

Neural coding merges sex and habitat chemosensory signals in an insect herbivore

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1 **Neural coding merges sex and habitat chemosensory signals in an**
2 **insect herbivore**

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13 **Running title:** Pheromone and plant odour coding

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17 **Keywords:** chemical communication; reproductive isolation; magic trait; intracellular
18 recordings; functional imaging; *Cydia pomonella*

19 **Abstract**

20 Understanding the processing of odour mixtures is a focus in olfaction research. Through
21 a neuroethological approach, we demonstrate that different odour types, sex and habitat
22 cues, are coded together in an insect herbivore. Stronger flight attraction of codling moth
23 males, *Cydia pomonella*, to blends of female sex pheromone and plant odour, compared
24 with single compounds, was corroborated by functional imaging of the olfactory centres in
25 the insect brain, the antennal lobes (AL). The macroglomerular complex (MGC) in the AL,
26 which is dedicated to pheromone perception, showed an enhanced response to blends of
27 pheromone and plant signals, while the response in glomeruli surrounding the MGC was
28 suppressed. Intracellular recordings from AL projection neurons that transmit odour
29 information to higher brain centres, confirmed this synergistic interaction in the MGC.
30 These findings underscore that, in nature, sex pheromone and plant odours are perceived

31 as an ensemble. That mating and habitat cues are coded as blends in the MGC of the AL
32 highlights the dual role of plant signals in habitat selection and in premating sexual
33 communication. It suggests that the MGC is a common target for sexual and natural
34 selection in moths, facilitating ecological speciation.

35 1. Introduction

36 Odours typically are blends of several chemicals, in specific proportions, and the olfactory
37 system decodes and discriminates these multidimensional signals rapidly and precisely. A
38 current question is how odour blends are represented in olfactory circuits and to what
39 extent the neural odour space reflects their ecological and evolutionary significance [1-4].

40 For reproduction, animals largely rely on two types of olfactory signals: sex pheromones
41 distinguish conspecific mates, and habitat odours signal food sources for adults and
42 offspring. Both sex and habitat odours are important mediators of premating reproductive
43 isolation and speciation [5-7] and the neural circuitry underlying the integration of these
44 two types of chemosensory cues is therefore an important target for sexual and natural
45 selection. The interaction of sexual and natural selection is thought to be a powerful
46 driver of speciation [8-10].

47 Insect herbivores are particularly suitable for studying the interaction between mating
48 and habitat cues, especially host plant odours, due to the importance of these signals for
49 their ecology and evolution. Host plant shifts have likely contributed to the remarkable
50 diversification of plant feeding insects [11,12] and most of these rely on sex pheromones
51 for mate finding [13,14].

52 Plant volatiles are recognized as sex pheromone modulators in many insect species
53 [15,16]. Although the behavioural interaction between pheromones and host plant
54 volatiles is well established, little is known about the neurophysiological correlates.
55 Research on the processing of odour blends in the primary olfactory centre in the brain,
56 the antennal lobe (AL), has focused mainly on sex pheromones or on plant volatiles, while
57 the combination of these two classes of compounds is being investigated only since
58 recently [17-19].

59 Separate investigation of pheromones and plant volatile stimuli has led to the idea of a
60 functional specialization of sensory processing in the AL and that these two odour classes
61 are represented in morphologically different regions of the AL of male moths. The
62 macroglomerular complex (MGC) is considered to be dedicated to pheromone coding and
63 the sexually isomorphic, ordinary glomeruli (OGs) to the coding of plant volatile
64 information [20]. Recent studies in the silk moth *Bombyx mori* and the noctuid moth
65 *Agrotis segetum*, however, do not corroborate a strict segregation of the two subsystems
66 and indicate that the MGC receives lateral input from the AL [17-19].

67 In the codling moth *Cydia pomonella* (Lepidoptera, Tortricidae), a reconstruction of the
68 glomerular structure of the AL, combined with electrophysiological recordings, suggested
69 significant cross-talk between the pheromone and general odour subsystems [21].
70 Codling moth is a key pest of apple and its sex pheromone and the behavioural role of
71 host plant volatiles have been carefully studied [22].

72 We investigated the neurophysiological mechanisms regulating the interaction between
73 female sex pheromone and behaviourally active host plant odorants, using functional
74 imaging of the AL and intracellular recordings (IR) of projection neurons (PNs) that
75 transmit olfactory signals to higher brain centres. The finding that the MGC is dedicated to
76 blends of social and environmental odours adds to our understanding of the role of
77 chemosensory cues in premating reproductive isolation and plant-insect ecology. It also
78 provides a new incentive for the refinement of sustainable insect control methods based
79 on behaviour-modifying chemicals.

80 **2. Materials and Methods**

81 **(a) Insects**

82 Experiments were done with 2- to 3-day-old unmated codling moth *Cydia pomonella*
83 (Lepidoptera, Tortricidae) males, which were reared for several generations on an
84 artificial diet (Andermatt Biocontrol, Grossdietwil, Switzerland). The males were kept at
85 70±5% RH, 23°C, under a 16L:8D photoperiod and they were fed with sugar water.

86 **(b) Odor stimuli**

87 Test odours included the main component of codling moth female sex pheromone,
88 codlemone, (*E,E*)-8,10-dodecadienol (>99.6% chemical and isomeric purity, Shin-Etsu
89 Chemical Co., Tokyo) and three plant volatiles, (*E*)- β -farnesene (93.4% pure), butyl
90 hexanoate (97.8%, both from Bedoukian Research Inc., Danbury, USA) and pear ester,
91 (*E,Z*)-2,4-decadienoate (87.4%, Sigma Aldrich).

92 For functional imaging and intracellular recordings, solutions of test compounds in 10 μ l
93 re-distilled hexane were applied on filter paper (0.5 x 1 cm), ca. 1 h before tests. After
94 the solvent evaporated during 1 min, one or two filter papers (compound blends) were
95 inserted into a Pasteur pipette. Codlemone was tested at amounts of 1 ng to 1 μ g, plant
96 compounds from 10 ng to 10 μ g, in decadic steps. A continuous charcoal-filtered and
97 moistened airstream (500 ml/min) passed through a glass tube (10 mm ID) over the
98 antenna. A stimulus controller (SFC-2/b, Syntech, Kirchzarten, Germany) injected a 0.5-s
99 puff (500 ml/min) through the pipettes into this glass tube. Odours were presented in
100 randomized order. Pipettes with filter paper loaded with 10 μ l of solvent were used as
101 control.

102 For behavioural tests, synthetic compounds were released from a piezo sprayer [23].
103 Compound dilutions were delivered at 10 $\mu\text{l}/\text{min}$ to a 20- μl glass capillary tube with a
104 drawn-out tip. A piezo-ceramic disc vibrated the capillary at ca. 100 kHz, producing an
105 aerosol, which evaporated a few cm downwind from the capillary tip at a constant rate
106 and known chemical purity. Codlemone was tested at 0.1 pg/min and plant compounds at
107 1 and 100 pg/min .

108 **(c) Behavioural assay**

109 Wind tunnel experiments were conducted according to Knight *et al.* [24]. A fan pulled air
110 through a charcoal filter, through a series of screens, at 0.25 m/s into the tunnel (1.6 x
111 0.6 x 0.6 m). Exhaust was expelled outside of the building. Room lighting was computer-
112 controlled to gradually decrease during a 60 min dusk period, between full light level
113 (1330 lux) and the dark period (25 lux). Ten batches of five moths were flown
114 consecutively to each lure, during the first 3 h of the scotophase. Male moth behaviour
115 was recorded for up to 6 min. The following types of behaviour were recorded: wing
116 fanning, take-off, upwind flight and contact with the screen. Proportional data were
117 adjusted with Bartlett's correction for small sample size. An angular transformation was
118 used to normalize proportional data prior to analysis of variance (ANOVA) (Statistix 9,
119 Analytical Software, Tallahassee, USA). An α -level of 0.05 was used to establish
120 significance, Tukey's method was used to compare means.

121 **(d) Functional imaging**

122 Individual moths were secured in a 1 ml plastic pipette, with the head protruding from the
123 narrow end, and fixed by dental wax (Surgident, Heraeus Kulzer Inc). The head capsule
124 was opened between the antenna and the eyes; muscle, glands, trachea, neural sheath
125 and the oesophagus were removed to expose the antennal lobes [25]. A calcium sensitive
126 dye (Calcium green-2-AM dye) was dissolved in 20% Pluronic F-127 in dimethyl sulfoxide
127 (Molecular Probes, Eugene, USA) and diluted in moth Ringer solution to 30 μM and then
128 applied to the brain, leaving the preparation in a dark and cold (5°C) environment for 3 h.

129 Recordings were made in vivo after incubation and washing, using an Olympus
130 microscope (20x air objective NA 0.50; filter settings: dichroic 500 nm, emission LP 515
131 nm). The preparation was illuminated at 475 nm. Stimulation started at frame 12 and
132 lasted 1 s. Images were binned twice (320 x 240 pixel) to increase signal-to noise ratio.
133 TILL Photonics imaging software (Gräfelfing, Germany) was used to record sequences of
134 40 frames (4 Hz, 200 ms exposure time) and noise was removed by a Gaussian filter. The
135 response magnitude was calculated as the average $\Delta F/F$ for each frame, where F was
136 estimated using a linear function fitted to the parts of the calcium fluorescence decay
137 curve outside the potential response. The onset of the signal was set to the time of the
138 first frame with a positive average $\Delta F/F$. For statistical analysis, a Kruskal-Wallis test was

139 followed by a Mann-Whitney U test with Holm-Bonferroni correction. A 3-D map of the
140 codling moth AL [21] was used to link the active area to AL glomeruli.

141 **(e) Intracellular recordings**

142 Insect preparation and recordings were done as described by Trona *et al.* [21]. During
143 recordings, the brain was super-fused with a pH 6.9 ringer solution delivered from a flow
144 system. A silver ground electrode was in contact with the ringer solution. Using a
145 micromanipulator, the AL was randomly penetrated with an electrode which was drawn
146 from a heated glass capillary (0.5 mm i.d., Sutter Instrument Co., Novato, USA) with the
147 tip filled with 1% neurobiotin (Vector Labs, Burlingame, USA) dissolved in 0.25 mM KCL
148 and the remaining part was filled with 1 mM KCl.

149 After recordings, the AL interneuron was stained with a depolarizing current (0.5-0.7 nA,
150 15 min). The brain was dissected from the head capsule and stained following the
151 protocol of Trona *et al.* [21]. Stained neurons were viewed in a laser scanning confocal
152 microscope (Zeiss LSM 510, Carl Zeiss, Jena, Germany) with a 40x1.4 oil-immersion DIC
153 objective. Alexa Fluor 488, fluorescein Avidin and Alexa Fluor 546 labelled structures were
154 excited with an argon laser 488 nm (with a 505 nm long-pass filter) and a HeNe laser
155 (with a 560 nm long-pass filter). Stacks of X-Y confocal images (1024 x 1024 pixel) were
156 scanned at 0.7 μ m step size.

157 Only complete recording sessions of the entire set of test stimuli were evaluated.
158 Responses were calculated from the number of net-spikes during 500 ms (number of
159 spikes 500 ms before stimulus onset subtracted from the number of spikes 500 ms after
160 stimulus onset). Net-spikes in response to control were subtracted from the net-spikes in
161 response to odour stimuli; blend responses were considered to be synergistic/suppressive
162 when the number of net-spikes in response to blends was significantly higher/lower than
163 the sum of net-spikes in response to the single compounds (G-test).

164 **Results**

165 **(a) Behavioural assay**

166 Blends of the main sex pheromone component, codlemone, and host plant volatiles
167 attracted significantly more codling moth males than single compounds (figure 1). All
168 three plant volatiles tested, (*E*)- β -farnesene, butyl hexanoate and pear ester, elicited
169 upwind orientation flights. Blending codlemone at 0.1 pg/min and plant volatiles at 100
170 pg/min significantly increased landings at the source, compared to codlemone alone
171 (figure 1).

172 (b) Functional imaging

173 Calcium signals revealed distinct glomerular activity patterns for each odorant tested
174 (figure 2). A threshold dose of codlemone (10 ng) elicited a significant response in the
175 MGC, including the cumulus (Cu) and nearby satellite glomeruli (20 and 37; figure 2b).
176 Plant volatiles alone did not elicit any response in the Cu, they instead activated satellite
177 glomeruli and glomeruli outside the MGC (figure 2c-e). A threshold dose of pear ester
178 (100 ng) was active in the satellite glomeruli 20 and 37, which also responded to
179 codlemone (figure 2c) plus glomerulus 11 outside the MGC.

180 Blends of 10 ng codlemone plus 100 ng of each plant volatile compound produced a
181 strong synergistic interaction in the Cu (figure 3a,e). This synergistic effect was not seen
182 at a 10-fold higher dose (figure 3a). Although several of the glomeruli surrounding the Cu
183 responded to plant volatiles and codlemone (figure 2b-e, 3e), there was no synergistic
184 interaction in these glomeruli: outside the Cu, the activity elicited by blends was
185 significantly lower than the sum of the activity elicited by the single compounds (figure
186 3b-d).

187 (c) Intracellular recordings

188 Figures 4 and 5 show the blend response of AL output neurons. Based on a dose-response
189 test with single compounds (figure 4a), codlemone and individual plant volatiles were
190 combined in a 1:10 ratio and 1:1000 ratio. The number of synergistic, suppressive and
191 additive responses of AL neurons to blends of codlemone and plant volatiles, in the Cu
192 and surrounding glomeruli is shown in figure 4b,c.

193 Analysis of 69 successful recordings demonstrates that odour blend interaction was not
194 merely additive ($p < 0.05$, G-test). Of the neurons showing a synergistic blend response,
195 52% responded to blends only, and not to single compounds. Suppressive responses
196 comprised both a decreased excitatory phase (53%) and complete response suppression
197 (47%) (figure 4b).

198 Twenty-nine neurons were successfully stained: 11 PNs arborizing in the Cu, 5 PNs in
199 satellite glomeruli surrounding the Cu, 10 PNs in glomeruli outside the MGC and, in
200 addition, 3 local interneurons (LNs). The Cu was innervated by uniglomerular PNs (figure
201 5a), and by one multiglomerular PN that also arborized in the satellite glomerulus 20
202 (figure 4d). Spike frequency histograms for selected PNs in response to compound blends
203 are shown in figure 5. A statistical comparison of the blend effects in stained PNs revealed
204 a significant difference: synergism occurred almost exclusively in the Cu, while blend
205 stimulation of glomeruli outside the MGC mostly had a additive or suppressive effect
206 (figures 4c, 5c).

207 Discussion

208 (a) Neural ensemble coding of sex pheromone and host plant odour in the MGC 209 of the male moth AL

210 Understanding how stimulation with a blend of odorants generates a unique perception in
211 the brain is a current research question. What adds to the complexity of olfactory coding
212 is the integration of separate, independent signals - sex and habitat odours - which are
213 together required to generate appropriate behavioural responses during mate-finding.

214 We combined functional imaging and intracellular recordings to study odour blend
215 processing in the codling moth *C. pomonella*, and show that the behavioural synergism
216 between sex pheromone and host plant odourants is mirrored neurophysiologically. The
217 MGC in the AL integrates signals from conspecific insects with habitat odours and
218 synergistic interactions between these two classes of odours occur both at the input and
219 output level. This demonstrates that processing of sex pheromone and plant volatiles,
220 which insects encounter as an ensemble in nature, does not employ functionally separate
221 pathways [17,18].

222 Blend enhancement and suppression in the AL may stem from odour interference in
223 antennal sensory neurons [19,26] and ultimately at the olfactory receptor level [27].
224 However, in codling moth, pheromone-plant volatile blends enhance the Cu response
225 while they simultaneously suppress surrounding glomeruli in a "center-surround" fashion.
226 Such complex coding may instead rely on lateral excitatory or inhibitory interconnections
227 between glomeruli through local interneurons (LNs) [2,28]. Functional studies of LNs will
228 be essential to understand olfactory processing in the AL.

229 Intracellular recordings of PNns, which connect the AL to higher brain centres, further
230 corroborate that the MGC processes blends of plant volatiles and sex pheromone.
231 Synergistic, blend-specific responses have been shown in the silk moth *B. mori* [17] and
232 in codling moth, where PNns innervate the Cu and satellite glomeruli of the MGC [21].

233 An antagonistic interaction modality was shown in the black cutworm *A. ipsilon*. A floral
234 volatile, which inhibits male attraction to pheromone, suppresses the pheromone
235 response in the AL [18] and in PNns innervating the MGC [19]. This suggests that odours
236 with different ecological roles may differently affect pheromone coding. A wiring diagram
237 of input and output signals in the codling moth AL, based on a more complete panel of
238 ecologically relevant odorants, from host and non-host plants or associated mutualistic
239 microorganisms [29,30], will reveal whether glomerulus morphology and position in the
240 AL correlates with the behavioural role of the respective key stimuli [31].

241 **(b) Behavioural and ecological physiology of pheromone-plant odour blend**
242 **perception**

243 Mate recognition in insects, and especially in habitat-specific plant-feeding species,
244 involves two main elements: sexual communication and recognition of larval and adult
245 food plants, which frequently serve as rendezvous sites. Both mate and host finding
246 largely rely on olfactory signals [14,32] which play a fundamental role in speciation
247 [6,33].

248 In the codling moth, host plant odour is part of the mate finding signal. The plant volatiles
249 chosen for this study are distinctive for the main hosts pear and apple, respectively. They
250 mediate female attraction for oviposition [29,34-37] and they synergize male attraction to
251 female sex pheromone. The MGC, in the olfactory centre of the moth brain, is the focal
252 point for processing blends of pheromone and these plant signals.

253 Speciation is thought to be facilitated by multiple-effect or "magic" traits, which are subject
254 to divergent selection and which contribute to nonrandom mating [9,10]. The MGC
255 interconnects mate and host choice and would accordingly be considered as a multiple-
256 effect trait. Host choice seemingly is under divergent selection in codling moth, which
257 forms distinct host races on apple, pear, walnut, plum and apricot. These differ in spring
258 emergence and diapause initiation, in close association with host flowering and fruit
259 maturation [38,39], and the genetically distinct walnut strain is adapted to toxic walnut
260 metabolites [40-42]. Females of several strains preferentially oviposit on their respective
261 host fruit [29,38].

262 A comparison of the female sex pheromones of closely related *Cydia* species further
263 corroborates the role of plant volatiles in reproductive isolation. Only few species share
264 the same pheromone, but these all feed on host plants belonging to different families. For
265 example, pea moth *C. nigricana* (Leguminosae) and pear moth *C. pyrivora* (*Pyrus*), the
266 sibling species of codling moth, use codlemone acetate (*E,E*)-8,10-dodecadienyl acetate,
267 which is a strong pheromone antagonist in codling moth males [43].

268 Pheromone and host odour communication is highly integrated also in other insects, for
269 example in *Drosophila* [44] and in bark beetles, where non-host volatiles, as opposed to
270 host volatiles, have an antagonistic effect on host and mate finding [45]. In the two
271 pheromone races of the European corn borer *Ostrinia nubilalis*, male preference for
272 females of the same race leads to premating isolation [46,47], which is reinforced by
273 preferential attraction to volatiles of their respective host plants, mugwort and maize
274 [48,49].

275 Ecological speciation, following host plant shifts, has likely contributed to the remarkable
276 diversity of phytophagous insects [11,33]. Our study provides physiological data that
277 suggest that mate recognition systems evolve in concert with chemosensory adaptation to

278 new hosts and ecological niches, and that sexual selection cannot be separated from
279 natural selection in male insect herbivores.

280 **(c) Practical implication**

281 Our knowledge of codling moth chemical ecology has led to the successful development of
282 species-specific and safe population control by pheromone-mediated mating disruption. In
283 spite of orchard applications on 200.000 ha [50], the behavioural mechanisms underlying
284 the disruption of mating are still under debate [51,52] and a better understanding of
285 them will give leads for improvement. Our study demonstrates that it will be useful to
286 consider the physiological and behavioural effect of plant volatiles on mating disruption,
287 since, in nature, pheromone and plant volatiles are perceived together.

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452 Figure Legends

453 **Figure 1.** Wind tunnel attraction of codling moth *C. pomonella* males ($n=50$) to the main
454 pheromone compound codlemone (released at 0.1 pg/min) and to plant volatiles butyl
455 hexanoate (*a*), (*E*)- β -farnesene (*b*), pear ester (*c*), at 1 pg/min and 100 pg/min. Grey
456 lines show attraction to 1:1000 blends of codlemone with these plant volatiles. Landings
457 at the source are significantly increased in response to each of these 2-component blends,
458 compared to pheromone alone (** $p < 0.001$, two-way ANOVA; butyl hexanoate
459 $F(4,45)=45.0$, β -farnesene $F(4,45)=23.75$, pear ester $F(4,45)=24.08$). Empty circles in
460 the codlemone response curve show significant differences between codlemone and single
461 plant volatiles alone ($p < 0.0001$, two-way ANOVA; butyl hexanoate $F(4,45)=23.35$, β -
462 farnesene $F(4,45)=53.96$, pear ester $F(4,45)=20.68$).

463 **Figure 2.** Calcium imaging of the codling moth male AL upon stimulation with single
464 odorants, sex pheromone (codlemone) and three plant volatiles. Dose-response
465 relationships of odor-evoked calcium signals, using an increasing dose of codlemone
466 ($n=19$), pear ester ($n=23$), β -farnesene ($n=14$) and butyl hexanoate ($n=19$) (*a*).
467 Glomerular activation patterns in response to 10 ng codlemone (*b*), to 100 ng of pear
468 ester (*c*), (*E*)- β -farnesene (*d*) and butyl hexanoate (*e*), respectively and in response to
469 the solvent (hexane) (*f*). Data points show means and standard errors (SEMs), glomeruli
470 numbers correspond to the 3D atlas of the codling moth AL [26].

471 **Figure 3.** Calcium imaging of the codling moth male AL following stimulation with 2-
472 component blends of sex pheromone (codlemone) and plant volatiles, butyl hexanoate,

473 pear ester and β -farnesene. Odour-evoked activity was measured in the cumulus (Cu)
474 and other responding glomeruli. Response in the Cu (*a*), showing a synergistic blend
475 interaction for 10:100 ng blends ($*p < 0.05$, $**p < 0.01$, Kruskal-Wallis test followed by
476 Mann-Whitney U-test with Holm-Bonferroni correction, $n = 30$ males). At a higher dose,
477 blends (100:1000 ng) were not significantly different from codlemone ($p = 0.36$, Kruskal-
478 Wallis test, $n = 30$ males). Response of glomeruli outside the cumulus (*b-d*) to plant
479 compounds, codlemone, their blends and the summed responses to single compounds
480 (Σ): butyl hexanoate, satellite glomerulus 20 and glomerulus 23 ($*p < 0.05$ and $**p < 0.01$,
481 $n = 26$) (*b*); pear ester, satellite glomeruli 20, 37 ($*p < 0.05$, $n = 30$) (*c*); β -farnesene,
482 satellite glomeruli 20, 21 ($***p < 0.001$ and $*p < 0.05$, one-sided t-test, $n = 31$) (*d*). Bars
483 show the standard error of the mean (SEM). Representative recording of codlemone, pear
484 ester and their blend (*e*). Glomeruli numbers correspond to the atlas of codling moth AL
485 [26].

486 **Figure 4.** Responses of AL neurons to single compounds and binary blends. Intracellular
487 recordings of AL neurons with increasing doses of codlemone ($n = 12$), butyl hexanoate
488 ($n = 10$), pear ester ($n = 11$) and β -farnesene ($n = 12$) (*a*). Histograms of synergistic,
489 suppressive and additive responses of 69 physiologically characterized interneurons to
490 blends of codlemone and plant volatiles (*b*). Number of synergistic, suppressive and
491 additive responses of neurons innervating Cu and glomeruli outside the MGC ($**p < 0.005$,
492 Chi2-test) (*c*). 3D-reconstruction of a multiglomerular PN innervating the Cu and the
493 satellite glomerulus 20, showing a synergistic response to a blend of codlemone and (E)-
494 β -farnesene. The horizontal bar shows the stimulus period (500 ms) (*d*).

495 **Figure 5.** Single confocal sections and spike frequency histograms (spikes/s) of
496 physiologically and morphologically characterized PNs in the codling moth male AL.
497 Synergistic responses of a PN innervating the Cu to blends of codlemone with pear ester
498 and β -farnesene (*a*). Synergistic responses of a multiglomerular PN, innervating the
499 satellite glomeruli 20 and 37, to blends of codlemone with pear ester and butyl hexanoate
500 (*b*). Suppressive responses of a PN innervating the glomerulus 14, to a blend of
501 codlemone and (E)- β -farnesene at different blend ratios (*c*). Confocal sections: entrance
502 of the antennal nerve (arrowheads), depth from anterior side of the AL (Z), scale bars (50
503 μm), glomeruli numbers correspond to the 3D AL atlas [Trona 2010]. Histograms:
504 stimulus period (bars, 500 ms).









