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Biosynthetic pathway of aliphatic formates via a Baeyer–Villiger oxidation in mechanism present in astigmatid mites

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Astigmatid mites depend on bioactive glandular secretions, pheromones, and defensive agents to mediate intra- and interspecies interactions. Aliphatic formates, such as (Z,Z)-8,11-heptadecadienyl formate (8,11-F17) and (Z)-8-heptadecenyl formate (8-F17), are rarely encountered natural products that are abundant in Sancassania sp. Sasagawa (Acari: Acaridae) mite secretions. Linoleic acid and oleic acid are predicted as key intermediates in the synthesis of the closely related aliphatic formates. To gain insight in this biosynthetic pathway, acarid mite feeding experiments were conducted using ¹³C-labeled precursors to precisely track incorporation. Analyses using ¹³C NMR spectroscopy demonstrated that the ¹³C-labeling pattern of the precursors was detectable on formates in exocrine secretions and likewise on fatty acids in total lipid pools. Curiously, the results demonstrated that the formates were biosynthesized without the dehomologation of corresponding fatty acids. Careful examination of the mass spectra from labeling experiments revealed that the carbonyl carbon of the formates is originally derived from the C-1 position of the fatty acids. Consistent with a Baeyer-Villiger oxidation reaction, labeling studies support the insertion of an oxygen atom between the carbonyl group and carbon chain. Empirical data support the existence of a Baeyer-Villiger monooxygenase responsible for the catalyzation of the Baeyer–Villiger oxidation. The predicted existence of a Baeyer–Villiger monooxygenase capable of converting aliphatic aldehydes to formates represents an exciting opportunity to expand the enzymatic toolbox available for controlled biochemical synthesis.

astigmatid mite | aliphatic formate | hydrocarbon | biosynthetic pathway | Baeyer–Villiger oxidation

Long-chain hydrocarbons in insects are key precursors to the construction of waterproof cuticles and further function in chemical communication, both within and between species (1). An array of hydrocarbons in astigmatid mites have also been characterized yet, unlike insects, comparatively short carbon chains ($<C_{20}$) predominate. One of the most widely distributed saturated hydrocarbons in the Astigmata is tridecane. With respect to unsaturated hydrocarbons, (*Z*)-8-heptadecane (8-C17) is widely distributed in Acaridae and Histiostomatidae, and (*Z*, *Z*)-6,9-heptadecadiene (6,9-C17) in Acaridae (2). As rarely detected chemicals in nature, 8-C17 serves as a sex pheromone in European house dust mites (*Dermatophagoides pteronyssinus*) (3), and similarly 6,9-C17 functions as an alarm pheromone in *Tortonia* sp., which attacks the nests of megachilid bees (*Osmia cornifons*) (4).

Linoleic acid [LA; (Z,Z)-9,12-octadecadienoic acid] has been experimentally proven to function as a precursor of 6,9-C17 in previous isotopic labeling studies with the mite (*Carpoglyphus lactis*) (5). *Carpoglyphus lactis* is able to synthesize LA de novo from a simple precursor (5, 6) and uses neral [(Z)-3,7-dimethyl-2,6-octadienal] as the alarm pheromone. Neral also reportedly functions as an alarm pheromone in several mite species and has been demonstrated to be a biosynthetic product of the mevalonate pathway in C. lactis (7). Mechanisms for lipid metabolism associated with mite secretory components have been gradually clarified (5-7). After being reported as an alarm pheromone of the mold mite (Tyrophagus putrescentiae) in 1975 (8), neryl formate [(Z)-3,7-dimethyl-2,6-octadienyl formate] has been subsequently identified as an alarm pheromone in several species of acaridae (2). Lardolure [(1R,3R,5R,7R)-1,3,5,7-tetramethyldecyl formate] has been identified as an aggregation pheromone in Lardoglyphus konoi (9, 10), and also shows induced activation in C. lactis, Aleuroglyphus ovatus, and T. putrescentiae (11). One of the lipidic components in mites, α -acaridial [2(E)-(4-methyl-3pentenyl)butenedial] has been demonstrated to function as a haptenic allergen that induced allergic contact dermatitis (12). The well-known house dust mites, D. pteronyssinus and Dermatophagoides farinare, are both major allergens, causing atopic diseases, such as asthma, rhinitis, and atopic dermatitis (13). Recent studies have shown that lipidic components, such as formate and hydrocarbon, function to attract the house dust mites and have the potential to be used for mite control (3, 13). Unlike more ubiquitous lipids, the biosynthetic pathways of chain-like formates present in mites have not been previously examined.

Significance

Aliphatic formates are commonly detected in astigmatid mite secretions and have been considered for roles in pheromone communication. Despite the natural occurrence of aliphatic formates, details surrounding the biosynthetic pathways remain unclear. To probe the biosynthetic mechanism, we performed ¹³C-incorporation experiments targeting two different chain-like aliphatic formates by feeding with suitable labeled precursors. Results support a mechanism whereby formates are generated via the Baeyer–Villiger oxidation of aldehydes as direct precursors. The enzyme that catalyzes this reaction has yet to be identified in animals, microbes, and plants. Results from the labeling studies inform approaches to identify the specific gene and encoded enzyme responsible for aliphatic formate biosynthesis, which will enable a facile dehomologation method in the field of organic synthesis.

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In our previous efforts, two formates, specifically (Z,Z)-8,11heptadecadienyl formate (8,11-F17) and (Z)-8-heptadecenyl formate (8-F17), were identified as the main secretion component of unidentified acarid mite (Sancassania sp. Sasagawa) (14). Based on the double-bond position, LA was hypothesized to be the biosynthetic precursor of 8,11-F17 and, similarly, oleic acid (OA) for 8-F17. If LA and OA are the biosynthetic precursors of the unsaturated hydrocarbons 6,9-C17 and 8-C17 in Sancassania sp. Sasagawa, new mechanistic insights involved in chain shortening and formate production are needed to understand the predictably unique steps involved. To provide a chemical framework for understanding the biosynthetic pathway of aliphatic formates, we performed experiments in which ¹³C-labeled precursors were administered to mites for the determination of ¹³C-labeling patterns of formate moiety using ¹³C NMR spectroscopy and gas chromatography-mass spectrometry (GC-MS). Results support an unusual biochemical mechanism in the generation of aliphatic formates, of which there are no previously reported cases. To the best of our knowledge, the enzyme that catalyzes this reaction is yet to be identified in animals, microbes, and plants.

Results

GC-MS Analysis Demonstrate Incorporation of ¹³C-Labeled Acetate into 8,11-F17 and 8-F17. Analyses of Sancassania sp. Sasagawa mite glandular secretions by GC-MS revealed two types of predominant hydrocarbons, specifically 6,9-C17 [retention time (RT) = 15.25 min and 8-C17 (RT = 15.32 min) and two types of formates, specifically 8,11-F17 (RT = 18.91 min), 8-F17 (RT = 18.95 min) as the main components (Fig. 1). Examination of the mass spectrum of 8,11-F17 revealed m/z 280 as a molecular ion (Fig. S1A). In contrast, the predicted molecular ion of 8-F17 was not clearly observable and instead the M⁺-HCOOH m/z 236 ion predominated (Fig. S1C). To gain insights into the origin of these molecules, Sancassania sp. Sasagawa mites were reared in an agar medium with dry yeast as the main ingredient in the presence of 10% [1-¹³C] acetate. To examine the pattern of ¹³C incorporation into formates 8,11-F17 and 8-F17, the mites secretions analyzed by GC-MS 7 d after the initiation cultivation with [1-¹³C] acetate. In the mass spectrum of 8,11-F17 derived from mites cultivated in unlabeled (control) media, the M⁺+3 ion $(m/z \ 283)$ was identified to be the highest molecular-weight isotopic parent ion detectable. However, when [1-13C] acetate was fed to the mites, the M⁺+10 ion (m/z 290) was identified to be the largest isotopic ion (Fig. S1B). Similarly, in the mass



Fig. 1. Mites (*Sancassania* sp. Sasagawa) contain abundant C17 unsaturated hydrocarbons and C17 aliphatic formates. Typical GC of extracts from *Sancassania* sp. Sasagawa. In order of GC retention time: (*Z*,*Z*)-6,9-heptadecadiene (6,9-C17), (*Z*)-8-heptadecene (8-C17), (*Z*,*Z*)-8,11-heptadecadienyl formate (8,11-F17), and (*Z*)-8-heptadecenyl formate (8-F17).

spectrum of 8-F17, the M⁺–HCOOH+2 ion (m/z 238) was identified to be the largest isotope ion in the control. However, when [1-¹³C] acetate was fed to the mites, the M⁺–HCOOH+8 ion (m/z 244) was identified to be the largest isotope ion (Fig. S1D). These results confirm that 1-¹³C acetate is incorporated into aliphatic formates, such as 8,11-F17 and 8-F17, 7 d after the administration experiment.

¹³C NMR Analysis of ¹³C-labeled LA and OA Yield Predictable Conserved Patterns of Chain Elongation. Mites cultivated in media including 10% [1-¹³C] acetate were separated from the culture media by suspension in saturated saline. The obtained mites of all developmental stages were immersed in hexane, and their total surface secretions were extracted. After mite bodies were isolated from hexane, total mite lipids were extracted using chloroform:methanol (2:1). After hydrolysis of the extracts by sodium methoxide, the products were treated with trimethylsilyldiazomethane to yield the corresponding fatty acid methyl esters. After isolation of LA methyl ester (LAME) and OA methyl ester (OAME) by a 10% (wt/wt) AgNO₃–SiO₂ column, each ester was analyzed by ¹³C NMR.

The ¹³C-isotopic abundances of each carbon atom of ¹³C-labeled LAME are summarized in Fig. S2 and Table S1. Chemical shifts for each carbon atom were assigned with reference to LAME ¹³C NMR data (5). Regarding the relative ¹³C-isotopic abundance values, the C-2 carbon atom was set as the standard signal and ¹³C-isotopic abundance was calculated for each carbon atom. In the mite lipid-derived ¹³C-labeled LAME, the odd-numbered carbon atoms 1, 3, 5, 7, 9, 11, 13, 15, and 17 showed strong signals from ¹³C. The individual levels of ¹³C-isotopic abundances of even-numbered carbon atoms were 750–1053%. Conversely, the ¹³C-isotopic abundances of even-numbered carbon atoms—that is, 2 (standard signal), 4, 6, 8, 10, 12, 14, 16, and 18—were low at –22 to 15%; these were therefore judged not to be enriched by ¹³C atoms.

The ¹³C-isotope abundances of each carbon atom of ¹³C-labeled OAME are summarized in Fig. S3 and Table S2. Chemical shifts of each carbon atom were assigned with reference to the previously described LAME ¹³C NMR data. Regarding calculations of the relative strength of the carbon signal, the C-2 carbon atom was set as the standard signal in the same manner as for LAME. In mite lipid-derived ¹³C-labeled OAME, odd-numbered carbon atoms 1, 3, 5, 7, 9, 11, 13, 15, and 17 were strongly labeled by ¹³C. The ¹³C-isotopic abundances of these nine carbon atoms were 689–919%. Conversely, ¹³C-isotopic abundances of even-numbered carbon atoms—that is, 2 (standard signal), 4, 6, 8, 10, 12, 14, 16, and 18—were low at –15 to 37%, and these were judged to not be enriched by ¹³C.

Contrasting Fatty Acids, ¹³C NMR Analysis of 8,11-F17 and 8-F17 **Demonstrate Carbonyl and Even** ¹³C-labeling of Carbons. After mite hexane extract was prepared as previously described, 8,11-F17 and 8-F17 were isolated via a 10% (wt/wt) AgNO₃-SiO₂ column, each ¹³C NMR spectrum was measured. ¹³C NMR spectra of chemically synthesized 8,11-F17 and 8-F17 were also measured as a control.

The ¹³C NMR spectra and data of ¹³C-labeled and ¹³C-unlabeled 8,11-F17 are summarized in Fig. 2 and Table S3. All chemical shifts of the carbon atoms were assigned with reference to ¹³C NMR data of the analogous molecules 6,9-C17 (5). The carbonyl carbon and the even-numbered carbons at 2, 4, 6, 8, 10, 12, 14, and 16 detected in mite-derived ¹³C-labeled 8,11-F17 were consistent with strong ¹³C labeling. The even-numbered carbons correspond to the C-1 carbon of the ¹³C-labeled acetyl units before the carbon chain is elongated. From the ¹³C NMR data, we conclude that $[1-^{13}C]$ acetate was incorporated into 8,11-F17. Conversely, odd-numbered carbons at 1, 3, 5, 7, 9, 11, 13,



Fig. 2. Mite incorporation of [1-¹³C] acetate into the aliphatic formate 8,11-F17 reveals even carbon chain ¹³C-labeling. ¹³C NMR spectra of ¹²C-unlabeled and ¹³C-labeled 8,11-F17. Filled circles indicate significantly ¹³C-enriched atoms. x: contaminant signal.

15, and 17 were not enriched with 13 C and no incorporation signal was detected.

⁻¹³C NMR spectra and data of ¹³C-labeled and ¹³C-unlabeled 8-F17 are summarized in Fig. 3 and Table S4. All chemical shifts of the carbon atoms are assigned with reference to 8,11-F17 (Table S3). Similar to ¹³C-labeled 8,11-F17, the signals from the carbonyl carbon and the even-numbered carbons at 2, 4, 6, 8, 10, 12, 14, and 16 in mite-derived ¹³C-labeled 8-F17 were consistent with ¹³C labeling. The ¹³C NMR data also confirmed the incorporation of $[1^{-13}C]$ acetate into 8-F17. Conversely, odd-numbered carbons at 1, 3, 5, 7, 9, 11, 13, 15, and 17 were not enriched with ¹³C and no incorporation signal was detected.

Incorporation of 1-¹³C-Labeled Stearic Acid Demonstrates Origins of 8,11-F17 and 8-F17 Carbonyl Carbons. To examine the incorporation and origin of ¹³C atoms into the formyl moiety present in 8,11-F17 and 8-F17, mites were fed on a mixture of dry yeast and $[1^{-13}C]$ stearic acid [1:4 (wt/wt)] every 4 d, and the secretions of mites were analyzed every 7 d. When raised with food containing no additional ¹³C-label, the abundance of the isotopic ion of 8,11-F17 (M⁺+1 ion, *m/z* 281) was 0.20 for M⁺ ion (*m/z* 280). A week after feeding the $[1^{-13}C]$ stearic acid, the values doubled. After 3 wk, they were 6-fold, and after an additional 3 wk they were 30-fold (Fig. 4). According to these data, it is possible that the ¹³C atoms were incorporated into the formyl moiety of 8,11-F17 molecules.

In the case of 8-F17, we did not determine whether ¹³C had been incorporated into the formyl moiety because there was a complete dissociation of H13COOH from the M+ ion, thus making the M^+ ion unobservable in electron ionization (EI) mode. However, the fact that no increases were observed in the amount of M⁺-HCOOH+1 ions (m/z 237) in 8-F17 is consistent with the lack of ¹³C incorporation anywhere else except the formyl moiety in the molecule (Fig. S4). To detect the molecular ion of 8-F17, we analyzed ¹³C-labeled and ¹³C-unlabeled 8-F17 by chemical ionization (CI). When raised with food not containing ¹³C-labeled stearic acid, the quasi-molecular ion of 8-F17 $(M^++H \text{ ion}, m/z 283)$ was detected (Fig. 5). Five weeks after feeding the [1-¹³C] stearic acid, $M^+ + H + 1$ ion (m/z 284) was detected together with the M⁺+H+1-H¹³COOH ion (m/z 237). These results are highly consistent with a single ¹³C atom incorporation specifically into the formyl moiety of the 8-F17.

In the case of 6,9-C17 and 8-C17 hydrocarbon biosynthesis, ${}^{13}CO_2$ would be dissociated from linoleyl aldehyde and oleyl aldehyde, respectively, during the decarbonylase reaction leading to cuticular hydrocarbon biosynthesis (15). By feeding with $[1^{-13}C]$ stearic acid, the corresponding hydrocarbons were not

labeled with ¹³C atom caused by loss of ¹³CO₂ (Fig. S5). Collectively, the current data are consistent with the ¹³C atom derived from $[1-^{13}C]$ stearic acid as selectively present in the formyl moiety and not significantly incorporated into other carbon positions.

Discussion

To elucidate the biosynthesis of aliphatic formates, specifically 8,11-F17 and 8-F17, in an arthropod model, ¹³C-labeled acetate was administered to mites (*Sancassania* sp. Sasagawa) following by targeted GC-MS and ¹³C NMR analyses. ¹³C was accurately incorporated into the formates 8,11-F17 and 8-F17, and the specific carbon location of its biosynthesis precursor LA and OA was determined. In the biosynthesis of fatty acids, the carbon chain is elongated by malonyl-CoA (C₂); this process is initiated with an acetyl-CoA (C₂). A result of incorporating [1-¹³C] acetate into mite lipids, odd-numbered ¹³C labeled LA and OA are generated through elongation and desaturation (Fig. 6). These data agree well with the results of a previous study, where ¹³C was inserted into an LA carbon chain in a regular manner by administering ¹³C-labeled glucose to *C. lactis* (5).

In contrast to established patterns for fatty acids, when the labeling pattern of formates 8,11-F17 and 8-F17 are considered, even-numbered carbon atoms are labeled with ¹³C in a regular manner. The most parsimonious hypothesis is that the carboxyl group of LA and OA has been decreased by one carbon. However, when we consider that a carbonyl carbon of formate also remains labeled with ¹³C, then it is further possible that formate is generated by an oxygen atom inserted between the fatty acid carboxyl carbon and the α carbon (Fig. 6). An alternative and more conservative hypothesis is that C18 fatty acids are converted to C17 molecules with primary alcohols through a process of α -oxidation and reduction, followed by the reconstitution of a C18 molecule by the conjugation to formate. Although biologically plausible, such a mechanism is highly disfavored considering empirical data that the carbonyl carbon of formate present in 8-F17 is concentrated with ¹³C (Fig. 5). Consequently, the currently proposed mechanism of aliphatic formate biosynthesis would proceed via Baeyer-Villiger oxidation following reduction of fatty acids (aliphatic acyl CoA) to aldehydes (Fig. 7). To verify that aldehydes are direct precursors, an isotopically labeled aldehyde was prepared for an administration experiment on mites. However, exogenous aldehyde feeding experiments resulted in significant miticide activity, thus forcing consideration of alternative experiments. Aldehydes



Fig. 3. Mite incorporation of $[1-^{13}C]$ acetate into the aliphatic formate 8-F17 also reveals even carbon chain ^{13}C -labeling. ^{13}C NMR spectra of ^{12}C -unlabeled and ^{13}C -labeled 8-F17. Filled circles indicate significantly ^{13}C -enriched atoms. \times : contaminant signal.



Fig. 4. Mites fed [1-¹³C] stearic acid as a precursor specifically retain ¹³C-labeled formate present in 8,11-F17. Mass spectra of ¹²C-unlabeled and ¹³C-labeled 8,11-F17 in El mode. Filled circles indicate significantly ¹³C-enriched atoms.

are known intermediates of hydrocarbons in insects (15–18), and our own studies make it increasingly clear that aldehydes are also intermediates of hydrocarbons in acaridae. Because mites (*Sancassania* sp. Sasagawa) secrete hydrocarbons and formates at the same time, it is hypothesized that two different types of enzymes act on aldehydes as a common substrate in the secretory glands and generate hydrocarbons and formates separately (Fig. 7).

Feeding experiments with [1-¹³C] stearic acid were carried out to determine if the formate 8,11-F17 is generated without the process of LA dehomologation. According to the previously described [1-¹³C] acetate administration experiment, because LA has been verified to be de novo biosynthesized, when $[1-^{13}C]$ stearic acid is taken into the body of mites, it should be unsaturated and generate [1-13C] LA. Given our hypothesis that 8,11-F17 is synthesized from LA, we predicted that [formyl-¹³C] 8,11-F17 (m/z 281) should be obtained. When mite secretions were analyzed by GC-MS, the strength of the M⁺+1 ion $(m/z \ 281)$ of 8,11-F17 increased over time during the administration experiment, and at 6 wk the ion strength was 30 times stronger compared with the control (Fig. 4). Given the rather surprising and strong 1-13C enrichment measured, we present these findings as evidence that 8,11-F17 is biosynthesized without going through the process of dehomologation, which-if predominant—would oppositely remove the 1-13C label present. We propose that this process similarly occurs for 8-F17, although we do present any parallel GC/EI-MS evidence for this example because of suboptimal ionization parameters. However, using an alternative soft CI approach, we analyzed 8-F17 after feeding mites with [1-13C] stearic acid and successfully obtained M++H+1

ion (m/z 284) derived from [formyl-¹³C] 8-F17 in mass spectrum (Fig. 5). This provides additional supporting evidence that 8-F17 is also biosynthesized without going through the process of dehomologation. In contrast, the findings that M⁺+1 parent ions of 6,9-C17 and 8-C17 do not increase are consistent with the generation of hydrocarbons (6,9-C17 and 8-C17) by decarbonylation of 1-¹³C-labeled aldehyde (Fig. S5). Considering all current results, we propose the involvement of a Baeyer–Villiger oxidation mechanism in the generation of an oxygen atom between the carbonyl group of aldehyde, the reduced product of LA (linoleoyl-CoA) and OA (oleoyl-CoA), and the carbon chain.

In 1948, indication of a biological Baeyer–Villiger oxidation was demonstrated for the first time during a series of microbiological studies on steroids (19). Since then, Baeyer–Villiger monooxygenase, which changes cyclic ketones to lactones and aliphatic and aromatic ketones to corresponding esters, has been predominantly found in microbes (20, 21). In plants, brassinolide has the highest physiological activity among brassinosteroid hormones and is essential in the promotion of extension growth; it is also known to be generated by the Baeyer–Villiger oxidation of castasterone, a biosynthetic intermediate (22, 23). To the best of our knowledge, we are unaware of enzymes capable of Baeyer–Villiger oxidations in animals. Baeyer–Villiger monooxygenases are useful





Fig. 6. Summary of measured ¹³C-labeling patterns present in fatty acids and formates, specifically 8,11-F17 and 8-F17, after feeding with [1-¹³C] acetate. Filled circles indicate significantly ¹³C-enriched atoms.

as biocatalysts, especially in the selective synthesis of natural products and pharmaceuticals, given the ability to produce optically active lactones or esters via the enantioselective oxidation of racemic ketones (21).

As a classic Baeyer-Villiger reaction of aliphatic and aromatic aldehydes using peroxide, carboxylic acids are selectively generated (24). However, in 2010 generation of formate from aliphatic and aromatic aldehvdes via a Baever-Villiger oxidation was demonstrated and enabled by a hypervalent difluoro- λ^3 -bromane in water (25). Although there are cases of formate being generated from aromatic aldehyde in enzymatic Baeyer-Villiger oxidations that use cyclohexanone monooxygenase and 4-hydroxyacetophenone monooxygenase of microbial origin (26-28), an enzyme that generates formate from aliphatic aldehyde has yet to be discovered. In organic chemistry, an efficient transformation of a primary alcohol to a one-carbon-shorter carboxylic acid using o-iodoxybenzoic acid and iodine has recently been reported (29). In the present study, the predicted arthropod enzyme would catalyze a novel dehomologation and generate a one-carbonshorter primary alcohol from an aldehyde via hydrolysis of formate. The substrate breath of substrate specificity of this enzyme still needs to be clarified; however, we expect useful applications requiring mild dehomologation in the synthesis of the labile bioactive molecules.

Methods

Chemical Analysis. Column chromatography was performed on Wakosil silica gel C-200 using the specified solvents. ¹H- and ¹³C NMR spectra were recorded on a Bruker Biospin AC400M spectrometer (400 MHz for ¹H and 100 MHz for ¹³C) using tetramethylsilane as the internal standard. GC-MS

was conducted using a Network GC System (6890N; Agilent Technologies) coupled with a mass-selective detector (5975 Inert XL; Agilent Technologies) operated at 70 eV using an HP-5MS capillary column (0.25 mm i.d. \times 30 m with 0.25-µm film thickness; Agilent Technologies). Helium was used as the carrier gas at a flow rate of 1.00 mL/min, with a splitless mode at a temperature set to increase from 60 (2 min) to 290 °C at a rate of 10 °C/min. The temperature was maintained at 290 °C for 5 min. Positive CI mass spectra were acquired on the same GC system coupled with mass spectrometer (JMS-700 MStation, JOEL) operated at 200 eV using an DB-5MS capillary column (0.25 mm i.d. \times 30 m with 0.25-µm film thickness; Agilent Technologies), with isobutane as reagent gas.

Mites. An unidentified *Sancassania* sp. Sasagawa strain (Acari: Acaridae) derived from hypopus attached to a Japanese rhinoceros beetle *Trypoxylus dichotomus* (L. 1771) originating from Kameoka city (Kyoto prefecture, Japan) was used in this study. Culture lines were established in our laboratory and maintained for several generations in a sterilized environment at 20 °C and 90% relative humidity in an agar medium, as reported previously (30). The internal transcribed spacer (ITS-II) region of this species consisted of 498, 496, or 492 bps (code name SS28, GenBank accession nos.: AB104962, AB104963, AB104964) as listed in Noge et al. (31).

Feeding Experiment with a ¹³C-Labeled Compound-Enriched Diet. For the feeding experiment, mites were fed on an agar medium with dry yeast and sodium [1-13C] acetate (>99% 13C enrichment; Sigma-Aldrich) [9:1 (wt/wt)] for 7 d. No adverse effect was observed with the addition of sodium acetate to the medium. The rate at which ¹³C atoms were incorporated into 8,11-F17 and 8-F17 was monitored by GC-MS on the second and seventh day. At each time interval, three mites were transferred to a conical glass insert (5 mm in diameter × 30 mm in height; Agilent Technologies) using a needle and soaked for 3 min in hexane (3 µL). The extract was directly subjected to GC-MS analysis. Another feeding experiment was conducted by using a mixture of dry yeast and [1-¹³C] stearic acid (99% ¹³C enrichment: Sigma-Aldrich) [1:4 (wt/wt)] for 6 wk. A set amount of feed (30-50 mg) was provided once every 4 d. The rate at which ¹³C atoms were incorporated into 8,11-F17 and 8-F17 was monitored by GC-MS in EI or CI mode at given time intervals over 7 d. Secretion by the mites was extracted with hexane and analyzed by GC-MS as described above.

Isolation of 8,11-F17 and 8-F17 Incorporated with Sodium [1-¹³C] Acetate. Mites of all developmental stages and both sexes were separated from the culture medium by suspension in a saturated saline. Medium-free mites (2.37 g) were then immersed in hexane (10 mL) for 10 min to obtain the secretions of the laterodorsal opisthonotal glands. After evaporation of the solvent, the extract was applied to a 10% (wt/wt) AgNO₃-SiO₂ column (2.8 g) and eluted with a mixture of ethyl acetate and hexane [1:10 (vol/vol)]. Purified 8,11-F17 (1 mg) and 8-F17 (<1 mg) were analyzed by ¹³C NMR spectroscopy.



Fig. 7. Proposed role of an arthropod-associated Baeyer–Villiger type oxidation in the biosynthesis of aliphatic formates (8,11-F17 and 8-F17) and hydrocarbons (6,9-C17 and 8-C17) via aldehydes as common precursors.

Isolation of LA and OA Incorporated with Sodium [1-13C] Acetate. After hexane extraction of the mites as described above, hexane-free mites were then extracted with chloroform:methanol [2:1 (vol/vol), 10 mL] for 3 d. After filtration, the filtrate was concentrated in vacuo and total lipids were hydrolyzed with sodium methoxide (100 mg) in methanol (5 mL) under reflux for 2 h. After acidifying the reaction mixture with 1N HCl at room temperature, ethyl acetate was added. The organic layer was successively washed with water and brine, dried over anhydrous Na₂SO₄, and concentrated in vacuo. The dry residue (17 mg) was dissolved with benzene:methanol [4:1 (vol/vol). 1 mL] and a trimethylsilyldiazomethane solution (0.6 M, 0.5 mL) was added to the solution at room temperature. After 30 min, the solvent was removed in vacuo, leaving fatty acid methyl esters (17 mg) in the form of oil. For isolation and purification of LAME and OAME, these esters were applied to a 10% (wt/wt) AgNO₃-SiO₂ column (1.7 g) and eluted with a mixture of ethyl acetate and hexane [1:10 (vol/vol)]. Purified LAME (3 mg) and OAME (3 mg) were analyzed, respectively, by ¹³C NMR spectroscopy. Samples of authentic unlabeled LAME and OAME were obtained from a commercial source for use as a standard (Sigma-Aldrich).

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Determination of ¹³C-Labeling Patterns. ¹³C NMR spectra of the compound labeled at the ¹³C atoms and of the naturally occurring compound were recorded under the same experimental conditions. Chemical shifts were assigned for every ¹³C NMR signal, and the signal integral for each respective carbon atom in the labeled compound was referenced to that of the compound with natural ¹³C abundance, thereby affording the relative ¹³C abundances for each position in the ¹³C-labeled compounds.

Chemical Identification. Two formates 8,11-F17 and 8-F17 were synthesized to establish unbiased structures (14). Two hydrocarbons, 6,9-C17 and 8-C17, have been previously identified (32, 33). Two fatty acids, LA and OA, were identified by comparison of mass spectra and GC retention times with authentic samples. A commercial calibration standard containing bacterial fatty acid methyl esters (Catalog No. 47080-U, Sigma-Aldrich) was used as the authentic sample.

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