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1 Origin of wiring specificity in an olfactory map: 2 dendrite targeting of projection neurons

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4 Kenneth Kin Lam Wong¹, Tongchao Li^{1,9}, Tian-Ming Fu^{2,†}, Gaoxiang Liu³, Cheng Lyu¹, Sayeh
5 Kohani¹, Qijing Xie¹, David J Luginbuhl¹, Srigokul Upadhyayula^{3,4,5}, Eric Betzig^{2,3,6}, Liquan
6 Luo^{1*}

7
8 ¹Department of Biology, Howard Hughes Medical Institute, Stanford University, Stanford,
9 United States

10 ²Howard Hughes Medical Institute, Janelia Research Campus, Ashburn, United States

11 ³Advanced Bioimaging Center, Department of Molecular and Cell Biology, University of
12 California, Berkeley, United States

13 ⁴Molecular Biophysics and Integrated Bioimaging Division, Lawrence Berkeley National
14 Laboratory, Berkeley, United States

15 ⁵Chan Zuckerberg Biohub, San Francisco, United States

16 ⁶Departments of Molecular and Cell Biology and Physics, Howard Hughes Medical Institute,
17 Helen Wills Neuroscience Institute, University of California, Berkeley, United States

18 *For correspondence: lluo@stanford.edu (L.L.)

19
20 ⁹Present Address:

21 ^aDepartment of Neurobiology and Department of Neurosurgery of Second Affiliated Hospital,
22 Zhejiang University School of Medicine, Hangzhou, P.R. China

23 ^bLiangzhu Laboratory, MOE Frontier Science Center for Brain Science and Brain-machine
24 Integration, State Key Laboratory of Brain-machine Intelligence, Zhejiang University, 1369
25 West Wenyi Road, Hangzhou 311121, China

26 ^cNHC and CAMS Key Laboratory of Medical Neurobiology, Zhejiang University, Hangzhou
27 310058, China

28 [†]Present Address: Princeton University, Princeton, United States

29 ABSTRACT

30 How does wiring specificity of neural maps emerge during development? Formation of the adult
31 *Drosophila* olfactory glomerular map begins with patterning of projection neuron (PN) dendrites
32 at the early pupal stage. To better understand the origin of wiring specificity of this map, we
33 created genetic tools to systematically characterize dendrite patterning across development at PN
34 type-specific resolution. We find that PNs use lineage and birth order combinatorially to build
35 the initial dendritic map. Specifically, birth order directs dendrite targeting in rotating and binary
36 manners for PNs of the anterodorsal and lateral lineages, respectively. Two-photon- and
37 adaptive optical lattice light-sheet microscope-based time-lapse imaging reveals that PN
38 dendrites initiate active targeting with direction-dependent branch stabilization on the timescale
39 of seconds. Moreover, PNs that are used in both the larval and adult olfactory circuits prune their
40 larval-specific dendrites and re-extend new dendrites simultaneously to facilitate timely olfactory
41 map organization. Our work highlights the power and necessity of type-specific neuronal access
42 and time-lapse imaging in identifying wiring mechanisms that underlie complex patterns of
43 functional neural maps.

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47 **INTRODUCTION**

48
49 Organization of neuronal connectivity into spatial maps occurs widely in the nervous systems
50 across species (Luo & Flanagan, 2007; Cang & Feldheim, 2013; Luo, 2021). For example, in the
51 retinotopic map of the visual system, neighboring neurons in the input field project axons to
52 neighboring neurons in the target field (Cang & Feldheim, 2013). Such a continuous
53 organization preserves spatial relationships of the visual world. Contrary to retinotopy, the
54 olfactory glomerular map consists of discrete units called glomeruli in which input neurons
55 connect with the cognate output neurons based on neuronal type rather than soma position
56 (Mombaerts et al., 1996; Gao et al., 2000; Vosshall et al., 2000). This discrete map represents a
57 given odor by the combinatorial activation of specific glomeruli. Whereas continuous maps are
58 readily built using gradients of guidance cues (Cang & Feldheim, 2013), how glomeruli are
59 placed at specific locations in discrete maps is less clear (Murthy, 2011). Understanding the
60 developmental origins of these neural maps is fundamental for deciphering the logic of their
61 functional organization through which information is properly represented and processed.

62 The adult *Drosophila* olfactory map in the antennal lobe (equivalent of the vertebrate
63 olfactory bulb) has proven to be a powerful model for studying mechanisms of wiring
64 specificity, thanks to the type-specific connections between the presynaptic olfactory receptor
65 neurons (ORNs) and the cognate postsynaptic projection neurons (PNs). Molecules and
66 mechanisms first identified in this circuit have been found to play similar roles in the wiring of
67 the mammalian brain (e.g., Hong et al., 2012; Berns et al., 2018; Pederick et al., 2021).
68 Assembly of the fly olfactory map begins with dendritic growth and patterning of PNs derived
69 primarily from the anterodorsal (adPNs) and lateral (lPNs) lineages and born with an invariant
70 birth order within each lineage (Jefferis et al., 2001, 2004; Marin et al., 2005; Yu et al., 2010;
71 Lin et al., 2012) (**Figures 1A and B**). This patterning creates a prototypic olfactory map, prior to
72 ORN axon innervation, indicative of the PN-autonomous ability to target dendrites into specific
73 regions. However, earlier studies could only unambiguously follow the development of one
74 single PN type – DL1 PNs (Jefferis et al., 2004). It remains unclear to date how the prototypic
75 olfactory map is organized and what cellular mechanisms PN dendrites use to achieve targeting
76 specificity (**Figure 1C₁₋₂**). The initial map formation is further complicated by circuit
77 remodeling during which embryonic-born PNs used in both the larval and adult circuits need to
78 reorganize their neurites (Marin et al., 2005). How embryonic-born PNs coordinate remodeling
79 with re-integration into the adult circuit is not known (**Figure 1C₃**).

80 Here, we set out to explore the origin of the olfactory map by performing a systematic
81 and comparative study of PN dendrite development at type-specific resolution *in vivo*, and two-
82 photon- and adaptive optical lattice light-sheet microscope-based time-lapse imaging of PN
83 dendrites in early pupal brain explants. Our study uncovers wiring logic that directs PN dendrites
84 to create an organized olfactory map, dendritic branch dynamics that lead to directional
85 selectivity, and a novel re-wiring mechanism that facilitates timely olfactory map formation.
86 These wiring strategies used in the initial map organization lay the foundation of precise synaptic
87 connectivity between PNs and ORNs in the final glomerular map.

88
89 **RESULTS**

90
91 **Overview of *Drosophila* olfactory circuit development at lineage-specific resolution**

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92 We first described the development of *Drosophila* olfactory circuit using pupal brains double-
93 labeled for adPNs and IPNs (**Figure 1D**; see genetic design in **Figure 2**). At the onset of
94 metamorphosis (0 hours after puparium formation; 0h APF), the adult-specific antennal lobe
95 (also referred to as ‘developing antennal lobe’) remained relatively small, located dorsolateral
96 and posterior to the larval-specific antennal lobe (also referred to as ‘degenerating antennal
97 lobe’) (**Figure 1D₁**). As PN dendrites continued to grow and innervate the developing antennal
98 lobe, its size increased considerably (**Figure 1D₁₋₃**). By 12h APF, PNs already appeared to be
99 sorting their dendrites into specific regions to form a prototypic map, as revealed by the
100 heterogeneous patterning of IPN dendrites (arrowheads in **Figure 1D₃**). From 21h to 50h APF,
101 dendrites of adPNs and IPNs gradually segregated and eventually formed intercalated but non-
102 overlapping glomeruli (**Figure 1D₄₋₅**). The development of the adult-specific antennal lobe
103 partially overlapped with the degeneration of the larval-specific antennal lobe, as indicated by
104 fragmentation of the larval-specific dendrites of embryonic-born PNs at 3h APF (**Figure 1D₂**).
105 This gross characterization at the resolution of two PN lineages was consistent with earlier
106 studies (Jefferis et al., 2004; Marin et al., 2005). However, the resolution was not sufficiently
107 high to answer the questions we raised in Introduction (**Figure 1C**).
108

109 **Expanded genetic toolkit for type-specific labeling of PNs during early pupal development**
110 To reveal how PN dendrites initiate olfactory map formation at high spatiotemporal resolution,
111 we needed genetic access to specific PN types during early pupal development. From our
112 recently deciphered single-cell PN transcriptomes (Xie et al., 2021), we searched for genetic
113 markers that are expressed strongly and persistently in single or a few PN types across pupal
114 development. This transcriptome-instructed search led to identification of *CR45223* (in place of
115 this non-coding gene, we used the adjacent *CG14322* that exhibits nearly identical expression
116 pattern), *lov*, and *tsh* (**Figures 2A and B; Figure 2 – figure supplement 1**).

117 Next, using CRISPR/Cas9, we generated knock-in transgenic QF2 expression driver lines
118 in which *T2A-QF2* (or *T2A-FLP* for intersection) was inserted immediately before the stop
119 codon of the endogenous gene (**Figure 2 – figure supplement 2**). The self-cleaving peptide T2A
120 allows QF2 to be expressed in the same pattern as the endogenous gene (Diao & White, 2012).
121 With these new *QF2* lines together with existing *GAL4* lines that label additional PN types (Xie
122 et al., 2019), we now have an expanded toolkit accessing PNs ranging from early- to late-born
123 PNs, from adPN to IPN lineages, and from PNs with neighboring glomerular projections to those
124 with distant projections in the adult antennal lobe (**Figures 2C and D**). As *QF2/QUAS* and
125 *GAL4/UAS* expression systems operate orthogonally to each other (Potter et al., 2010), we
126 crossed our *QF2* lines with existing *GAL4* lines for simultaneous labeling of distinct PN types in
127 the same brain (see inset in **Figure 2C**). This combinatorial use of driver lines permitted
128 comparative analyses of development of distinct PN types with minimal biological and technical
129 variations.

130 To limit driver expression only in PNs, we applied intersectional logic gates (AND and
131 NOT gates) using our newly generated conditional reporters genetically encoding either
132 mGreenLantern, Halo tags, and/or SNAP tags (Kohl et al., 2014; Sutcliffe et al., 2017; Campbell
133 et al., 2020) (**Figures 2E and F; Figure 2 – figure supplement 3**). These reporters can be
134 broadly used in other systems. Finally, we used MARCM (Lee & Luo, 1999) to label PNs that
135 remain inaccessible due to lack of drivers (**Figure 2C**; discussed in **Figure 3**).
136

137 **Early larval-born adPN dendrites initially share similar targeting regions**

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138 Using the new genetic tools, we first re-visited the dendrite development of DL1 PNs—the first
139 larval-born adPN type—using pupal brains double-labeled for DL1 PNs (labeled by *71B05-*
140 *GAL4*) and adPNs (**Figure 3A**). Consistent with our previous study (Jefferis et al., 2004), DL1
141 PNs already showed robust dendritic growth at the wandering third instar larval stage (**Figure 3**
142 – **figure supplement 1A**). At 0h APF, DL1 PN dendrites extended radially outwards from the
143 main process, reaching nearly the entire developing antennal lobe and often overshooting it
144 (white arrowheads in **Figure 3A₁**), likely surveying the surroundings. By 6h APF, most of the
145 dendrites already occupied the dorsolateral (DL) corner of the antennal lobe (**Figure 3A₂**). As
146 the antennal lobe continued to grow, this dorsolateral positioning of the DL1 PN dendrites
147 remained largely unchanged (**Figure 3A₃₋₆**). From 21h APF onwards, the dendrites underwent
148 progressive refinement: they were restricted into a smaller area by 30h APF (**Figure 3A₄₋₅**), and
149 eventually formed a compact, posterior glomerulus by 50h APF (**Figure 3A₆** showing a single *z*
150 section).

151 To assess whether other PN types follow the same developmental trajectory, we next
152 examined *CG14322+* PNs, which include DL1 PNs and DA3 PNs—the first and second larval-
153 born adPN types, respectively. In the same brain, we also labeled with a different fluorophore
154 DC2 PNs—the third larval-born adPN type (**Figure 3B**). The dendritic pattern of DL1/DA3 PNs
155 appeared indistinguishable from that of DL1 PNs from 0h to 12h APF (compare yellow channel
156 of **Figure 3B₁₋₃** with **Figure 3A₁₋₃**), suggesting that DL1 and DA3 PN sent dendrites to the same
157 region in the antennal lobe. We began to see differences in 21h APF pupal brains in which
158 DL1/DA3 PN dendrites not only occupied the dorsolateral region but also spread ventrally
159 (white arrowhead in **Figure 3B₄**; compare with **Figure 3A₄**). The more ventrally targeted
160 dendrites likely belong to DA3 PNs. This suggests that ~21h APF marks the beginning of
161 dendritic segregation of DL1 and DA3 PNs. By 30h APF, DL1 and DA3 dendrites were clearly
162 separable (**Figure 3B₅**), which respectively formed more posteriorly and anteriorly targeted
163 glomeruli at 50h APF (**Figure 3B₆**; see single *z* sections in **Figure 3 – figure supplement 1C**).
164

165 Next, we focused on the third-born—DC2 PNs labeled by *91G04-GAL4* (**Figure 3B**).
166 This *GAL4* labeled additional embryonic-born adPNs from 0h to 6h APF, but the expression in
167 these PNs diminished afterwards. As embryonic-born adPNs do not have any dendrites in the
168 developing antennal lobe at 0h APF (discussed in **Figure 7**), dendrites found in the antennal lobe
169 should belong to the larval-born DC2 PNs. Like DL1/DA3 PNs, DC2 PNs initiated radial
170 dendritic extension across the antennal lobe at 0h APF (**Figure 3B₁**; **Figure 3 – figure**
171 **supplement 1B**). Notably, DL1/DA3 and DC2 PN dendrites exhibited substantial overlap from
172 0h to 12h APF and shared a similar targeting region at the dorsolateral corner from 6h to 12h
173 APF (**Figure 3B₁₋₃**). It was not until 21h APF that DL1, DA3, and DC2 dendrites began to
174 segregate from each other along both medial-lateral and anterior-posterior axes (**Figure 3B₄₋₅**).
175 By 50h APF, DC2 glomerulus was separated from DL1/DA3 glomeruli by intermediate
176 glomeruli (**Figure 3B₆**).

177 In summary, dendrites of consecutively larval-born DL1, DA3, and DC2 adPNs (here
178 collectively named ‘early larval-born adPNs’) develop in a similar fashion and share a similar
179 targeting region at early pupal stages (0–12h APF). This is then followed by their segregation
180 into distinct regions close to their adult glomerular positions during mid-pupal stages (21–50h
APF).

181

182 **Larval-born adPNs with distant birth order send dendrites to distinct regions**

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183 The analysis of early larval-born adPNs (**Figures 3A and B**) led us to hypothesize that larval-
184 born adPNs might use their birth order to coordinate dendrite targeting during early pupal stages.
185 If this were true, we would expect dendrites of larval-born adPNs with distant birth order to
186 occupy distinct regions. To test this hypothesis, we compared dendrite targeting regions of early
187 larval-born adPNs with those of later-born adPNs.

188 We first examined DC3/VA1d adPNs (referred to as ‘early-middle larval-born adPNs’)
189 using *Mz19-GAL4* (**Figure 3C**). This *GAL4* is expressed in 3 PN types from 24h APF to
190 adulthood: DC3 adPNs, VA1d adPNs, and DA1 IPNs (Jefferis et al., 2004). To distinguish
191 adPNs from IPNs, we previously adopted a FLP-out strategy labeling *Mz19*⁺ PNs with either
192 GFP or RFP based on their lineages and studied dendrite segregation and refinement during mid-
193 pupal stages (Li et al., 2021) (**Figure 3C4-7**). However, the weak *GAL4* expression before 24h
194 APF prevented us from visualizing any dendrites at earlier stages. To overcome this, we
195 incorporated Halo and SNAP chemical labeling (Kohl et al., 2014) in place of the
196 immunofluorescence approach. This modification substantially extended the detection to
197 developmental stages as early as 12h APF (**Figure 3C1**). We found that, from 12h to 21h APF,
198 DC3/VA1d PN dendrites targeted the ventrolateral (VL) corner of the antennal lobe (**Figure**
199 **3C1-4**). Thus, early (DL1/DA3/DC2) and early-middle (DC3/VA1d) larval-born adPN dendrites
200 occupy distinct regions at 12h APF.

201 As we did not have reliable drivers to access other later-born PNs at early pupal stages,
202 we turned to MARCM (Lee & Luo, 1999) to generate heat shock-induced single-cell clones of
203 PNs born at different times (**Figure 3 – figure supplement 2**). We used *GH146-GAL4(IV)*, a PN
204 driver that labels the majority of PN types, including later-born adPNs (**Figure 3 – figure**
205 **supplement 2D-E**), with a tight temporal control of heat shock and analyzed heat shock-induced
206 animals that were among the first to form puparium to minimize the effects of unsynchronized
207 development among individual animals (see **Materials and Methods** for details). These
208 optimizations permitted a systematic clonal analysis at higher PN type-specific resolution that
209 correlates with birth time. Applying heat shocks at specific times, we accessed four groups of
210 larval-born adPNs: (1) first-born (DL1), (2) early-born (DL1, DA3, DC2 and D), (3) middle-born
211 (VM7v, VM7d, VM2, DM6 and VA1v), and (4) late-born (DM6, VA1v, DL2v, DL2d). We note
212 that DM6 and VA1v PNs belonged to both groups of middle- and late-born adPNs, reflecting the
213 nature of short birth timing differences among these later-born PNs. Using this strategy, we
214 could also label IPNs born at different times (**Figure 3 – figure supplement 2F**).

215 Clonal analysis revealed that, at 12h APF, the first-born DL1 adPNs sent dendrites to the
216 dorsolateral corner of the antennal lobe as expected (**Figure 3D1-3**). By contrast, dendrites of
217 middle larval-born adPNs occupied a large region on the medial/dorsomedial (M/DM) side
218 (**Figure 3D4-6**). The dendritic arborization patterns of these PNs varied widely, most likely
219 because they belonged to different PN types. Intriguingly, late larval-born adPN dendrites
220 targeted the peripheral, dorsomedial (abbreviated as pDM) corner where the staining of the pan-
221 neuropil marker N-Cadherin was relatively weak (**Figure 3D7-9**). The weak staining on the
222 periphery implies that this area is less populated by PN dendrites (the major constituent of the
223 antennal lobe neuropil at this stage), possibly because (1) this area is not innervated by many
224 PNs and/or (2) the dendrites of late-born PNs innervate later and remain less elaborate than
225 earlier-born PNs (we will explore this later).

226 Together, our data (**Figure 3A-D**) suggest that larval-born adPNs with adjacent birth
227 order send dendrites to similar regions of the developing antennal lobe whereas those with
228 distant birth order send dendrites to distinct regions (**Figure 3E_{2,3}**). We therefore propose that, at

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229 early pupal stages, larval-born adPNs are grouped by their birth order, and those belonging to the
230 same group share similar dendrite targeting specificity (**Figure 3E1**). Notably, the birth order of
231 the examined PNs does not specify dendrite targeting randomly (**Figure 3E4**). Rather, the
232 stereotyped dendritic pattern in the prototypic map correlates with the birth order in an organized
233 manner (rotating clockwise in the right hemisphere when viewed from the front; anti-clockwise
234 in the left: early↔ DL; early-middle↔VL; middle↔M/DM; late↔pDM). One can therefore infer
235 at least the approximate birth order of a larval-born adPN based on its initial dendrite targeting,
236 and *vice versa*.

237

238 **Timing of larval-born adPN dendrite targeting depends on birth order**

239 Having provided evidence for birth order-dependent spatial patterning of larval-born adPN
240 dendrites, we next asked whether the timing of dendritic extension and targeting is also
241 influenced by birth order. We noticed that the extent of dendritic innervation of 0h APF first-
242 born DL1 adPNs resembled that of 6h APF middle-born adPNs (compare **Figure 3 – figure**
243 **supplement 3A₁₋₄** with **Figure 3 – figure supplement 3B₅₋₈**). Such a resemblance was also seen
244 between 0h APF middle-born and 6h APF late-born adPNs (compare **Figure 3 – figure**
245 **supplement 3B₁₋₄** with **Figure 3 – figure supplement 3C**). Quantitative analyses of the
246 exploring volume of dendrites and the number of terminal branches showed that, at 0h APF, DL1
247 PN dendrites were more elaborate than middle-born PN dendrites (**Figure 3 – figure**
248 **supplement 3F**). By 6h APF, the middle-born appeared to catch up, showing an extent of
249 innervation comparable to DL1 PNs.

250 We next examined when the dendrites reach their targeting regions. We found that
251 whereas early larval-born adPNs (DL1, DA3, DC2) concentrated their dendrites to the
252 dorsolateral corner by 6h APF (**Figure 3B₂**; **Figure 3 – figure supplement 3A₅₋₈**), later-born
253 PNs concentrated their dendrites to the medial/dorsomedial or peripheral dorsomedial side at 12h
254 APF (**Figure 3D₄₋₉**; **Figure 3 – figure supplement 3B_{5-8-C}**). Thus, our results suggest larval-
255 born adPN dendrites innervate and pattern the antennal lobe using a ‘first born, first developed’
256 strategy.

257

258 **Contribution of lineage to early PN dendritic patterning**

259 Both lineage and birth order of PNs contribute to the eventual glomerular choice of their
260 dendrites (Jefferis et al., 2001). What is the involvement of lineage in the prototypic map
261 formation? Do IPN dendrites pattern the developing antennal lobe following similar rules as
262 adPNs? To characterize IPN dendrite development at type-specific resolution, we used *tsh-GAL4*
263 to genetically access DA1/DL3 IPNs, and MARCM clones of IPNs as a complementary approach
264 (**Figure 4**). We focused on the dendritic patterns of *tsh*+ DA1/DL3 IPNs from 0h to 12h APF as
265 *tsh-GAL4* labeled additional PNs from 21h APF onwards (**Figure 4A₄₋₆**; **Figure 4 – figure**
266 **supplement 1B₄₋₆**; **Figure 4 – figure supplement 2**; **Figure 2 – figure supplement 1**).

267 Examination of pupal brains double-labeled with DA1/DL3 IPNs (referred to as ‘middle
268 larval-born IPNs’) and DL1/DA3 adPNs revealed that, like the early larval-born adPNs, dendritic
269 growth of DA1/DL3 IPNs was evident by the wandering third instar larval stage (**Figure 4 –**
270 **figure supplement 1A**). At this stage, most DA1/DL3 IPN dendrites innervated the antennal
271 lobe and intermingled with those of DL1/DA3 adPNs. From 0h to 12h APF, despite a high
272 degree of overlap among those dendrites that explored the surroundings, DA1/DL3 IPN dendrites
273 primarily targeted an area ventrolateral to those of DL1/DA3 adPNs (**Figure 4A₁₋₃**). Such a
274 spatial distinction was also observed between middle larval-born adPNs and IPNs in 0h and 6h

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275 APF pupal brains where occasionally single-cell clones from both lineages were simultaneously
276 generated by MARCM (**Figure 3 – figure supplement 3D_{1-4, 7-10}**). Thus, at least some adPNs
277 and IPNs sort their dendrites into distinct regions very early on regardless of birth timing.

278 Next, we used MARCM to ask if IPNs born earlier and later than DA1/DL3 IPNs would
279 send dendrites to regions different from that of DA1/DL3 IPNs. We found that dendrites of early-
280 born IPNs primarily occupied the medial/dorsomedial side of the antennal lobe (**Figure 4B₁₋₃**);
281 we note that adPNs born at the same time sent dendrites to the dorsolateral side (see yellow
282 arrowhead in **Figure 4B₃**). Also, in contrast to the ventrolateral targeting of middle-born IPN
283 dendrites, late-born IPNs sent dendrites to the dorsomedial corner (**Figure 4B₄₋₆**). Like larval-
284 born adPNs, late-born IPNs innervated the antennal lobe later than earlier-born IPNs (**Figure 3 –**
285 **figure supplement 3D₇₋₁₂–E, G**).

286 These data suggest that, at early pupal stages, IPN dendrites pattern the developing
287 antennal lobe following similar rules as larval-born adPNs: grouping by birth order; adjacent
288 birth order → similar dendrite targeting; distant birth order → distinct dendrite targeting; ‘first
289 born, first developed’. However, unlike the correlation of birth order and target positions in a
290 rotational manner for adPNs (**Figure 3E**), the IPN dendritic map formation appears binary:
291 early↔M/DM; middle↔VL; late↔DM (**Figure 4C**). Our type-specific characterization
292 corroborated with the gross examination of the IPN dendrites as previously reported (Jefferis et
293 al., 2004): at 12h APF, IPN dendrites mostly occupied the opposite corners along the
294 dorsomedial-ventrolateral axis, leaving the middle of the axis largely devoid of IPN dendrites
295 (arrowheads in **Figure 1D₃**).

296 In summary, lineage and birth order of larval-born PNs contribute to their dendrite
297 targeting in a combinatorial fashion (**Figure 4D**). The wiring logic of PN dendrites in the
298 developing antennal lobe can therefore be represented by [lineage, birth order] = dendrite
299 targeting; one can deduce the unknown if the other two are known.

300 **An explant system for time-lapse imaging of PN development at early pupal stages**

301 So far, we have identified wiring logic governing the initial dendritic map formation (**Figures 3**
302 and **4**). To examine dendrite targeting at higher spatiotemporal resolution, we established an
303 early-pupal brain explant culture system based on previous protocols (Özel et al., 2015;
304 Rabinovich et al., 2015; Li and Luo, 2021; Li et al., 2021), and performed single- or dual-color
305 time-lapse imaging with two-photon microscopy as well as adaptive optical lattice light-sheet
306 microscopy (AO-LLSM) (**Figure 5A–C**). The following lines of evidence support that our
307 explant system recapitulates key features of *in vivo* olfactory circuit development.

308 First, during normal development, the morphology of the brain lobes changes from
309 spherical at 0h APF to more elongated rectangular shapes at 15h APF (Rabinovich et al., 2015).
310 After 22h *ex vivo* culture, the spherical hemispheres of brains dissected at 3h APF became more
311 elongated, mimicking ~15h APF *in vivo* brains characterized by the separation of the optic lobes
312 from the central brain (**Figure 5D**).

313 Second, dual-color, two-photon imaging of PNs every 20 min for 22h revealed that IPNs
314 in 3h APF brains initially produced dynamic but transient dendritic protrusions in many
315 directions, followed by extensive innervation into the antennal lobe (arrowheads in **Figure 5E₁₋₃**;
316 **Figure 5 – video 1**). In higher brain centers, IPN axons clearly showed direction-specific
317 outgrowth of collateral branches into the mushroom body calyx as well as forward extension into
318 lateral horn (arrows in **Figure 5E₃**), thus resembling *in vivo* development (**Figure 1 – figure**
319 **supplement 2**).

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321 Third, larval-specific dendrites observed in 0h APF brains cultured for 12h *ex vivo*
322 (orange arrowhead in **Figure 5F4**) were no longer seen in those cultured for 24h *ex vivo* (**Figure**
323 **5F5**), indicative of successful pruning and clearance of larval-specific dendrites. Also, the size of
324 the developing antennal lobe in the brains cultured for 24h *ex vivo* increased considerably
325 (**Figure 5F5**). These imply that olfactory circuit remodeling (degeneration of larval-specific
326 processes and growth of adult-specific processes) proceeds normally, albeit at a slower rate
327 (compare with **Figure 5F1-3**).

328 Fourth, dendrites from genetically identified DL1 and DA1/DL3 PNs targeted to their
329 stereotyped locations in the antennal lobe in 0h APF brains cultured for 24h *ex vivo*, (**Figure**
330 **5G**), mimicking *in vivo* development (**Figure 4A**).

331 Finally, segregation of dendrites of PNs targeting to neighboring proto-glomeruli could
332 be recapitulated in brains dissected at 24h APF and cultured for 8h (**Figure 5 – figure**
333 **supplement 1; Figure 5 – video 2**). Specifically, despite constant dynamic interactions among
334 dendrites that explore the surroundings (arrowheads in **Figure 5 – figure supplement 1A2-4**),
335 DC3/VA1d and DA1 PNs exhibited a 1–2 μ m increase in the distance between centers of the two
336 dendritic masses and a substantial decrease in the overlap of their core targeting regions (**Figure**
337 **5 – figure supplement 1B–D**). Taken together, these data support that the explant culture and
338 imaging system established here reliably captures key neurodevelopmental events starting from
339 early pupal stages.

340
341 **Single-cell, two-photon imaging reveals active dendrite targeting**
342 Our observation in fixed brains revealed that dendrites of DL1 adPNs transition from a uniform
343 extension in the antennal lobe at 0h APF to concentration at the dorsolateral corner of the
344 antennal lobe at 6h APF (**Figure 3A**). To identify mechanisms of dendrite targeting specificity
345 that could be missed in static developmental snapshots, we performed two-photon time-lapse
346 imaging of single-cell MARCM clones of DL1 PNs in 3h APF brains (**Figure 6; Figure 6 –**
347 **figure supplement 1; Figure 6 – video 1**). Although we did not have a counterstain outlining
348 the antennal lobe, we could use the background signals to discern the orientation of DL1 PNs in
349 the brain (**Figure 6 – figure supplement 1A**). The final targeting regions relative to the antennal
350 lobe revealed by *post hoc* fixation and immunostaining confirmed proper dendrite targeting
351 (yellow arrowhead in **Figure 6A10**; **Figure 6 – figure supplement 1B–C**).

352 Using DL1 PN in **Figure 6A** (pseudo-colored in yellow; **Figure 6 – video 1**) as an
353 example, we observed that the PN initially extended dendrites in every direction (**Figure 6A1-3**),
354 like what we observed in fixed tissues (**Figure 3A1**). The first sign of active targeting emerged at
355 2h 20min *ex vivo* when DL1 PN began to generate long, albeit transient, dendritic protrusions in
356 the dorsolateral direction; these selective protrusions were more prominent at 3h *ex vivo*
357 (arrowheads in **Figure 6A4–6**). The dorsolateral targeting continued to intensify, leading to the
358 formation a highly focal dendritic mass seen at 13h *ex vivo* (arrowhead in **Figure 6A8**). As the
359 dendrites reached the dorsolateral corner and explored locally, the change in shape appeared less
360 pronounced (**Figure 6A9**).

361 To quantitatively characterize the active targeting process, we categorized the bulk
362 dendritic masses emanated from the main process according to their targeting directions: DL,
363 DM, VM, and VL (**Figure 6B**). During the initial phase, the percentage of dendritic volume in
364 each direction varied from 10% to 40% (**Figures 6C and D**), indicative of active exploration
365 with little targeting specificity. Despite these variations, the total amount of dendritic mass seen
366 in the VM direction over the entire imaging time (area under graph of **Figure 6C**) was the

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367 smallest across all samples examined (**Figure 6E**). The initial phase of exploration in every
368 direction was followed by a ~4-hour transitional phase during which DL1 PNs predominantly
369 extended dendrites in 2 of the 4 directions (**Figure 6C; Figure 6 – figure supplement 1D–E**).
370 One of the 2 directions was always DL whereas the other was either DM or VL but never VM.
371 At the final phase, DL1 PN dendrites always preferred DL out of the 2 available directions.
372 Lastly, we analyzed the bulk dendritic movements. We defined bulk extension and retraction
373 events when dendrites respectively extended and retracted more than 2 μ m between two
374 consecutive time frames. The analyses showed a striking shift from frequent extension and
375 retraction towards stabilization, reflecting the pre- and post-targeting dynamics respectively
376 (**Figure 6F and G**).

377 Hence, long-term two-photon imaging of single-cell DL1 PNs revealed that dendrite
378 targeting specificity increases over time via active targeting in specific direction and stepwise
379 elimination of unfavorable trajectory choices (see summary in **Figure 7F_{1–3}**).
380

381 **AO-LLSM imaging suggests a cellular mechanism underlying dendrite targeting specificity**
382 To capture fast dynamics of single dendritic branches, we performed dual-color adaptive optical
383 lattice sheet microscopy (AO-LLSM) imaging (Chen et al., 2014; Wang et al., 2014; Liu et al.,
384 2018) of PNs every 30 seconds for 15 minutes, following a protocol we recently established (Li
385 et al., 2021; Li & Luo, 2021). We selected 3h, 6h, and 12h APF pupal brains double-labeled with
386 DL1 PNs and bulk adPNS (**Figure 7A–C; Figure 7 – videos 1–3**). The labeling of adPNS with
387 GFP outlined PN cell bodies and the developing antennal lobe but not the degenerating one,
388 presumably because the GFP in larval-specific dendrites was quickly quenched upon glial
389 phagocytosis (Marin et al., 2005).

390 In the 15-min imaging window, we observed 4 types of terminal branches regardless of
391 neuronal types or developmental stages: (1) stable branch that existed throughout the entire
392 imaging time, (2) transient branch that was produced and eliminated within the imaging window,
393 (3) emerging branch that was produced after imaging began, and (4) retracting branch that was
394 eliminated within the imaging period (**Figure 7 – figure supplement 1A**). To examine if
395 terminal branch dynamics exhibit any directional preference, we assigned the branches according
396 to their targeting directions (**Figure 7D**). Extension and retraction events were defined when
397 speed exceeded 0.5 μ m/min. Terminal branches were selected for analyses as branches closer to
398 the main process were too dense to resolve. **Figure 7D_{1–3}** showed the dynamics of ~15 randomly
399 selected terminal branches in each direction from the representative 3h, 6h, and 12h APF DL1
400 PNs (**Figure 7A–C**).

401 Quantitative analyses revealed that at 3h APF, DL1 PNs constantly produced, eliminated,
402 extended, and retracted dendritic branches (**Figure 7A, Figure 7D₁, Figure 7 – video 1**). Even
403 stable branches were not immobile. Rather, they spent comparable amounts of time extending
404 and retracting at ~1.5 μ m/min (**Figure 7 – figure supplement 1A₁, 1B**). Transient, emerging,
405 and retracting branches had similar, but more variable speed, ranging from 1 to 2.5 μ m/min.
406 Although there was no correlation between targeting direction and frequency/speed of
407 extension/retraction, the number of stable branches in the VM direction was significantly lower
408 than in other directions across all 3h DL1 PN samples examined (**Figure 7E₁**). This suggests that
409 even though dendritic branches were developed in every direction at the early stages, those
410 branches in the VM direction were short-lived and might be eliminated by retraction. The
411 direction-dependent stability/lifespan of dendritic branches on the timescale of seconds

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412 uncovered from AO-LLSM imaging explains why bulk dendrites in unfavorable trajectories
413 failed to persist in long-term two-photon imaging.

414 From 6h to 12h APF, DL1 PNs no longer manifested direction-specific branch
415 de/stabilization (**Figure 7B–C**, **Figure 7D_{2–3}**, **Figure 7 – videos 2–3**). At the same
416 developmental stage, stable branches in one direction appeared indistinguishable from those in
417 other directions in terms of abundance, frequency, and speed (**Figure 7D_{2–3}**, **Figure 7 – figure**
418 **supplement 1C–D**). This suggests that the entire dendritic mass tends to stay in equilibrium
419 upon arrival at target regions. At 12h APF, the abundance of stable branches of DL1 PNs was the
420 highest (**Figure 7D–E₁**). Also, the stable branches of 12h APF DL1 PNs moved at a significantly
421 lower speed (~1 μ m/min) (**Figures 7E₂**) and spent more time being stationary than those at 3h
422 and 6h (**Figure 7 – figure supplement 1B–D**). The reduced branch dynamics at 12h APF is
423 consistent with observations from two-photon imaging showing fewer bulk extension/retraction
424 events in the final phase of targeting (**Figure 6E–F**). Despite the slowdown, dendritic
425 arborization was evident in terminal branches of 12h APF DL1 PNs (**Figure 7 – figure**
426 **supplement 1E**), indicating that PN dendrites are transitioning from simple to complex branch
427 architectures. Although it remains unclear if there is a causal relationship between reduced
428 branch dynamics and increased structural complexity, we propose that both contribute to the
429 sustentation of dendrite targeting specificity.

430 In summary, AO-LLSM imaging reveals that PNs selectively stabilize branches in the
431 direction towards the target and destabilize those in the opposite direction, providing a cellular
432 basis of dendrite targeting specificity. Upon arrival at the target, the specificity is sustained
433 through branch stabilization in a direction-independent manner (summarized in **Figure 7F_{4–7}**).
434

435 **Embryonic-born PNs timely integrate into adult olfactory circuit by simultaneous dendritic 436 pruning and re-extension**

437 In earlier sections, we uncovered wiring logic of larval-born PN dendritic patterning and cellular
438 mechanisms of dendrite targeting specificity used to initiate olfactory map formation (**Figures 3–**
439 **7**). In this final section, we focused on embryonic-born PNs, which participate in both larval and
440 adult olfactory circuits by reorganizing their processes (Marin et al., 2005). Our previous study
441 demonstrates that embryonic-born PNs prune their larval-specific dendrites during early
442 metamorphosis (Marin et al., 2005) (**Figure 1D_{1–3}**). Here, we examined when and how
443 embryonic-born PNs re-extend dendrites used in the adult olfactory circuit.

444 It is known that γ neurons of *Drosophila* mushroom body (γ Kenyon cells) and sensory
445 Class IV dendritic arborization (C4da) neurons prune their processes between 4h and 18h APF
446 and show no signs of re-extension at 18h APF (Lee et al., 2000; Watts et al., 2003; Lee et al.,
447 2009). Do embryonic-born adPNS follow a similar timeframe? We first examined developing
448 brains double-labeled for embryonic-born DA41/VA6/VA2 adPNS (collectively referred to as
449 '*lov*+' PNs) and early larval-born DC2 adPNS (**Figure 8A**; **Figure 8 – figure supplement 1**).
450 We found that, by 12h APF, *lov*+' PNs already sent adult-specific dendrites to a region
451 ventromedial to DC2 PN dendrites (green arrowhead in **Figure 8A₃**). This implies that *lov*+' PNs
452 have already caught up with DC2 PNs on dendrite development at this stage, and re-extension of
453 *lov*+' PN dendrites must have happened even earlier. Indeed, we observed *lov*+' PN dendrites
454 innervated the developing antennal lobe extensively at 6h APF (**Figure 8A₂**). Such innervation
455 was not observed at 0h APF (**Figure 8A₁**). After 12h APF, the time course of *lov*+' PN dendrite
456 development was comparable to that of DC2 PNs (**Figure 8A_{4–6}**).

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457 To characterize dendritic re-extension at single-cell resolution, we developed a sparse,
458 stochastic labeling strategy to label single *lov*+ PNs (Figure 8B). We found that *lov*+ PNs
459 produced nascent branches from the main process dorsal to larval-specific dendrites as early as
460 3h APF (Figure 8C₂₋₃; arrowheads in Figure 8C₆₋₇). At 6h APF, when larval-specific dendrites
461 were completely segregated from *lov*+ PNs, robust extension of adult-specific dendrites was
462 seen across the developing antennal lobe (Figure 8C₄). These data indicate that *lov*+ PNs re-
463 extend their adult-specific dendrites at a more dorsal location before the larval-specific dendrites
464 are completely pruned.

465 Do other embryonic-born PNs prune and re-extend their dendrites simultaneously? Like
466 *lov* drivers, *Mz612-GAL4* labels embryonic-born PNs, one of which is VA6 PN (Marin et al.,
467 2005). In 3h APF brains co-labeled for *Mz612*+ and *lov*+ PNs, we could unambiguously access
468 3 single embryonic-born PN types: (1) *lov*+ *Mz612*– PN, (2) *lov*– *Mz612*+ PN, and (3) *lov*–
469 *Mz612*+ PN (Figure 8 – figure supplement 2A–B). Tracing of individual dendritic branches
470 showed that all these PNs already re-extended dendrites to varying extents prior to separation of
471 larval-specific dendrites from the rest of the processes (Figure 8 – figure supplement 2C).
472 Thus, concurrent pruning and re-extension apply to multiple embryonic-born PN types.

473 To capture the remodeling at higher temporal resolution, we performed two-photon time-
474 lapse imaging of single embryonic-born PNs labeled by *Split7-GAL4* (Figure 8D, Figure 8 –
475 video 1, Figure 8 – figure supplement 3). This *GAL4* labels one embryonic-born PN (either
476 VA6 or VA2 PN) at early pupal stages but eight PN types at 24h APF (Xie et al., 2021). Initially
477 (3h APF + 0h *ex vivo*), no adult-specific dendrites were detected in live *Split7*+ PNs (Figure
478 8D₁). The following ~3h *ex vivo* saw thickening of the main process (arrowhead in Figure 8D₃).
479 From 4h *ex vivo* onwards, re-extension occurred in the presumed developing antennal lobe
480 located dorsal to larval-specific dendrites (arrowheads in Figure 8D₄₋₈; see traces in Figure
481 8D₉). Live imaging of *Split7*+ PNs also revealed that fragmentation of larval-specific dendrites
482 occurred at the distal ends (Figure 8 – figure supplement 3B₁₋₅), and the process leading to
483 larval-specific dendrites gradually disappeared as pruning approached completion (Figure 8 –
484 figure supplement 3B₆₋₁₀). These observations suggest that pruning of embryonic-born PN
485 dendrites is not initiated by severing at the proximal end. Distal-to-proximal pruning, rather than
486 in the reversed direction, further supports concurrent but spatially segregated pruning and re-
487 extension processes.

488 It has been shown that dendritic pruning of embryonic-born PNs requires ecdysone
489 signaling in a cell-autonomous manner (Marin et al., 2005). We asked if the re-extension process
490 also depends on ecdysone signaling. We expressed a dominant negative form of ecdysone
491 receptor (EcR-DN) in most PNs (including *lov*+ PNs) and monitored the development of *lov*+
492 PN dendrites (Figure 8 – figure supplement 4). We found that inhibition of ecdysone signaling
493 by *EcR-DN* expression not only suppressed pruning, but also blocked re-extension. This is
494 consistent with a previous study reporting the dual requirement of ecdysone signaling in the
495 pruning and re-extension of *Drosophila* anterior paired lateral (APL) neurons, although, unlike
496 embryonic-born PNs, APL neurons prune and re-extend processes sequentially (at 6h and 18h
497 APF, respectively) (Mayseless et al., 2018). We currently could not distinguish if the lack of re-
498 extension is due to defective pruning, or if ecdysone signaling controls pruning and re-extension
499 independently.

500 Taken together, our data demonstrate that embryonic-born PNs prune and re-extend
501 dendrites simultaneously at spatially distinct regions, and that both processes require ecdysone
502 signaling (Figure 8E). Such a ‘multi-tasking’ ability explains how embryonic-born PNs can re-

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503 integrate into the adult olfactory circuit and engage in its prototypic map formation in a timely
504 manner.

505
506 **DISCUSSION**
507

508 **Wiring logic for the prototypic olfactory map**

509 Prior to this study, no apparent logic linking PN lineage, birth order, and adult glomerular
510 position has been found. Our systematic analyses of dendritic patterning at the resolution of
511 specific PN types across development identified wiring logic underlying the spatial organization
512 of the prototypic olfactory map (**Figures 3 and 4**).

513 We found that PNs of a given lineage are grouped by birth order, and PNs of the same
514 group share similar dendrite targeting specificity and timing. The strong correlation found
515 between birth order and transcriptomic similarity among 0h APF adPNS (Xie et al., 2021)
516 provides a molecular basis for executing this logic. In addition, we found that dendrites of adPNS
517 and IPNs respectively pattern the antennal lobe in rotating and binary manners following birth
518 order. Certain lineage-specific transcription factors are known to instruct dendrite targeting
519 (Komiyama et al., 2003; Komiyama & Luo, 2007; Li et al., 2017; Xie et al., 2022), which may
520 account for why the adPN and IPN dendritic maps differ. Cellular interactions among PN
521 dendrites likely contribute further to the resulting olfactory map, given the robust dendritic
522 dynamics seen during circuit formation (**Figures 5–8**). Our new tools for labeling and genetic
523 manipulation of distinct PN types (**Figure 2**) will now enable in-depth investigations into the
524 potential PN dendrite-dendrite interactions and molecular mechanisms leading to the initial map
525 organization.

526
527 **Wiring logic evolves as development proceeds**

528 After the initial map formation at 12h APF, dendrite positions in the antennal lobe could change
529 substantially in the next 36 hours (for example, see DC2 PNs in **Figure 3B_{4–6}** and DA1 and
530 VA1d/DC3 PNs in **Figure 3C_{4–7}**). These changes occur when dendrites of PNs with neighboring
531 birth order begin to segregate and when ORN axons begin to invade the antennal lobe.

532 Accordingly, the oval-shaped antennal lobe turns into a globular shape (30–50h APF; **Figure**
533 **3C_{6–7}**). These PN-autonomous and non-autonomous changes likely mask the initial wiring logic,
534 explaining why previous studies, which mostly focused on examining the final glomerular
535 targets in adults (Jefferis et al., 2001), have missed the earlier organization. Interestingly, the
536 process of PN dendritic segregation coincides with the peak of PN transcriptomic diversity at
537 24h APF (Li et al., 2017; Xie et al., 2021).

538 Recent proteomics and genetic analyses have indicated that PN dendrite targeting is
539 mediated by cell-surface proteins cooperating as a combinatorial code (Xie et al., 2022). The
540 evolving wiring logic, which is consistent with the stepwise assembly of olfactory circuit (Hong
541 & Luo, 2014), suggests the combinatorial codes are not static. We propose that PNs use a
542 numerically simpler code for initial dendrite targeting (e.g., 18 adPN types being sorted into four
543 groups with distinct targeting specificity; **Figure 3E**). Following the expansion of transcriptomic
544 diversity, PNs acquire a more complex code mediating dendritic segregation of neighboring PNs
545 and matching of PN dendrites and ORN axons. Functional characterization of differentially
546 expressed genes between 12h and 24h APF PNs may provide molecular insights into how the
547 degree of discreteness in the olfactory map arises.

548

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549 **Selective branch stabilization as a cellular mechanism for dendrite targeting**

550 Utilizing an early pupal brain explant culture system coupled with two-photon and AO-LLSM
551 imaging (**Figure 5**), we presented the first time-lapse videos following dendrite development of a
552 specific PN type – DL1 PNs (**Figures 6 and 7**). We found that DL1 PN dendrites initiate active
553 targeting towards their dorsolateral target with direction-dependent branch stabilization. This
554 directional selectivity provides a cellular basis of the emerging targeting specificity of PN
555 dendrites at the beginning of olfactory map formation.

556 Although selective branch stabilization as a mechanism to achieve axon targeting
557 specificity has been described in neurons in the vertebrate and invertebrate systems (e.g., Yates
558 et al., 2001; Li et al., 2021), our time-lapse imaging showed, for the first time to our knowledge,
559 that selective branch stabilization is also used to achieve dendrite targeting specificity.
560 Furthermore, AO-LLSM imaging revealed that selective stabilization and destabilization of
561 dendritic branches occur on the timescale of seconds. As the rate of olfactory circuit
562 development in the brain explants was slower than normal development (**Figure 5F**), we might
563 have captured PN dendritic dynamics in slow motion. Using AO-LLSM for high spatiotemporal
564 resolution imaging, we just begin to appreciate how fast PN dendrites are coordinating trajectory
565 choices with branch stabilization to make the appropriate decision. Having characterized the
566 dendritic branch dynamics of the wild-type DL1 PNs, we have set the stage for future studies
567 addressing how positional cues and the downstream signaling instruct wiring, and whether other
568 PN types follow similar rules as DL1 PNs.
569

570 **Simultaneous pruning and re-extension as novel remodeling mechanism for neuronal
571 remodeling**

572 Our data on embryonic-born adPN dendrite development reveals a novel mode of neuronal
573 remodeling during metamorphosis (**Figure 8**). In mushroom body γ neurons and body wall
574 somatosensory neurons, two well-characterized systems, larval-specific neurites are first pruned,
575 followed by re-extension of adult-specific processes (Watts et al., 2003; Williams and Truman,
576 2005; Yaniv and Schuldiner, 2016). However, embryonic-born adPNs prune larval-specific
577 dendrites and re-extend adult-specific dendrites simultaneously but at spatially separated regions.
578 Such spatial segregation suggests that regional external cues could elicit compartmentalized
579 downstream signals leading to opposite effects on the dendrites. Subcellular
580 compartmentalization of signaling and cytoskeletal organization has been observed in diverse
581 neuron types across species (Rolls et al., 2007; Kanamori et al., 2013; O'Hare et al., 2022).

582 Why do embryonic-born adPNs ‘rush’ to re-extend dendrites? During normal
583 development, it takes at least 18 hours for embryonic-born adPNs to produce and properly target
584 dendrites (growth at 3–6h APF, initial targeting at 6–12h APF, and segregation at 21–30h APF).
585 Given that the dendritic re-extension of embryonic-born PNs is ecdysone dependent (**Figure 8 –**
586 **figure supplement 4**), if the PNs did not re-extend dendrites at 3h APF, they would have to wait
587 for the next ecdysone surge at ~20h APF (Thummel, 2001), which might be too late for their
588 dendrites to engage in the prototypic map formation. Thus, embryonic-born PNs develop a
589 remodeling strategy that coordinates with the timing of systemic ecdysone release. By
590 simultaneous pruning and re-extension, embryonic-born adPNs timely re-integrate into the adult
591 prototypic map that readily serves as target for subsequent ORN axon innervation.

592 In conclusion, our study highlights the power and necessity of type-specific neuronal
593 access and time-lapse imaging to identify wiring logic and mechanisms underlying the origin of
594 an olfactory map. Applying similar approaches to other developing neural maps across species

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595 should broaden our understanding of the generic and specialized designs that give rise to
596 functional maps with diverse architectures.

597

598 MATERIALS AND METHODS

599

600 *Drosophila* stocks and husbandry

601 Flies were maintained on standard cornmeal medium at 25°C. Fly lines used in this study
602 included *GH146-FLP* (Hong et al., 2009), *QUAS-FRT-stop-FRT-mCD8-GFP* (Potter et al.,
603 2010), *UAS-mCD8-GFP* (Lee & Luo, 1999), *UAS-mCD8-FRT-GFP-FRT-RFP* (Stork et al.,
604 2014), *VT033006-GAL4* (Tirian & Dickson, 2017), *Mz19-GAL4* (Jefferis et al., 2004), *91G04-*
605 *GAL4* (Jenett et al., 2012), *Mz612-GAL4* (Marin et al., 2005), *71B05-GAL4* (Jenett et al., 2012),
606 *Split7-GAL4* (Xie et al., 2021), *QUAS-FLP* (Potter et al., 2010) and *UAS-EcR.B1-AC655.F645A*
607 (Cherbas et al., 2003). The following GAL4 lines were obtained from Bloomington *Drosophila*
608 Stock Center (BDSC): *tsh-GAL4* (BDSC #3040) and *lov-GAL4* (BDSC #3737).

609 The following two stocks were used for MARCM analyses: (1) *UAS-mCD8-GFP*, *hs-*
610 *FLP*; *FRT*^{G13}, *tub-GAL80*; *GH146-GAL4*, and (2) *FRT*^{G13}, *UAS-mCD8-GFP* (Lee & Luo,
611 1999).

612 The following lines were generated in this study: *UAS-FRT*^{I0}-*stop-FRT*^{I0}-*3xHalo7-CAAX*
613 (on either II or III chromosome), *UAS-FRT-myr-4xSNAPf-FRT-3xHalo7-CAAX* (III), *UAS-FRT-*
614 *myr-mGreenLantern-FRT-3xHalo7-CAAX* (II), *QUAS-FRT-stop-FRT-myr-4xSNAPf* (III), *run-*
615 *T2A-FLP* (X), *acj6-T2A-FLP* (X), *acj6-T2A-QF2* (X), *CG14322-T2A-QF2* (III) and *lov-T2A-*
616 *QF2* (II).

617

618 *Drosophila* genotypes

619 **Figure 1D, Figure 1 – figure supplement 1, Figure 1 – figure supplement 2:** *run-T2A-FLP*/+;
620 *UAS-mCD8-FRT-GFP-FRT-RFP*/+; *VT033006-GAL4*/+

621 **Figure 3A:** *acj6-T2A-QF2*/+; *GH146-FLP*, *QUAS-FRT-stop-FRT-mCD8-GFP/UAS-FRT*^{I0}-
622 *stop-FRT*^{I0}-*3xHalo7-CAAX*; *71B05-GAL4*/+

623 **Figure 3B, Figure 3 – figure supplement 1C:** *GH146-FLP/UAS-FRT*^{I0}-*stop-FRT*^{I0}-*3xHalo7-*
624 *CAAX*; *91G04-GAL4/CG14322-T2A-QF2*, *QUAS-FRT-stop-FRT-myr-4xSNAPf*

625 **Figure 3C:** *acj6-T2A-FLP*/+; *Mz19-GAL4*; *UAS-FRT-myr-4xSNAPf-FRT-3xHalo7-CAAX*/+

626 **Figure 3D, Figure 3 – figure supplement 2, Figure 3 – figure supplement 3:** *UAS-mCD8-*
627 *GFP*, *hs-FLP*/+; *FRT*^{G13}, *tub-GAL80/FRT*^{G13}, *UAS-mCD8-GFP*; *GH146-GAL4* (IV)/+

628 **Figure 3 – figure supplement 1A:** *GH146-FLP/UAS-FRT*^{I0}-*stop-FRT*^{I0}-*3xHalo7-CAAX*;
629 *71B05-GAL4*/+

630 **Figure 3 – figure supplement 1B:** *GH146-FLP/UAS-FRT*^{I0}-*stop-FRT*^{I0}-*3xHalo7-CAAX*;
631 *91G04-GAL4*/+

632 **Figure 4A, Figure 4 – figure supplement 1:** *GH146-FLP*, *UAS-FRT*^{I0}-*stop-FRT*^{I0}-*3xHalo7-*
633 *CAAX/tsh-GAL4*; *CG14322-T2A-QF2*, *QUAS-FRT-stop-FRT-myr-4xSNAPf*/+

634 **Figure 4B:** *UAS-mCD8-GFP*, *hs-FLP*/+; *FRT*^{G13}, *tub-GAL80/FRT*^{G13}, *UAS-mCD8-GFP*;
635 *GH146-GAL4* (IV)/+

636 **Figure 4 – figure supplement 2:** *acj6-T2A-FLP*/+; *tsh-GAL4*, *UAS-mCD8-FRT-GFP-FRT-RFP*

637 **Figure 5E, Figure 5 – video 1:** *run-T2A-FLP*/+; *UAS-FRT-myr-mGreenLantern-FRT-3xHalo7-*
638 *CAAX*/+; *VT033006-GAL4*/+

639 **Figure 5F:** *UAS-mCD8-GFP*/+; *VT033006-GAL4*/+

640 **Figure 5G1:** *GH146-FLP/UAS-FRT*^{I0}-*stop-FRT*^{I0}-*3xHalo7-CAAX*; *71B05-GAL4*/+

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641 **Figure 5G2: GH146-FLP/tsh-GAL4; UAS-FRT¹⁰-stop-FRT¹⁰-3xHalo7-CAAX/+**
642 **Figure 5 – figure supplement 1, Figure 5 – video 2: acj6-T2A-FLP/+; Mz19-GAL4/UAS-FRT-**
643 **myr-mGreenLantern-FRT-3xHalo7-CAAX**
644 **Figure 6A, Figure 6 – figure supplement 1, Figure 6 – video 1: UAS-mCD8-GFP, hs-FLP/+;**
645 **FRT^{G13}, tub-GAL80/FRT^{G13}, UAS-mCD8-GFP;; GH146-GAL4 (IV)/+**
646 **Figure 7A–C, Figure 7 – figure supplement 1, Figure 7 – videos 1–3: acj6-T2A-QF2/+,**
647 **GH146-FLP, QUAS-FRT-stop-FRT-mCD8-GFP/UAS-FRT¹⁰-stop-FRT¹⁰-3xHalo7-CAAX;**
648 **71B05-GAL4/+**
649 **Figure 8A, Figure 8 – figure supplement 1: GH146-FLP, QUAS-FRT-stop-FRT-mCD8-**
650 **GFP/lov-T2A-QF2; UAS-FRT¹⁰-stop-FRT¹⁰-3xHalo7-CAAX/91G04-GAL4**
651 **Figure 8C: GH146-FLP/lov-GAL4; UAS-FRT¹⁰-stop-FRT¹⁰-3xHalo7-CAAX/QUAS-FRT-stop-**
652 **FRT-myr-4xSNAPf**
653 **Figure 8D, Figure 8 – figure supplement 3, Figure 8 – video 1: UAS-mCD8-GFP/+; Split7-**
654 **GAL4 (i.e. FlyLight SS01867: 72C11-p65ADZp; VT033006-ZpGDBD)/+**
655 **Figure 8 – figure supplement 2: GH146-FLP, QUAS-FRT-stop-FRT-mCD8-GFP/lov-T2A-**
656 **QF2, Mz612-GAL4; UAS-FRT¹⁰-stop-FRT¹⁰-3xHalo7-CAAX/+**
657 **Figure 8 – figure supplement 4A: lov-T2A-QF2, QUAS-FLP/+; VT033006-GAL4/UAS-mCD8-**
658 **FRT-GFP-FRT-RFP**
659 **Figure 8 – figure supplement 4B: lov-T2A-QF2, QUAS-FLP/UAS-EcR-DN; VT033006-**
660 **GAL4/UAS-mCD8-FRT-GFP-FRT-RFP**

661

662 MARCM clonal analyses

663 MARCM clonal analyses have been previously described (Lee & Luo, 1999). Larvae of the
664 genotype *UAS-mCD8-GFP, hs-FLP/+; FRT^{G13}, tub-GAL80/FRT^{G13}, UAS-mCD8-GFP;; GH146-*
665 *GAL4/+* were heat shocked at 37°C for 1 hour. To label the first-born DL1 PNs, heat shock was
666 applied at <24h after larval hatching (ALH). MARCM clones of early, middle and late larval-
667 born PNs were generated by applying heat shocks at 48h, 72h and >96h ALH, respectively. As
668 larvae developed at different rates (Tennessen & Thummel, 2011), we reasoned that even if we
669 could collect 0h–2h ALH larvae, their development might have varied by the time of heat shock.
670 To minimize the effects of unsynchronized development, we selected those heat-shocked larvae
671 that were among the first to form puparia and collected these white pupae in a ~3-hour window
672 for the clonal analyses.

673

674 Transcriptomic analyses

675 Transcriptomic analyses have been described previously (Xie et al., 2021). tSNE plots and dot
676 plots were generated in Python using PN single-cell RNA sequencing data and code available at
677 https://github.com/Qijing-Xie/FlyPN_development.

678

679 Generation of T2A-QF2/FLP lines

680 To generate a *T2A-QF2/FLP* donor vector for *acj6* (we used the same strategy for *run*, *CG14322*
681 and *lov*), a ~2000-bp genomic sequence flanking the stop codon of *acj6* was PCR amplified and
682 introduced into *pCR-Blunt II-TOPO* (ThermoFisher Scientific #450245), forming *pTOPO-acj6*.
683 To build *pTopo-acj6-T2A-QF2*, *T2A-QF2* including *loxP*-flanked 3xP3-RFP was PCR amplified
684 from *pBPGUw-HACK-QF2* (Addgene #80276), followed by insertion into *pTOPO-acj6* right
685 before the stop codon of *acj6* by DNA assembly (New England BioLabs #E2621S). To generate
686 *T2A-FLP*, we PCR amplified *FLP* from genomic DNA of *GH146-FLP* strain. *QF2* in *pTopo-*

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687 *acj6-T2A-QF2* was then replaced by *FLP* through DNA assembly. Using CRISPR Optimal
688 Target Finder (Gratz et al., 2014), we selected a 20-bp gRNA target sequence that flanked the
689 stop codon and cloned it into *pU6-BbsI-chiRNA* (Addgene #45946). If the gRNA sequence did
690 not flank the stop codon, silent mutations were introduced at the PAM site of the donor vector by
691 site-directed mutagenesis. Donor and gRNA vectors were co-injected into *Cas9* embryos in-
692 house or through BestGene.

693

694 **Generation of FLP-out reporters**

695 To generate *pUAS-FRT¹⁰-stop-FRT¹⁰-3xHalo7-CAAX*, *FRT¹⁰-stop-FRT¹⁰* was PCR amplified
696 from *pUAS-FRT¹⁰-stop-FRT¹⁰-mCD8-GFP* (Li et al., 2021) and inserted into *pUAS-3xHalo7-*
697 *CAAX* (Addgene #87646) through NotI and DNA assembly.

698 To generate *pUAS-FRT-myr-4xSNAPf-FRT-3xHalo7-CAAX*, we first PCR amplified *myr-*
699 *4xSNAPf* from *pUAS-myr-4xSNAPf* (Addgene #87637) using *FRT*-containing primers. *FRT-myr-*
700 *4xSNAPf-FRT* was then introduced into *pCR-Blunt II-TOPO*, forming *pTOPO-FRT-myr-*
701 *4xSNAPf-FRT*. Using NotI-containing primers, *FRT-myr-4xSNAPf-FRT* was PCR amplified and
702 subcloned into *pUAS-3xHalo7-CAAX* through NotI.

703 To generate *pUAS-FRT-myr-mGreenLantern-FRT-3xHalo7-CAAX*, we first PCR
704 amplified *mGreenLantern* from *pcDNA3.1-mGreenLantern* (Addgene #161912). Using *MluI* and
705 *XbaI*, we replaced *4xSNAPf* in *pUAS-myr-4xSNAPf* with *mGreenLantern* to build *pUAS-myr-*
706 *mGreenLantern*. *myr-mGreenLantern* was PCR amplified with the introduction of *FRT*
707 sequence, followed by insertion into *pCR-Blunt II-TOPO*. Using the NotI-containing primers,
708 *FRT-myr-mGreenLantern-FRT* was PCR amplified and subcloned into *pUAS-3xHalo7-CAAX*
709 through NotI.

710 To generate *pQUAS-FRT-stop-FRT-myr-4xSNAPf*, we first PCR amplified *FRT-stop*
711 from *pJFRC7-20XUAS-FRT-stop-FRT-mCD8-GFP* (Li et al., 2021) and inserted it into *pTOPO-*
712 *FRT-myr-4xSNAPf-FRT* through DNA assembly to form *pTOPO-FRT-stop-FRT-myr-4xSNAPf-*
713 *FRT*. Using NotI-containing forward and KpnI-containing reverse primers, *FRT-stop-FRT-myr-*
714 *4xSNAPf* was PCR amplified and subcloned into *p10XQUAST*. *p10XQUAST* was generated using
715 *p5XQUAS* (Addgene #24349) and *p10xQUAS-CsChrimson* (Addgene #163629).

716 *attP24* and *86Fb* landing sites were used for site-directed integration.

717

718 **Immunofluorescence staining and confocal imaging**

719 Fly brain dissection for immunostaining and live imaging has been described (Wu & Luo, 2006).
720 Briefly, brains were dissected in phosphate buffered saline (PBS) and fixed with 4%
721 paraformaldehyde in PBS for 20 minutes on a nutator at room temperature. Fixed brains were
722 washed with 0.1% Triton X-100 in PBS (PBST) for 10 min twice. After blocking with 5%
723 normal donkey serum in PBST for 1 hour at room temperature, the brains were incubated with
724 primary antibodies overnight at 4°C. After PBST wash, brains were incubated with secondary
725 antibodies (1:1000; Jackson ImmunoResearch) in dark for 2 hours at room temperature. Washed
726 and mounted brains were imaged with confocal laser scanning microscopy (ZEISS LSM 780;
727 LSM 900 with Airyscan 2). Images were processed with ImageJ. Neurite tracing images were
728 generated using Simple Neurite Tracer (SNT) (Arshadi et al., 2021). Primary antibodies used
729 included chicken anti-GFP (1:1000; Aves Lab #GFP-1020), rabbit anti-DsRed (1:500; TaKaRa
730 #632496), rat anti-Cadherin DN (1:30; Developmental Studies Hybridoma Bank DSHB DN-
731 Ex#8 supernatant), and mouse anti-Bruchpilot (1:30; DSHB nc82 supernatant).

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733 **Chemical labeling**

734 Chemical labeling of *Drosophila* brains has been described (Kohl et al., 2014). Janelia Fluor (JF)
735 Halo and SNAP ligands (stocks at 1 mM) were gifts from Dr. Luke Lavis (Grimm et al., 2017,
736 2021).

737 Fixed brains were washed with PBST for 5 min, followed by incubation with Halo and/or
738 SNAP ligands (diluted in PBS) for 45 min at room temperature. Brains were then washed with
739 PBST for 5 min, followed by blocking and immunostaining if necessary. For co-incubation of
740 Halo and SNAP ligands, JF503-cpSNAP (1:1000) and JF646-Halo (1:1000) were used.
741 Alternatively, JFX650-SNAP (1:1000) and JFX554-Halo (1:10000) were used. When only Halo
742 ligands were needed, either JF646-Halo or JF635-Halo (1:1000) was used.

743 For live brain imaging, dissected brains were incubated with Halo ligands diluted in
744 culture media (described below) for 30 min at room temperature. For two-photon imaging,
745 JF570-Halo was used at 1:5000. For AO-LLSM imaging, following JF646-Halo incubation at
746 1:1000, the brains were incubated with 1 μ M Sulforhodamine 101 (Sigma) for 5 min at room
747 temperature. The brains were then briefly washed with culture media before imaging.

748

749 **Brain explant culture setup and medium preparation**

750 Brain explant culture setup was modified based on Li et al., 2021; Li & Luo, 2021. A Sylgard
751 plate with a thickness of ~ 2 millimeters was prepared by mixing base and curing agent at 10:1
752 ratio (DOW SYLGARD™ 184 Silicone Elastomer Kit). The mixture was poured into a 60 mm x
753 15 mm dish in which it was cured for two days at room temperature. Once cured, the plate was
754 cut into small squares (~15 mm x ~15 mm). Indentations were created based on the size of an
755 early pupal brain using a No.11 scalpel. Additional slits were made around the indentations for
756 attaching imaginal discs which served as anchors to hold the brain position. A square Sylgard
757 piece was then placed in a 60 mm x 15 mm dish or on a 25-mm round coverslip in preparation
758 for two-photon/AO-LLSM imaging.

759 Culture medium was prepared based on published methods (Rabinovich et al., 2015; Li
760 and Luo, 2021; Li et al., 2021). The medium contained Schneider's *Drosophila* Medium
761 (ThermoFisher Scientific #21720001), 10% heat-inactivated Fetal Bovine Serum (ThermoFisher
762 Scientific #16140071), 10 μ g/mL human recombinant insulin (ThermoFisher Scientific
763 #12585014; stock = 4 mg/mL), 1:100 Penicillin-Streptomycin (ThermoFisher Scientific
764 #15140122). For 0h–6h APF brain culture, 0.5 mM ascorbic acid (Sigma #A4544; stock
765 concentration = 50 mg/mL in water) was included. 20-hydroxyecdysone (Sigma #H5142; stock
766 concentration = 1 mg/mL in ethanol) was used for 0h–6h and 12h brain explants at 20 μ M and 2
767 μ M, respectively. Culture medium was oxygenated for 20 minutes before use.

768

769 **Single- and dual-color imaging with two-photon microscopy**

770 Single- and dual-color imaging of PNs were performed at room temperature using a custom-built
771 two-photon microscope (Prairie Technologies) with a Chameleon Ti:Sapphire laser (Coherent)
772 and a 16X water-immersion objective (0.8 NA; Nikon). Excitation wavelength was set at 920 nm
773 for GFP imaging, and at 935 nm for co-imaging of mGreenLantern and JF570-Halo. z-stacks
774 were obtained at 4- μ m increments (10- μ m increments for **Figure 5 – video 1**). Images were
775 acquired at a resolution 1024 x 1024 pixel² (512 x 512 for **Figure 5 – video 1**), with a pixel
776 dwell time of 6.8 μ s and an optical zoom of 2.1, and at a frequency every 20 minutes for 8–23
777 hours.

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779 **Dual-color imaging with AO-LLSM**

780 For AO-LLSM based imaging, the excitation and detection objectives along with the 25-mm
781 coverslip were immersed in ~40 mL of culture medium at room temperature. Explant brains held
782 on Sylgard plate were excited simultaneously using 488 nm (for GFP) and 642 nm (for JF-646)
783 lasers operating with ~2–10 mW input power to the microscope (corresponding to ~10–50 μ W at
784 the back aperture of the excitation objective). An exposure time of 20–50 msec was used to
785 balance imaging speed and signal-to-noise ratio (SNR). Dithered lattice light-sheet patterns with
786 an inner/outer numerical aperture of 0.35/0.4 or 0.38/0.4 were used. The optical sections were
787 collected by an axial step size of 250 nm in the detection objective coordinate, with a total of 81–
788 201 steps (corresponding to a total axial scan range of 20–50 μ m). Emission light from GFP and
789 JF-646 was separated by a dichromatic mirror (Di03-R561, Semrock, IDEX Health & Science,
790 LLC, Rochester, NY) and captured by two Hamamatsu ORCA-Fusion sCMOS cameras
791 simultaneously (Hamamatsu Photonics, Hamamatsu City, Japan). Prior to the acquisition of the
792 time series data, the imaged volume was corrected for optical aberrations using two-photon guide
793 star based adaptive optics method (Chen et al., 2014; Wang et al., 2014; Liu et al., 2018). Each
794 imaged volume was deconvolved using Richardson-Lucy algorithm on HHMI Janelia Research
795 Campus' or Advanced Bioimaging Center's computing cluster
796 (<https://github.com/scopetools/cudadecon>, <https://github.com/abcucberkeley/LLSM3DTools>)
797 with experimentally measured point spread functions obtained from 100 or 200 nm fluorescent
798 beads (Invitrogen FluoSpheresTM Carboxylate-Modified Microspheres, 505/515 nm, F8803,
799 FF8811). The AO-LLSM was operated using a custom LabVIEW software (National
800 Instruments, Woburn, MA).

801

802 **Statistics**

803 For data analyses, t-test and one-way ANOVA were used to determine *p* values as indicated in
804 the figure legend for each graph, and graphs were generated using Excel. Exact *p* values were
805 provided in Source Data files.

806

807 **Material and data availability**

808 All reagents generated in this study are available from the lead corresponding author without
809 restriction. Figure 3 - Source Data 1, Figure 5 - Source Data 1, Figure 6 - Source Data 1, and
810 Figure 7 - Source Data 1 contain the numerical and statistical data used to generate the figures.

811

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819

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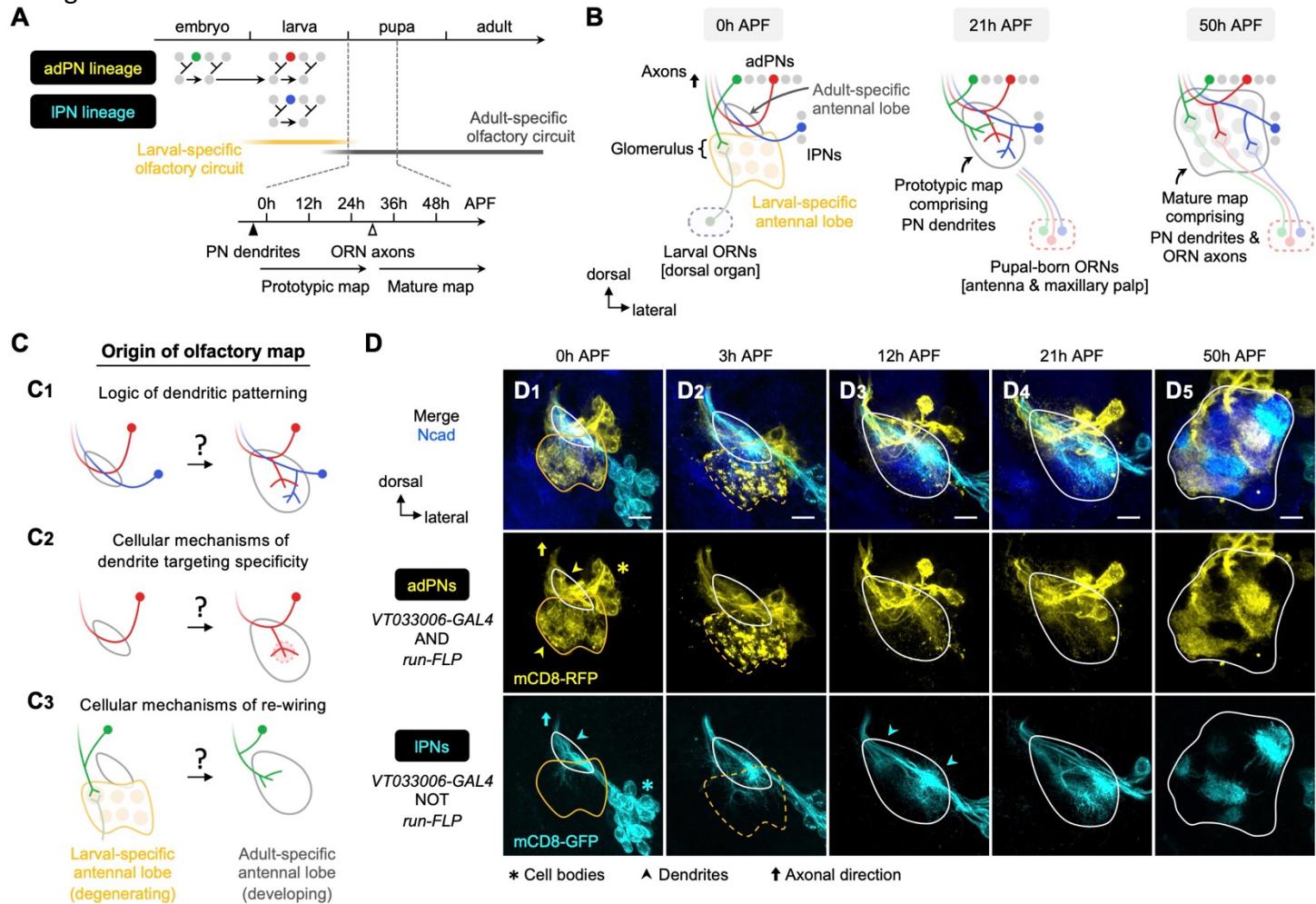


Figure 1. Organization and development of the adult olfactory circuit in *Drosophila*.

(A, B) Timeline (A) and schematic illustration (B) of *Drosophila* olfactory circuit development. Green, red, and blue circles denote the birth of embryonic-born adPN, larval-born adPN, and larval-born IPN, respectively. At the onset of metamorphosis, larval-specific olfactory circuit degenerates; larval ORNs die while embryonic-born adPNs prune their larval-specific processes and re-extend new processes into the adult-specific olfactory circuit. In the adult-specific olfactory circuit, PN dendrites extend first and form a prototypic map. This is followed by extension of ORN axons and synaptic partner matching between cognate PN dendrites and ORN axons to form a mature map. Solid and open arrowheads in A indicate innervation timing of PN dendrites and ORN axons, respectively.

(C) Overview of this study investigating the logic of dendritic patterning (C₁; see **Figures 3 and 4**) as well as cellular mechanisms of dendrite targeting specificity (C₂; see **Figures 6 and 7**) and re-wiring (C₃; see **Figure 8**) that contribute to the developmental origin of the adult *Drosophila* olfactory map.

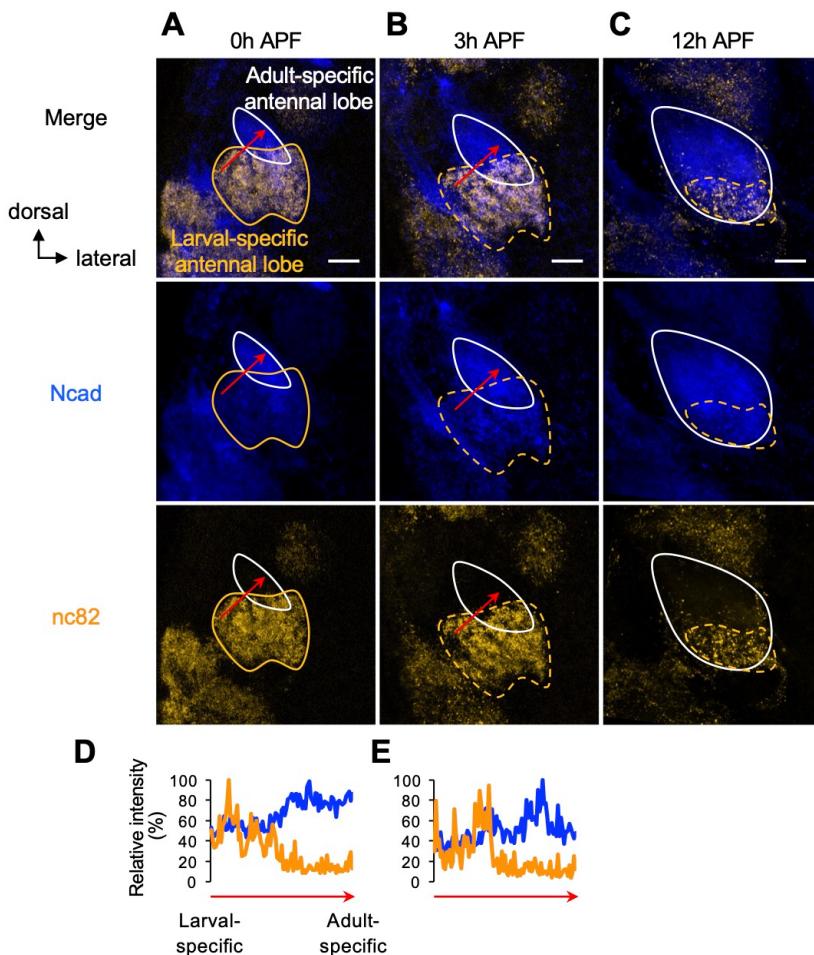
(D) Staining of fixed brains at indicated stages showing dendrite development of adPNs (*VT033006+ run+*; labeled in yellow) and IPNs (*VT033006+ run-*; labeled in cyan). As *run-FLP* is expressed before 0h APF in adPN but not IPN neuroblasts, we can use it to label adPNs and IPNs with two distinct colors using an intersectional reporter (see **Materials and Methods** for the genotype). Yellow arrowheads in D₁ mark larval- and adult-specific dendrites of adPNs in larval- and adult-specific antennal lobes, respectively. Cyan arrowheads in D₃ denote specific targeting of IPN dendrites at the opposite ends of the dorsomedial-ventrolateral axis.

Common notations in this study: Unless otherwise indicated, all images in this and subsequent figures are partial z projections of confocal stacks. Antennal lobe neuropils are revealed by N-Cadherin (Ncad; in blue) staining. Adult-specific (developing) antennal lobe is outlined with white solid line. Larval-specific antennal

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lobe is outlined with orange line (dashed line used to denote the degeneration stage) and is distinguished from the developing antennal lobe by the more intense nc82 staining as shown in **Figure 1 – figure supplement 1** (nc82 channel not shown here). Asterisks (*) indicate PN cell bodies, which are outside the antennal lobe neuropil (and sometimes appear on top because of the z-projections). Arrowheads mark PN dendrites. Arrows mark PN axons projecting towards higher olfactory centers (see **Figure 1 – figure supplement 2** for PN axons at their targets in the mushroom body and lateral horn). h APF: hours after puparium formation; h ALH: hours after larval hatching. DL: dorsolateral; DM: dorsomedial; VM: ventromedial; VL: ventrolateral. Scale bar = 10 μ m.

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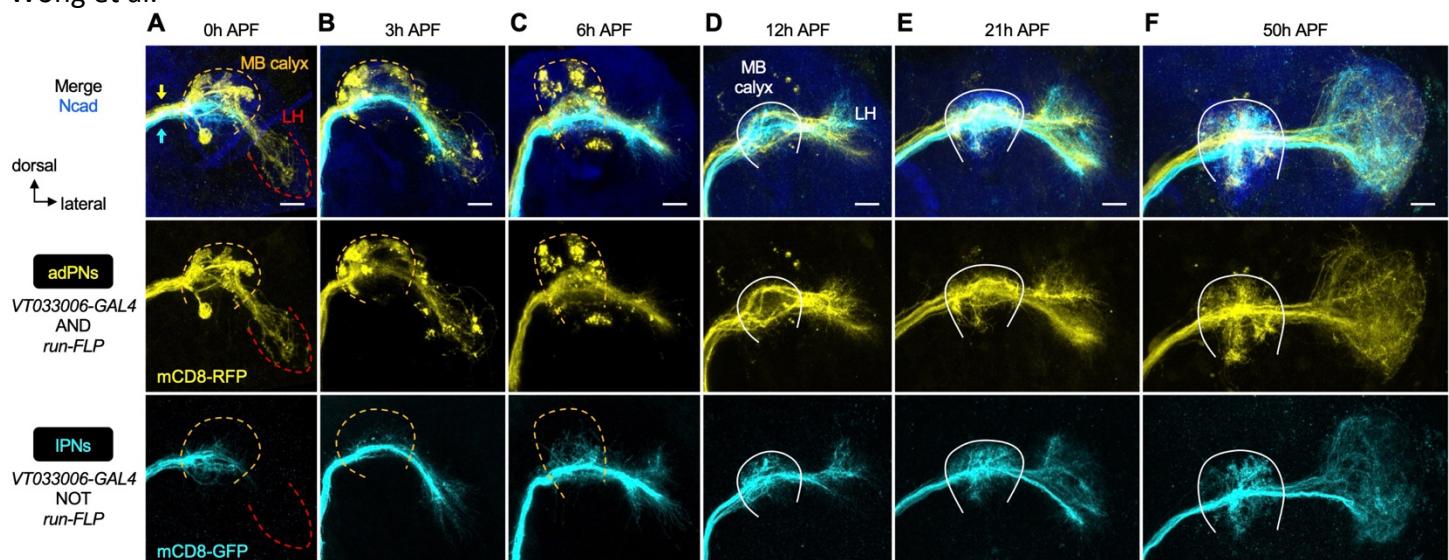


Figure 1 – figure supplement 2. PN axon development across pupal stages.

(A–F) Staining of fixed brains at indicated stages showing axon development of adPNs (*VT033006+ run+*; labeled in yellow) and IPNs (*VT033006+ run-*; labeled in cyan). Yellow and cyan arrows in A indicate the segregation of adPN and IPN axons along the inner antennocerebral tract. MB: mushroom body; LH: lateral horn. MB calyx (where PN axons and Kenyon cells of the mushroom body form synapses) and LH neuropils (where PN axons form synapses with their postsynaptic target neurons) are outlined as follows. In A–C, orange dashed line denotes the degeneration of larval-specific MB calyx. In A, larval-specific LH located more ventrally is outlined by red dashed line. In D–F, the developing adult-specific MB calyx is outlined by white solid line and adult-specific LH is to the right of the calyx.

See Figure 1 legend for common notations.

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A

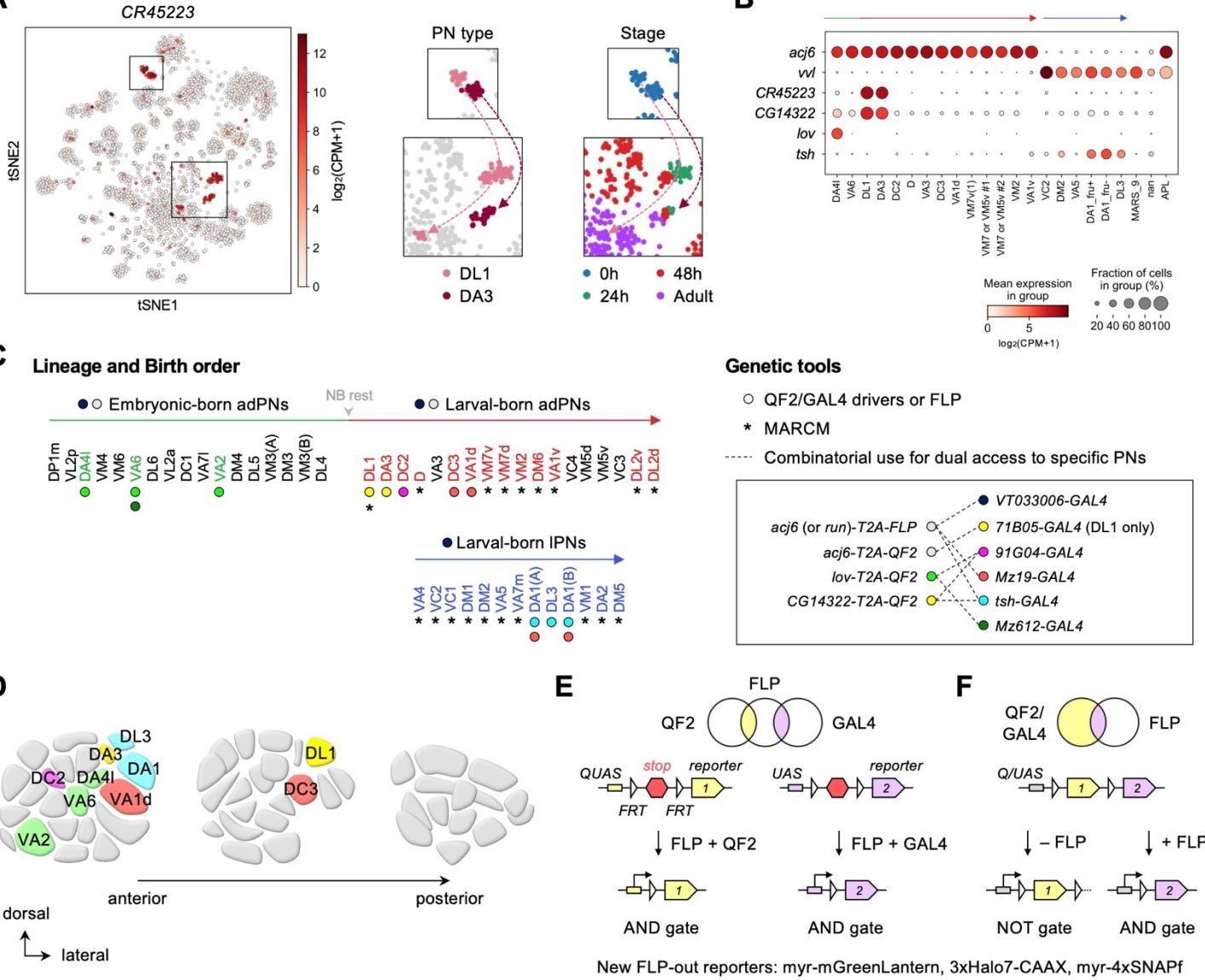


Figure 2. Expanded genetic toolkit for dual-color, type-specific labeling of PNs.

(A) tSNE plot of PN single-cell transcriptomes, color-coded according to *CR45223* expression level in $[\log_2(\text{CPM}+1)]$, where CPM stands for transcript counts per million reads. Zoom-in of boxes in the tSNE plot (left) are shown on the right, color-coded according to PN types and developmental stages.

(B) Dot plot showing the expression of *acj6*, *vvl*, *CR45223*, *CG14322*, *lov*, and *tsh* in 0h APF PNs arranged according to their birth order and lineage (green: embryonic-born adPNs; red: larval-born adPNs; blue: larval-born lPNs). Unit of expression is $[\log_2(\text{CPM}+1)]$ as in (A).

(C) Birth orders of adPNs and iPNS [summarized from (Lin et al., 2012; Yu et al., 2010)] and genetic tools used to access them. **Left:** Accessible PN types are colored. Circles beneath the PN types denote *QF2/GAL4* drivers used to access them. Asterisks beneath the PN types denote access by MARCM. Grey arrowhead marks neuroblast (NB) rest. **Right:** Genetic tools. Inset shows combinatorial use of *QF2/FLP* and *GAL4* (linked by dashed lines) for comparative analyses of dendrite development of two groups of PNs in the same animal.

(D) Schematic of glomerular projections of *QF2/GAL4*-accessible PNs in the adult antennal lobe. Indicated glomeruli are color-coded based on the genetic tools used to access them. See the color code in **Figure 2C**. **(E, F)** Schematic of intersectional logic gates for dual-color labeling of PNs. See **Figure 2 – figure supplement 2** for newly generated FLP-out reporters.

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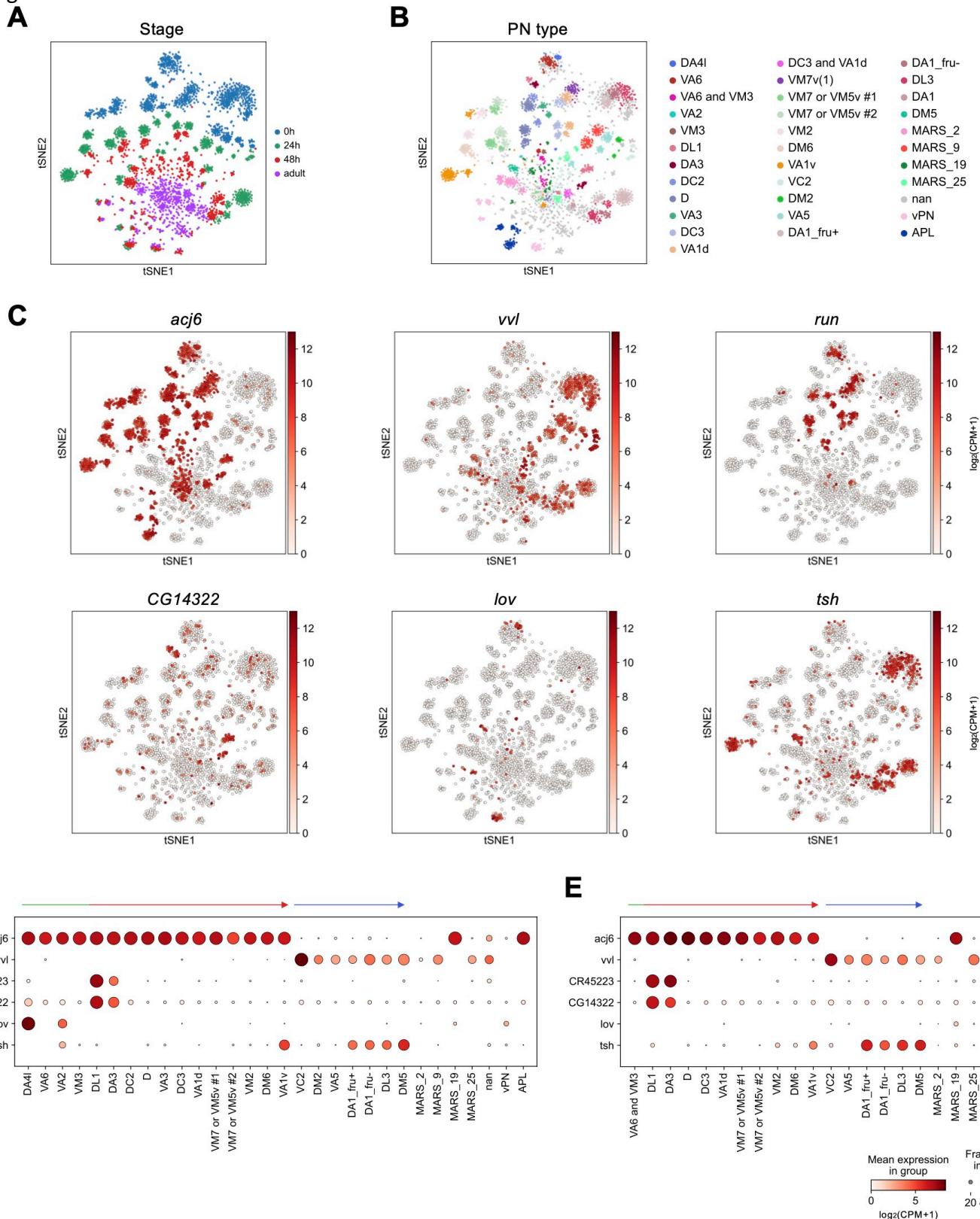


Figure 2 – figure supplement 1. Expression of PN marker genes across development.

(A–C) tSNE plots of PN transcriptomes, color-coded according to developmental stages (A), PN types (B), or the expression levels of indicated gene [$\log_2(\text{CPM}+1)$] (C) using scRNA-seq data from Xie et al., 2021. (D, E) Dot plot showing the expression levels of *acj6*, *vvl*, *CR45223*, *CG14322*, *lov*, and *tsh* in PNs [$\log_2(\text{CPM}+1)$] at 24h APF (D) and 48h APF (E).

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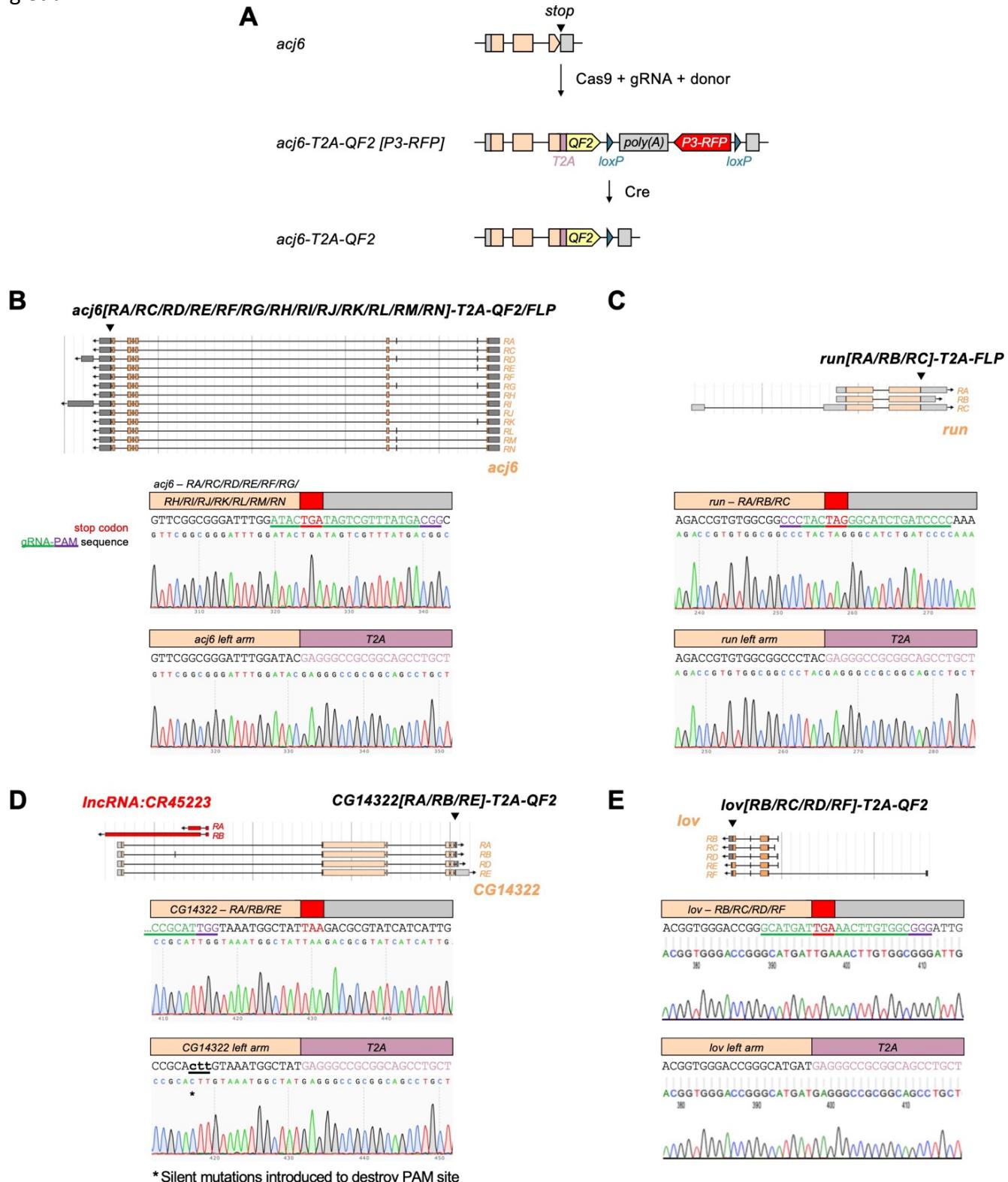
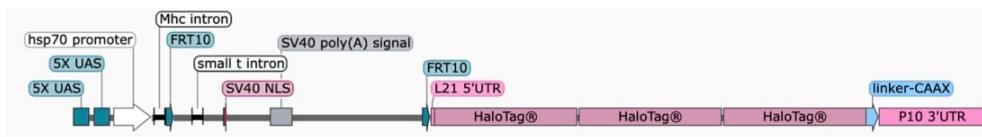


Figure 2 – figure supplement 2. Generation of T2A-QF2/FLP transgenic flies by CRISPR/Cas9.
(A) Schematic of generation of transgenic driver lines by CRISPR/Cas9. *acj6-T2A-QF2* is shown as an example.
(B-E Top) Transcripts of *acj6* (B), *run* (C), *CG14322* (D), and *lov* (E) visualized using FlyBase JBrowse.
(B-E Bottom) Targeted insertion of T2A-QF2/FLP right before the stop codon of the endogenous gene. Stop codon and gRNA-PAM sequence are color-coded as indicated.

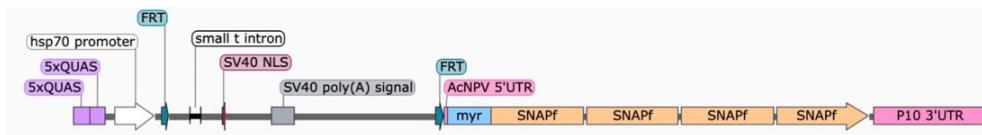
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- Single-color FLP-out reporters

UAS-FRT¹⁰-stop-FRT¹⁰-3xHalo7-CAAX

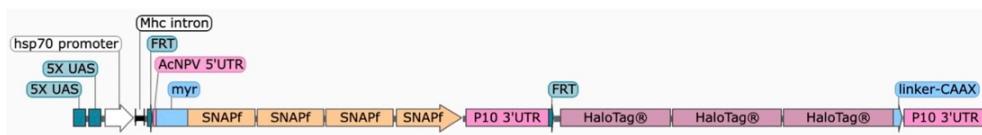


QUAS-FRT-stop-FRT-myr-4xSNAPf

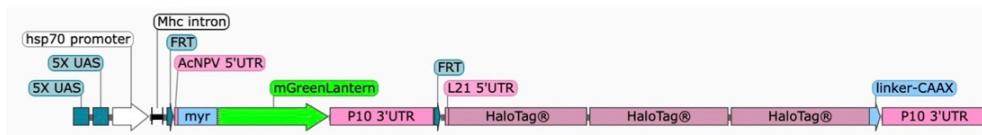


- Dual-color FLP-out reporters

UAS-FRT-myr-4xSNAPf-FRT-3xHalo7-CAAX



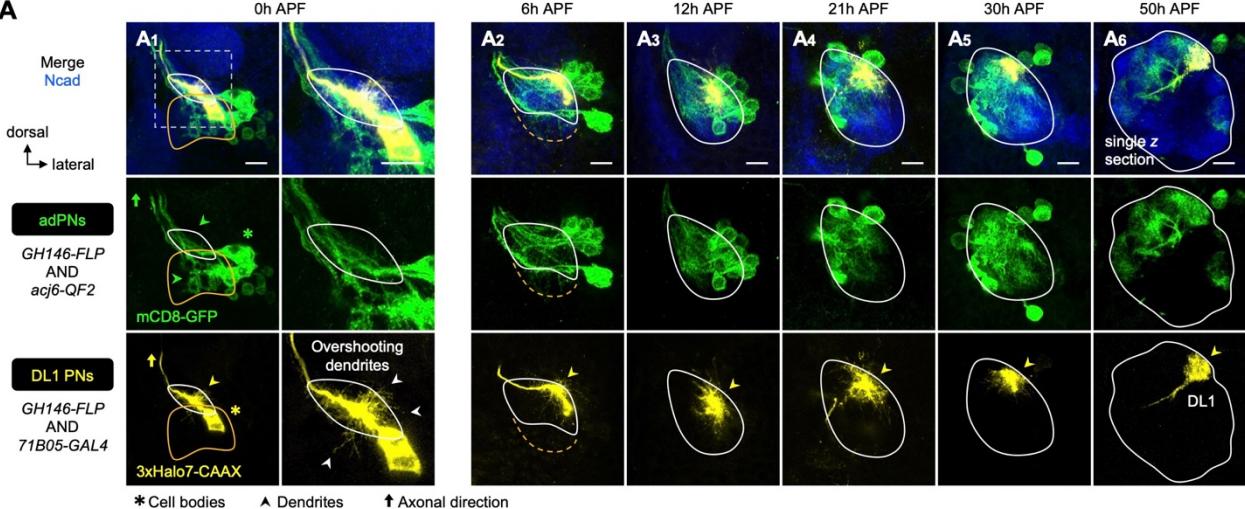
UAS-FRT-myr-mGreenLantern-FRT-3xHalo7-CAAX



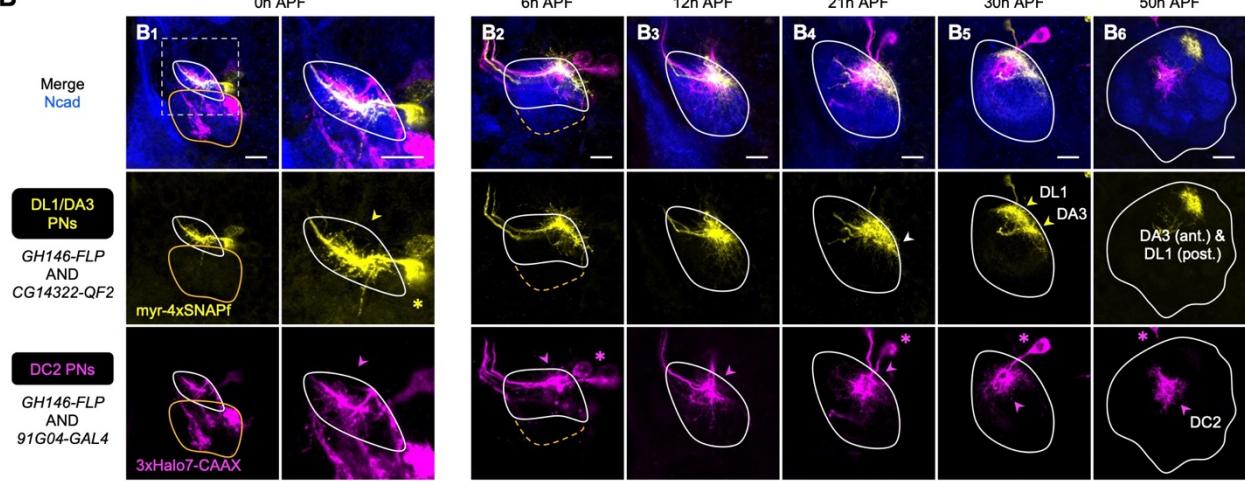
7
8
9 **Figure 2 – figure supplement 3. Design of single- and dual-color FLP-out reporters.**

10 Images, created with SnapGene, show four newly generated *Q/UAS*-based single- and dual-color FLP-out
11 reporters.
12

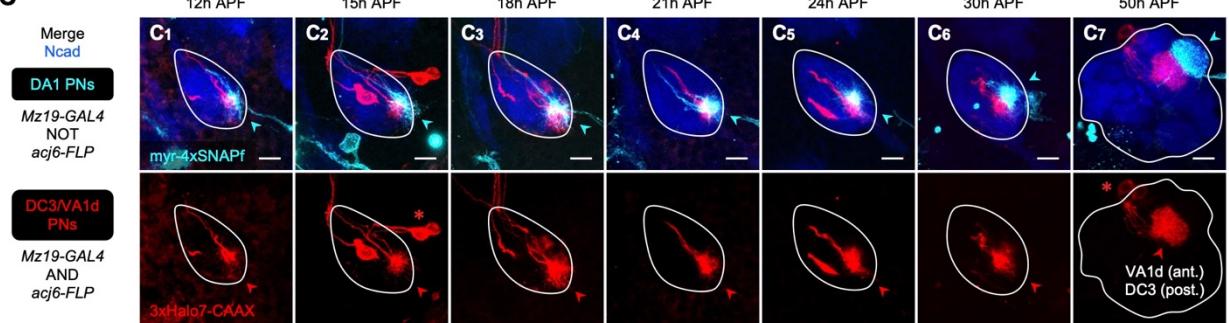
Wong et al. A



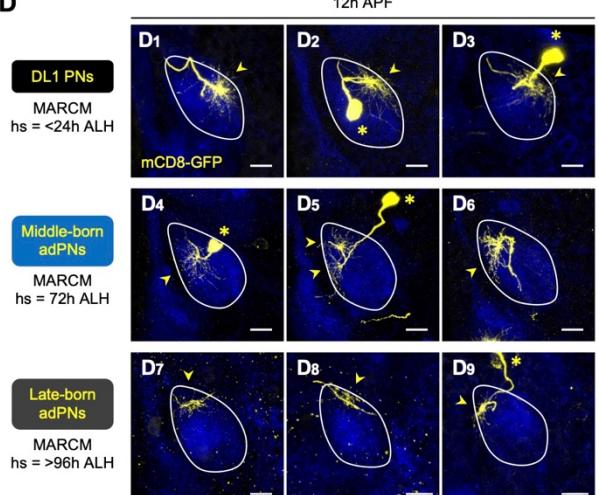
B



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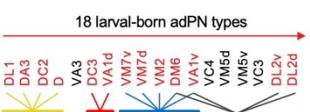


D



E Wiring logic of larval-born adPN dendrites during development

1) Grouping of PNs by birth order



2) Adjacent birth order → similar dendrite targeting



3) Distant birth order → distinct dendrite targeting



4) Birth order correlates with dendrite targeting in a rotating** manner.



5) First-born, first-developed

* We do not see a correlation between birth order and cell body position.
** Clockwise in the right hemisphere; anti-clockwise in the left

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Figure 3. Birth order-dependent spatial patterning of adPN dendrites in the developing antennal lobe.

(A) Confocal images of fixed brains at indicated stages showing dendrite development of adPNs (*acj6*⁺; labeled in green) and DL1 adPNs (*71B05*⁺; labeled in yellow). Right column of **A**₁ shows zoom-in of the dashed box.

The labeling of *acj6*⁺ adPNs outlines the developing antennal lobe and is used in dual-color AO-LLSM imaging later (see **Figure 7A–C**). White arrowheads in **A**₁ mark dendrites overshooting the antennal lobe.

(B) Confocal images of fixed brains at indicated stages showing dendrite development of DL1/DA3 adPNs (*CG14322*⁺; labeled in yellow) and DC2 adPNs (*91G04*⁺; labeled in magenta). As *91G04-GAL4* labels some embryonic-born PNs from 0–6h APF, their neurites are found in the larval-specific antennal lobe (**B**_{1,2}). Right column of **B**₁ shows zoom-in of the dashed box. White arrowhead in **B**₄ denotes the more ventrally targeted DL1/DA3 dendrites.

(C) Confocal images of fixed brains at indicated stages showing dendrite development of DC3/VA1d adPNs (*Mz19*⁺ *acj6*⁺; labeled in red) and DA1 IPNs (*Mz19*⁺ *acj6*⁻; labeled in cyan).

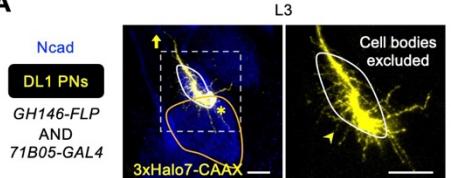
(D) Confocal images of single-cell MARCM clones (in yellow) of DL1 PNs (**D**_{1–3}), middle larval-born adPNs (**D**_{4–6}) and late larval-born adPNs (**D**_{7–9}) in 12h APF pupal brains, generated by heat shocks (hs) at indicated times.

(E) Summary of wiring logic of larval-born adPN dendrites to form olfactory map in the 12h APF developing antennal lobe.

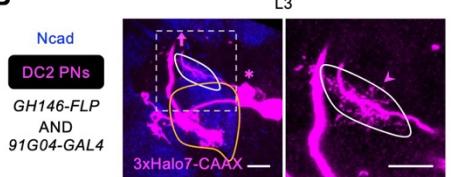
See Figure 1 legend for common notations.

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A



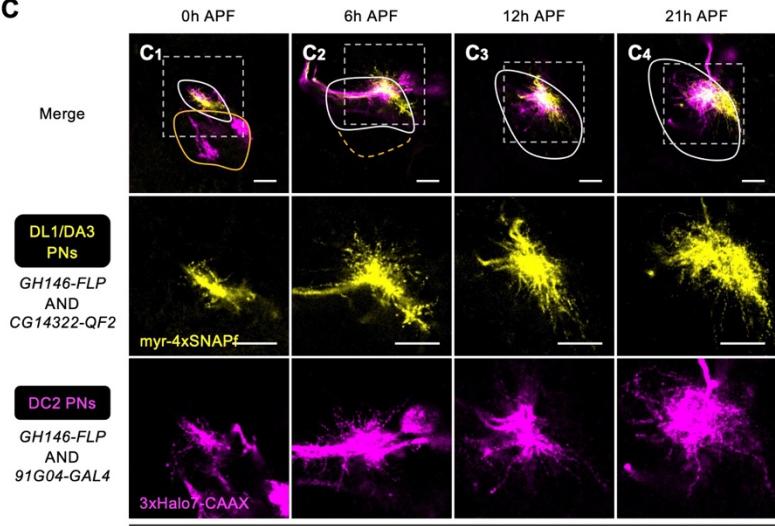
B



dorsal
↑ lateral

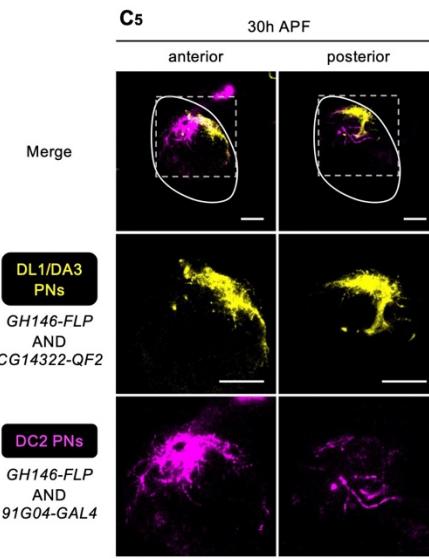
* Cell bodies
▲ Dendrites
↑ Axonal direction

C

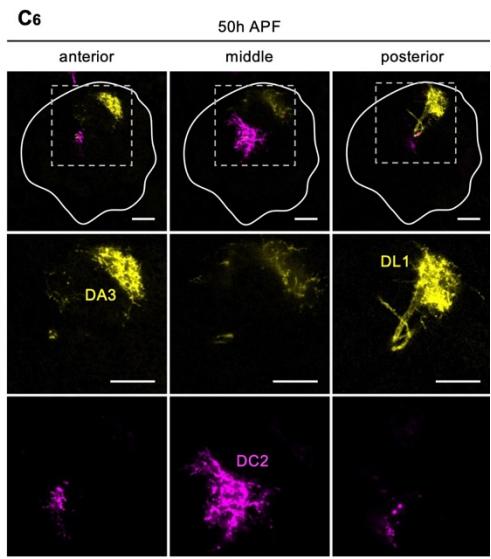


Single z sections of Figure 3B

C5



C6



Single z sections of Figure 3B

3

4

Figure 3 – figure supplement 1. Dendrite development of early larval-born PNs.

(A) Dendritic extension of DL1 PNs (*71B05*); labeled in yellow) across the developing antennal lobe at the wandering third instar larval stage (L3). Zoom-in of the dashed box shown on the right.

(B) Dendritic extension of DC2 PNs (*91G04*; labeled in yellow) across the developing antennal lobe at L3. Zoom-in of the dashed box shown on the right.

(C) Single z sections of Figure 3B showing dendrite development of DL1/DA3 adPNs (*CG14322*; labeled in yellow) and DC2 adPNs (*91G04*; labeled in magenta).

See Figure 1 legend for common notations.

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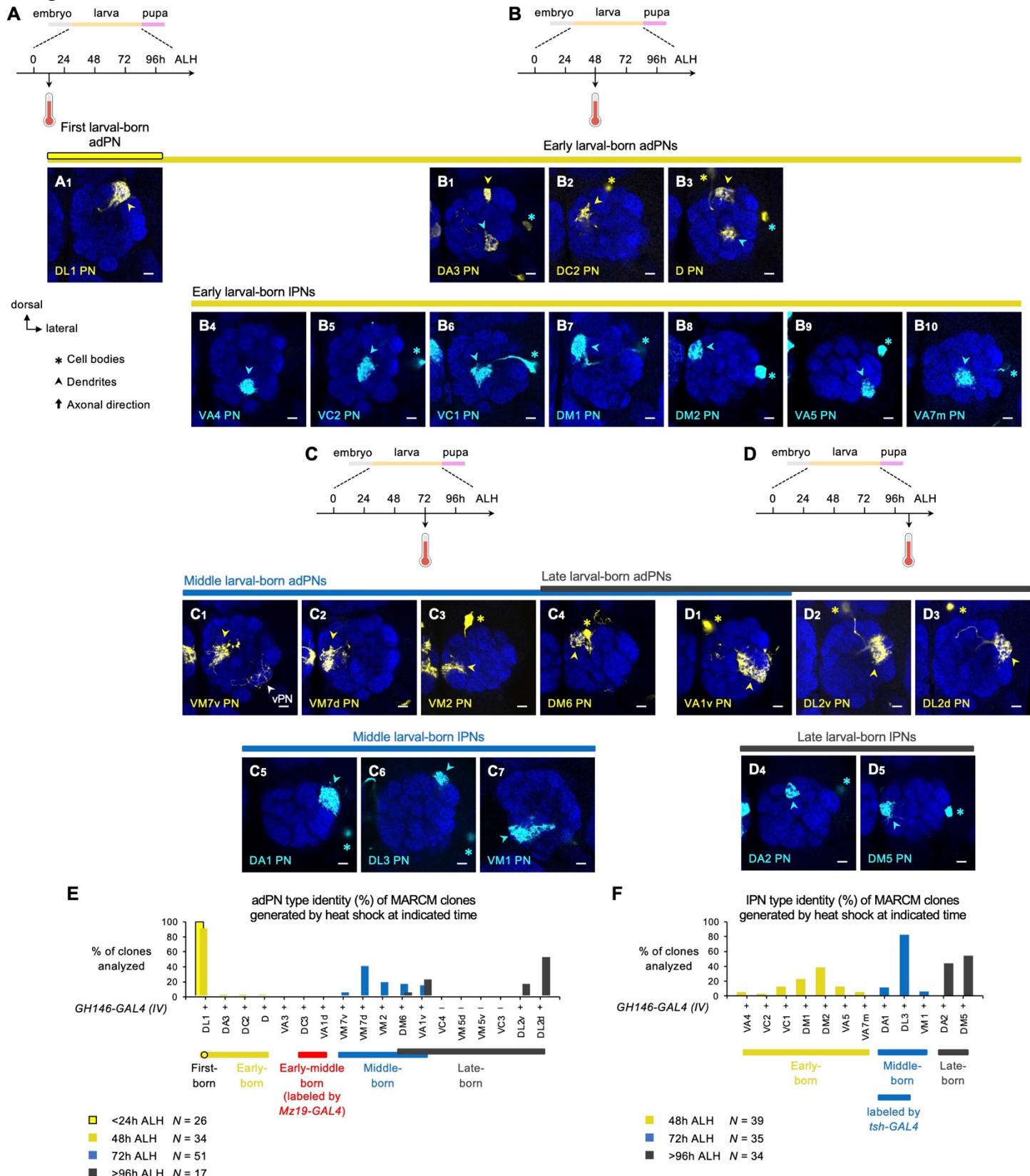


Figure 3 – figure supplement 2. MARCM-labeled single-cell PNs of indicated lineages in adult brains.

(A) MARCM clone of DL1 PN (in yellow) generated by heat shock at <24h ALH.

(B) MARCM clones of early larval-born PNs (B₁₋₃: adPNs in yellow; B₄₋₁₀: IPNs in cyan) generated by heat shock at 48h ALH. In B₁, single-cell clone of the adPN lineage and that of the IPN lineage, corresponding to

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i0 DA3 PN (yellow arrowhead) and VA5 PN (cyan asterisk and arrowhead), were simultaneously generated. In **B₃**,
i1 single-cell adPN and IPN, corresponding to D PN (yellow asterisk and arrowhead) and VA7m PN (cyan asterisk
i2 and arrowhead), were simultaneously generated.

i3 **(C)** MARCM clones of middle larval-born PNs (**C₁₋₄** and **D₁**: adPNs in yellow; **C₅₋₇**: IPNs in cyan) generated
i4 by heat shock at 72h ALH. In **C₁**, white arrowhead mark processes of vPN clone that do not belong to VM7v
i5 PN.

i6 **(D)** MARCM clones of late larval-born PNs (**C₄** and **D₁₋₃**: adPNs in yellow; **D₄₋₅**: IPNs in cyan) generated by
i7 heat shock at >96h ALH.

i8 **(E)** Percentage bar graph showing the adPN type identity of MARCM clones generated by heat shock at
i9 indicated times. Sample size *N* indicates the number of clones analyzed.

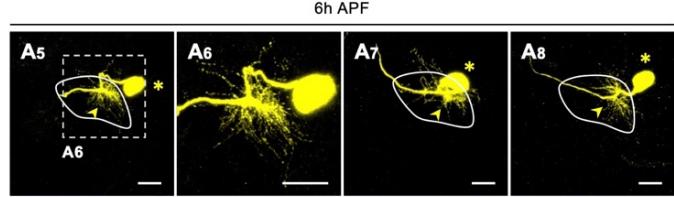
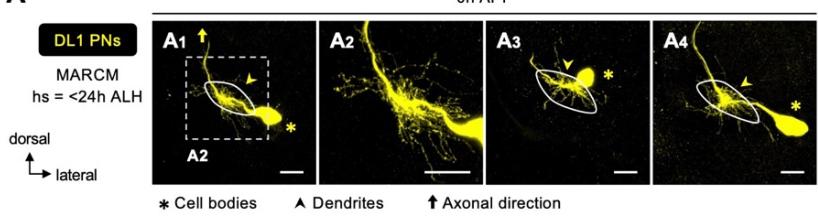
i0 **(F)** Percentage bar graph showing the IPN type identity of MARCM clones generated by heat shock at indicated
i1 times. Sample size *N* indicates the number of clones analyzed.

i2 See Figure 1 legend for common notations.

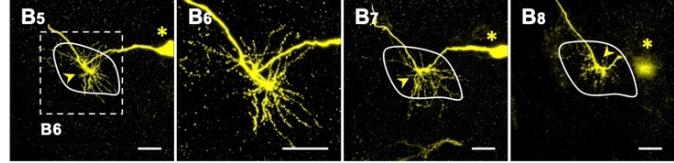
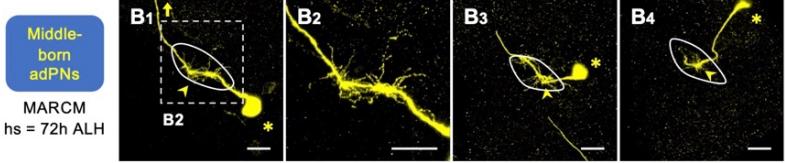
i3

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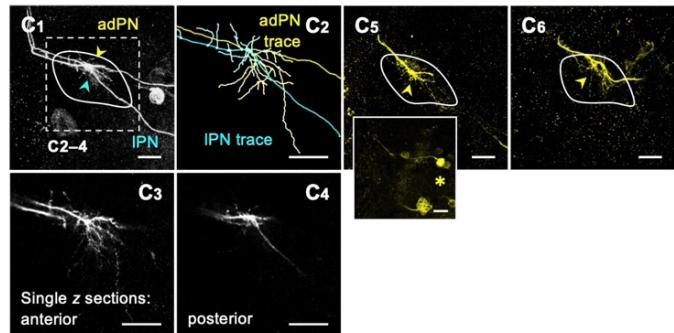
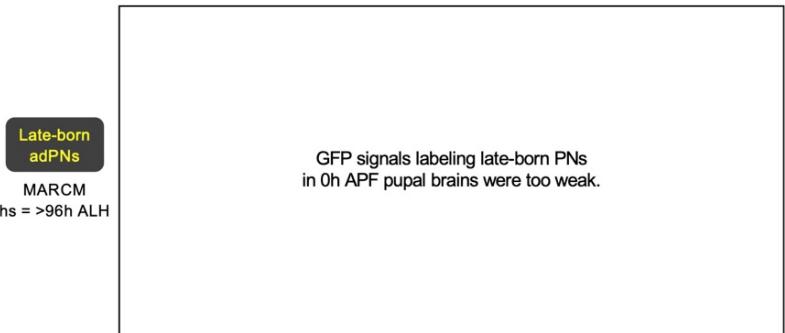
A



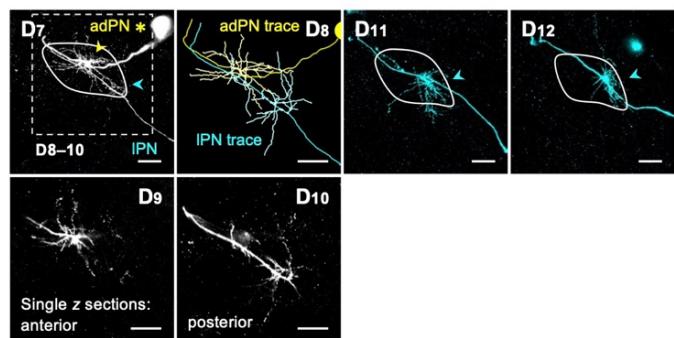
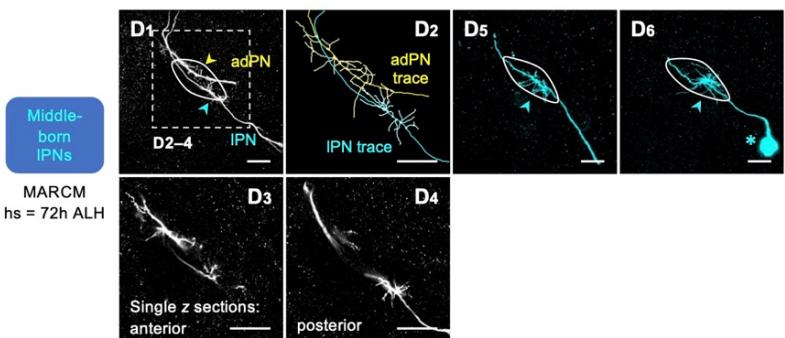
B



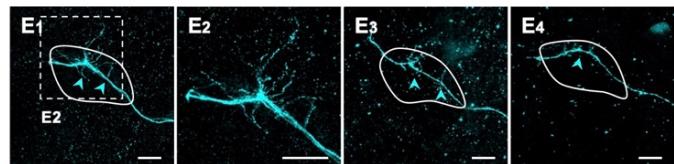
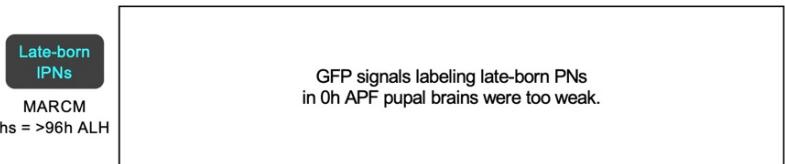
C



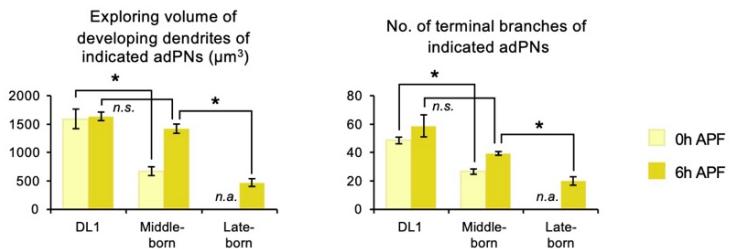
D



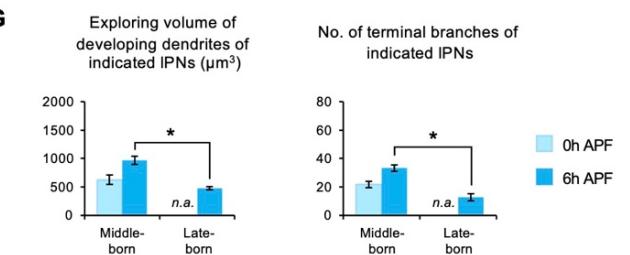
E



F



G



'4

'5 Figure 3 – figure supplement 3. Dendrite development of DL1, middle larval-born, and late larval-born PNs at early stages.

'6 Adult-specific antennal lobes (revealed by high Ncad staining; not shown) are outlined by white solid line.

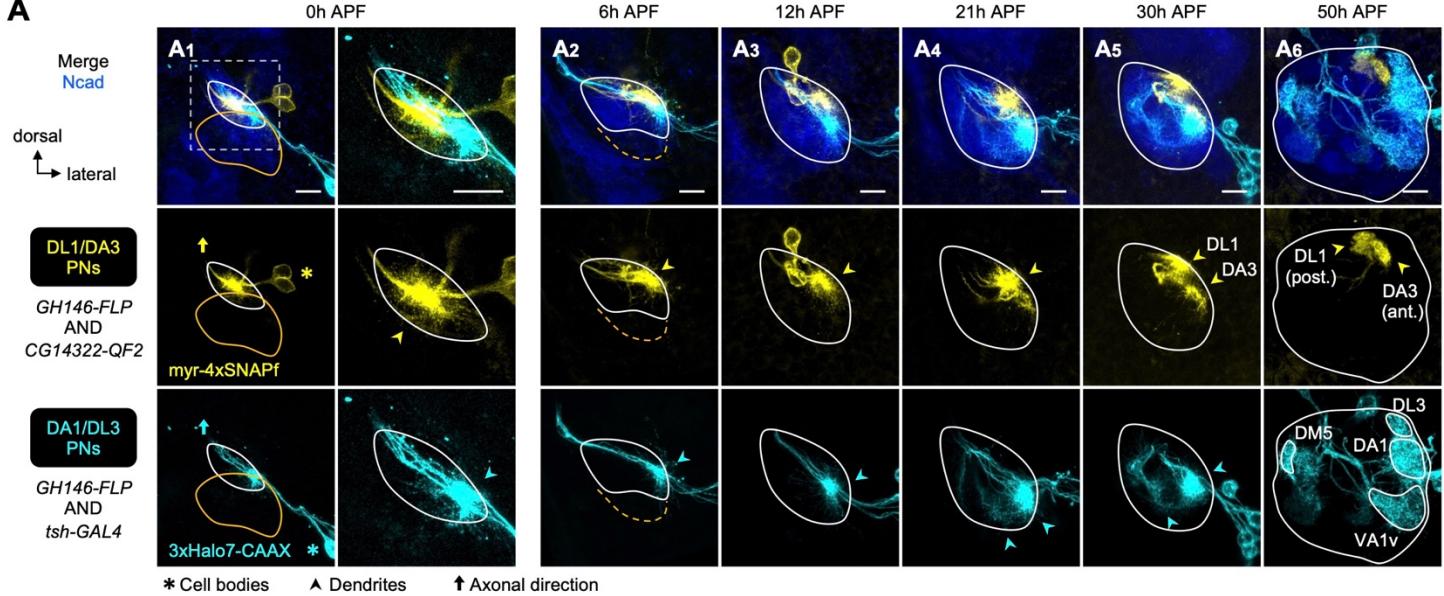
'7 (A) MARCM clones of DL1 PNs (in yellow), generated by heat shock at <24h ALH, in 0h APF (A₁₋₄) and 6h APF (A₅₋₈) pupal brains.

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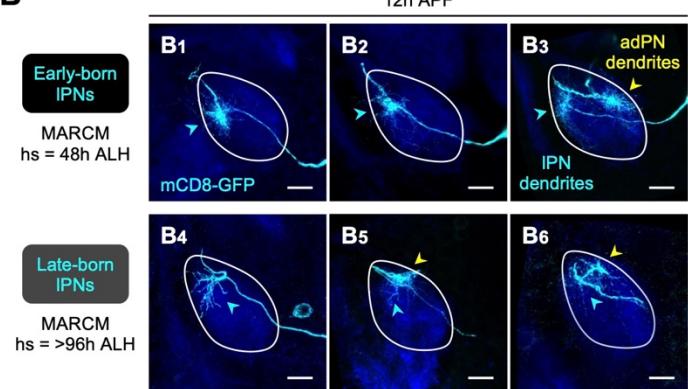
0 (B) MARCM clones of middle larval-born adPNs (in yellow), generated by heat shock at 72h ALH, in 0h APF
1 (**B₁₋₄**) and 6h APF (**B₅₋₈**) pupal brains.
2 (C) MARCM clones of late larval-born adPNs (in yellow), generated by heat shock at >96h ALH, in 6h APF
3 pupal brains. In **C₁**, single-cell MARCM clones of adPN and IPN lineages were simultaneously labeled. **C₂**
4 shows neurite tracing of adPN (in yellow) and IPN (in cyan) in **C₁**. Single *z* sections of **C₁** are shown in **C₃₋₄**.
5 Small inset below **C₅** reveals the cell body position.
6 (D) MARCM clones of middle larval-born IPNs (in cyan), generated by heat shock at 48h ALH, in 0h APF (**D₁₋₆**)
7 and 6h APF (**D₇₋₁₂**) pupal brains. In **D₁₋₄** and **D₇₋₁₀**, single-cell adPN and IPN were simultaneously labeled. **D₂**
8 and **D₈** shows neurite tracing of **D₁** and **D₇**, respectively (adPN in yellow; IPN in cyan). Single *z* sections of **D₁**
9 are shown in **D₃₋₄**, and those of **D₇** are shown in **D₉₋₁₀**.
10 (E) MARCM clones of late larval-born IPNs (in cyan), generated by heat shock at >96h ALH, in 6h APF pupal
11 brains.
12 (F–G) Quantification of exploring volume of developing dendrites of indicated PNs (F: adPNs; G: IPNs) at 0h
13 and 6h APF (left). Quantification of number of terminal branches of indicated PNs (F: adPNs; G: IPNs) at 0h
14 and 6h APF (right). Error bars, SEM; *t* test; *, $p < 0.05$; n.s., $p \geq 0.05$. SEM, standard error of the mean; n.s., not
15 significant; n.a., not applicable.
16

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A

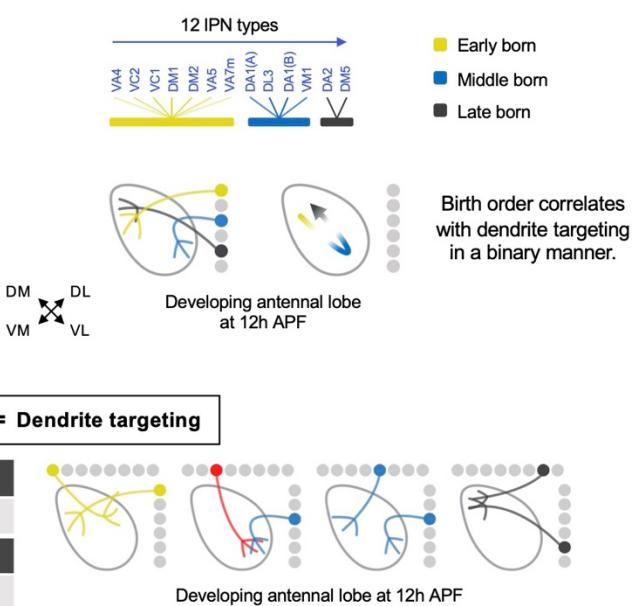


B



C

Wiring logic of IPN dendrites during development



D

[Lineage; Birth Order] = Dendrite targeting

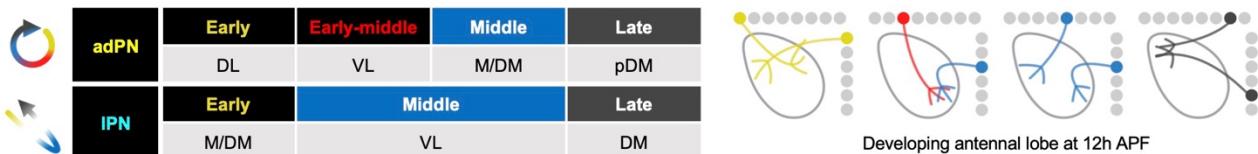


Figure 4. Birth order-dependent spatial patterning of IPN dendrites in the developing antennal lobe.
(A) Confocal images of fixed brains at indicated stages showing dendrite development of DL1/DA3 adPNs (*CG14322*⁺; labeled in yellow) and DA1/DL3 IPNs (*tsh*⁺; labeled in cyan). Right column of A₁ shows zoom-in of the dashed box.
(B) MARCM clones (in cyan) of early (B₁₋₃) and late (B₄₋₆) larval-born IPNs in 12h APF pupal brains, generated by heat shocks (hs) at indicated times. In B₃, B₅, and B₆, single-cell clones of adPN (yellow arrowheads) and IPN (cyan arrowheads) lineages were simultaneously labeled.
(C) Summary of wiring logic of larval-born IPN dendrites to form olfactory map in the 12h APF developing antennal lobe.
(D) Summary of determination of dendrite targeting of larval-born PNs by lineage and birth order.
See Figure 1 legend for common notations.

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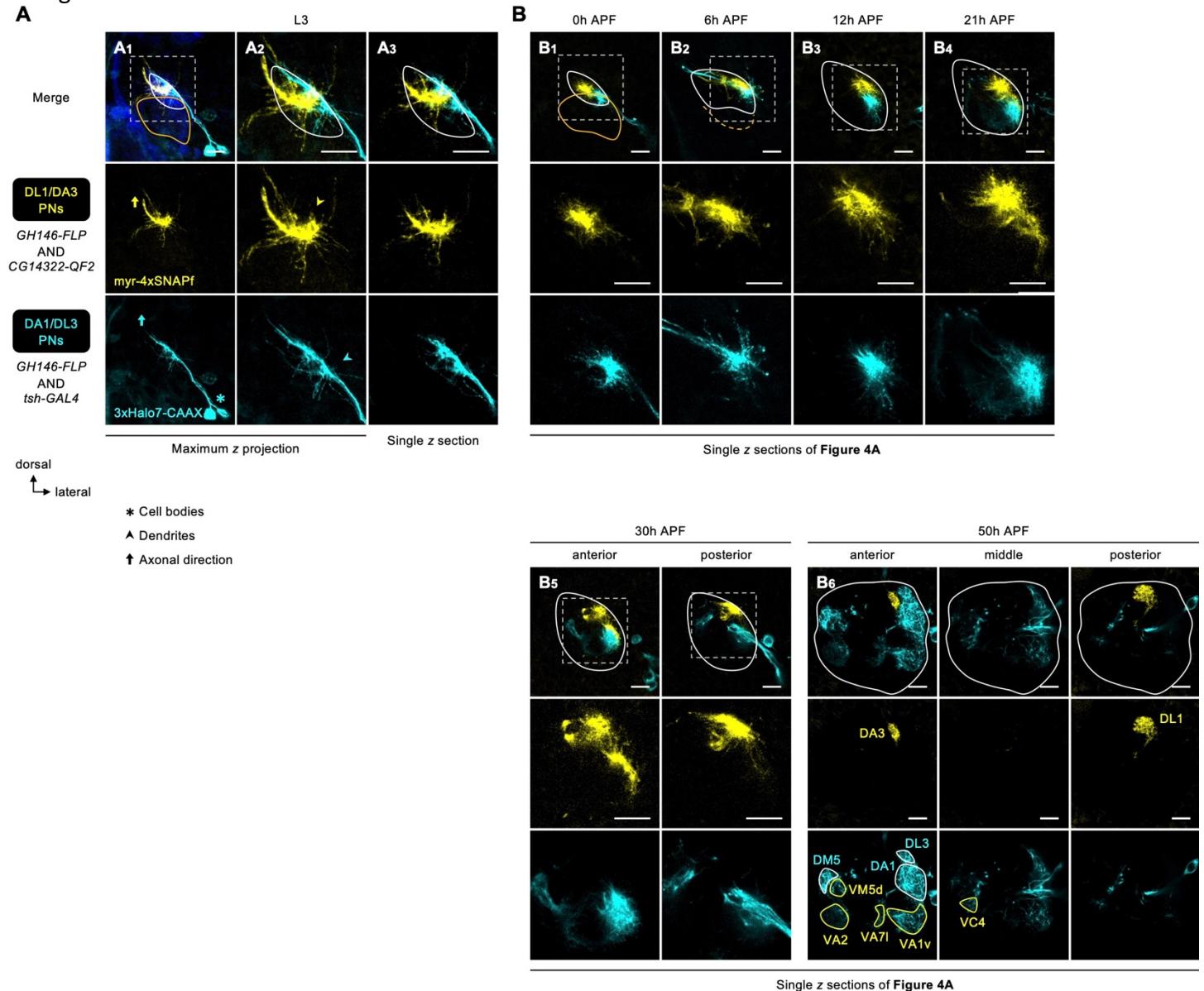


Figure 4 – figure supplement 1. Dendrite development of DL1/DA3 and DA1/DL3 PNs.
 (A) Dendritic extension of DL1/DA3 PNs (*CG14322*+/−; labeled in yellow) and DA1/DL3 PNs (*tsh*+/−; labeled in cyan) across the developing antenna lobe at the wandering third instar larval stage (L3).
 (B) Single *z* sections of Figure 4A showing dendrite development of DL1/DA3 PNs (*CG14322*+/−; labeled in yellow) and DA1/DL3 PNs (*tsh*+/−; labeled in cyan). In B6, glomeruli innervated by *tsh*+ adPN and *tsh*+ IPN dendrites are outlined in yellow and cyan, respectively.
 See Figure 1 legend for common notations.

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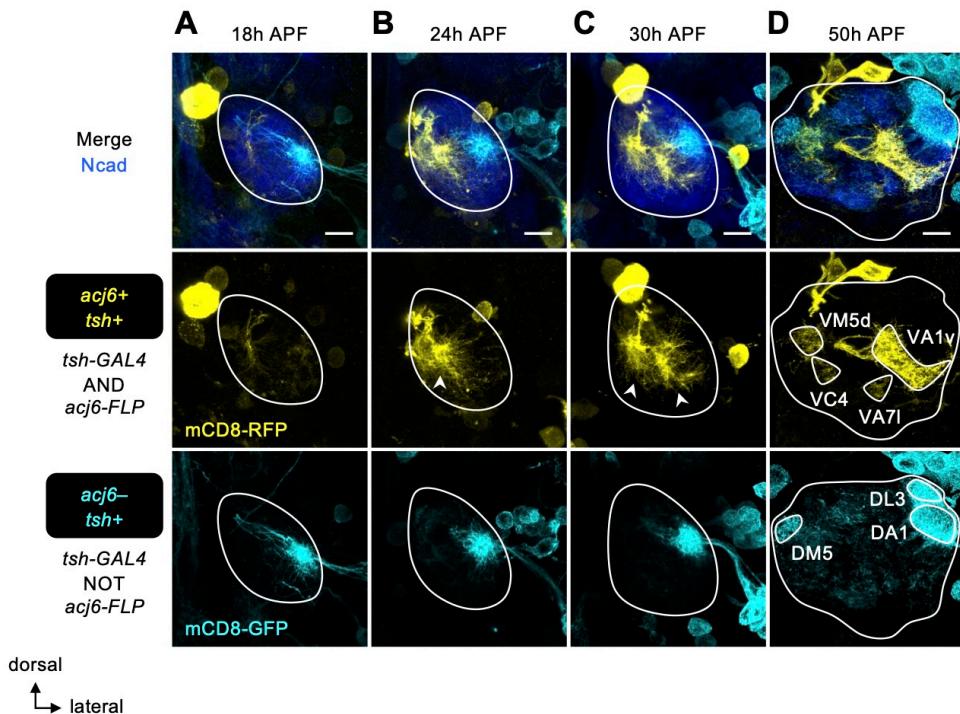


Figure 4 – figure supplement 2. Expression patterns of *tsh* in the developing antennal lobe during mid-pupal stages.

(A) At 18h APF, *tsh* is only expressed in PNs of the lateral lineage (*acj6-*; in cyan).

(B–D) From 24h APF onwards, *tsh* is expressed in some adPNs (*acj6+*; in yellow), consistent with the transcriptome data (Figure 2B, Figure 2 – figure supplement 1C–E). *tsh-GAL4* seems to weakly label local interneurons at 50h APF (*acj6-*; in cyan).

See Figure 1 legend for common notations.

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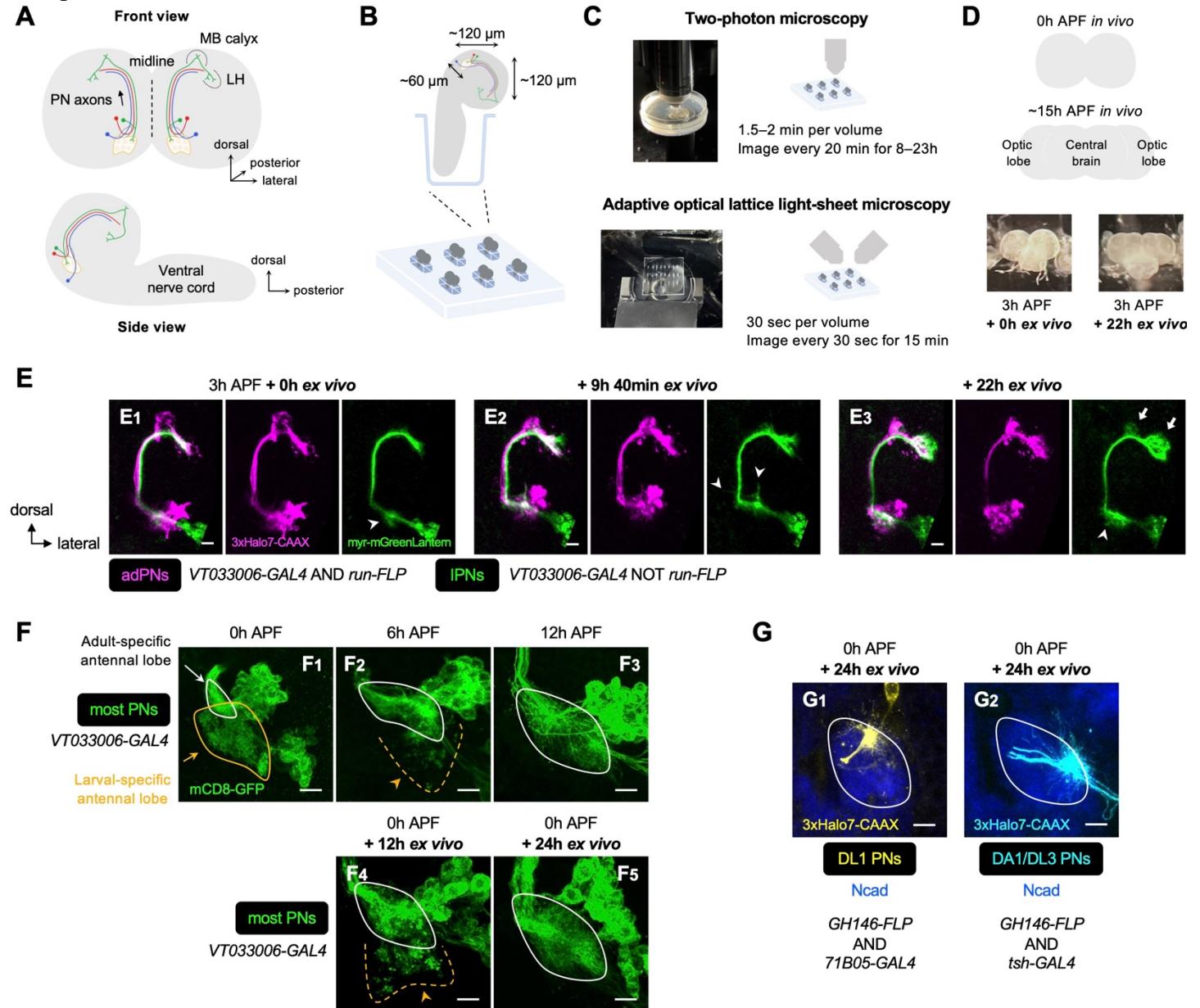


Figure 5. Establishment of explant system for time-lapse imaging of olfactory map formation.

(A) Schematic of anatomical organization of the olfactory circuit in early pupal brain (0–3h APF). Green, red, and blue denote embryonic-born adPN, larval-born adPN and larval-born IPN, respectively. MB: mushroom body; LH: lateral horn.

(B) Schematic of explant culture system for early pupal brains. Wells created in the Sylgard plate from which brains were imbedded are shown in blue.

(C) Schematic of explant culture and imaging system for early pupal brains.

(D) **Top:** Schematic of morphological changes of brain lobes from 0h to ~15h APF during normal development. **Bottom:** Morphologies of a brain explant dissected at 3h APF and cultured for 0h *ex vivo* and that cultured for 22h *ex vivo*.

(E) Two-photon time-lapse imaging of adPNs (*VT033006+ run+*; labeled in magenta) and IPNs (*VT033006+ run-*; labeled in green) in pupal brain dissected at 3h APF and cultured for 0–22h *ex vivo*. Arrowheads mark dynamic but transient dendritic protrusions of IPNs in E_{1,2}, and extensive dendritic innervation of IPNs in E₃. Arrows in E₃ mark axonal innervation of IPNs in mushroom body calyx and lateral horn.

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7 (F) Confocal images of antennal lobes labeled by *VT033006*⁺ PNs (in green) at 0h (**F**₁), 6h (**F**₂) and 12h (**F**₃)
8 APF *in vivo*. Confocal images of antennal lobes labeled by *VT033006*⁺ PNs in pupal brains dissected at 0h APF
9 and cultured for 12h (**F**₄) and 24h (**F**₅) *ex vivo*.

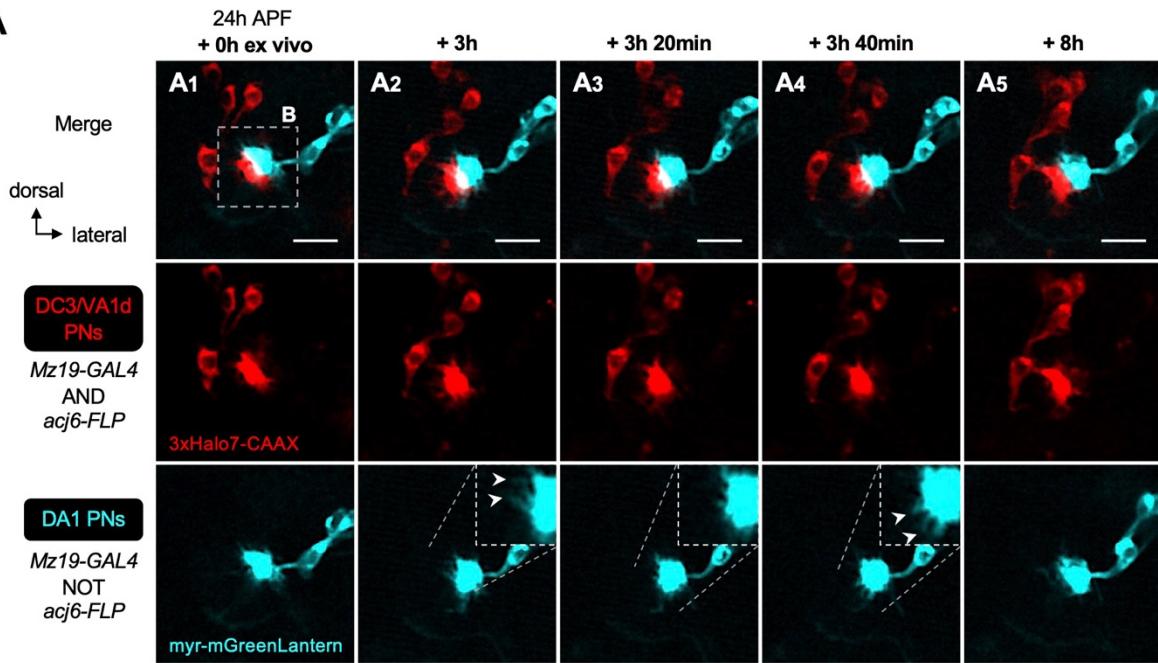
10 (G) Dendrite targeting regions of DL1 PNs (*71B05*⁺; in yellow; **G**₁) and DA1/DL3 PNs (*tsh*⁺; in cyan; **G**₂) in
11 the antennal lobes in pupal brains dissected at 0h APF and cultured for 24h *ex vivo*. Antennal lobes are revealed
12 by N-Cadherin (Ncad; in blue) staining.

13 See Figure 1 legend for common notations.

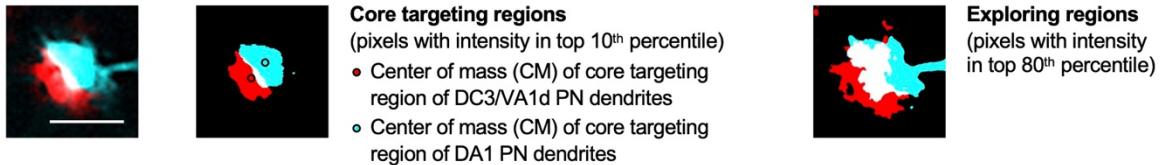
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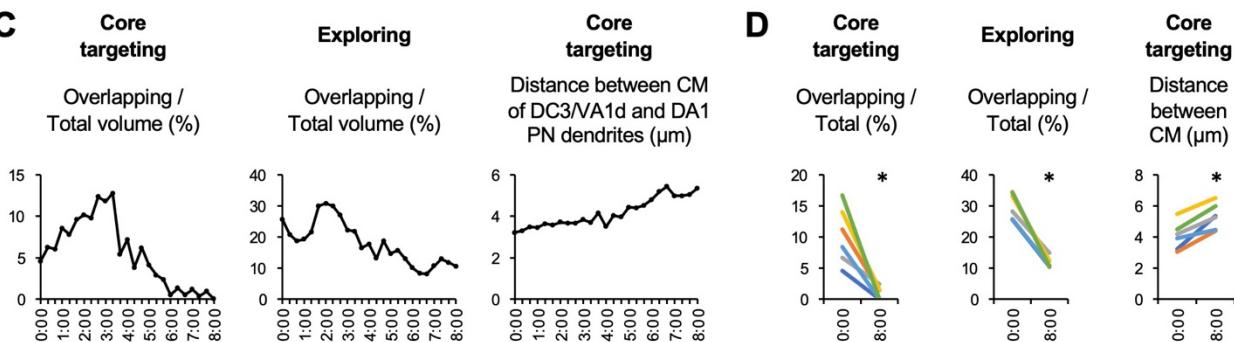
A



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C



D

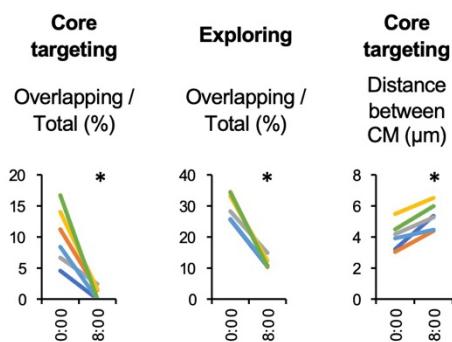


Figure 5 – figure supplement 1. Dendritic segregation of DC3/VA1d adPNs and DA1 IPNs targeting neighboring proto-glomeruli.

(A) Two-photon time-lapse imaging of DC3/VA1d adPN (*Mz19*+ *acj6*+; labeled in red) and DA1 IPN (*Mz19*+ *acj6*–; labeled in red) dendrites in pupal brain dissected at 24h APF and cultured for 8h *ex vivo*. Insets in A₂₋₄ show the zoom-in. Arrowheads in A₂₋₄ indicate the disappearance (compare A₂ with A₃) and extension (compare A₄ with A₃) of dendrites.

(B) Core targeting region of PN dendrites is defined using pixels with intensity in the top 10th percentile. Red and cyan circles mark the centers of mass of the core targeting regions of DC3/VA1d and DA1 PN dendrites, respectively. Exploring region of PN dendrites is defined using pixels with intensity in the top 80th percentile.

(C) **Left:** Ratio of overlapping to total core targeting volume (in percentage) across the 8-h imaging period. **Middle:** Ratio of overlapping to total exploring volume (in percentage) across the 8-h imaging period. **Right:** Distance between centers of mass of DC3/VA1d and DA1 core targeting regions across the 8-h imaging period. Sample size $N = 1$. Timestamp 00:00 refers to HH:mm; H, hour; m, minute.

(D) **Left:** Ratio of overlapping to total core targeting volume (in percentage) at 0h and 8h *ex vivo*. **Middle:** Ratio of overlapping to total exploring volume (in percentage) at 0h and 8h *ex vivo*. **Right:** Distance between

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'2 centers of mass of DC3/VA1d and DA1 core targeting regions at 0h and 8h *ex vivo*. Sample size $N = 6$. Error
'3 bars, standard error of the mean; t test; *, $p < 0.05$. Timestamp 00:00 refers to HH:mm; H, hour; m, minute.

'4

'5 **Figure 5 – video 1. Two-photon time-lapse imaging of PN development.**

'6 See **Figure 5E** for details. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

'7

'8 **Figure 5 – video 2. Two-photon time-lapse imaging of PN dendritic segregation.**

'9 See **Figure 5 – figure supplement 1** for details. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute;
'0 s, second.

'1

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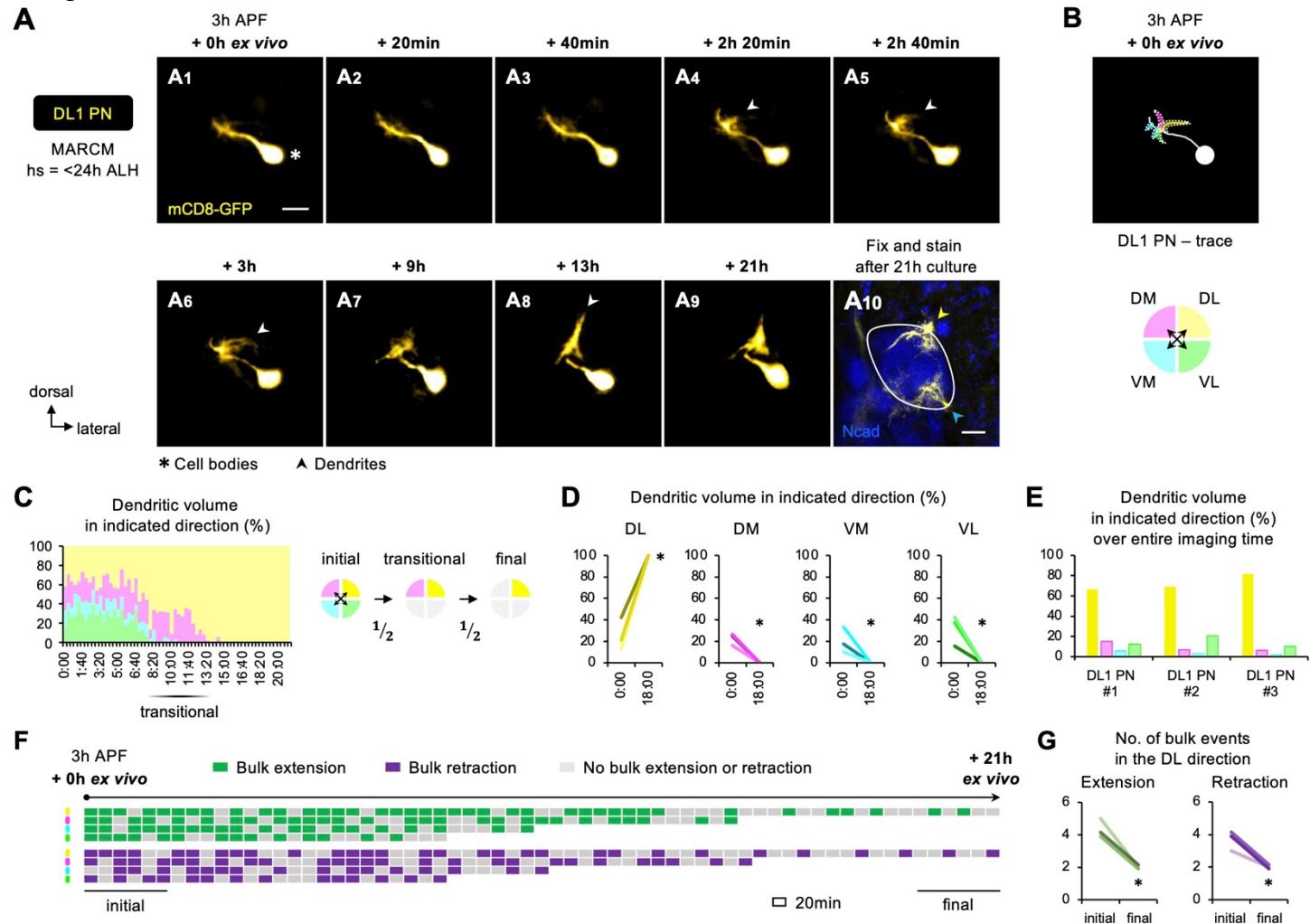


Figure 6. Two-photon time-lapse imaging reveals active dendrite targeting.

(A) Two-photon time-lapse imaging of MARCM-labeled DL1 PN (pseudo-colored in yellow) in a brain dissected at 3h APF and cultured for 21h *ex vivo* (A₁₋₉). Arrowheads in A₄₋₆ denote protrusions of dendritic branches towards the dorsolateral direction. After 21h culture, explant was fixed and immune-stained for N-Cadherin (Ncad; in blue) to outline the developing antennal lobe (A₁₀). Yellow and cyan arrowheads indicate DL1 PN dendrites and processes of other *GH146*⁺ cells, respectively.

(B) Neurite tracing of DL1 PN at the beginning of live imaging (3h APF + 0h *ex vivo*). Dendrites are categorized based on the directions to which they extend and color-coded accordingly.

(C) Left: Quantification of the percentage of dendritic volume in indicated direction during the time-lapse imaging period reveals a transitional phase during which dendrites were found in only two out of the four directions. Right: Schematic of the initial, transitional and final phases during the course of targeting. ‘½’ denotes the reduction of available trajectory directions by half. Timestamp 00:00 refers to HH:mm; H, hour; m, minute.

(D) Quantification of the percentage of DL1 PN dendritic volume in indicated direction in 3h APF cultured brains at the beginning (0h *ex vivo*) and at/near the end of imaging (18h *ex vivo*). DL1 PN sample size = 3. *t* test; *, *p* < 0.05. Timestamp 00:00 refers to HH:mm; H, hour; m, minute.

(E) Quantification of the percentage of sum of DL1 PN dendritic volume in indicated directions throughout the entire imaging time. DL1 PN sample size = 3.

(F) Bulk dendrite dynamics of DL1 PN in **Figure 6A**. Each row represents bulk dendrite dynamics in the indicated direction (color coded as in **Figure 6B**) across the 22-h imaging period. Each block represents a 20-min window. Bulk extension (in green) and retraction (in magenta) events are defined as dendrites extending

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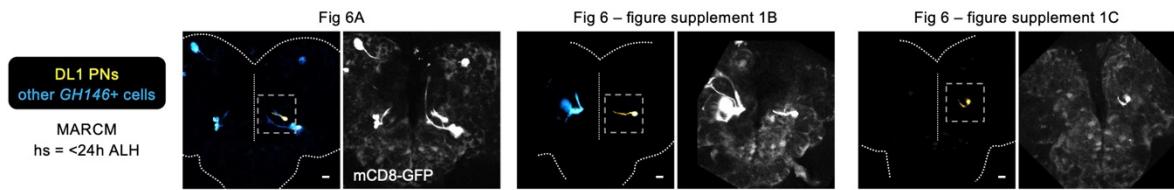
15 and retracting more than 2 μ m between two consecutive time windows. The first and last 6 consecutive
16 windows refer to the initial and final phases of imaging.

17 (G) Quantification of number of bulk extension and retraction events in the dorsolateral direction during the
18 initial and final phases of imaging. DL1 PN sample size = 3. *t* test; *, $p < 0.05$.

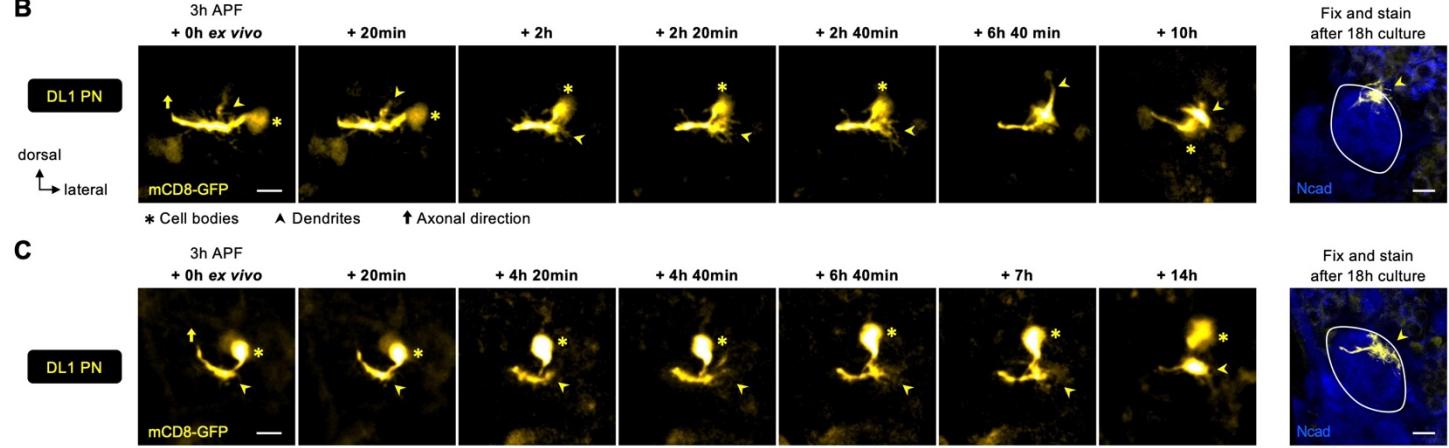
19

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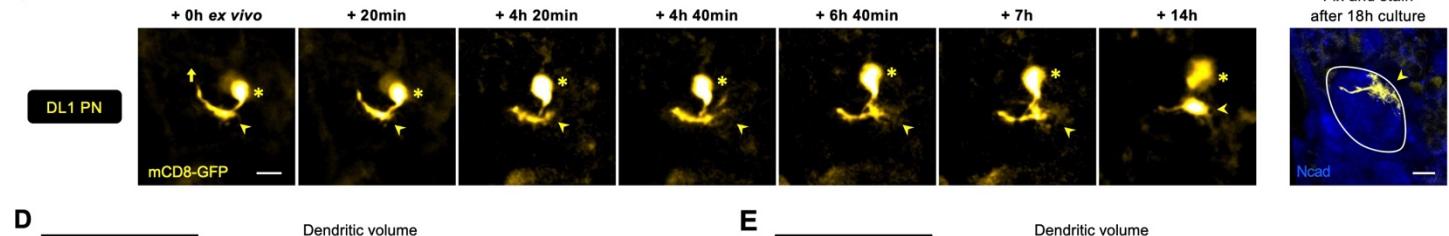
A



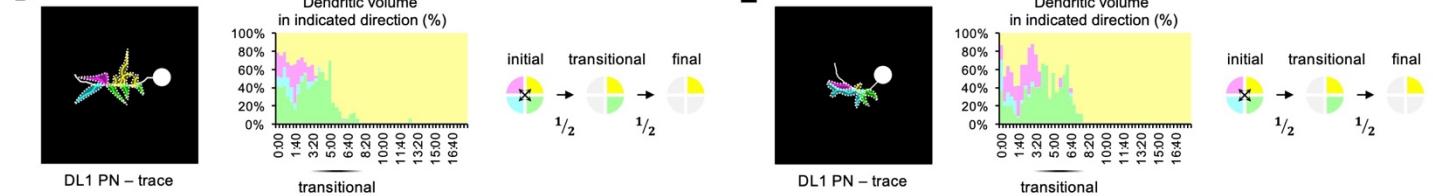
B



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E

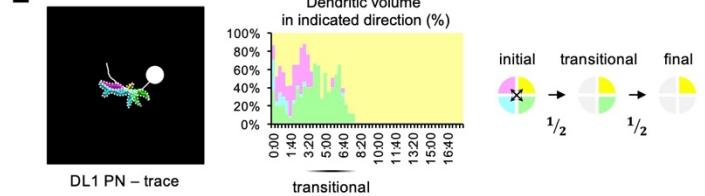


Figure 6 – figure supplement 1. Two-photon time-lapse imaging of DL1 PNs.

(A) Two-photon images of MARCM-labeled DL1 PNs (pseudo-colored in yellow) and other *GH146*+ cells (pseudo-colored in blue) in pupal brains dissected at 3h APF. Zoom-in time-lapse images of the dashed boxes are shown in **Figure 6A**, **Figure 6 – figure supplement 1B**, **C**. Background signals (in grey) are used to discern the orientation of DL1 PN in the brain.

(B, C) Two-photon time-lapse imaging of MARCM-labeled DL1 PNs (pseudo-colored in yellow) in additional pupal brains dissected at 3h APF and cultured for 18h *ex vivo*. After culture, explant was fixed and immunostained for N-Cadherin (Ncad; in blue) to outline the developing antennal lobe.

(D, E) Neurite tracing of DL1 PN (**Figure 6 – figure supplement 1B**, **C**) at the beginning of live imaging (3h APF + 0h *ex vivo*) and quantification of the percentage of dendritic volume in indicated direction during the time-lapse imaging period. During the transitional period, dendrites are only found in the dorsolateral and ventrolateral directions. Timestamp 00:00 refers to HH:mm; H, hour; m, minute.

Figure 6 – video 1. Two-photon time-lapse imaging of DL1 PN dendrites.

See **Figure 6A** for details. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

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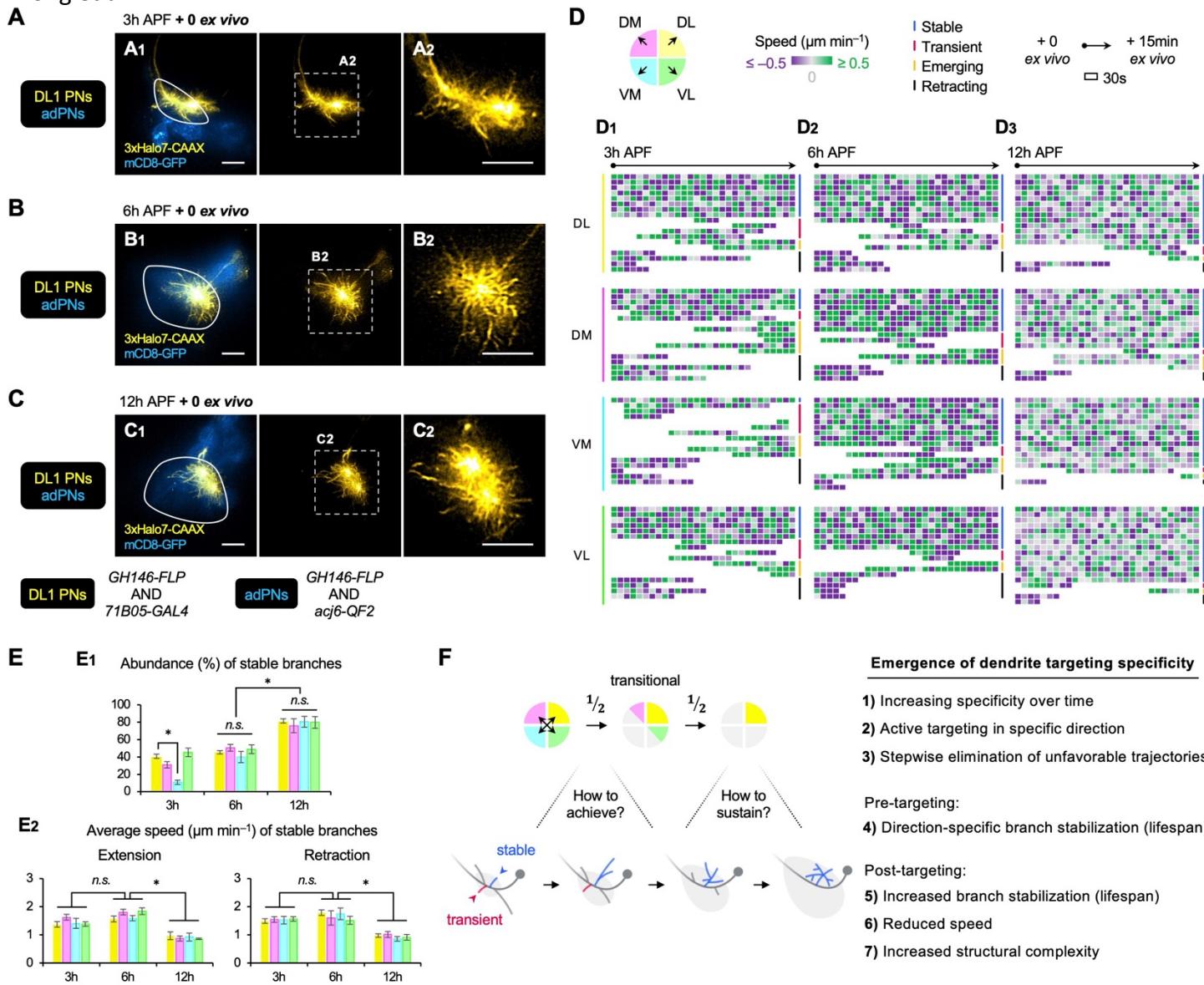


Figure 7. AO-LLSM time-lapse imaging reveals cellular mechanisms of dendrite targeting specificity.

(A–C) AO-LLSM imaging of DL1 PNs (*71B05*⁺; labeled in yellow) and adPNs (*acj6*⁺; labeled in blue) in cultured brains dissected at 3h (A), 6h (B), and 12h (C) APF. Zoom-in, single z-section images of A₁, B₁, and C₁ (outlined in dashed boxes) are shown in A₂, B₂, and C₂, respectively.

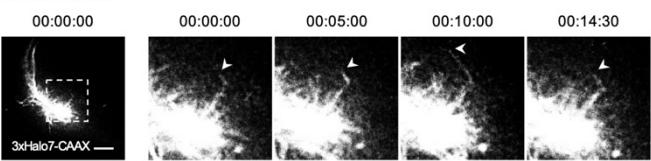
(D) Single dendritic branch dynamics of DL1 PN at 3h (D₁), 6h (D₂), and 12h (D₃) shown in **Figure 7A–C**. Terminal branches are analyzed and categorized based on the directions in which they extend. Their speeds are color-coded using dual purple-grey-green gradients (negative speeds, retraction; positive speeds, extension). Individual branches are also assigned into four categories: stable, transient, emerging, and retracting (color coded on the right; see **Figure 7 – figure supplement 1A**). Each block represents a 30s-window. Each row represents individual branch dynamics across the 15-min imaging period.

(E) Quantification of the abundance (in percentage) of DL1 PN stable branches in indicated direction at 3h, 6h, and 12h (E₁). Average speed of DL1 PN stable branches in indicated direction at 3h, 6h, and 12h (E₂). DL1 PN sample size: 3h = 4; 6h = 3; 12h = 3. Error bars, SEM; *t* test; One-way ANOVA; *, *p* < 0.05; n.s., *p* ≥ 0.05. SEM, standard error of the mean; n.s., not significant.

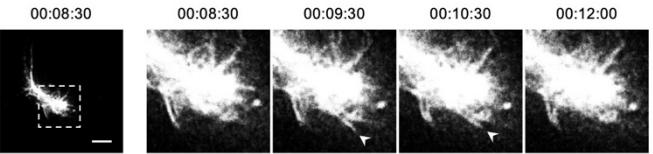
(F) Summary of mechanisms underlying the emergence of dendrite targeting specificity revealed by two-photon and AO-LLSM imaging of DL1 PN dendrites.

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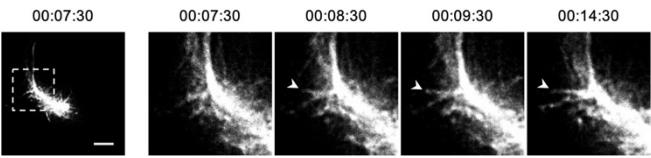
A 1. Stable branch



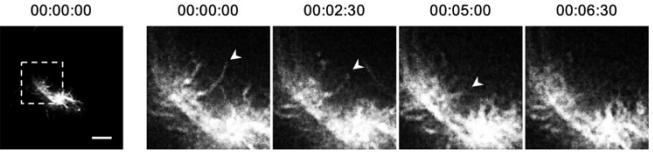
2. Transient branch



3. Emerging branch

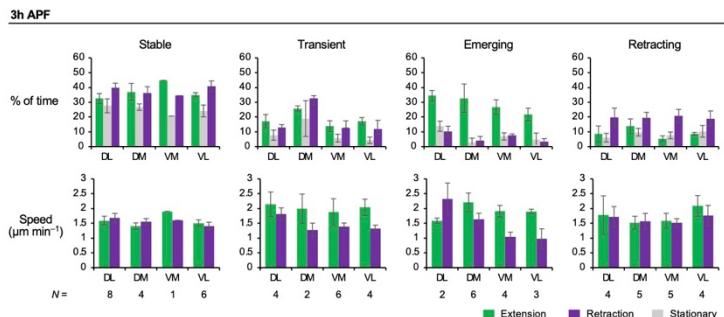


4. Retracting branch

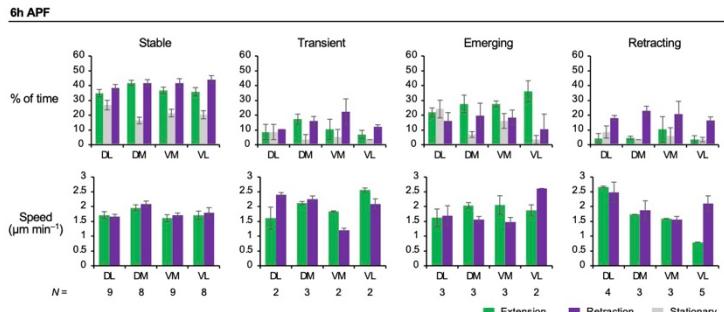


dorsal
↑ lateral

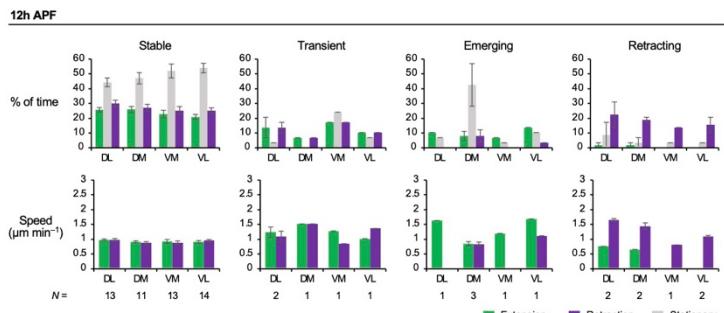
B



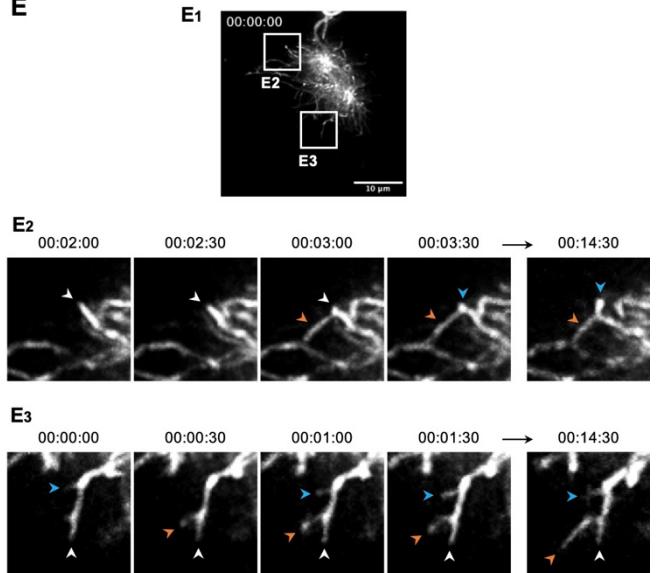
C



D



E



.7
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Figure 7 – figure supplement 1. Analyses of DL1 PN dendritic branches captured by AO-LLSM imaging.

(A) Categorization of branches into **(1)** stable, **(2)** transient, **(3)** emerging, and **(4)** retracting branches.

Representative branches of each category are shown. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

(B–D) **Top:** Quantification of the percentage of time given branches spent on extending, retracting, and being stationary. **Bottom:** Extension/retraction speeds of DL1 PN stable, transient, emerging, and retracting branches

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in indicated directions at 3h APF (**B**), 6h APF (**C**), and 12h APF (**D**). Data are analyzed using DL1 PNs in **Figure 7A–C**. *N* indicates branch number.

(E) Time-lapse AO-LLSM imaging of 12h APF DL1 PNs (**E**₁; **Figure 7C**) reveals terminal branch arborization. **E**₂ and **E**₃ are selected time-lapse images of zoom-in of indicated boxes in **E**₁. White arrowheads point to the terminal branch of interest, and colored arrowheads point to secondary branches produced from the branch of interest. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

Figure 7 – video 1. AO-LLSM time-lapse imaging of 3h DL1 PN dendrites.

See **Figure 7A** for details. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

Figure 7 – video 2. AO-LLSM time-lapse imaging of 6h DL1 PN dendrites.

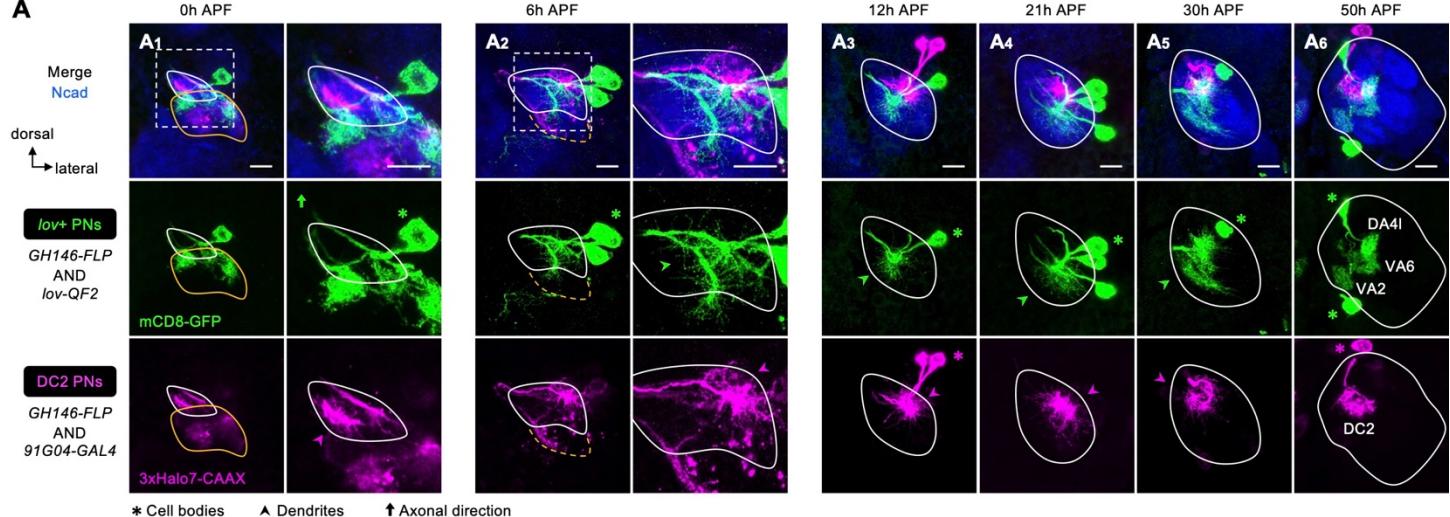
See **Figure 7B** for details. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

Figure 7 – video 3. AO-LLSM time-lapse imaging of 12h DL1 PN dendrites.

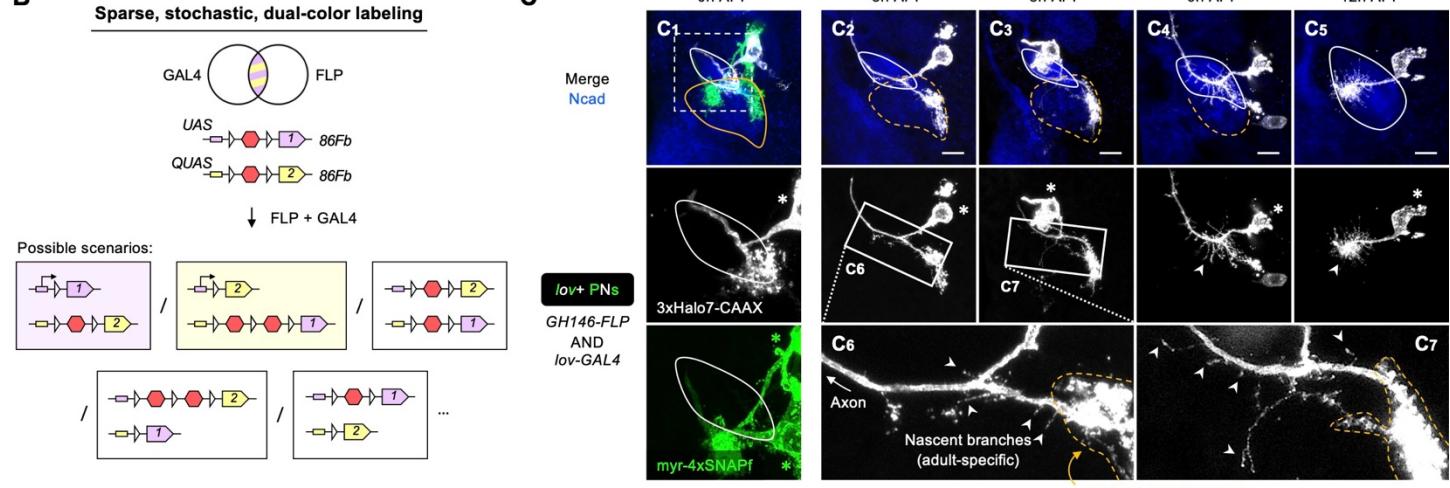
See **Figure 7C** for details. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

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A



B



D

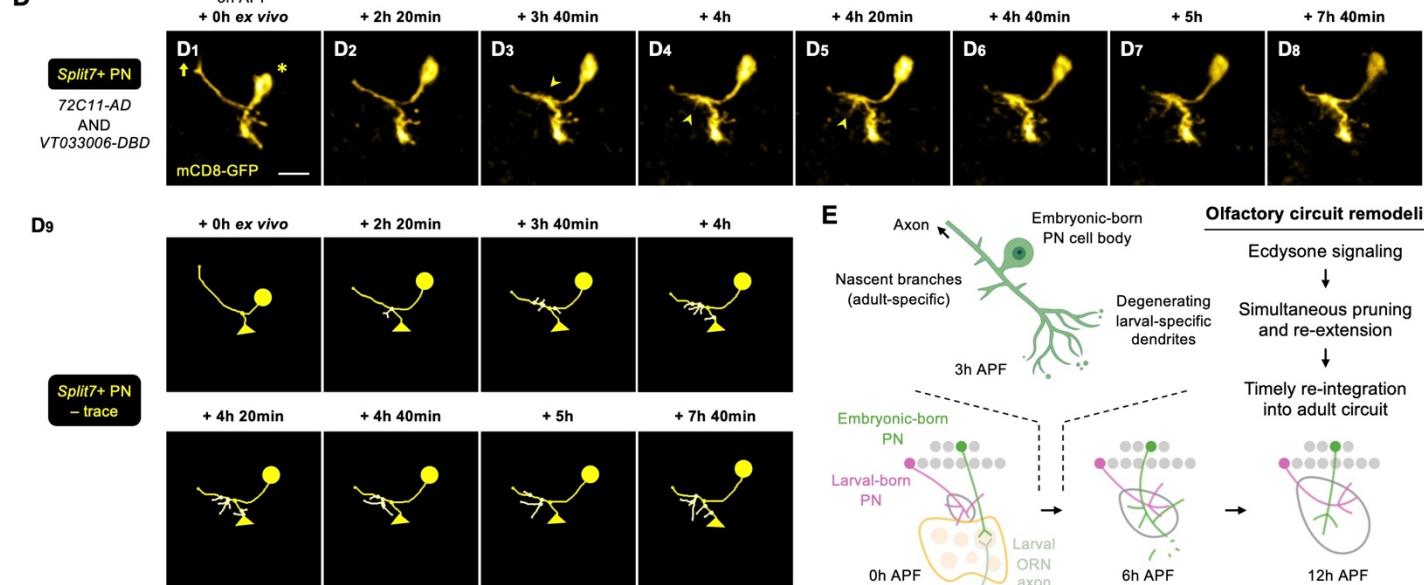
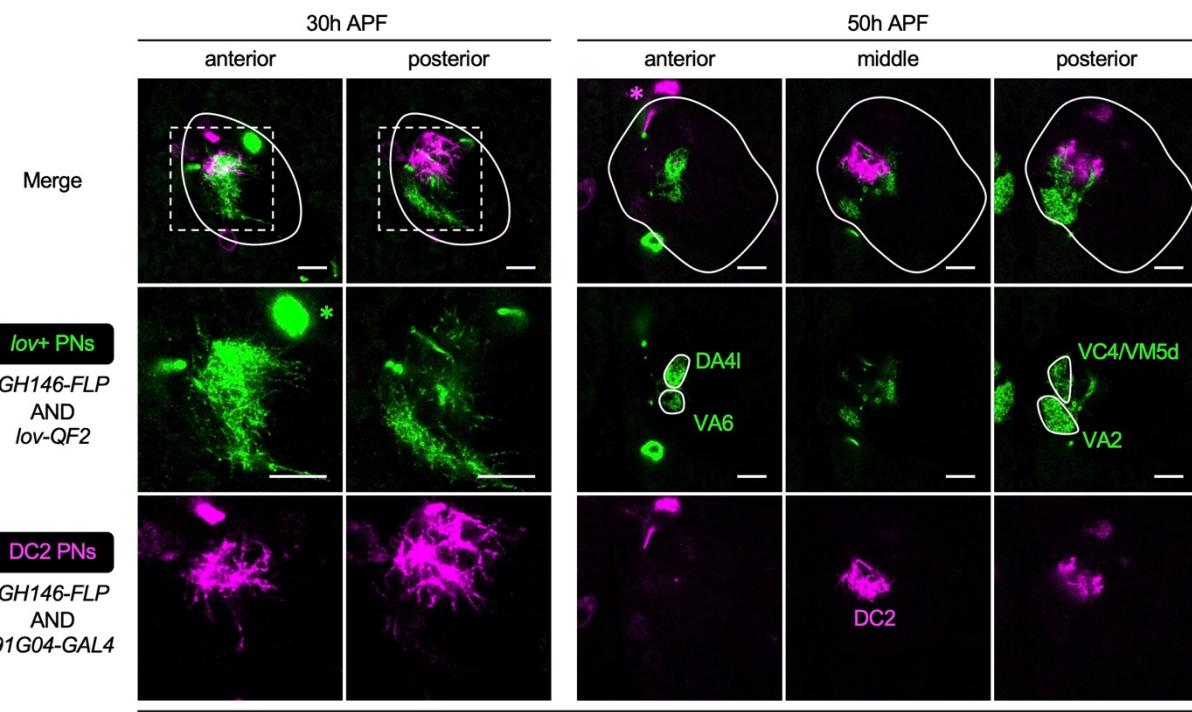
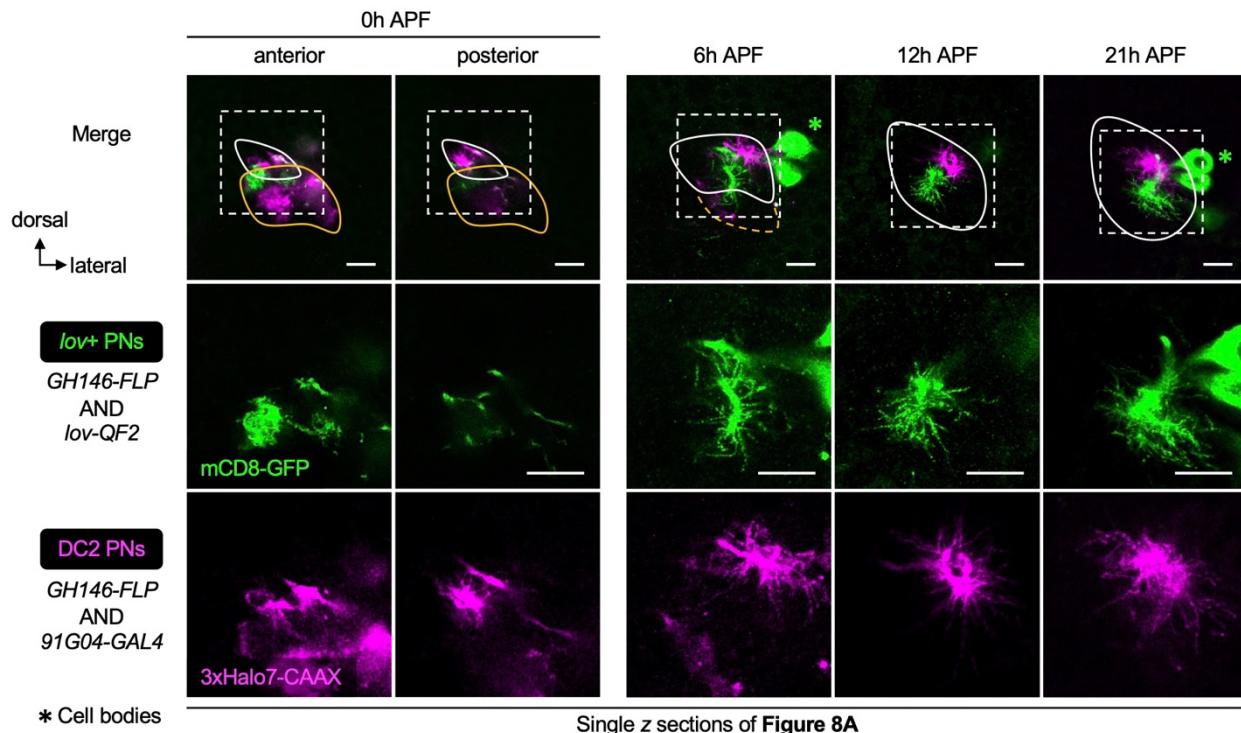


Figure 8. Embryonic-born PNs timely participate in olfactory map formation via simultaneous pruning and re-extension.

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'5 (A) Confocal images of fixed brains at indicated stages showing dendrite development of *lov*⁺ PNs (embryonic-
'6 born; labeled in green) and *91G04*⁺ DC2 PNs (larval-born; labeled in magenta). As *91G04-GAL4* also labels
'7 some embryonic-born PNs from 0–6h APF, their processes are found in the larval-specific antennal lobe (A_{1,2}).
'8 Right columns of A_{1,2} show zoom-in of the dashed boxes. Green arrowhead in A₂ indicates robust dendrite re-
'9 extension of embryonic-born PNs across the developing antennal lobe at 6h APF.
'0 (B) Schematic of the sparse, stochastic, and dual-color labeling strategy. Following *FLP* and *GAL4* co-
'1 expression, random recombination of *FRT* sites yields expression of either reporter 1 or 2, or no labeling at all.
'2 (C) Sparse labeling of *lov*⁺ PNs (labeled in green; single-cell *lov*⁺ PNs in grey) at indicated developmental
'3 stages. C₆ and C₇ are zoom-in images of the rectangular boxes in C₂ and C₃, respectively. Arrowheads indicate
'4 nascent, adult-specific dendrites. Larval-specific dendrites are outlined by dashed orange lines. Arrows indicate
'5 axons projecting towards high brain centers.
'6 (D) Two-photon time-lapse imaging of a single embryonic-born PN (*Split7*⁺; pseudo-colored in yellow) in a
'7 brain dissected at 3h APF and cultured for 23h *ex vivo*. Arrowhead in D₃ denote the thickening of the main
'8 process. Arrowheads in D_{4,5} denote dendritic protrusions dorsal to larval-specific dendrites. D₉ show neurite
'9 tracing of the embryonic-born PN. Triangles in D₉ indicate the degenerating larval-specific dendrites.
'0 (E) Schematic summary of remodeling of embryonic-born PN dendrites. Following simultaneous pruning and
'1 re-extension, embryonic-born PNs timely integrates into adult olfactory circuit and, together with larval-born
'2 PNs, participate in the prototypic map formation.
'3

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15
16 **Figure 8 – figure supplement 1. Dendrite development of *lov*⁺ embryonic-born PNs.**

17 Single z sections of Figure 8A showing the dendrite development of *lov*⁺ PNs (embryonic-born; labeled in
18 magenta) and 91G04+ DC2 PNs (larval-born; labeled in green).

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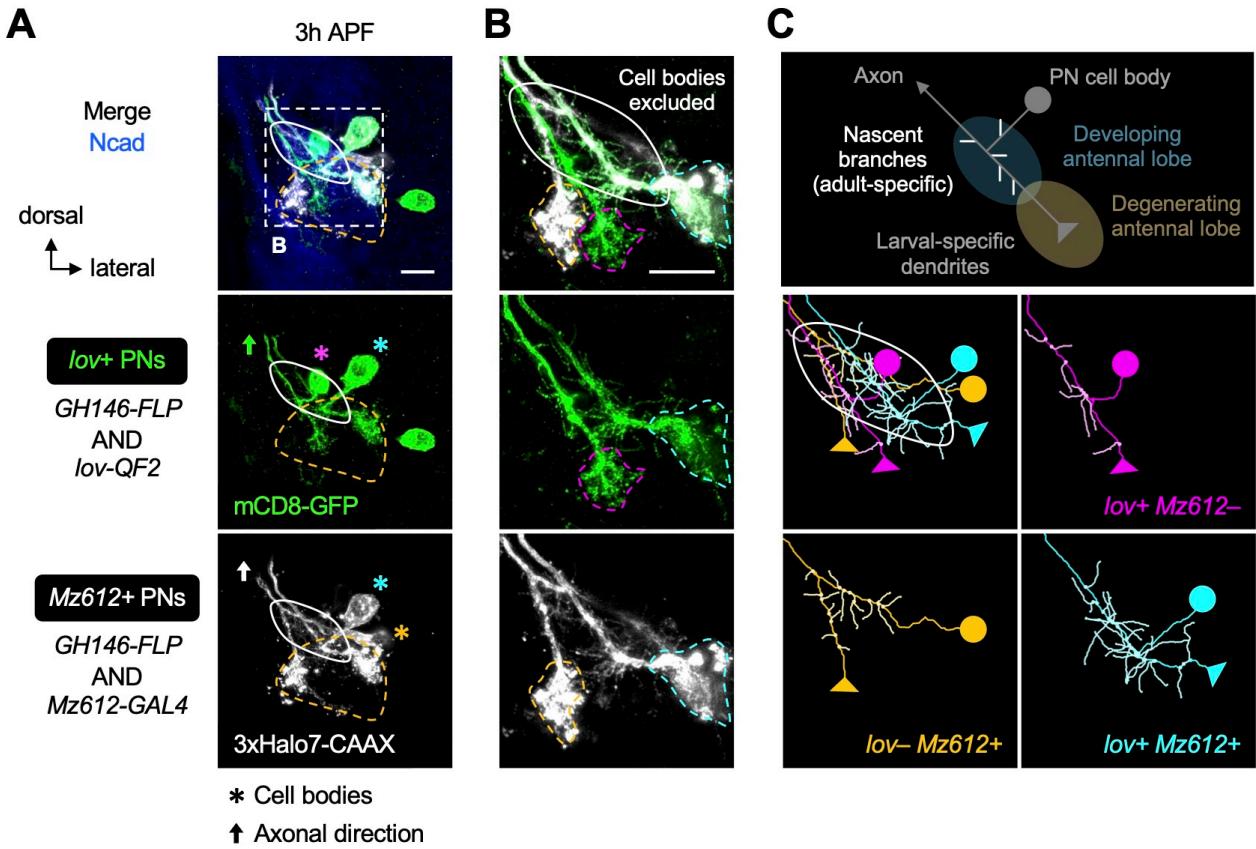


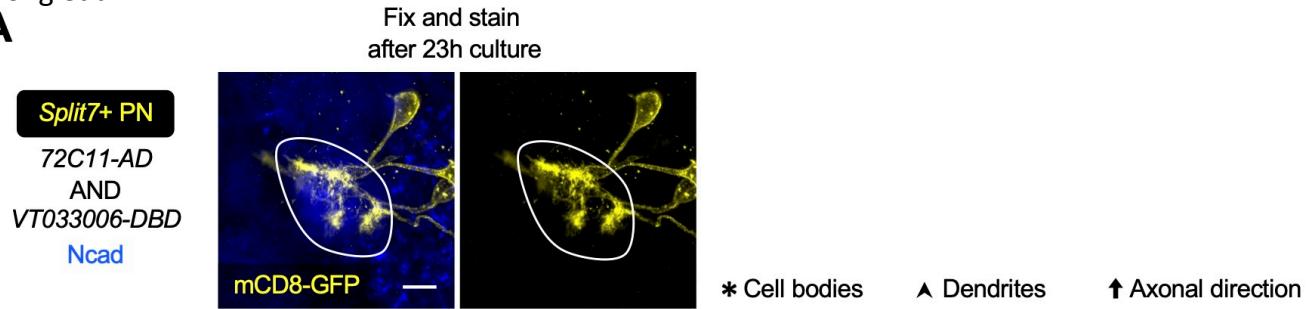
Figure 8 – figure supplement 2. Dendrite re-extension of *lov*+ and *Mz612*+ embryonic-born PNs.

(A, B) Dendritic re-extension of *lov*+ (labeled in green) and *Mz612*+ (labeled in grey) PNs. B is the zoom-in of dashed box in A. Magenta, cyan, and orange asterisks in A indicate cell bodies of *lov*+ *Mz612*– PN, *lov*+ *Mz612*+ PN, and *lov*– *Mz612*+ PN, respectively. Using the same color code, their larval-specific dendrites are outlined with dashed lines in B.

(C) Schematic of co-existence of larval- and adult-specific dendrites of an embryonic-born PN (top row), and neurite tracing of the three embryonic-born PNs (middle and bottom rows).

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A



B

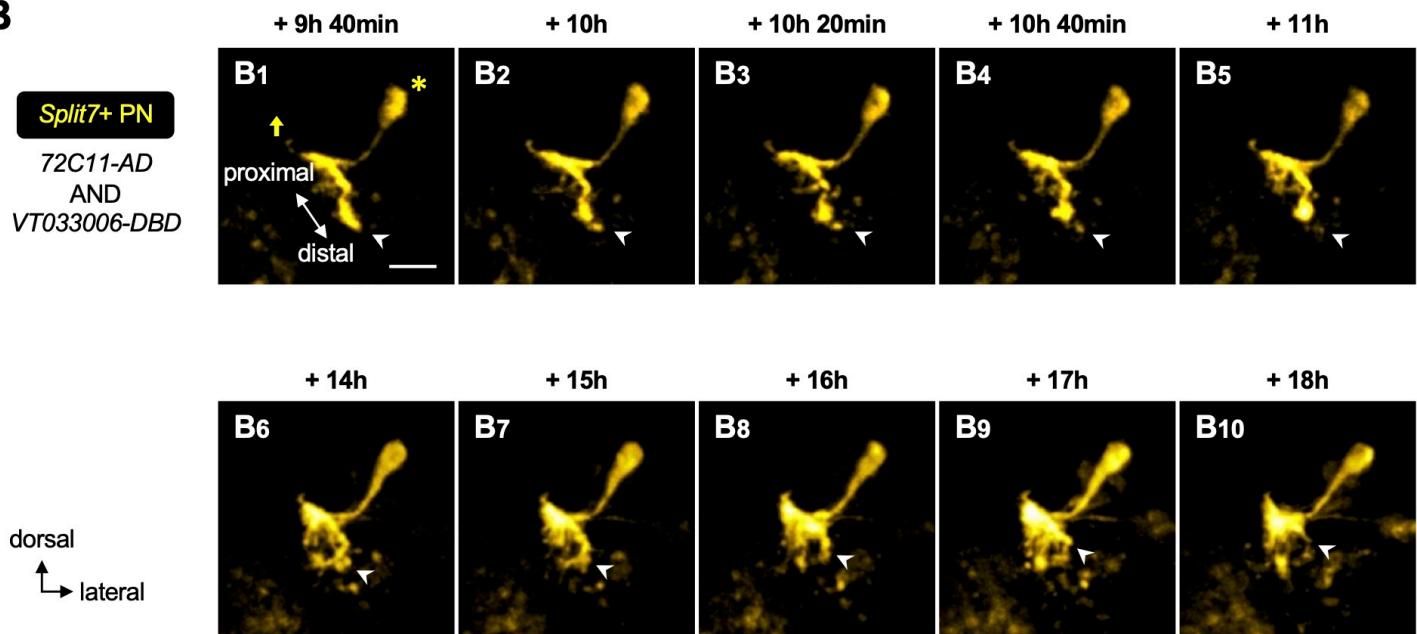


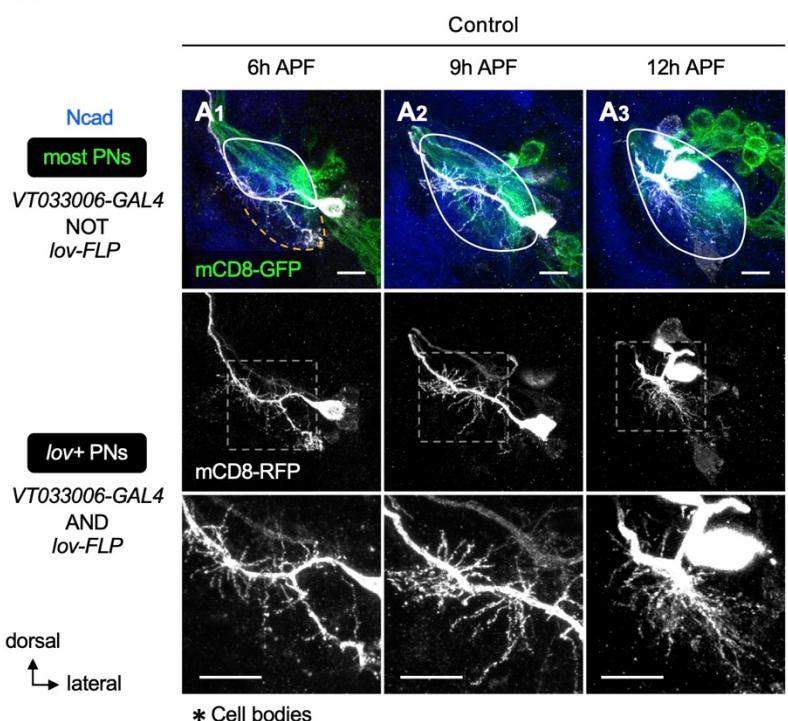
Figure 8 – figure supplement 3. Two-photon time-lapse imaging of *Split7*+ PN dendrites.

(A) Confocal image of *Split7*+ PNs (labeled in yellow; **Figure 8D**) after 23h culture. N-Cadherin (Ncad; in blue) staining outlines the developing antennal lobe. *Split7-GAL4* is expressed in more than one PN type at later stages.

(B) Two-photon time-lapse imaging of a *Split7*+ PN showing distal-to-proximal pruning of larval-specific dendrites.

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A



B

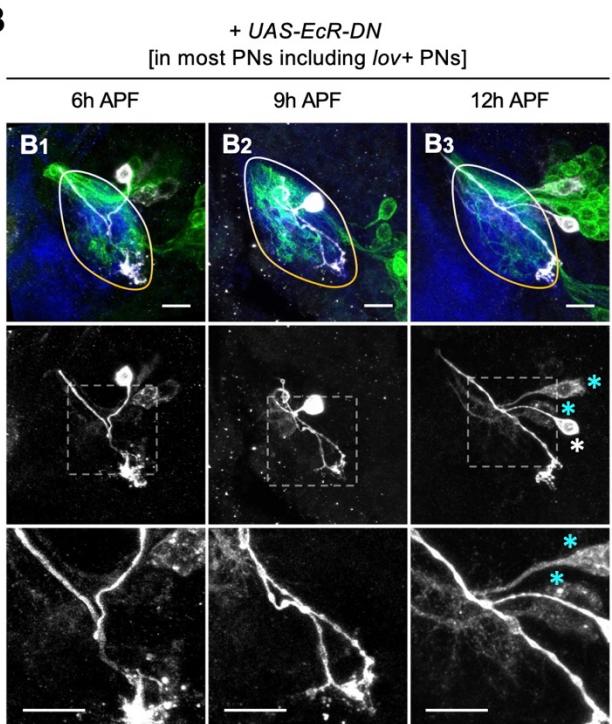


Figure 8 – figure supplement 4. Dual requirement of ecdysone signaling in pruning and re-extension of embryonic-born PN dendrites.

(A) Normal dendrite development seen in control *lov*+ embryonic-born PNs (*VT033006*+ *lov*); labeled in grey). Other PNs (referred to as most PNs; *VT033006*+ *lov*-) are labeled in green. Bottom row shows zoom-in of the dashed boxes.

(B) Expression of a dominant negative form of ecdysone receptor (*EcR-DN*) in most PNs including *lov*+ PNs suppresses both pruning and re-extension of *lov*+ PN dendrites. Similar results were seen in multiple biological samples ($N \geq 3$ for each stage per genotype). Cyan asterisks in B3 mark *lov*+ cells that had weak *VT033006-GAL4* and thereby weak *EcR-DN* expression. These cells appeared to still elaborate dendrites, suggestive of a dose-dependent effect of *EcR-DN*. White asterisk in B3 mark a *lov*+ cell with strong *VT033006-GAL4* expression. The presumed fused larval- and adult-specific antennal lobes are outlined with white-orange gradient line.

Figure 8 – video 1. Two-photon time-lapse imaging of *Split7*+ PN dendrites.

See Figure 8D for details. Timestamp 00:00:00 refers to HH:mm:ss; H, hour; m, minute; s, second.

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