Biosynthesis of the insect pheromone (S)-4-methyl-3-heptanone

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Using stable isotope-labelled probes and mass spectrometry, the insect pheromone (*S*)-4-methyl-3-heptanone is shown to be biosynthesised from three propionate units following a polyke-tide/fatty acid-type metabolic route.

Simple 3-ketones (Fig. 1) are common secondary metabolites in insects and other arthropods, where they serve a range of communicatory and ecological functions.¹ Ketones **1–5** have all been identified in various exocrine glands of ants, with (*S*)-4-methyl-3-heptanone (**1**) exhibiting particularly widespread taxonomic distribution. Ketone **1** is usually located in the mandibular glands and serves as an alarm pheromone, but, in at least one species (*Aphaenogaster albisetosus*), it is stored in the poison gland and is used to coordinate nestmate recruitment to food sources.² In addition to its role as a pheromone, ketone **1** is employed by opilionids (Arachnida) as a defensive allomone against ants.³ Moreover, in the interaction between the ant *Paraponera clavata* and its parasite *Apocephalus paraponerae* (Diptera), there is evidence that **1** functions as a kairomone (a semiochemical that disfavours the emitter and benefits another organism).⁴

In comparison with plants and microorganisms, very little is known about the biosynthesis of secondary metabolites in insects. The route to 3-ketone 1 and its relatives, for example, has never been investigated. It has been proposed that these simple alkanones are aceto/propiogenins.⁵ Indeed, it is easy to see how all the structures in Fig. 1 can be assembled from the condensation of acetate and/or propionate units. A potential route to 4-methyl-3-heptanone (1) is shown in Scheme 1. A starter unit of propionyl– SEnz is condensed with methylmalonate to yield diketide **6**. Following total reduction of the β -keto group (by the action of a putative ketoreductase, dehydratase and hydrogenase), a second methylmalonate is incorporated to give triketide **8**. Hydrolysis of the thioester and decarboxylation then results in the production of methyl ketone **1**. Thus, although **1** has only eight carbons, it is assembled from three C₃ units.

Here, we report a study on the biosynthesis of **1** in the ant *Harpegnathos saltator*⁶ using stable isotope-labelled probes together with mass spectrometric (MS)-based detection. [${}^{2}H_{3}$]Methylmalonic acid and [${}^{2}H_{3}$]methyl[1,3- ${}^{13}C_{2}$]malonic acid were synthesised and introduced into the diet of *H. saltator.*[†] GC/MS



analysis of the mandibular gland contents from individual treated ants revealed clear incorporation of labelling. Combining MS scans over the entire GC peak of 4-methyl-3-heptanone resulted in a spectrum containing ions from a number of isotopomers (Fig. 2). A maximum of nine ²H atoms, in multiples of three, were incorporated into ketone **1**, a result consistent with the biogenetic origin outlined in Scheme 1. As a consequence of the higher volatility of $[^{2}H_{9}]$ -**1** over isotopomers bearing fewer deuterium atoms, it was possible to achieve partial GC resolution of this species, such that



Scheme 1 Proposed biosynthetic route to **1**: (i) methylmalonate, $-CO_2$; (ii) reduction, dehydration, reduction; (iii) methylmalonate, $-CO_2$; (iv) thioester hydrolysis, $-CO_2$.



Fig. 2 Molecular ion region of the mass spectrum of ketone 1 from *H*. *saltator* following exposure to $[{}^{2}H_{3}]$ methylmalonic acid. M⁺⁺ for unlabelled 1 (*m*/*z* 128) is accompanied by signals due to incorporation of one, two and three $[{}^{2}H_{3}]$ methyl groups.

a pure spectrum of $[{}^{2}H_{9}]$ -1 could be obtained. Similarly, upon treatment of the ants with $[{}^{2}H_{3}]$ methyl $[1,3{}^{-13}C_{2}]$ malonic acid, a clean spectrum of $[{}^{2}H_{9}][{}^{13}C_{2}]$ -1 was recorded (Fig. 3). Comparison of the *m*/*z* values for fragment ions from unlabelled [Fig. 3(A)] and labelled [Fig. 3(B) and (C)] 1 revealed a deuterium labelling pattern consistent with the structures in Fig. 3. Moreover, α -cleavage either side of the C=O group uniquely identified C3 as the site of one of the 13 C labels in Fig. 3(C). Strictly, the second 13 C could possibly have resided at C5 or C6, as the fragmentation of ketone 1 did not distinguish between these positions. Only C5, however, exhibits a



Fig. 3 Mass spectra of ketone 1 from *H. saltator*: (A) without treatment; (B) following exposure to $[^{2}H_{3}]$ methylmalonic acid; (C) following exposure to $[^{2}H_{3}]$ methyl $[1,3^{-13}C_{2}]$ malonic acid. The asterisks on the structure indicate ${}^{13}C_{2}$

1,3-relationship with two $[^{2}H_{3}]$ groups, retaining the relative position of labels seen in the malonate precursor. Thus, it is highly probable that the second ^{13}C was located at C5.

The labelling patterns observed in **1** (Fig. 3) demonstrate that this ketone is produced from three propionate building blocks, with loss of C1 from one C₃ unit. These results provide the first evidence to support the proposed biosynthetic route shown in Scheme 1 and demonstrate that 4-methyl-3-heptanone is a product of polyketide/ fatty acid-type metabolism. The notion that related ketones are also produced by this general route is supported by the observation that labelling from $[^{2}H_{3}]$ methylmalonic acid was incorporated into 4-methyl-3-hexanone (**2**), a trace component in the mandibular glands of *H. saltator*. In this case, the Me branch and C1 were labelled with $[^{2}H_{3}]$, but the C6 Me group remained unlabelled (data not shown). This result is consistent with a mixed acetate/ propionate origin for **2**, where C5 and C6 stem from acetate and the remaining carbons are propionate-derived.

In summary, we have shown that simple ketones (Fig. 1) can be synthesised by insects, using the polyketide/fatty acid pathway, and stored in exocrine glands for use as semiochemicals.

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Notes and references

† Labelled methylmalonic acids were synthesised from [1,3-¹³C₂]- or unlabelled dimethyl malonate and [²H₃]iodomethane in NaOMe/MeOH, followed by saponification. Aqueous solutions of the labelled probes (5 µl at 0.1 g ml⁻¹) were injected into crickets (pre-paralysed by *H. saltator* venom) and the ant colonies fed on a diet of three treated crickets per week. After three weeks, callow worker ants were dissected and their mandibular glands extracted individually in dichloromethane (10 µl) before GC/MS analysis. Approximately 10% of samples revealed incorporation into **1**.

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