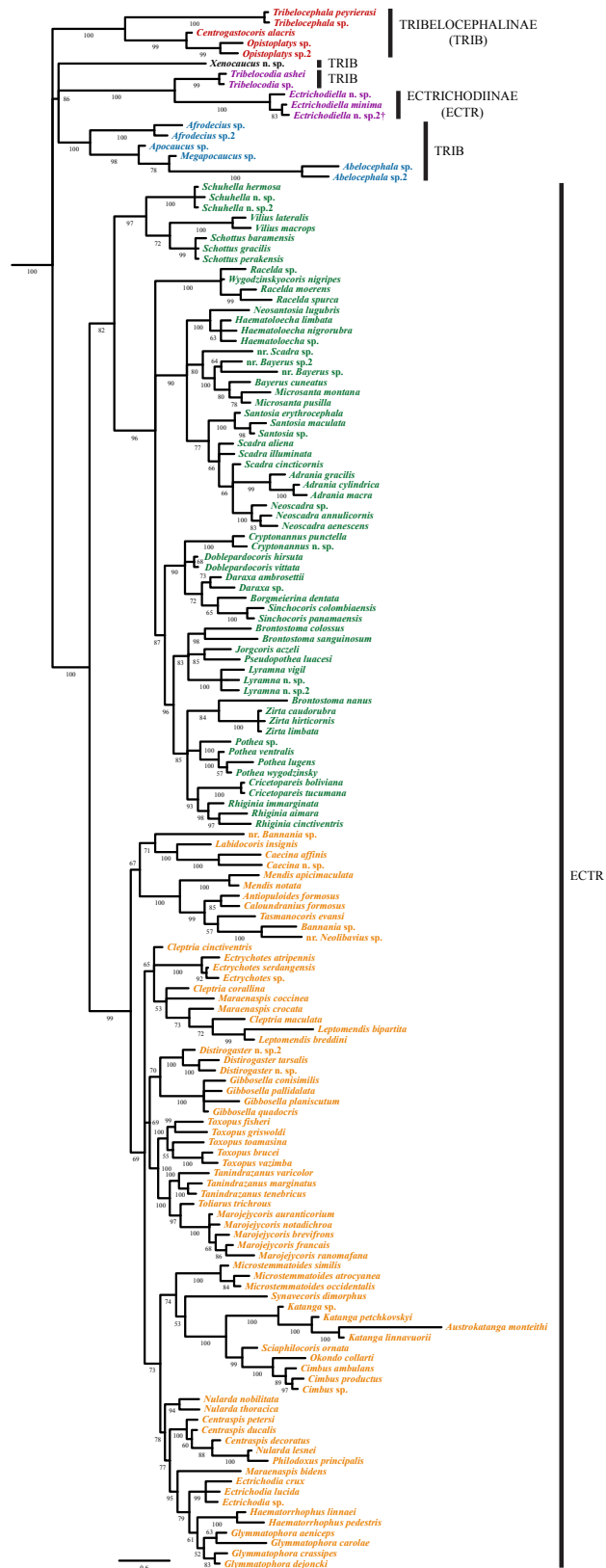
Supplementary material 4.8. Combined morphological parsimony phylogenetic analyses of 156 taxa (outgroups removed from figure). Terminal taxon names are colored by clades recovered in the combined morphological and molecular ML best tree in Fig. 4.2 for comparison. Bootstrap values  $\geq 50$  are reported. A. Strict consensus tree from 664 parsimonious trees (length = 10,458 steps; RI = 0.527; CI = 0.288) based on an equal weights TNT analysis. B. Single most parsimonious tree (length = 753.08685 steps; RI = 0.519; CI = 0.285) based on an implied weights ( $k = 3.000000$ ) TNT analysis.



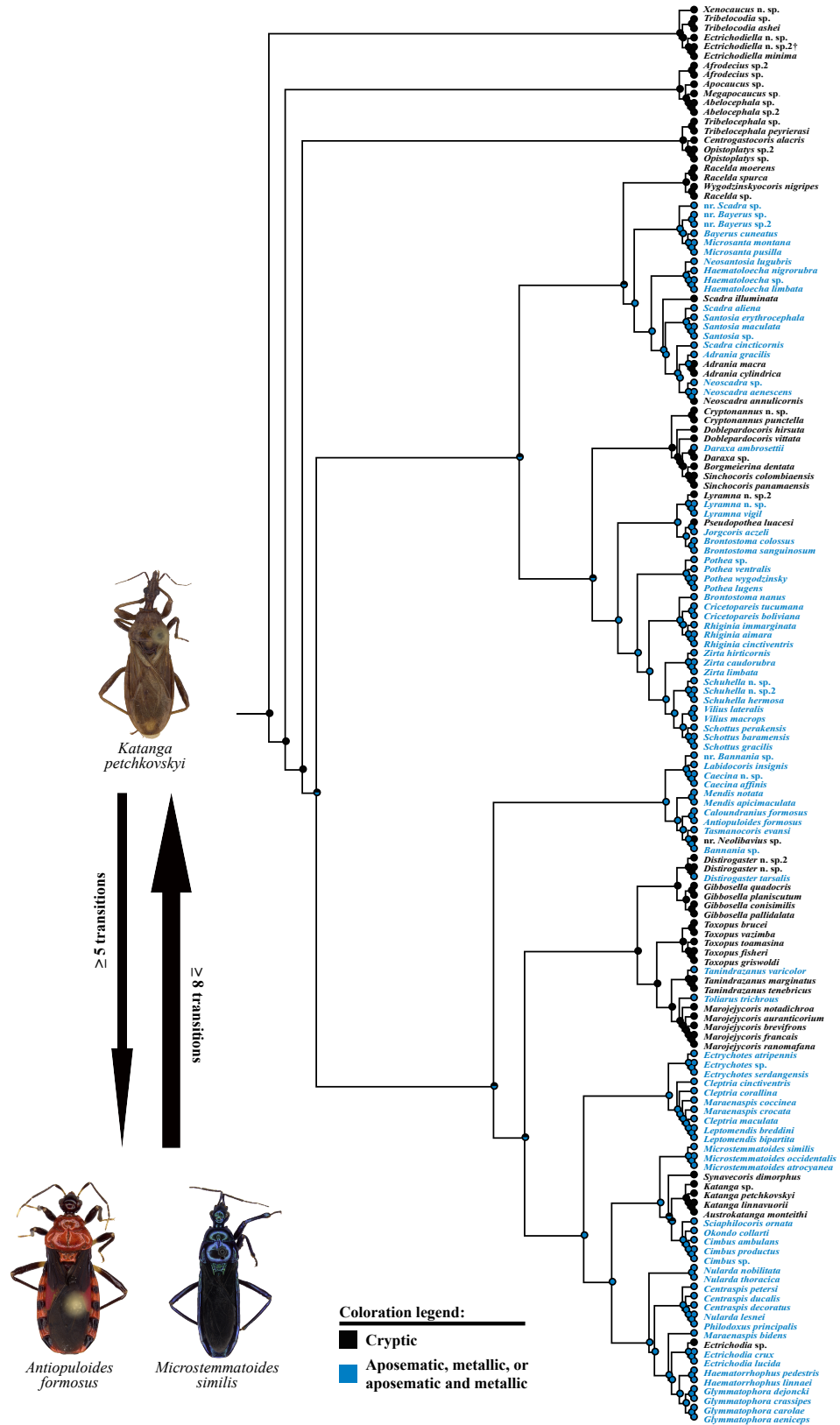
Supplementary material 4.9. Combined morphological parsimony phylogenetic analyses of 156 taxa (outgroups removed from figure). Terminal taxon names are colored by clades recovered in the combined morphological and molecular ML best tree in Fig. 4.2 for comparison. Bootstrap values  $\geq 50$  are reported. A. Single most parsimonious tree (length = 652.35368 steps; RI = 0.520; CI = 0.285) based on an implied weights ( $k = 4.266835$ ) TNT analysis. B. Strict consensus tree from three parsimonious trees (length = 274.39610 steps; RI = 0.526; CI = 0.288) based on an implied weights ( $k = 18.980060$ ) TNT analysis.



Supplementary Material 4.10. Bayesian MCMC majority-rule consensus tree based on of 156 taxa (outgroups removed from figure) and a combined morphological and molecular dataset. Molecular data were divided into three partitions and models of sequence evolution: 16S, 28S D2, 28S, D3–D5 (GTR+ $\Gamma$ +I); 18S, COI codon positions 1 and 2, Wg (GTR+ $\Gamma$ +I); COI codon position 3 (HKY+ $\Gamma$ +I). Morphological data were analyzed under Lewis' (2001) Mkv model of morphological evolution. Terminal taxon names are colored by clades recovered in the combined morphological and molecular ML best tree in Fig. 4.2 for comparison. Posterior probabilities  $\geq 50$  are reported.



Supplementary material 4.11. Parsimony ancestral state reconstruction on the ML phylogram (Fig. 4.2) for Tribelocephalinae and Ectrichodiinae taxa (tree converted to cladogram for visual) and color character 198. Terminal taxa coded as cryptic (black) or either aposematic, metallic, or aposematic and metallic (blue).



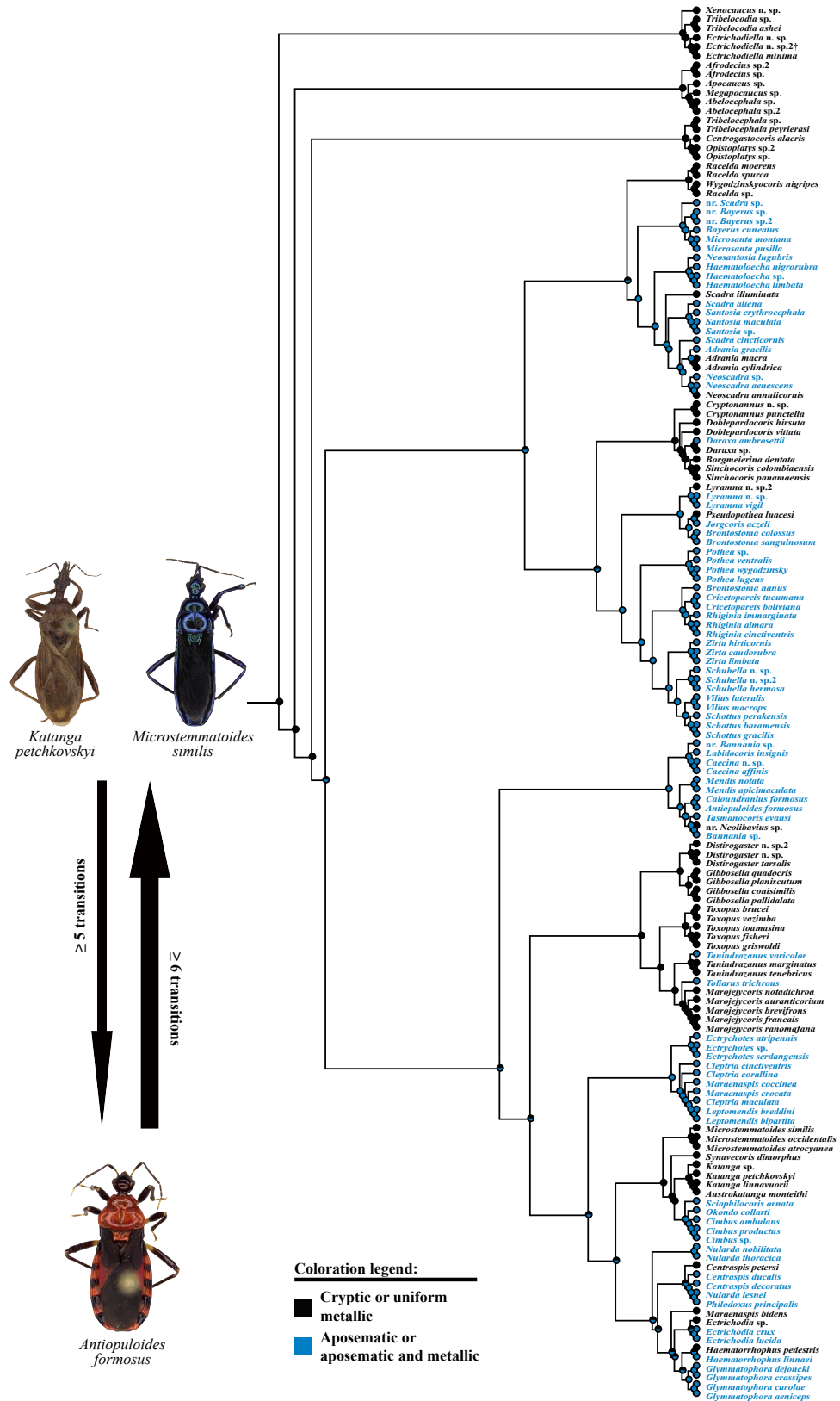


Supplementary material 4.12. Figshare data repository link to BayesTraits MultiState ML and MCMC ancestral state reconstruction output for color character 198 on ML phylogram and ultrametric trees. The gray panel lists internal node IDs and subtending terminal taxa. The blue panel are results from ML ancestral state reconstruction. The yellow panel are results from MCMC ancestral state reconstruction. Abbreviations: Ultra 0, ultrametric tree generated with penalty set to 0; Ultra 10, ultrametric tree generated with penalty set to 10; Ultra 10000, ultrametric tree generated with penalty set to 10000.

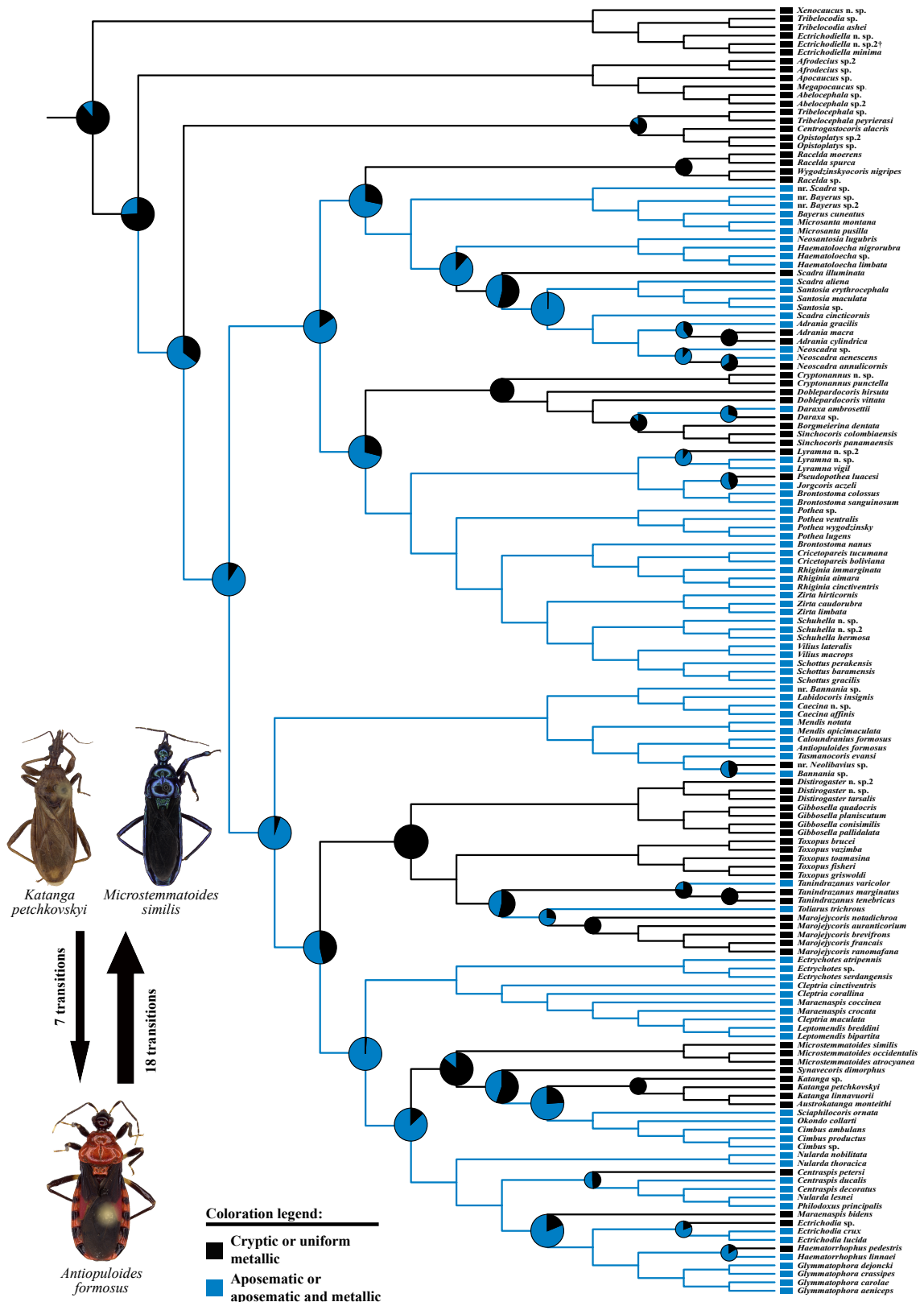
Figshare data repository link: <https://figshare.com/s/f0d20be1ac405a64a727>

Figshare doi: 10.6084/m9.figshare.2908249

Supplementary material 4.13. Parsimony ancestral state reconstruction on the ML phylogram (Fig. 4.2) for Tribelocephalinae and Ectrichodiinae taxa (tree converted to cladogram for visual) and color character 199. Terminal taxa coded as cryptic (black) or either aposematic, metallic, or aposematic and metallic (blue).



Supplementary material 4.14. BayesTraits MultiState ML ancestral state reconstruction of color character 199 on the ML phylogram (Fig. 4.2) for Tribelocephalinae and Ectrichodiinae taxa (tree converted to cladogram for visual). Terminal taxa coded as cryptic or uniform metallic (black) or aposematic or aposematic and metallic (blue). Pie charts at select nodes show probability values from the ML reconstructions; branches are colored to reflect the highest probability of a color state at that branch.



Supplementary material 4.15. Figshare data repository link to BayesTraits MultiState ML and MCMC ancestral state reconstruction output for color character 199 on ML phylogram and ultrametric trees. The gray panel lists internal node IDs and subtending terminal taxa. The blue panel are results from ML ancestral state reconstruction. The yellow panel are results from MCMC ancestral state reconstruction. Abbreviations: Ultra 0, ultrametric tree generated with penalty set to 0; Ultra 10, ultrametric tree generated with penalty set to 10; Ultra 10000, ultrametric tree generated with penalty set to 10000.

Figshare data repository link: <https://figshare.com/s/f81b72ae4f384d8daace>

Figshare doi: 10.6084/m9.figshare.2909611

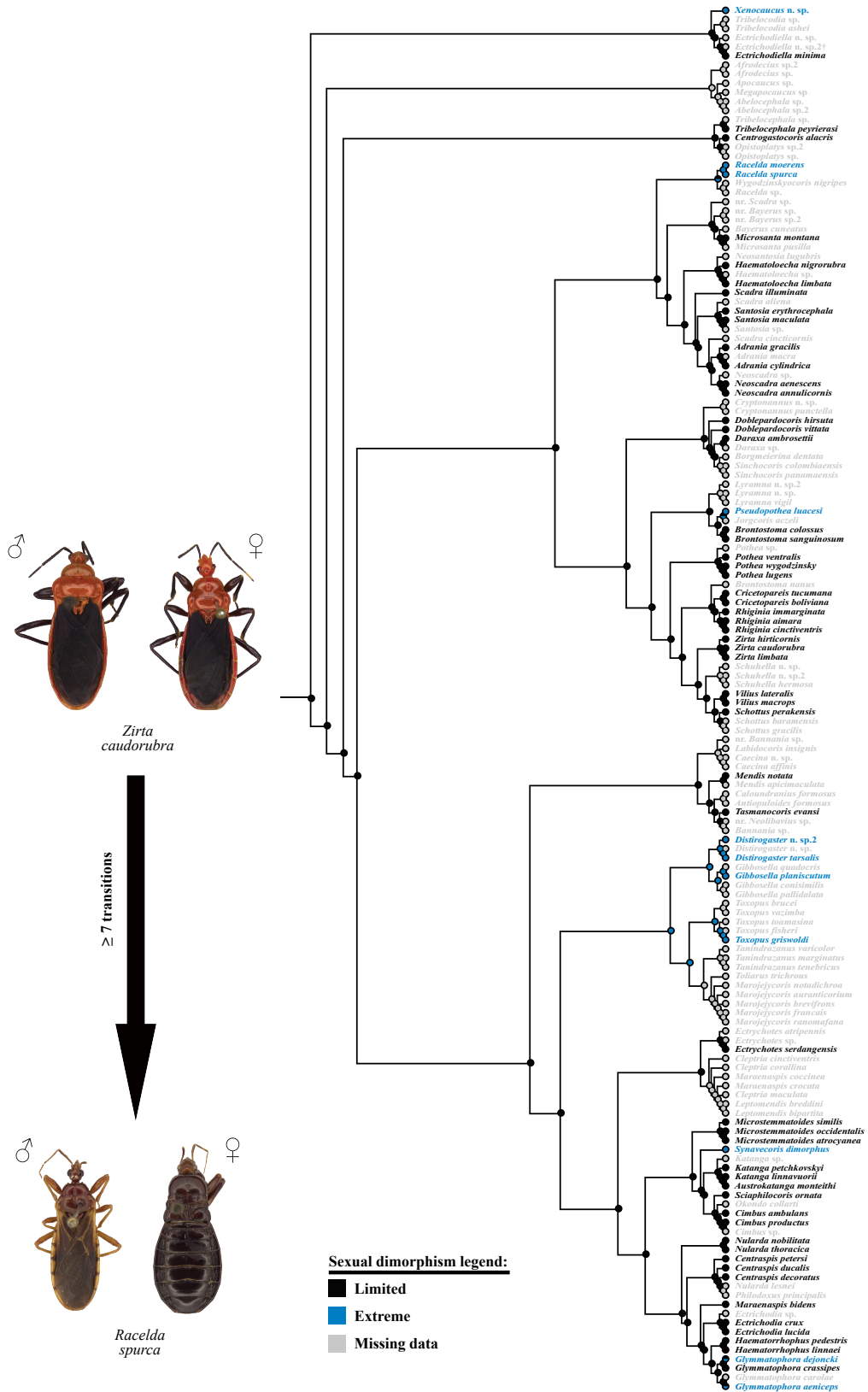
Supplementary material 4.16. Phylogenetic logistic regression parameter estimates for the effects of log-transformed male body length on coloration in 133 Ectrichodiinae and Tribelocephalinae species. Cryptic and uniform metallic coloration is treated as one state with aposematic and aposematic and metallic color patterns as a separate state. Parameters are phylogenetic signal ( $a$ ), intercept ( $b_0$ ), and the regression coefficient ( $b_1$ ), and p-values  $\leq 0.05$  are indicated in bold. Abbreviations: Phylo, phylogram; Ultra0, ultrametric tree generated with penalty set to 0; Ultra10, ultrametric tree generated with penalty set to 10; Ultra10000, ultrametric tree generated with penalty set to 10000; Par, parameter; SE, standard error; BS, bootstrap; C, converged; NC, non-converged.

Tree/dataset	Par	Estimate	SE	p-value	BS mean (C and NC) <sup>a</sup>	BS confidence interval (C and NC) <sup>a</sup>	BS p-value (C and NC) <sup>a</sup>	BS mean (C only)	BS confidence interval (C only)
Phylo C+UM									
	$a$	-1.0391			-1.4282	(-2.6531, -0.4579)	<b>0.0052</b>	-1.3942	(-2.6406, -0.4433)
	$b_0$	-0.0466	0.4660	0.9205	-0.0419	(-2.4529, 2.6352)	0.8712	-0.0340	(-2.2389, 2.5531)
	$b_1$	0.6073	0.2262	<b>0.0082</b>	0.3744	(-2.5873, 2.8295)	0.4488	0.3830	(-2.4233, 2.5408)
Ultra0 C+UM									
	$a$	-0.6724			-0.9606	(-3.3316, 0.4277)	<b>0.0184</b>	—	—
	$b_0$	-0.3662	0.5515	0.5078	-0.3904	(-1.6487, 0.9075)	0.5128	—	—
	$b_1$	0.6804	0.2264	<b>0.0032</b>	0.7669	(0.2764, 1.4656)	<b>0.0016</b>	—	—
Ultra10 C+UM									
	$a$	-0.0837			-0.0510	(-3.0062, 0.9890)	<b>0.0192</b>	-0.5191	(-2.9789, 0.9509)
	$b_0$	-0.2572	0.6808	0.7063	-0.2964	(-1.6978, 1.2196)	0.7000	-0.2884	(-1.6878, 1.2196)
	$b_1$	0.6209	0.2245	<b>0.0065</b>	0.7305	(0.2352, 1.3738)	<b>0.0064</b>	0.7300	(0.2355, 1.3701)
Ultra10000 C+UM									
	$a$	-0.0832			-0.4968	(-2.8611, 0.9529)	<b>0.0164</b>	-0.4983	(-2.8414, 0.9391)
	$b_0$	-0.2572	0.6908	0.7063	-0.2817	(-1.6744, 1.2164)	0.6944	-0.2815	(-1.6710, 1.2164)
	$b_1$	0.6208	0.2245	<b>0.0065</b>	0.7266	(0.2371, 1.3525)	<b>0.0048</b>	0.7268	(0.2380, 1.3525)

<sup>a</sup>Parametric bootstrapping was performed to obtain means and confidence intervals, as well as test the null hypotheses that there is no phylogenetic signal in the residuals, ( $H_0: a = -4$ , 1-tailed test) and that the regression coefficient equals 0 ( $H_0: b_1 = 0$ , 2-tailed test).

Supplementary material 4.17. Parsimony ancestral sexual dimorphism reconstruction on the ML phylogram (Fig. 4.2) for Tribelocephalinae and Ectrichodiinae taxa (tree converted to cladogram for visual). Sexual dimorphism for terminal taxa coded as limited (black) or extreme (blue). Missing data (gray) for terminals is indicated.





Supplementary material 4.18. Figshare data repository link to BayesTraits MultiState ML and MCMC ancestral state reconstruction output for sexual dimorphism character 200 on ML phylogram and ultrametric trees. The gray panel lists internal node IDs and subtending terminal taxa. The blue panel are results from ML ancestral state reconstruction. The yellow panel are results from MCMC ancestral state reconstruction. Abbreviations: Ultra 0, ultrametric tree generated with penalty set to 0; Ultra 10, ultrametric tree generated with penalty set to 10; Ultra 10000, ultrametric tree generated with penalty set to 10000.

Figshare data repository link: <https://figshare.com/s/433681e7784d456e9a8d>

Figshare doi: 10.6084/m9.figshare.2912692

## Conclusion

The research conducted in this dissertation has significantly advanced our understanding of Ectrichodiinae biodiversity, natural history, and evolution, as well as Reduviidae phylogeny and classification. Prior to this work, there was a lack of thorough taxonomic treatments and phylogenetic hypotheses for the subfamily. No tribal-level classification existed, and relationships between genera had not been cladistically investigated, with the exception of one genus-level phylogeny of the New World fauna. Faunistic checklists and descriptions of single taxa based on few specimens of a single sex have dominated the Ectrichodiine literature.

In Chapter 1, Ectrichodiinae predatory behaviors were reviewed and supplemented with observations on *Rhiginia cinctiventris* (Stål), 1872, in Costa Rica. The predator-prey association between Ectrichodiinae and millipedes was first documented almost a century ago. However, feeding behaviors have since received little attention, rendering this intriguing phenomenon largely inaccessible. About 2% of Ectrichodiinae species have been documented to prey on millipedes, leaving ample room to further test the claim that Ectrichodiinae, as a group, specialize on millipedes. Ectrichodiinae appear to have a preference for benzoquinone-producing juliform millipedes (Spirostreptida and Spirobolida), with Polydesmida rarely attacked. Furthermore, Ectrichodiinae insert maxillary and mandibular stylets at the millipede's intersegmental membranes on the ventral and ventrolateral trunk area or between the head and collum. Whether these areas are attacked in relation to the location of the millipede nervous system or defense glands remains to be investigated. Lastly, immature and adult Ectrichodiinae exhibit both solitary and communal predation. The majority of observations suggest that immatures typically exhibit communal predation, while adults usually engage in solitary predation.

The second chapter highlights our incomplete knowledge of the Madagascan invertebrate fauna, especially for the Ectrichodiinae. However, this situation is improving as a result of large

taxonomic surveys, such as the one that enabled this taxonomic project. Prior to the dissertation, only 10 described Madagascan Ectrichodiinae species were known. Here, 63 new species and three new genera of Ectrichodiinae that are endemic to the island are described. *Toxopus* Bergroth, 1905, *Gibbosella* Chłond, 2010, *Gibbosella elongata* Chłond, 2010, *Glymmatophora crassipes* Horváth, 1914, and *Maraenaspis bidens* (Reuter), 1887, are redescribed, with the first description of males for each taxon. *Cleptria signoretii* Reuter, 1887, is transferred to *Toxopus*.

Undoubtedly, the ~80 my isolation of Madagascar from other landmasses, along with geologic and climatic factors within Madagascar and throughout the Indian Ocean, has influenced the evolution of the island's biota. However, few systematic analyses incorporating modern divergence dating and biogeographic analyses have focused on Madagascan insects. The drastic increase of the described diversity in Chapter 2 presented an opportunity to investigate the timing and origin of Madagascan Ectrichodiinae using phylogenetic, divergence dating, and biogeographic analyses in Chapter 3. Based on the phylogenetic results, Madagascan Ectrichodiinae do not form a monophyletic group, with different clades closely related to Afrotropical and Oriental lineages. Ectrichodiinae are shown to have colonized the island within the last ~68 my via transoceanic dispersal at least twice from the Oriental region and once from the Afrotropical region. Some biogeographic models also infer a single out-of-Madagascar dispersal event to the Afrotropical region. Dispersal from the Oriental region to Madagascar may have been facilitated by the Seychelles-Mascarene Plateau and volcanic islands in the Indian Ocean. Factors facilitating dispersal from the Afrotropical region to Madagascar are more difficult to determine, but aerial dispersal via stepping-stone islands in the Mozambique Channel is the most probable hypothesis when accounting for ancient oceanographic reconstructions.

In the final chapter, the the evolution of sexual dimorphism, aposematic coloration, and the relationship between coloration and body size in Ectrichodiinae is investigated. Investigating

aposematic coloration in a phylogenetic context has been limited compared to the numerous published microevolutionary studies in animals. Few phylogenetic comparative studies in vertebrates and Lepidoptera have shown a positive correlation of body size and aposematic coloration. However, one lepidopteran study found no correlation between these two traits, a result that may be confounded by larval gregariousness. Adult Ectrichodiinae are typically solitary and afford an opportunity to test for a coloration-body size relationship. The evolution of extreme sexual dimorphism is also explored in a phylogenetic context. Aposematic coloration is shown to have evolved from a cryptic ancestor early in the evolutionary history of Ectrichodiinae+ Tribelocephalinae, with subsequent reversals and regains. A positive association between aposematic coloration and body length is recovered when phylogenetic logistic regressions are conducted on ultrametric trees but not on phylograms, which may be an artifact of the method. Extreme sexual dimorphism is shown to evolve multiple times from a more limited condition. Thus, careful consideration must be given to the use of female morphology in phylogenetic investigations and taxonomic decisions. Lastly, a new classification based on phylogenetic results is proposed for Ectrichodiinae.