

# Biogenesis of sex pheromones in the female olive fruit-fly

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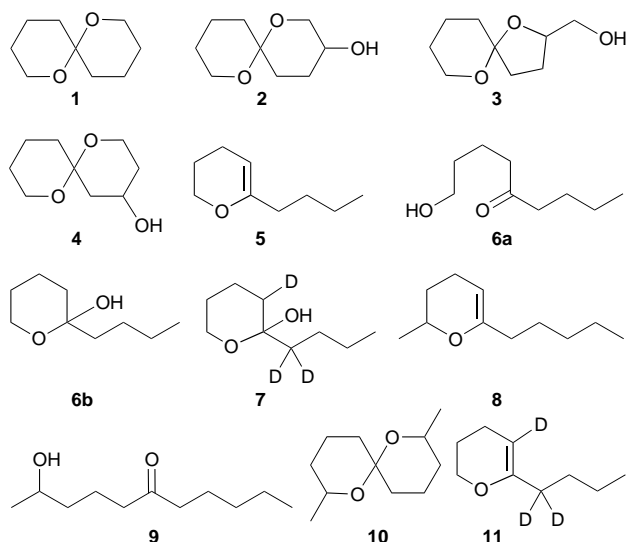
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**A likely pathway to the sex pheromones of *Bactrocera oleae* (olive fruit-fly) is presented, based mainly on feeding experiments with deuterium labelled precursors.**

There is little knowledge of the chemistry, enzymology or molecular biology of the biosynthesis of fruit-fly pheromones which could be linked with the known chemistry<sup>1,2</sup> of these species to facilitate species-specific monitoring and control. We now describe experiments with the pestilent species, *Bactrocera oleae* (olive fruit-fly),<sup>1,3</sup> that permit a proposal for the biogenesis of the (female) pheromone, which is essentially one component.<sup>4</sup> This suggests the operation of a single, major biosynthetic pathway for study, and possible disruption.

The major component of the pheromone is racemic 1,7-dioxaspiro[5.5]undecane **1**<sup>5</sup> which is accompanied by low levels (*ca.*



3%) of hydroxy derivatives **2–4**.<sup>6</sup> We selected 6-*n*-butyl-3,4-dihydro-2*H*-pyran **5**<sup>7</sup> for investigation as an advanced precursor of **1**<sup>8</sup> for the following reasons. Although **5** has not been identified in the olive fruit-fly, it co-occurs with **1** in *B. cacuminata*,<sup>9</sup> in which keto alcohol **6a** also occurs; dehydration of the corresponding hemiketal **6b** would afford **5**. Secondly, 2-methyl-6-*n*-pentyl-3,4-dihydro-2*H*-pyran **8**<sup>10</sup> and keto alcohol **9** accompany isomers of 2,8-dimethyl-1,7-dioxaspiro[5.5]undecane **10** in *B. halfordiae* and *B. kraussii*,<sup>2</sup> so that the nexus that applies to **5** and **1** also applies to **8** and **10**. Appropriate side-chain hydroxylation of these dihydropyrans (**5** and **8**) followed by cyclisation would yield the corresponding spiroacetals **1** and **10**.

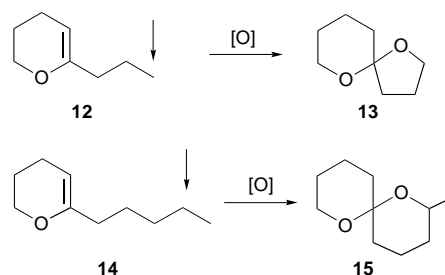
A [<sup>2</sup>H<sub>3</sub>]-labelled analogue of **5**, *viz* **11**, was prepared from [<sup>2</sup>H<sub>6</sub>]-propanone,<sup>11</sup> and essentially complete deuteration at the indicated positions in **11** was confirmed by <sup>1</sup>H and <sup>13</sup>C NMR and mass spectral analyses.‡ Examination by GC–MS of the

rectal glandular contents of female olive flies, subsequent to the receipt of diet-administered **11**, established that **1** was now significantly <sup>2</sup>H-enriched, as were the accompanying hydroxy-spiroacetals **2–4**, and the fragmentation patterns confirmed that deuterium was located at the anticipated positions  $\alpha$  to the spiro centre. In this and in other cases described here, variation in isotopic composition for **1** was observed by progressive mass spectral examination from the fore- to the centre of the GC peak. More highly deuterated species elute earlier.<sup>11</sup> These examinations showed at least 40% of **1** incorporated deuterium. It is possible that *in vivo* hydration of **5** to form the hydroxy ketone **6a** (or the corresponding cyclic hemiketal **6b**) provided the real substrate for methyl oxidation. Thus, **7** (a deuterium analogue of **6**) was synthesised and administered and resulted in efficient formation of labelled **1**, indicating that **6** is also a possible penultimate precursor of **1**.

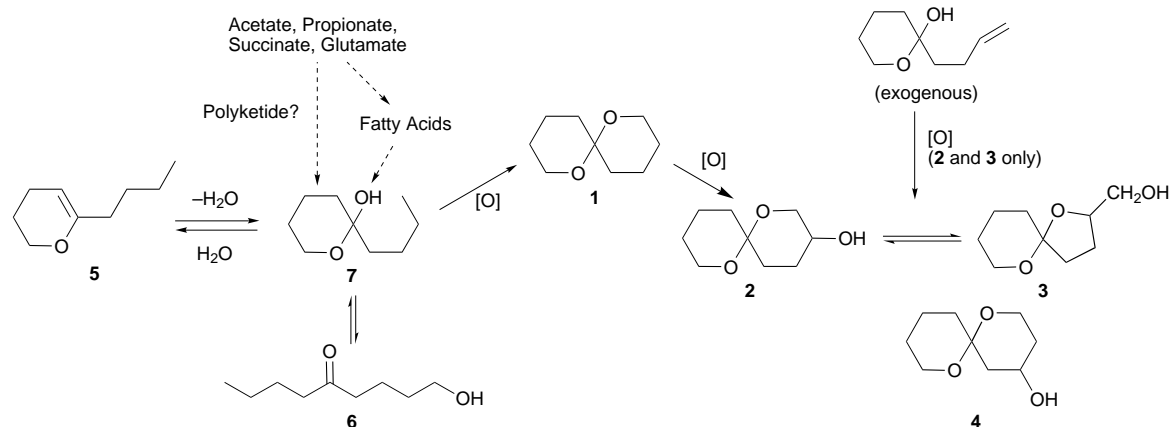
To explore the nature and specificity of the presumed side-chain oxidation, dihydropyrans **12** and **14** were also administered to female olive flies and inefficient oxidation (compared with **11**) occurred, to provide low levels of the known spiroacetals **13** and **15**,<sup>2</sup> respectively (Scheme 1). To the best of our knowledge these have never been detected in olive fruit-flies under natural conditions. Labelled alkenes **16** and **17**, which correspond to possible precursors of the hydroxy-spiroacetals **2** and **3**, were also administered to whole female *B. oleae*. Analyses confirmed that deuterium from **16** or, more effectively, **17** was specifically incorporated into **2** and **3**, but not **4** or the parent spiroacetal **1**. This outcome is shown in Scheme 2.

The proportion of labelled **2** and **3** relative to **1** and **4** increased to a level twenty times higher when **17**, but not **16**, was administered. This supports the view that the penultimate step in the biosynthesis of **1** in *B. oleae* is  $\omega$ -oxidation of **6**, followed by cyclisation. Significantly, the results for **12**, **14**, **16** and **17** demonstrate that exogenous compounds can access the enzymes of the biosynthetic sequence, a crucial feature if enzyme inhibitors<sup>12</sup> are to be devised for pest control.

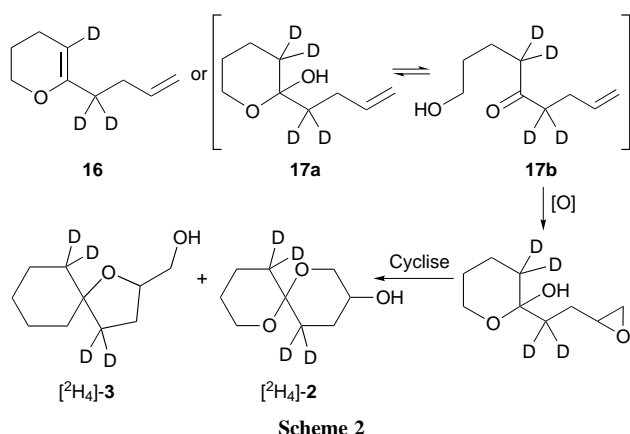
With respect to the origin of the minor co-occurring hydroxy-spiroacetals **2–4**, we considered these might arise from hydroxylation of intact, initially formed spiroacetal **1**. To test this proposal, [<sup>5,5,11,11</sup>-<sup>2</sup>H<sub>4</sub>]-**1** was synthesised<sup>13</sup> and administered, and GC–MS analysis confirmed the efficient formation



Scheme 1



Scheme 3



Scheme 2

of labelled 2–4. The enantiomeric compositions of the hydroxy spiroacetals from this trial group closely matched those from the control group.<sup>13,14</sup> In contrast to this, the enantiomeric profile of the hydroxy derivatives resulting from administration of 16 or 17 was quite different. For example, 3 from the control group was predominantly (2*R*,5*R*) and (2*R*,5*S*) (20% ee), but was predominantly (2*S*,5*S*) and (2*S*,5*R*) (60% ee) for those derived from 17.<sup>13</sup> This indicates that epoxidation of an unsaturated precursor is an unlikely natural route to 2 and 3. We believe that 2–4 are hydroxylation products of intact 1 (or a chemically equivalent species).<sup>15</sup>

In summary, the above results are consistent with  $\omega$ -hydroxylation of 6 (or possibly 5) followed by dehydrative spirocyclisation to yield 1. The remarkable, highly regioselective oxidation (compare results from 5, 12 and 14) of a remote Me or CH<sub>2</sub> group is reminiscent of similar oxidations mediated by cytochrome P450s in other eucaryotic systems.<sup>16</sup> This class of enzymes is known to occur in insects, mediating important transformations such as pesticide detoxification.<sup>17</sup>

With this framework established (Scheme 3), we are now directing attention to the origin of 6, the likely precursor of 1, and to establishing the generality of the biosynthetic relationship between keto alcohols and the corresponding spiro acetals, e.g. 9 and 10. Keto alcohols 6 and 9 could conceivably arise by either a fatty acid or polyketide pathway and experiments to differentiate these are being conducted. In this context, it is of interest that labelled nonanoic acid and 5-oxononanoic acid were not incorporated into 1 in *B. oleae*. The isolation and characterisation of the likely P450 enzyme in *B. oleae* and *B. cacuminata* are also being pursued.

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## Notes and References

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‡ All new compounds exhibited appropriate <sup>1</sup>H and <sup>13</sup>C NMR spectra and high resolution mass spectra.

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