

1 **The *Drosophila melanogaster* pheromone Z4-11Al is**  
2 **encoded together with habitat olfactory cues and mediates**  
3 **species-specific communication**

4 Sebastien Lebreton<sup>1</sup>, Felipe Borrero-Echeverry<sup>2</sup>, Francisco Gonzalez<sup>1</sup>, Marit Solum<sup>1</sup>, Erika  
5 Wallin<sup>3</sup>, Erik Hedenström<sup>3</sup>, Bill S. Hansson<sup>4</sup>, Anna-Lena Gustavsson<sup>5</sup>, Marie Bengtsson<sup>1</sup>,  
6 Göran Birgersson<sup>1</sup>, William B. Walker<sup>1</sup>, Hany Dweck<sup>6</sup>, Paul G. Becher<sup>1</sup>, Peter Witzgall<sup>1\*</sup>

7 <sup>1</sup>Department of Plant Protection Biology, Swedish University of Agricultural Sciences, Box  
8 102, 23053 Alnarp, Sweden

9 <sup>2</sup>Biological Control Laboratory, Colombian Corporation of Agricultural Research Research,  
10 AA 240142 Las Palmas, Bogota, Colombia

11 <sup>3</sup>Department of Chemical Engineering, Mid Sweden University, Holmgatan 10, 85170  
12 Sundsvall, Sweden

13 <sup>4</sup>Department of Evolutionary Neuroethology, Max Planck Institute for Chemical Ecology,  
14 Hans-Knoell-Strasse 8, 07745 Jena, Germany

15 <sup>5</sup>Department of Medical Biochemistry and Biophysics, Karolinska Institutet, Scheeles väg  
16 2, 17165 Solna, Sweden

17 <sup>6</sup>Department of Molecular, Cellular, and Developmental Biology, Yale University, New  
18 Haven, CT 06520, USA

19 \*Correspondence: Peter Witzgall, SLU, Box 102, 23053 Alnarp, Sweden  
20 phone +46 70 2426939, peter.witzgall@slu.se

21 **Abstract**

22 Mate recognition in animals evolves during niche adaptation and involves habitat and  
23 social olfactory signals. *Drosophila melanogaster* is attracted to fermenting fruit for  
24 feeding and egg-laying. We show that, in addition, female flies release a pheromone (Z)-  
25 4-undecenal (Z4-11Al), that elicits flight attraction in both sexes. The biosynthetic  
26 precursor of Z4-11Al is the cuticular hydrocarbon (Z,Z)-7,11-heptacosadiene (7,11-HD),  
27 which is known to afford reproductive isolation between the sibling species *D.*  
28 *melanogaster* and *D. simulans*. A pair of alternatively spliced receptors, Or69aB and  
29 Or69aA, is tuned to Z4-11Al and to food olfactory cues, respectively. These receptors are

30 co-expressed in the same olfactory sensory neurons, and feed into a neural circuit  
31 mediating species-specific, long-range communication: the close relative *D. simulans*,  
32 which shares food resources and co-occurs with *D. melanogaster*, does not respond. That  
33 Or69aA and Or69aB have adopted dual olfactory traits highlights the interplay of habitat  
34 and social signals in mate finding. These olfactory receptor genes afford a collaboration  
35 between natural and sexual selection, which has the potential to drive phylogenetic  
36 divergence.

## 37 **Keywords**

38 pheromone, kairomone, olfaction, reproductive isolation

## 39 **Significance Statement**

40 Volatile insect sex pheromones carry a message over a distance, they are perceived by  
41 dedicated olfactory receptors, and elicit a sequence of innate behaviours. Pheromones  
42 mediate specific mate recognition, but are embedded in and perceived together with  
43 environmental olfactory cues. We have identified the first long-range, species-specific  
44 pheromone in *Drosophila melanogaster*. A pair of spliced olfactory receptors, feeding into  
45 the same neural circuit, has developed a dual affinity to this pheromone and kairomones,  
46 encoding adult and larval food. Blends of this pheromone and kairomone specifically  
47 attract *D. melanogaster*, but not the close relative *D. simulans*. This becomes an  
48 excellent paradigm to study the interaction of social signals and habitat olfactory cues in  
49 premating reproductive isolation and phylogenetic divergence.

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## 51 **Introduction**

52 Volatile insect pheromones transmit species-specific messages and elicit long-range flight  
53 attraction. Premating communication with pheromones facilitates and accelerates mate-  
54 finding, and reduces predation risk and energy expenditure, which is particularly adaptive  
55 in short-lived insects (1,2).

56 Sexual communication subserves mate-finding and ultimately reproduction. Female  
57 insects search for larval food and oviposition sites soon after mating, and both sexes

58 forage to offset the nutritional cost of reproduction. The search for mates and food is  
59 accordingly interconnected, and so is the response to sex and habitat olfactory signals.  
60 The sensory drive hypothesis predicts that mate recognition in animals evolves during  
61 niche adaptation and that premating sexual communication involves olfactory  
62 specialization to both social signals and habitat cues (3,4).

63 Pheromones are released into an atmosphere that is filled with environmental, habitat-  
64 related olfactory cues, some of which signal mating sites and food sources. The response  
65 to sex pheromones and food or habitat odourants (kairomones) is under sexual and  
66 natural selection, respectively. Pheromones and kairomones are always perceived as an  
67 ensemble in a natural context and this leads to an interaction of sexual and natural  
68 selection during adaptive divergence of sexual signaling, which is thought to facilitate  
69 premating reproductive isolation (3-7).

70 Olfactory sexual communication is studied at cellular and molecular resolution in the fruit  
71 fly *Drosophila melanogaster*, but volatile pheromones encoding species-specific, long-  
72 range mate recognition have not yet been found. *Drosophila* is attracted to yeast and  
73 fruit odorants for feeding, mating and oviposition (8-10) and the interconnection between  
74 perception of pheromones and food semiochemicals is a current research theme (11, 12).  
75 For example, the male-produced sex pheromone cVA and food stimuli are integrated to  
76 coordinate feeding, courtship behavior and oviposition site selection (13-16). Perception  
77 of cVA is a current and outstanding paradigm for studying the molecular and neuronal  
78 logic of innate, olfactory-mediated reproductive behavior (15, 17, 18). cVA and other  
79 known olfactory pheromones are active during courtship, and since they are all shared  
80 with other *Drosophila* species, they cannot account for species-specific communication  
81 (19, 20, 21).

82 *D. melanogaster* and *D. simulans* are sibling species with no gene flow between them  
83 (22). Interspecific matings of *D. melanogaster* with *D. simulans*, or other closely related  
84 species are inhibited by the female-produced cuticular hydrocarbon (Z,Z)-7,11-  
85 heptacosadiene (7,11-HD), which is perceived through gustatory receptors at close range  
86 (23-25). 7,11-HD or other non-volatile cuticular hydrocarbons convey species-specificity,  
87 but they are not volatile and cannot account for long-range communication and flight  
88 attraction. This raises the question whether *Drosophila* uses, in addition, volatile  
89 pheromone signals that mediate specific mate recognition at a distance.

90 We have identified the first long-range, species-specific pheromone in *D. melanogaster*. A  
91 pair of spliced olfactory receptors, feeding into the same neural circuit, has developed a  
92 dual affinity to this pheromone and to environmental semiochemicals, encoding adult and  
93 larval food. A blend of this pheromone and a food odourant specifically attracts *D.*  
94 *melanogaster*, but not the close relative *D. simulans*. This becomes an excellent paradigm  
95 for studying the interaction of social signals and habitat olfactory cues in premating  
96 reproductive isolation and phylogenetic divergence.

## 97 **Results**

### 98 ***Drosophila melanogaster* females produce a suite of volatile aldehydes.**

99 A focus in *Drosophila* pheromone research has been on cuticular hydrocarbons, which are  
100 active during close-range courtship. Our scope was to investigate volatile compounds  
101 encoding long-range communication. We therefore collected volatile compounds released  
102 by *D. melanogaster* flies in a glass aeration apparatus and found 16 aliphatic aldehydes,  
103 according to chemical analysis by gas chromatography-mass spectrometry (GC-MS).

104 Males and females shared saturated aldehydes with a carbon chain length of C7 to C18,  
105 but mono-unsaturated aldehydes were released by females only (Fig. 1A; Table 1). The  
106 most abundant compound was identified as (Z)-4-undecenal (Z4-11Al) and synthesized.

107 GC-MS analysis showed that Z4-11Al was present also in cuticular extracts of females,  
108 although in lower amounts ( $0.27 \pm 0.12$  ng/female,  $n = 5$ ) than in headspace collections  
109 ( $3.0 \pm 0.81$  ng/female,  $n = 10$ ;  $P < 0.01$  Mann-Whitney test). Cuticular profiles of  
110 *Drosophila* flies have been investigated, but Z4-11Al or other aldehydes have not been  
111 reported (19, 26, 27).

112 The sister species *D. simulans* did not release Z4-11Al, nor other monounsaturated  
113 aldehydes (Fig. 1A). Unlike *D. melanogaster*, *D. simulans* does not produce (Z,Z)-7,11-  
114 heptacosadiene (7,11-HD) (23). This led us to hypothesize that the production of mono-  
115 unsaturated aldehydes with a double bond in position 4 was linked to oxidation of di-  
116 unsaturated cuticular hydrocarbons. Oxidation of 7,11-HD is expected to generate two  
117 saturated aldehydes, heptanal and hexadecanal, and two unsaturated aldehydes, Z4-11Al  
118 and (Z)-4-eicosenal (Fig. 1B). This was experimentally verified by applying 100 ng  
119 synthetic 7,11-HD to a glass vial. After 60 min,  $1.92 \pm 0.42$  ng Z4-11Al were retrieved ( $n$   
120 = 3). Based on the cuticular hydrocarbon profile of *D. melanogaster* (24), 26 aldehydes  
121 are expected to be formed by oxidation, 16 of which were found in our headspace  
122 analysis (Table 1), others may have been below detection level.

123 Next, single sensillum electrophysiological recordings (SSR) from all basiconic, trichoid,  
124 coeloconic, and intermediate olfactory sensilla in *D. melanogaster* (Fig. 1C) and GC-  
125 coupled SSR recordings (GC-SSR) from ab9 sensilla (Fig. 1D) showed that Z4-11Al  
126 strongly activates ab9A olfactory sensory neurons (OSNs). A weaker response from ab4A  
127 neurons to Z4-11Al probably reflects the sensitivity of Or7a (expressed in ab4A) to  
128 aldehydes, such as the leaf volatile (E)-2-hexenal or bombykal, a lepidopteran  
129 pheromone compound (28-30).

130 **The olfactory receptor Or69aB responds to Z4-11Al.**

131 ab9A OSNs express the olfactory receptor (Or) Or69a (31). We therefore screened ab9A  
132 OSNs with known ligands of Or69a (30) and Z4-11Al. In the *D. melanogaster* strains  
133 Canton-S and Zimbabwe, the monoterpene (*R*)-carvone elicited the strongest response  
134 from ab9A, although the response to Z4-11Al was not significantly different. In *D.*  
135 *simulans*, Z4-11Al elicited a significantly lower response than (*R*)-carvone (Fig. 2A).

136 The Or69a gene encodes two proteins, Or69aA and Or69aB, as a result of alternative  
137 splicing (Fig. 2D), which occurred prior to the split of the *D. obscura* and *D. melanogaster*  
138 groups (32, 33). Heterologous co-expression of both Or69a splice variants in ab3A  
139 (*Δhalo*) empty neurons (34) produced a response similar to native ab9A OSNs; whereas  
140 individual expression revealed distinct response profiles for Or69aA and Or69aB (Fig. 2 A  
141 and B). Or69aB responds best to both isomers of carvone, followed by Z4-11Al.

142 Carvone and Z4-11Al, which seem unlike at first glance, share a structural motif, a  
143 carbonyl functional group with an equidistant double bond in position 4 (Fig. 2 B and C).  
144 Different ligands, upon binding to the same Or, are thought to adopt a complementary  
145 bioactive conformation. The strain energy required for any compound to assume a steric  
146 conformation that aligns with an active ligand should typically not exceed 5 kcal/mol  
147 (35). Conformational analysis showed that Z4-11Al aligns with (*R*)-carvone, which elicited  
148 the strongest Or69aB response, at a strain energy cost of only 1.5 kcal/mol. Or69aA, on  
149 the other hand, is tuned to terpenoid alcohols and responded significantly less to Z4-  
150 11Al. The most active ligands (*S*)-terpineol, (*S*)- and (*R*)-linalool, which again share the  
151 functional group and a double bond in position 4, align at 3.0 kcal/mol (Fig. 2 B and C).  
152 Conformational analysis confirms that the most active ligands for Or69aA and Or69aB,  
153 respectively, are structurally related.

154 **Z4-11Al elicits upwind flight attraction in *D. melanogaster*, but not in *D.***  
155 ***simulans*.**

156 Z4-11Al elicited upwind flight and landing at the source, in cosmopolitan Dalby and  
157 Canton-S strain *D. melanogaster* males and females. In contrast, males of the Zimbabwe  
158 strain and *D. simulans* were not attracted (Fig. 3 A and B). This shows that Z4-11Al, in  
159 addition to its precursor 7,11-HD (Fig. 1) participates in sexual isolation between *D.*  
160 *melanogaster* and its sister species *D. simulans* (23), and between cosmopolitan and  
161 African *D. melanogaster* strains (36-38).

162 Moreover, admixture of Z4-11Al eliminated *D. simulans* attraction to the yeast volatile  
163 (*R*)-linalool (39) (Fig. 3A). *D. melanogaster* and its sister species *D. simulans* co-occur in  
164 the same habitat and use partially overlapping food resources. The very rarely occurring  
165 hybrid matings are sterile; the antagonistic interaction between pheromone and food  
166 stimulus is therefore adaptive and substantiates a contributing role of Or69a in

167 reproductive isolation. A tentative explanation for the significantly reduced attraction  
168 response of *D. melanogaster* females to the blend of Z4-11Al and linalool, compared to  
169 males (Fig. 3A) is that females would avoid food resources and rendezvous sites  
170 overcrowded with other females.

171 The response magnitude of wild-type flies to Z4-11Al released at a rate of 10 ng/min was  
172 similar to the upwind flight response to vinegar headspace, when the main compound  
173 acetic acid was released at a 170-fold amount (Fig. 3A; 36). This illustrates the high  
174 responsiveness and sensitivity of *D. melanogaster* males and females to Z4-11Al.

175 Finally, we used tetanus toxin mutants to verify that Or69a encodes Z4-11Al. Upwind  
176 flight attraction to Z4-11Al or linalool was significantly reduced when Or69a OSNs were  
177 disrupted (Fig. 3B). In summary, Z4-11Al is a powerful attractant that enables specific  
178 mate recognition at a distance. Its heterospecific role was conserved even in blends with  
179 the food attractant linalool.

## 180 **Conclusions**

181 Z4-11Al is the first species-specific, long-range sex pheromone of *D. melanogaster*. It is  
182 produced by females and perceived by Or69aB in both sexes. The precursor of Z4-11Al is  
183 the cuticular hydrocarbon 7,11-HD, which mediates isolation between *D. melanogaster*  
184 and its sister species *D. simulans* during courtship (23-25).

185 In addition to pheromone, Or69aB and its twin receptor Or69aA bind kairomonal  
186 terpenoids, such as linalool or terpineol, which are found in both fruit and yeast  
187 headspace. Citrus peel, a preferred oviposition substrate (8), and baker's yeast which  
188 grows on ripe fruit and elicits fly attraction and oviposition (9) are sources of all main  
189 ligands of Or69aA and Or69aB (39, 41).

190 Combined pheromone and food odour tuning in the two Or69a splice variants underscores  
191 the tie between sexual and natural selection during the evolution of specific mate  
192 communication (3, 4). Flies releasing pheromone tint pervading habitat and food  
193 odorants and thus shape and foreground a communication channel that facilitates mate  
194 finding. This is particularly adaptive in *Drosophila* when mating sites, fruit and berries,  
195 are abundant and widely spread.

196 A twin receptor, emerging from the Or69a splice event (32,33), facilitated adaptive  
197 changes in ligand tuning, without compromising the established functional role of the  
198 Or69a channel. Functional divergence has apparently been biased towards structurally  
199 related ligands (Fig. 2 B and C; 30) and ecologically relevant odorant signals. This is

200 expected; conservative diversification of the splice variants is constrained to a behavioral  
201 theme, since both Ors feed into one OSN.

202 Olfactory representations of other *Drosophila* Ors involved in food and pheromone  
203 perception project through separate channels to the LH, where third-order neurons  
204 partially overlap and integrate (15). In stark contrast, Or69a is the first olfactory gene  
205 known to encode dual olfactory traits. Or69aA and Or69aB co-express in the same ab9A  
206 OSNs (31) and thus achieve a coordination of mating and food stimuli already in first  
207 order neurons, at the antennal periphery. This makes Or69a a target for selection during  
208 phylogenetic divergence. The tuning range of Ors evolves more rapidly than hardwired  
209 neural circuits in higher brain centres (42) and selection pressure is further relaxed  
210 following a splice event. Differential tuning of Or69a in closely related cosmopolitan and  
211 African strains of *D. melanogaster* (Fig. 2 A) corroborates this idea.

212 Tuning changes in the two splice forms of Or69a are restricted with respect to the  
213 behavioral and ecological role of their ligands, since they both feed into a neural circuit  
214 mediating sexual and habitat attraction. The two splice forms provide, on the other hand,  
215 degrees of freedom during adaptive divergence, since they allow fly populations to adopt  
216 new kairomone or pheromone signals; alteration of either one produces a new  
217 communication channel. Reproductive isolation may arise as a byproduct (43-45) and the  
218 Or69a gene therefore has the potential to drive speciation (46, 47). Species in the *D.*  
219 *melanogaster* and *D. obscura* groups provide a rich substrate for studying Or ligand  
220 evolution and its consequences on disruptive selection on ecological interactions and  
221 mate choice.

## 222 **Methods**

### 223 **Insects**

224 Canton-S, Zimbabwe (S-29; Bloomington #60741) and Dalby-HL (Dalby, Sweden) (48)  
225 strains of *D. melanogaster* were used as wild type flies for behavioral experiments.  
226 Canton-S was used for comparison with knockouts of the same background. Further tests  
227 were done with the sister species *D. simulans*.

228 We used the Or69a-Gal4/UAS TeTx, tetanus toxin knockout line to verify the role of  
229 Or69a in flight attraction to Z4-11Al. Canton-S/UAS TeTx (Bloomington #28838 and  
230 28997) and Canton-S/Or69a-Gal4 (Bloomington #10000) were used as parental controls.

231 Flies were reared on a standard sugar-yeast-cornmeal diet at room temperature (19 to  
232 22°C) under a 16:8-h L:D photoperiod. Newly emerged flies were anesthetized under CO<sub>2</sub>  
233 and sexed under a dissecting microscope. Virgin flies were identified by the presence of

234 meconium, and were kept together with flies of the same sex. Flies were kept in 30-ml  
235 Plexiglas vials with fresh food. Experiments were done with 3- to 5-d-old flies.

## 236 **Chemicals**

237 (*Z*)-4-undecenal (*Z*4-11Al) and (*E*)-4-undecenal (*E*4-11Al) were synthesized. A short  
238 description follows below, for a complete account of the chemical synthesis, see  
239 Supplemental Information.

240 (*Z*)-4-Undecenoic acid was synthesized via a modified version of Wube et al. (49) in 80%  
241 stereoisomeric purity. Esterification under acidic conditions with sulfuric acid in methanol  
242 resulted in 80% *Z*-isomer and a 93% yield over two steps. Stereoisomeric purity was  
243 controlled with NMR and GC-FID by comparing analysis for acid and ester, the  
244 appearance of a small quartet, in the NMR spectra, at 1.96 indicates the presence of *E*-  
245 isomer. Gas chromatographic separation on a polar Varian factorFOUR vf-23ms of *Z*- and  
246 *E*-ester proved that the stereochemistry was not affected by the acidic conditions during  
247 esterification. Methyl (*Z*)-4-undecenoate was purified on regular silica gel and on silver  
248 nitrate impregnated silica gel to obtain a stereoisomeric purity of 98.6%. Methyl (*Z*)-4-  
249 undecenoate was reduced to (*Z*)-4-undecenol with lithium aluminum hydride in  
250 diethylether and oxidized to *Z*4-11Al with Dess-Martin periodinane in dichloromethane.

251 A modified version of Virolleaud's (50) metathesis was used to produce (*E*)-4-undecenoic  
252 acid in a 56% yield (87.5% of the *E*-isomer). (*E*)-4-undecenoic acid was esterified under  
253 the same conditions as the (*Z*)-acid, without isomerisation of the double bond (according  
254 to GC-FID and <sup>1</sup>H-NMR). The methyl-(*E*)-4-undecenoate was reduced to the alcohol with  
255 lithium aluminum hydride in diethylether and purified on silver nitrate impregnated silica  
256 gel to obtain a purity of 99.8% of the (*E*)-isomer, which was oxidized with Dess-Martin  
257 periodinane in dichloromethane to obtain *E*4-11Al.

258 Commercially available compounds were: (*R*)-carvone (97% chemical purity, CAS #6485-  
259 40-1, Firmenich), (*S*)-carvone (98%, CAS #2244-16-8, Firmenich), (*S*)-terpineol (97%,  
260 CAS #10482-56-1, Aldrich), (*S*)-linalool (97%, CAS #126-91-0, Firmenich), (*R*)-linalool  
261 (97%, CAS #126-90-9, Firmenich), citronellol (99%, CAS #106-22-9, Aldrich), geraniol  
262 (98%, CAS #106-24-1, Aldrich), 3-octanol (99%, CAS #589-98-0, Aldrich), decanol  
263 (99%, CAS #112-30-1, Fluka), 11-Al (99%, CAS #112-44-7, Aldrich).

## 264 **Odor collection and chemical analysis**

265 Twenty *D. melanogaster* (Dalby), *D. melanogaster* (Canton) (*n* = 10) or 20 *D. simulans*  
266 (*n* = 10) unmated female or unmated male flies were placed in a glass aeration  
267 apparatus designed for collection of airborne pheromone (effluvia) (51). The flies were  
268 held in a glass bulb with a narrow open outlet (ø 1 mm), which prevented them from  
269 escaping. A charcoal-filtered air flow (100 mL/min) passed over the flies during 75 min.



270 Fly effluvia were collected on the glass surface, breakthrough was monitored by attaching  
271 a 10-cm glass capillary ( $\varnothing$  1 mm) attached to the outlet. After 75 min, flies were  
272 removed, 100 ng of heptadecyl acetate (internal standard) was deposited in the glass  
273 bulb, which was then rinsed with 50  $\mu$ l hexane, the solvent was concentrated to 10  $\mu$ l in  
274 Francke vials.

275 Cuticular extracts ( $n = 5$ ) were obtained by dropping 20 *D. melanogaster* females for 5  
276 min in 400  $\mu$ l hexane containing 100 ng heptadecyl acetate. After 5 min, the extracts  
277 were transferred to Francke vials and concentrated to 10  $\mu$ l before analysis. Fly extracts  
278 and volatile collections were stored at -20°C.

279 Oxidation of (*Z,Z*)-7,11-heptacosadiene (7,11-HD) was analysed by dropping 100 ng  
280 synthetic 7,11-HD into a 1.5-mL glass vial, at 19°C. Vials were rinsed with 10  $\mu$ l of  
281 hexane, which contained 100 ng heptadecyl acetate as an internal standard, after 15, 30,  
282 45, 60 and 75 min ( $n = 3$ ).

283 Samples were analysed by combined gas chromatography and mass spectrometry (GC-  
284 MS; 6890 GC and 5975 MS, Agilent technologies Inc., Santa Clara, CA, USA). Two  $\mu$ l  
285 were injected (injector temperature 225°C) splitless (30 s) into fused silica capillary  
286 columns (60 m x 0.25 mm), coated with HP-5MS UI (Agilent Technologies Inc.,  $d_f = 0.25$   
287  $\mu$ m) or DB-wax (J&W Scientific, Folsom, CA, USA,  $d_f = 0.25$   $\mu$ m), that were temperature-  
288 programmed from 30 to 225°C at 8°C/min. Helium was used as mobile phase at 35 cm/s.  
289 The MS operated in scanning mode over  $m/z$  29-400. Compounds were tentatively  
290 identified based on their mass spectra and Kovats retention indices, using custom and  
291 NIST (Agilent) libraries, followed by comparison with authentic standards.

## 292 **Behavioural assays**

293 Upwind flight behavior was observed in a glass wind tunnel (30 x 30 x 100 cm) equipped  
294 with a piezo sprayer (40). The flight tunnel was lit diffusely from above, at 13 lux,  
295 temperature ranged from 22 to 24°C, relative humidity from 38% to 48% and charcoal  
296 filtered air, at a velocity of 0.25 m/s, was produced by a fan (Fischbach GmbH,  
297 Neunkirchen, Germany). Compounds were delivered from the centre of the upwind end of  
298 the wind tunnel via a piezo-electric micro-sprayer (52). Forty flies were flown individually  
299 to each treatment. "Attraction" was defined as upwind flight, directly from a release tube  
300 at the end of the tunnel over 80 cm towards the odor source, followed by landing.

301 Unmated fed, 3-d-old Dalby wild-type males and females, *D. melanogaster* Zimbabwe  
302 strain males and *D. simulans* males were flown towards (*Z*)-4-undecenal (released at 10  
303 ng/min), (*R*)-linalool (10 ng/min) and the blend of (*Z*)-4-undecenal and (*R*)-linalool (10  
304 ng/min, each).

305 **Heterologous expression of Or69aA and Or69aB**

306 Or69aA and Or69aB receptors were cloned from antennae of *D. melanogaster*, Dalby line  
307 (53). Briefly, cDNA was generated from RNA extracts of antennae of 100 males and  
308 females using standard procedures. Or69a variants were PCR amplified with the following  
309 primers: Or69aA\_5': GTCATAGTTGAAACCAGGATGCAGTTGC, Or69aB\_5':  
310 ATAATTCAGGACTAGATGCAGTTGGAGG, Or69aAB\_3':  
311 TGCACCTTTGCCCTTTTATTTAAGGGAC.

312 The splice variants were amplified with unique 5' primers and a common 3' primer,  
313 reflective of genomic structure at this locus. These primers encompass the entire open  
314 reading frame of the receptor variants, and are located partially upstream and  
315 downstream of the start and stop codons. PCR amplicons were gel-purified and cloned  
316 into the pCR8/GW/Topo-TA Gateway entry vector (Thermo-Fisher Scientific, Waltham,  
317 MA, USA) according to standard procedure, with vector inserts sequenced to confirm  
318 fidelity of Or sequence. Or inserts were subsequently transferred to pUAS.g-HA.attB (54)  
319 with LR Clonase II enzyme (Thermo-Fisher Scientific), according to manufacturers  
320 protocol; vector inserts were sequenced to confirm fidelity of Or sequence.

321 Mini-prep purified pUAS.g-HA.attB plasmid with Or69aA or Or69aB insert were delivered  
322 to Best Gene Inc. (Chino Hills, CA USA) for generation of transgenic *D. melanogaster*  
323 flies. Using the PhiC31 targeted genomic-integration system (54) vectors with Or69aA or  
324 Or69aB were injected into the following fly strain, for integration on the 3<sup>rd</sup> chromosome:  
325 M{3xP3-RFP.attP}ZH-86Fb (with M{vas-int.Dm}ZH-2A) (Bloomington *Drosophila* Stock  
326 Number: 24749). For expression of single receptor variants in the empty neuron system,  
327 Or69a transgenes were crossed into the  $\Delta halo$  background to give genotype: *w*;  
328  $\Delta halo/Cyo$ ; UAS-DmelOr69a(A or B), and these flies were crossed to flies with genotype:  
329 *w*;  $\Delta halo/Cyo$ ; DmelOr22a-Gal4, as described previously (53). Experimental  
330 electrophysiology assays were performed on flies with genotype: *w*;  $\Delta halo$ ; UAS-  
331 DmelOr69a(A or B)/DmelOr22a-Gal4.

332 For co-expression of Or69aA and Or69aB in the same empty neurons, a second fly-line  
333 with Or69aB was generated with Or69aB present on the X-chromosome. The same UASg-  
334 HA.attB:Or69aB plasmid generated previously was injected into the following fly strain:  
335 *y,w*, P{CaryIP}su(Hw)attP8 (Bloomington *Drosophila* Stock Number: 32233). The Or69aB  
336 transgene was crossed into the DmelOr22a-Gal4 line in  $\Delta halo$  background to give  
337 genotype: UAS-DmelOr69aB;  $\Delta halo/Cyo$ ; DmelOr22a-Gal4; these flies were crossed to  
338 flies with genotype: *w*;  $\Delta halo/Cyo$ ; UAS-DmelOr69aA. Experimental electrophysiology  
339 assays were performed on flies with genotype: UAS-DmelOr69aB/*w*;  $\Delta halo$ ; UAS-  
340 DmelOr69aA/DmelOr22a-Gal4.

### 341 **Conformational analysis**

342 MacroModel version 11.0 (Schrodinger LLC, New York, NY, USA) in the Maestro Version  
343 10.4.017 were used to build, minimize and to perform conformational analysis of Z4-  
344 11Al, (*R*)-carvone, (*S*)-terpineol and (*R*)-linalool, using default settings (OPLS3 as force  
345 field, water as the solvent and mixed torsional/low-mode sampling method). The  
346 assumed bioactive conformations of the conformationally more flexible compounds, Z4-  
347 11Al and (*R*)-linalool, were based on the position of the shared functional groups in the  
348 conformationally more restricted compounds, (*R*)-carvone and (*S*)-terpineol. The  
349 carbonyl and the double bond atoms were kept fixed during minimization of the proposed  
350 bioactive conformation of Z4-11Al; the alcohol functional group and the double bond were  
351 kept fixed in (*R*)-linalool. The strain energies, the energy cost for adopting proposed  
352 bioactive conformations, were then calculated as the difference between the lowest  
353 energy conformations and the assumed bioactive conformation.

### 354 **Electrophysiological recordings**

355 Single sensillum recordings (SSR) were done as described earlier (19). Unmated males  
356 were restrained in 100- $\mu$ l pipette tips, with half of the head protruding, the third antennal  
357 segment or palps were placed on a glass microscope slide and held by dental wax. For  
358 the initial screening, all basiconic, trichoid, coeloconic, and intermediate sensilla (31)  
359 were localized in *D. melanogaster* (Canton-S strain) males, under a binocular at 1000x  
360 magnification. Further recordings were made from small basiconic ab9 sensilla, in *D.*  
361 *melanogaster* (Canton-S and Zimbabwe strains) and in *D. simulans* males, and from large  
362 basiconic ab3 sensilla in mutant *D. melanogaster*, where Or69aA and Or69aB were  
363 heterologously expressed (see above).

364 Tungsten electrodes (diameter 0.12 mm, Harvard Apparatus Ltd, Edenbridge, United  
365 Kingdom) were electrolytically sharpened with a saturated KNO<sub>3</sub> solution. The recording  
366 electrode was introduced with a DC-3K micromanipulator equipped with a PM-10 piezo  
367 translator (Märzhäuser Wetzler GmbH, Germany) at the base of the sensilla. The  
368 reference electrode was inserted into the eye. The signal from olfactory sensory neurons  
369 (OSNs) was amplified with a probe (INR-02; Syntech), digitally converted by an IDAC-4-  
370 USB (Syntech) interface, and analyzed with Autospike software v. 3.4 (Syntech). Neuron  
371 activities were recorded during 10 s, starting 2 s before odor stimulation. Neuron  
372 responses were calculated from changes in spike frequency, during 500 ms before and  
373 after odor stimulation.

374 Odorants were diluted in redistilled hexane, 10  $\mu$ g of test compounds in 10  $\mu$ l hexane  
375 were applied to filter paper (1 cm<sup>2</sup>), kept in Pasteur pipettes. The test panel contained  
376 the most active ligands known for Or69a (30) and several aldehydes. Diagnostic  
377 compounds for confirmation of sensillum identity were 2-phenyl ethanol (ab9) and 2-  
378 heptanone (ab3). Control pipettes contained solvent only. Puffs (2.5 ml, duration 0.5 s)

379 from these pipettes, produced by a stimulus controller (Syntech GmbH, Kirchzarten,  
380 Germany), were injected into a charcoal-filtered and humidified airstream (0.65 m/s),  
381 which was delivered through a glass tube to the antenna.

382 For GC–SSR recordings, GC columns and the temperature programmes were the same as  
383 for the GC–MS analysis. At the GC effluent, 4 psi of nitrogen was added and split 1:1 in a  
384 3D/2 low dead volume fourway-cross (Gerstel, Mühlheim, Germany) between the flame  
385 ionization detector and the antenna. Towards the antenna, the GC effluent capillary  
386 passed through a Gerstel ODP-2 transfer line, that tracked GC oven temperature, into a  
387 glass tube (30 cm x 8 mm ID), where it was mixed with charcoal-filtered, humidified air  
388 (20°C, 50 cm/s).

### 389 **Statistical analysis**

390 Generalized linear models (GLM) with a Bernoulli binomial distribution were used to  
391 analyse wind tunnel data. Landing at source and sex were used as the target effects.  
392 *Post-hoc* Wald pairwise comparison tests were used to identify differences between  
393 treatments. For all the electrophysiological tests, differences in spike activity derived from  
394 SSRs were analyzed with Kruskal Wallis H test followed by pairwise comparisons with  
395 Mann Whitney U *post hoc* test. All statistical analysis were carried out using R (R Core  
396 Team 2013) and SPSS Version 22 (IBM Corp.).

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## 560 **Author Contributions**

561 S.L., P.G.B. and P.W. conceived the study. F.B.-E. and M.S. carried out behavioural  
562 studies, F.G. and W.B.W produced empty neuron flies, F.G. and H.D. did single sensillum  
563 recordings, supervised by B.S.H, E.W. and E.H. synthesized the compound, A.-L.G.  
564 calculated the conformational analysis, M.B., G.B. and S.L. performed chemical analysis,  
565 P.W. wrote the paper with input from all co-authors.



566 **Legends**

567 **Figure 1. Headspace analysis of *Drosophila* females and males by GC-MS**  
568 **and electrophysiological screening of the candidate pheromone**  
569 **compound Z4-11Al on male antennae.**

570 (a) Chromatograms of headspace collections from *D. melanogaster* females (lilac traces;  
571 upper trace: amplified signal; lower trace: entire chromatogram), males (blue trace), and  
572 *D. simulans* females (green trace). The headspace of *D. melanogaster* females contains  
573 16 yet undescribed compounds: heptanal (1), octanal (2), (Z)4-nonenal (3), nonanal (4),  
574 (Z)4-undecenal (Z4-11Al) (5), undecanal (6), dodecanal (7), (Z)4-tridecanal (8),  
575 tridecanal (9), tetradecanal (10), pentadecanal (11), (Z)4-hexadecenal (12), hexadecanal  
576 (13), (Z)4-octadecenal (14), octadecanal (15) and (Z)4-eicosenal (16) (see Table 1).  
577 Female-specific compounds are coloured, the most abundant cuticular hydrocarbon, 7-  
578 tricosene (17) is shown for reference, the internal standard (IS) was heptadecyl acetate.  
579 Inset: mass spectrum of the most abundant female-specific compound Z4-11Al.

580 (b) Oxidation of the most abundant female cuticular hydrocarbon (Z,Z)-7,11-  
581 heptacosadiene (7,11-HD), affording two saturated and two unsaturated aldehydes,  
582 heptanal, hexadecanal, Z4-11Al and (Z)4-eicosenal.

583 (c) Single sensillum recordings (SSR) from all *D. melanogaster* olfactory sensory neurons  
584 (OSNs) with Z4-11Al (error bars show SEM;  $n = 5$ ).

585 (d) SSR coupled to GC (GC-SSR), showing a response of ab9A to three different amounts  
586 of Z4-11Al.

587 **Figure 2. SSR-Response of Or69a splice variants to ten odorants, in**  
588 **native ab9A OSNs and in ab3A OSNs, following heterologous expression.**

589 (a) SSR from ab9A OSNs, in *D. melanogaster* (Canton-S, Zimbabwe) and *D. simulans*  
590 males, which natively express both splice variants Or69aA and Or69aB. Responses were  
591 normalized to the most active odorant.

592 (b) SSR from ab3A OSNs in *D. melanogaster*, heterologously expressing Or69aA and  
593 Or69aB, together and singly. Test panel includes the known most active ligands for Or69a  
594 (27) and three aldehydes. Cross-hatched bars indicate behaviorally active compounds  
595 (Fig. 3). Bars followed by different letters indicate statistically significant differences for  
596 each fly type ( $P < 0.05$ ; Mann-Whitney test,  $n = 5$  for ab9A,  $n = 10$  for ab3A).

597 (c) Key ligands for Or69aA, (S)-terpineol (1) and (R)-linalool (4), and for or Or69aB, (R)-  
598 carvone (8) and Z4-11Al (10). Alignment of these ligands illustrates shared structural  
599 motifs.

600 (d) Alternative splicing of Or69a, where coloured boxes A and B show unique exons  
601 encoding the splice products; dark boxes show shared exons, generating co-expression of  
602 Or69aA and Or69aB in the same neurons in ab9 sensilla in *D. melanogaster*.

603 **Figure 3. Z4-11Al mediates long range attraction in *D. melanogaster*.**

604 (a) Upwind flights to 10 ng/min of Z4-11Al and (*R*)-linalool, followed by landing at the  
605 source, in *D. melanogaster* (Dalby) males and females, in *D. melanogaster* (Zimbabwe)  
606 males and *D. simulans* males. Lower case letters indicate statistical differences between  
607 test insect strains and species, for each treatment. Asterisks indicate significant  
608 differences between treatments ( $n = 40$ ,  $P < 0.001$ ; binomial GLMs followed by post-hoc  
609 Wald pairwise comparison tests).

610 (b) Upwind flights to 10 ng/min of Z4-11Al and (*R*)-linalool in *D. melanogaster* (Canton-  
611 S) males expressing a tetanus toxin in OSNs expressing Or69a, and in the parental lines.  
612 Letters indicate statistical differences within treatments ( $n = 40$ ,  $P < 0.001$ ; binomial GLM,  
613 followed by *Post-hoc* Wald pairwise comparison tests).

614

615 **Table 1. Saturated and unsaturated aldehydes found in headspace**  
616 **collections of *D. melanogaster* (Dalby) females and males.**

Compound	Females (%±SD, n=10)	Males (%±SD, n=10)
heptanal	3.1 ± 0.8	9.9 ± 4.8
octanal	tr <b>a</b>	0.2 ± 0.5
nonanal	4.1 ± 0.9	8.7 ± 7.9
(Z)-4-nonenal	tr	- <b>b</b>
undecanal	tr	-
(Z)-4-undecenal	23.3 ± 1.8	-
dodecanal	7.9 ± 1.3	-
tridecanal	tr	-
(Z)-4-tridecenal	0.4 ± 0.9	-
tetradecanal	11.2 ± 0.8	4.9 ± 4.1
pentadecanal	3.2 ± 0.9	2.4 ± 2.1
hexadecanal	27.8 ± 3.3	69.1 ± 9.7
(Z)-4-hexadecenal	2.9 ± 0.3	-
octadecanal	4.7 ± 1.0	4.4 ± 3.9
(Z)-4-octadecenal	3.0 ± 0.6	-
(Z)-4-eicosenal	5.6 ± 1.7	-

617

618 **a** - traces

619 **b** - not found

## 620 **Supplementary Information: Chemical Synthesis**

621 Dry THF and dry Et<sub>2</sub>O was obtained from a solvent purification system (Activated alumina  
622 columns, Pure Solv PS-MD-5, Innovative technology, Newburyport, USA) and used in the  
623 reactions when dry conditions were needed. All other chemicals were used without  
624 purification. Reactions were performed under Argon atmosphere unless otherwise stated.  
625 Flash chromatography was performed on straight-phase silica gel (Merck 60, 230–400  
626 mesh, 0.040–0.063 mm, 10–50 g/g of product mixture) employing a gradient technique  
627 with an increasing concentration (0–100 %) of distilled ethyl acetate in distilled  
628 cyclohexane. In cases of very polar products chromatography was continued with ethanol  
629 in ethyl acetate (0–20 %). Thin-layer chromatography (TLC) was performed to monitor  
630 the progress of the reaction on silica gel plates (Merck 60, precoated aluminium foil),  
631 using ethyl acetate (40 %) in cyclohexane as an eluent, and plates were developed by  
632 means of spraying with vanillin in sulfuric acid and heating at 120°C. Purity of the  
633 product was checked with gas chromatography (GC) analysis on a Varian 3300 GC  
634 instrument equipped with a flame ionization detector (FID) using a capillary column  
635 Equity-5 (30 m x 0.25 mm i.d, d<sub>f</sub> = 0.25 μm, with nitrogen (15 psi) as carrier gas and a  
636 split ratio of 1:20). The oven temperature was programmed at 50°C for 5 min followed by  
637 a gradual increase of 10°C min<sup>-1</sup> to reach a final temperature of 300°C. An Agilent 7890  
638 GC equipped with a polar capillary column FactorFOUR vf-23ms (30 m x 0.25 mm i.d., d<sub>f</sub>  
639 = 0.25 μm) coupled to an Agilent 240 ion-trap MS detector for separation of some  
640 isomeric intermediates. The injector was operated in split mode (1:20) at 275°C, and a  
641 helium flow rate of 1 ml min<sup>-1</sup> and a transfer line temperature of 280 °C. The analyses  
642 were performed in the external ionisation configuration. EI spectra were recorded with a  
643 mass range of m/z 50–300 at fast scan rate. Nuclear magnetic resonance (NMR) spectra  
644 were recorded on a Bruker Avance 500 (500 MHz <sup>1</sup>H, 125.8 MHz <sup>13</sup>C) spectrometer using  
645 CDCl<sub>3</sub> as solvent and internal standard.

646 **(Z)-4-Undecenoic acid.** NaHMDS (6.78 mmol, 1 M in hexane) was added dropwise,  
647 during 30 min, to a suspension of (3-carboxypropyl)triphenylphosphonium bromide (1.45  
648 g, 3.39 mmol) in THF (25 mL). The mixture was stirred for 2 h then cooled to 0°C on  
649 ice/water bath, and heptanal (0.387g, 3.39 mmol) in THF (2.5 mL) was added slowly  
650 during 15 min. The mixture was stirred for 5 h at 0°C then allowed to reach room  
651 temperature over night. The reaction was quenched with H<sub>2</sub>O (20 mL) and the organic  
652 solvent was evaporated. The remaining water phase was extracted with Et<sub>2</sub>O (3 x 20  
653 mL), the obtained organic phases discarded and the basic aqueous phase was acidified  
654 with HCl (2M) until pH 1 and extracted with Et<sub>2</sub>O (3 x 20 mL). The combined organic  
655 phases were dried over MgSO<sub>4</sub> (anhydr.) and the solvent evaporated off. The obtained  
656 crude product was dissolved in pentane, cooled at -18°C and filtered to remove the  
657 precipitated OPPh<sub>3</sub> followed by evaporation of the solvent to result in 0.547 g of a yellow  
658 oil (87.5% yield). <sup>1</sup>H-NMR: 5.52–5.30 (m, 2H), 2.35 (m, 4H), 2.04 (q, J = 6.5 Hz, 1.6H,

659 *Z*-isomer), 1.96 (q,  $J=6.5$  Hz, 0.4H, *E*-isomer), 1.37–1.19 (m, 8H), 0.89 (t,  $J=7$  Hz, 3H)  
660 ppm. The NMR data is in accordance with data previously reported (46, 52). The  
661 relationship by integration between protons at 2.04 and 1.95 indicates approximately a  
662 *Z*:*E*-ratio of 80:20 which is supported by GC-MS analysis on a Varian factorFOUR vf-23ms  
663 column. The obtained crude product was used in the next step without further  
664 purification.

665 **Methyl (*Z*)-4-undecenoate.** (*Z*)-4-Undecenoic acid (0.547 g, 2.97 mmol) from above  
666 was dissolved in methanol (15 mL) and 7 drops of concentrated H<sub>2</sub>SO<sub>4</sub> were added  
667 followed by heating at 70°C over night. The mixture was allowed to reach room  
668 temperature and the methanol was evaporated and the remaining crude product was  
669 dissolved in Et<sub>2</sub>O (15 mL). The organic phase was washed with H<sub>2</sub>O (3 x 10 mL) and brine  
670 (2 x 10 mL), dried over Na<sub>2</sub>SO<sub>4</sub> (anhydr.) and solvent evaporated resulting in 0.547 g of  
671 a yellow oil (92.8% yield). GC-MS (FactorFour vf-23ms) shows a *Z*:*E*-ratio of 80:20. <sup>1</sup>H-  
672 NMR(CDCl<sub>3</sub>): 5.4 (m, 2H), 3.67 (s, 3H), 2.3 (m, 4H), 2.03 (q,  $J = 6.5$  Hz, 1.6H, *Z*-  
673 isomer), 1.96 (q,  $J=6.5$  Hz, 0.4H, *E*-isomer), 1.33–1.21 (m, 8H), 0.89 (t,  $J=6.5$  Hz, 3H)  
674 ppm (no data found in the literature). <sup>13</sup>C-NMR(CDCl<sub>3</sub>): 134.2, 119.9, 32.3, 31.9, 29.5,  
675 29.3, 27.43, 22.7, 14.1 ppm. <sup>13</sup>C-NMR data similar to reported (53). Proton NMR shows a  
676 80:20 *Z*:*E*-ratio between the diastereomers. Enrichment of the *Z*-isomer on AgNO<sub>3</sub> (10%)  
677 impregnated silica resulted in 63 mg of 98.6:1.4 *Z*:*E*-ratio according to GC-FID analysis  
678 on the vf-5 column as the diastereoisomeric purity was not possible to measure when  
679 using <sup>1</sup>H-NMR.

680 **(*Z*)-4-Undecenol.** Methyl (*Z*)-4-undecenoate (63 mg, 0.32 mmol) was dissolved in Et<sub>2</sub>O  
681 (5 mL) and LiAlH<sub>4</sub> (2 spatel tips) was added followed by stirring at room temperature for  
682 30 min. HCl (2 M, 2 mL) was added to quench the reaction and the mixture was extracted  
683 with Et<sub>2</sub>O (2 x 3 mL), the combined organic layer was dried over MgSO<sub>4</sub> (anhydr.) and  
684 solvent was evaporated. Purification with flash chromatography on SiO<sub>2</sub> resulted in 37  
685 mg. <sup>1</sup>H-NMR(CDCl<sub>3</sub>): 5.43–5.32 (m, 2H), 3.67 (m, 2H), 2.16–2.10 (m, 2H), 2.08–2.02  
686 (m, 2H), 1.69–1.60 (m, 2H), 1.39–1.22 (m, 8H), 0.89 (t,  $J=6.5$  Hz, 3H) ppm. NMR-data  
687 were similar to Kim and Hong (54) and Davis and Carlsson (55). Diastereomeric purity  
688 was checked with GC-FID before next step.

689 **(*Z*)-4-Undecenal.** (*Z*)-4-Undecenol (37 mg, 0.22 mmol) in DCM (3 mL) was added to  
690 Dess-Martin periodinane (0.140 g, 0.33 mmol) in DCM (0.5 mL). After 50 min, NaOH  
691 (2M, 10 mL) was added to quench the reaction. The two layers were separated and the  
692 aqueous phase was extracted with Et<sub>2</sub>O (3 x 10 mL), the combined organic layers were  
693 washed with NaOH (2M, 10 mL), dried over MgSO<sub>4</sub> (anhydr.) and solvent was evaporated  
694 resulting in 30 mg of a yellow oil (81% yield). The crude product was purified with  
695 Kugelrohr distillation at bp 65–70 °C (1.6 mbar), resulted in 17 mg. <sup>1</sup>H-NMR(CDCl<sub>3</sub>): 9.77  
696 (s, 1H), 5.48–5.22 (m, 2H), 2.47 (t,  $J = 7$  Hz, 2H), 2.37 (q,  $J = 7$  Hz, 2H), 2.04 (q,  $J =$   
697 7Hz, 2H), 1.37–1.23 (m, 8H), 0.88 (t,  $J = 7$  Hz, 3H) ppm. <sup>13</sup>C-NMR(CDCl<sub>3</sub>): 202.1, 131.8,

698 127.0, 43.9, 31.8, 29.5, 29.0, 27.2, 22.6, 20.1, 14.1 ppm. Both  $^1\text{H}$ - and  $^{13}\text{C}$ -NMR data  
699 were in accordance with published results (56, 57). Analysis on GC-MS (FactorFour vf-  
700 23ms) resulted in a 98.6:1.4 *Z*:*E*-ratio, the *E*-isomer could not be detected by  $^1\text{H}$ -NMR.

701 **(*E*)-4-Undecenoic acid.** 4-Pentenoic acid (0.5 g, 5 mmol) and 1-octene (2.8 g, 25  
702 mmol) was dissolved in DCM (50 mL) and Grubbs II catalyst (85 mg, 0.1 mmol) was  
703 added and the reaction was refluxed. After 7 h was a second portion of Grubbs II catalyst  
704 (85 mg, 0.1 mmol) added and the reaction refluxed for 16 h. Reaction was allowed to  
705 reach room temperature and the solvent was evaporated. The obtained crude product  
706 was dissolved in Et<sub>2</sub>O (50 mL) and filtered through a short pad of silica gel. The product  
707 was purified with flash chromatography by gradient elution (0–100% EtOAc in *c*-hexane  
708 followed by 0–10% EtOH in EtOAc) resulting in 0.52 g oil (56% yield).  $^1\text{H}$ -NMR(CDCl<sub>3</sub>):  
709 5.51–5.33 (m, 2H), 2.41 (q, *J*=7 Hz, 2H), 2.32 (q, *J*=7 Hz, 2H), 2.04 (q, *J*=6.5 Hz, 0.25  
710 H, *Z*-isomer), 1.97 (q, *J*=6.5 Hz, 1.75H, *E*-isomer), 1.37–1.22 (m, 9H), 0.88 (t, *J*=7.5  
711 Hz, 3H) ppm. The relation between proton at 2.04 and 1.97 reveals a 87.5:12.5 *E*:*Z*-  
712 ratio. The isolated product was used in the next step without further purification.

713 **Methyl (*E*)-4-Undecenoate.** (*E*)-4-Undecenoic acid (0.52 g, 2.82 mmol) was dissolved  
714 in methanol (25 mL) and a catalytic amount H<sub>2</sub>SO<sub>4</sub> was added and the mixture was  
715 refluxed over night. After evaporation of the solvent the crude product was dissolved in  
716 Et<sub>2</sub>O (10 mL) and washed with H<sub>2</sub>O (20 mL). The aqueous phase was extracted with Et<sub>2</sub>O  
717 (2 × 25 mL), the combined organic layer was washed with H<sub>2</sub>O (20 mL) and brine (20  
718 mL), dried over MgSO<sub>4</sub> (anhydr.) and evaporation of solvent resulted in 0.439 g (78%  
719 yield).  $^1\text{H}$ -NMR(CDCl<sub>3</sub>): 5.51–5.33 (m, 2H), 3.67 (s, 3H), 2.40–2.27 (m, 4H), 1.96 (q,  
720 *J*=6.5 Hz, 2H), 1.38–1.21 (m, 8H), 0.88 (t, *J*=6.5 Hz, 3H) ppm. Purification with flash  
721 chromatography resulted in 0.401g (71.7% yield). GC-FID showed the same  
722 stereoisomeric ratio as for the acid above.

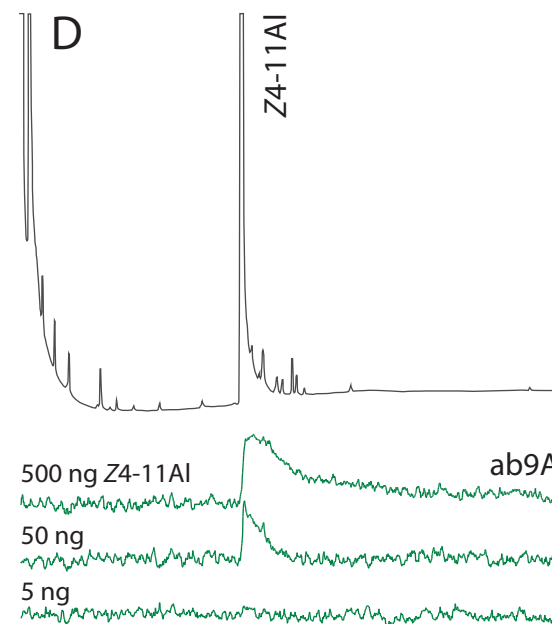
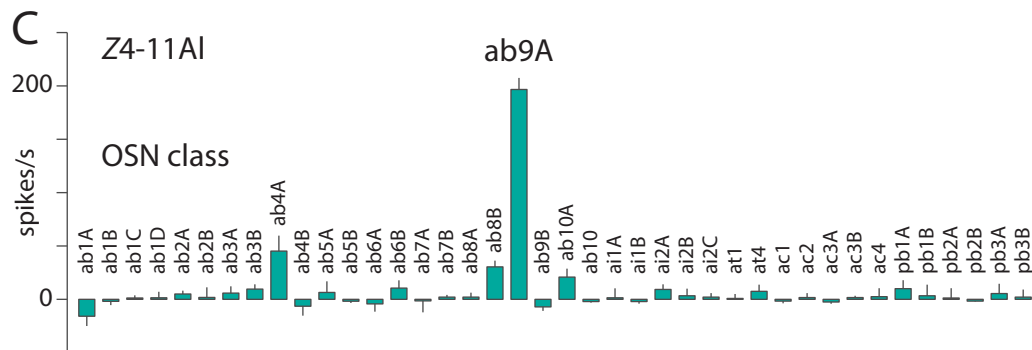
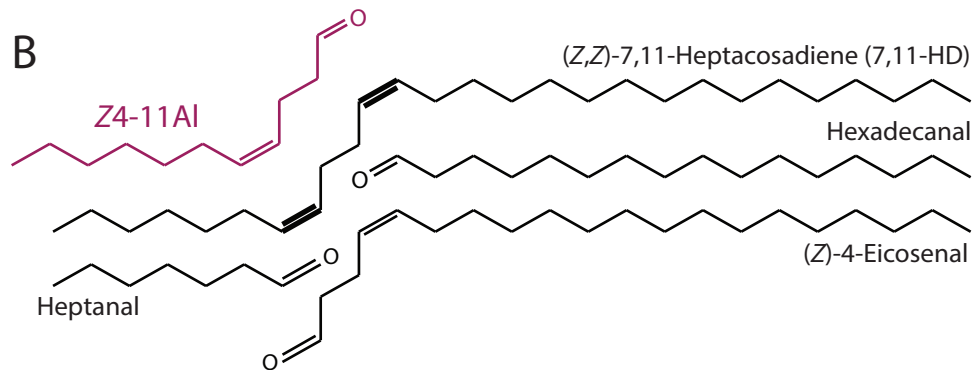
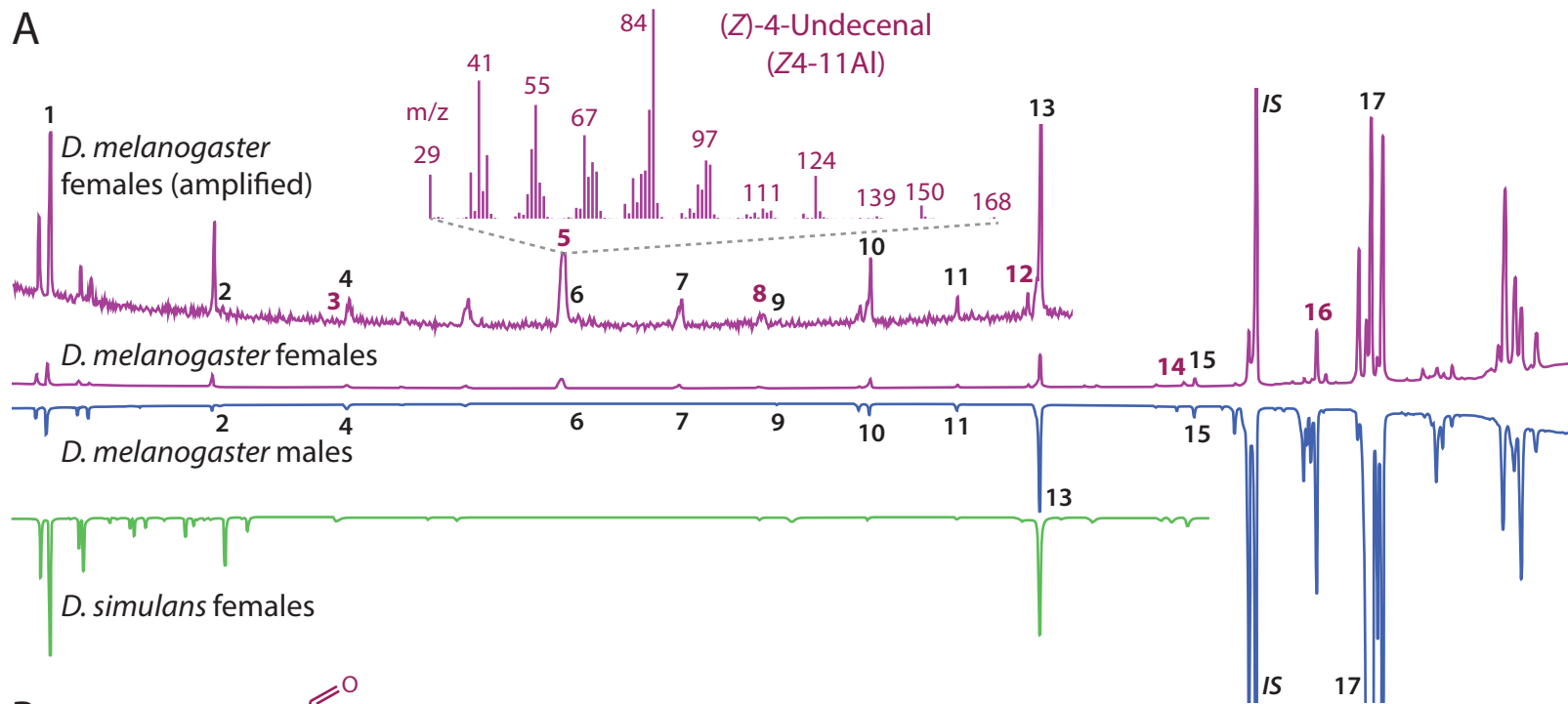
723 **(*E*)-4-Undecen-1-ol.** LiAlH<sub>4</sub> (0.055 g, 1.46 mmol) was added to methyl (*E*)-4-  
724 undecenoate (0.145 g, 0.73 mmol) dissolved in Et<sub>2</sub>O (5 mL). After 30 minutes was HCl (2  
725 M, 5 mL) added to quench the reaction. The acidic water phase was extracted with Et<sub>2</sub>O  
726 (3 × 10 mL) and the combined organic layers were dried over MgSO<sub>4</sub> (anhydr.) and  
727 evaporation of solvent resulted in 0.104 g (99% yield). Enrichment of the *E*-isomer with  
728 medium pressure liquid chromatography (MPLC) on AgNO<sub>3</sub> (10% impregnated) silica  
729 resulted in 30 mg of a clear oil (>99.8 % *E*).  $^1\text{H}$ -NMR(CDCl<sub>3</sub>): 5.43 (m, 2H), 3.65 (m,  
730 2H), 2.08 (q, *J*=7 Hz, 2H), 1.97 (q, *J*=7 Hz, 2H), 1.63 (pent, 2H), 1.35–1.21 (m, 9H),  
731 0.88 (t, *J*=6.5 Hz, 3H) ppm.  $^{13}\text{C}$ -NMR(CDCl<sub>3</sub>): 134.4, 131.3, 129.4, 62.6, 32.6, 32.5,  
732 31.7, 29.6, 29.5, 28.9, 28.8, 22.6, 14.1 ppm. All NMR-data were in accordance with  
733 previous published data (58).

734 **(*E*)-4-Undecenal.** Dess-Martin Periodinane (0.110 g, 0.26 mmol) was added to (*E*)-4-  
735 undecen-1-ol (0.030 g, 0.22 mmol) in DCM (4 mL). NaOH (2 M, 10 mL) was added after  
736 1 h to quench reaction. The aqueous phase was extracted with Et<sub>2</sub>O (3 × 10 mL) and the

737 combined organic layers were dried over MgSO<sub>4</sub> (anhydr.) and evaporation of the solvent  
738 resulted in 30 mg (98% yield). Purification of the crude with Kugelrohr distillation at 65°C  
739 (2 mbar) resulted in 10 mg of product (33% yield, 97% chemical purity, 3% undecenal).  
740 <sup>1</sup>H-NMR(CDCl<sub>3</sub>): 9.76 (t, *J*=1.5 Hz, 1H), 5.50–5.36 (m, 2H), 2.48 (d of t, *J*=7.5, 1.5 Hz,  
741 2H), 2.33 (q, *J*=7 Hz, 2H), 1.97 (q, *J*=6.5 Hz, 2H), 1.32–1.19 (m, 8H), 0.87 (t, *J*=6.5 Hz,  
742 3H). <sup>13</sup>C-NMR (CDCl<sub>3</sub>): 202.5, 132.2, 127.6, 43.6, 32.5, 31.7, 29.4, 28.8, 25.2, 22.6,  
743 14.1. The NMR-data were in accordance with published data (57, 58).

744 (*Z*)-4-Undecenoic acid was synthesized via a modified version of Wube and Hübner (46) in  
745 80% stereoisomeric purity. Esterification under acidic conditions with sulfuric acid in  
746 methanol resulted in 80% *Z*-isomer and a 93% yield over two steps. Stereoisomeric  
747 purity was controlled with NMR and GC-FID by comparing analysis for acid and ester, the  
748 appearance of a small quartet, in the NMR spectra, at 1.96 indicates the presence of *E*-  
749 isomer. Gas chromatographic separation on a polar Varian factorFOUR vf-23ms of *Z*- and  
750 *E*-ester proved that the stereochemistry was not affected by the acidic conditions during  
751 esterification. Methyl-(*Z*)-4-undecenoate was purified on regular silica gel and on silver  
752 nitrate impregnated silica gel to obtain a stereoisomeric purity of 98.6 %. Methyl-(*Z*)-4-  
753 undecenoate was reduced to (*Z*)-4-undecenol with lithium aluminum hydride in  
754 diethylether and oxidized to (*Z*)-4-undecenal with Dess-Martin periodinane in  
755 dichloromethane.

756 A modified version of Virolleaud's (47) metathesis was used to produce the (*E*)-4-  
757 undecenoic acid and in a 56 % yield (87.5% of the *E*-isomer). (*E*)-4-undecenoic was  
758 esterified under the same conditions as the (*Z*)-acid and once again there was no  
759 isomerisation of the double bond (according to GC-FID and <sup>1</sup>H-NMR. The methyl-(*E*)-4-  
760 undecenoate was reduced to alcohol with LiAlH<sub>4</sub> in diethylether and purified on silver  
761 nitrate impregnated silica gel to obtain a purity of 99.8 % of the (*E*)-isomer, which was  
762 oxidized with Dess-Martin periodinane in dichloromethane to obtain the wanted (*E*)-4-  
763 undecenal.

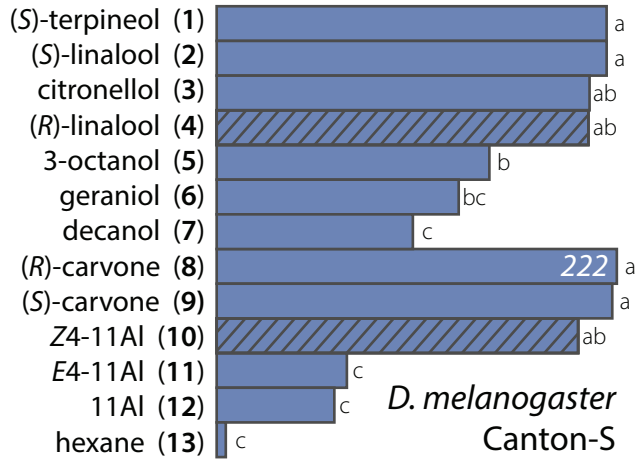




**A**

**Native ab9A OSN Response**

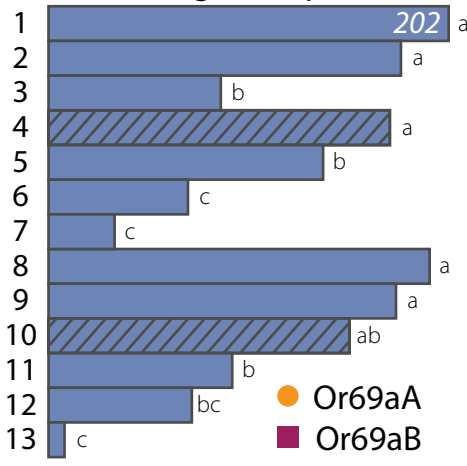
● Or69aA and ■ Or69aB



**B**

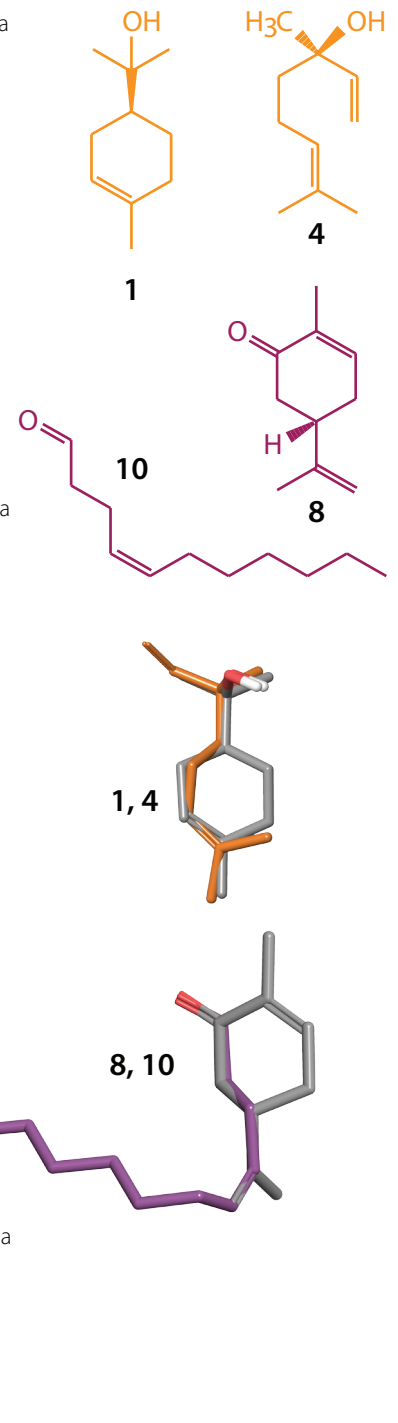
**ab3A( $\Delta$ halo) OSN Response**

(heterologous expression)



**C**

**Main Ligands**



**D**

