Lawrence Berkeley National Laboratory

Lawrence Berkeley National Laboratory

Title JGI Fungal Genomics Program

Permalink https://escholarship.org/uc/item/02q3q97r

Author Grigoriev, Igor V.

Publication Date 2011-08-10

JGI Fungal Genomics Program

Igor V. Grigoriev¹

¹Lawrence Berkeley National Laboratory

August 2011

The work conducted by the U.S. Department of Energy Joint Genome Institute is supported by the Office of Science of the U.S. Department of Energy under Contract No. DE-AC02-05CH11231

DISCLAIMER

This document was prepared as an account of work sponsored by the United States Government. While this document is believed to contain correct information, neither the United States Government nor any agency thereof, nor The Regents of the University of California, nor any of their employees, makes any warranty, express or implied, or assumes any legal responsibility for the accuracy, completeness, or usefulness of any information, apparatus, product, or process disclosed, or represents that its use would not infringe privately owned rights. Reference herein to any specific commercial product, process, or service by its trade name, trademark, manufacturer, or otherwise, does not necessarily constitute or imply its endorsement, recommendation, or favoring by the United States Government or any agency thereof, or The Regents of the University of California. The views and opinions of authors expressed herein do not necessarily state or reflect those of the United States Government or any agency thereof or The Regents of the University of California.



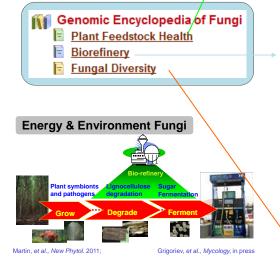
JGI Fungal Genomics Program

Contact: Igor Grigoriev <ivgrigoriev@lbl.gov>

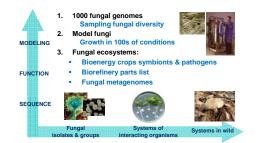


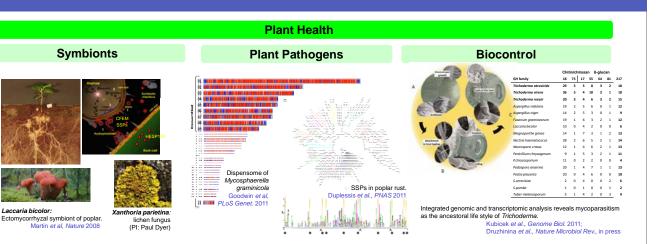
Abstract

Genomes of energy and environment fungi are in focus of the Fungal Genomic Program at the US Department of Energy Joint Genome Institute (JGI). Its key project, the Genomics Encyclopedia of Fungi, targets fungi related to plant health (symbionts, pathogens, and biocontrol agents) and biorefinery processes (cellulose degradation, sugar fermentation, industrial hosts), and explores fungal diversity by means of genome sequencing and analysis. Over 60 fungal genomes have been sequenced by JGI to date and released through MycoCosm (www.jgi.doe.gov/fungi), a fungal webportal, which integrates sequence and functional data with genome analysis tools for user community. Sequence analysis supported by functional genomics leads to developing parts list for complex systems ranging from ecosystems of biofuel crops to biorefineries. Recent examples of such 'parts' suggested by comparative genomics and functional analysis in these areas are presented here.



Future Grand Challenges





Biorefinery Lignocellulose Degradation **Xylose Fermenters** Thermophiles Orthologs Thermophilic biomass-degrading fungi Comparative genomics and Myceliophthora thermophila and Thielavia transcriptomics of x terrestris are the first with completely STREETSTEET and xylose growers revealed finished genomes, new candidate for cell candidate genes for factories with secreted thermostable S.cereviseae strain enzymes. CAZy and FOLy in Basidiomycetes improvement. Berka, Tsang et al., subm Tandem duplication Convergent evolution of brown rot and ECM from white rot ancestor. of peroxidases in Wohlbach, et al. Release of reducing sugars from alfalfa straw by extracellular enzymes PNAS 2011 Trametes. Eastwood et al., Science 2011 Riley et al .. in prep

Fungal Diversity

MycoCosm: 80+ fungal genomes

fungi **Genome-Centric View Comparative View** www.iai.doe.aov/funa ----OTHERDY 0

Genome-centric View

HAIT TOTAL OF SHEET AND CLAIMER AND AND AND

Supports genome analysis, functional genomics, user data deposition & curation

JGIS Whome Altycocosm Revolution Allog



Comparative View

Enables analysis of groups of related fungi

