

1 **A locomotor neural circuit persists and functions similarly in larvae and adult *Drosophila***

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12 **Abstract**

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14 Individual neurons can undergo drastic structural changes, known as neuronal remodeling or structural
15 plasticity. One example of this is in response to hormones, such as during puberty in mammals or
16 metamorphosis in insects. However, in each of these examples it remains unclear whether the remodeled
17 neuron resumes prior patterns of connectivity, and if so, whether the persistent circuits drive similar
18 behaviors. Here, we utilize a well-characterized neural circuit in the *Drosophila* larva: the Moonwalking
19 Descending Neuron (MDN) circuit. We previously showed that larval MDN induces backward crawling, and
20 synapses onto the Pair1 interneuron to inhibit forward crawling (Carreira-Rosario et al., 2018). MDN is
21 remodeled during metamorphosis and regulates backward walking in the adult fly. We investigated whether
22 Pair1 is remodeled during metamorphosis and functions within the MDN circuit during adulthood. We
23 assayed morphology and molecular markers to demonstrate that Pair1 is remodeled during metamorphosis
24 and persists in the adult fly. In the adult, optogenetic activation of Pair1 resulted in arrest of forward
25 locomotion, similar to what is observed in larvae. MDN and Pair1 are also synaptic partners in the adult,
26 showing that the MDN-Pair1 interneuron circuit is retained in the adult following hormone-driven pupal
27 remodeling. Thus, the MDN-Pair1 neurons are an interneuronal circuit – i.e. a pair of synaptically connected
28 interneurons – that persists through metamorphosis, taking on new input/output neurons, yet generating
29 similar locomotor behavior at both stages.
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31 Introduction

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33 Large-scale changes in neuronal morphology and function occur during mammalian puberty (Barendse et al.,
34 2018; Mills et al., 2016; Sisk and Zehr, 2005), as well as several neurobiological disorders including depression
35 (Patel et al., 2019), or chronic pain (Kuner and Flor, 2017). Similarly, major changes in neuronal numbers and
36 type occur as a result of insect metamorphosis (Kanamori et al., 2015; Truman and Reiss, 1976; Yaniv and
37 Schuldiner, 2016). Despite these changes, there are documented cases of individual insect neurons persisting
38 from larval to adult stages. In *Drosophila*, individual motor and sensory neurons have been shown to persist
39 throughout metamorphosis and undergo dramatic remodeling (Consoulas et al., 2002, 2000; Yaniv and
40 Schuldiner, 2016; Yu and Schuldiner, 2014). Similar findings have been reported for the insect mushroom
41 body, where Kenyon cells partners (projection neurons, DANs) exist at both larval and adult stages (Li et al.,
42 2020; Marin et al., 2005). Yet, it remains unclear whether the remodeled neurons re-establish connectivity
43 with the identical neurons in the larva and adult.

44 During *Drosophila* metamorphosis the animal changes from a crawling limbless larva to a walking six-
45 legged adult (Riddiford, 1980; Riddiford et al., 2003). Despite the obvious differences, some behaviors are
46 similar: both larvae and adults undergo forward locomotion in search of food, backward locomotion in
47 response to noxious stimuli, and pausing in between antagonistic behaviors (Carreira-Rosario et al., 2018). We
48 and others identified a neuron that, when activated, can trigger backward locomotion in both larvae and
49 adults (Bidaye et al., 2014; Carreira-Rosario et al., 2018; Sen et al., 2017), despite the obvious differences in
50 limbless and six-legged locomotion. This neuron, named Moonwalker/Mooncrawler Descending Neuron
51 (MDN) is present in two bilateral pairs per brain lobe, with all four MDNs having similar synaptic partners,
52 and all four MDNs capable of eliciting backward larval locomotion in larvae (Carreira-Rosario et al., 2018).
53 Larval MDNs induce backward locomotion via the coordinate arrest of forward locomotion followed by the
54 initiation of backward locomotion. Halting forward locomotion is done via activation of the Pair1 descending
55 interneuron, which inhibits the A27h premotor neuron, to prevent it from inducing forward locomotion
56 (Carreira-Rosario et al., 2018). Activating backward locomotion is likely to be due, in part, to MDN activation
57 of the A18b premotor neuron, which is specifically active during backward locomotion (Carreira-Rosario et
58 al., 2018). Thus, MDN-Pair1 are synaptically coupled members of a locomotor circuit in the *Drosophila* larva.

59 Here we follow our previous work showing that MDN is remodeled during metamorphosis and persists
60 into the adult (Carreira-Rosario et al., 2018) by asking: Is the MDN partner neuron Pair1 also maintained in
61 the adult? Does the adult Pair1 induce an inhibition in forward locomotion, similar to its role in larvae? And,
62 are the adult Pair1 and MDN synaptically coupled? We find that all of these questions are answered in the
63 affirmative, showing that the core MDN-Pair1 decision-making circuit (a pair of synaptically-connected
64 interneurons) persists from larva to adult, despite profound remodeling during metamorphosis, and that this
65 circuit coordinates forward/backward locomotion in both larvae and adults.

66 Results

67 The Pair1 neuron persists from larval to adult stages

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69 To determine if Pair1 neurons were present in the adult, we mapped expression of a Pair1-Gal4 line (*R75C02-
70 Gal4*) from early larval to adult stages. We identified the larval Pair1 neurons based on their characteristic cell
71 body position in the medial subesophageal zone (SEZ), dense local ipsilateral dendritic arborizations (defined
72 as dendritic based on enrichment for post-synapses in the TEM reconstruction of the larval Pair1 neuron;
73 [Figure 1 – supplement 1](#)), and contralateral axons descending into the ventral nerve cord (VNC) in an
74 extremely lateral axon tract (Carreira-Rosario et al., 2018). Using the Pair1-Gal4 line, we could identify Pair1
75

76 neurons with this morphology at 28h and 96h after larval hatching (ALH; [Figure 1A,B](#)). The Pair1 neuron cell
77 bodies and proximal neurites could still be observed at 24 hrs after pupal formation (APF), but virtually all of
78 the dendritic processes and descending axonal process had been pruned ([Figure 1C](#), only one neuron
79 labeled). This is expected, given that many or all neurons undergo axon/dendrite remodeling during
80 metamorphosis (Kanamori et al., 2015; Truman and Reiss, 1976; Yaniv and Schuldiner, 2016). At 48 hrs APF,
81 Pair1 neurons exhibited dendritic branching in the SEZ and a descending axon into the VNC, regaining
82 morphological features similar to that of larval Pair1 neurons ([Figure 1D](#)). The axon innervated the T1
83 (prothoracic) neuropil and descended further down the VNC. These morphological features were maintained
84 into the adult fly, where we could trace the Pair1 axon to primarily innervate the T1 neuropil ([Figure 1E](#)),
85 with less extensive innervation of the mesothoracic (T2) and metathoracic (T3) neuropils.

86 Although we can use the Pair1-Gal4 line to track neurons with Pair1 morphological features from
87 larva to adult, it remains possible that the Gal4 line switches off in Pair1 and switches on in a similar
88 descending neuron at a stage in between those we assayed. To conclusively demonstrate that the larval Pair1
89 neuron survives into adulthood, we used a genetic technique to permanently label or “immortalize” the larval
90 Pair1 neurons and assay for their presence in the adult brain. Briefly, the method achieves spatial specificity
91 by using Pair1-Gal4 to drive UAS-FLP which removes a stop cassette from nSyb-FRT-stop-FRT-LexA
92 resulting in permanent LexA expression in Pair1-Gal4 neurons; it achieves temporal specificity (e.g. labeling
93 only larval Pair1-Gal4+ neurons) by using a heat inducible KD recombinase to “open” the lexAop-
94 KDRTstopKDRT-HA reporter (see Methods for additional details). Thus, a heat shock will permanently
95 label all Pair1-Gal4+ neurons at the time of heat shock. We immortalized Pair1 neurons in the larva, and
96 assayed expression in the adult, and observed the two bilateral Pair1 neurons, based on characteristic medial
97 SEZ cell body position, local ipsilateral arbors and contralateral descending axons that preferentially innervate
98 the prothoracic neuropil ([Figure 1F](#)). Pair1 innervation is clearer in neurons immortalized during larval
99 stages, which reduces the off-target neuron expression in the adult VNC, and reveals an greatly enriched level
100 of innervation in the T1 neuropil ([Figure 1F](#)).

101 The Pair1-Gal4 line is expressed in several off-target neurons in addition to Pair1. One of these, a
102 sensory neuron from the proboscis, can be reduced from the adult Pair1 pattern by removing the proboscis a
103 day prior to analysis (see Methods) but is present at the 48 hr APF timepoint ([Figure 1D, E](#)). In addition,
104 there are off-target neurons that innervate all three thoracic neuropils (T1-T3), obscuring Pair1 innervation
105 ([Figure 1E](#)). We took advantage of the sparse labeling of the immortalization genetics and found brains that
106 maintained preferential targeting of Pair1 to the prothoracic neuropil but lacked T1-T3 off-target innervation,
107 confirming that they are indeed off-target neurons ([Figure 1F-F''](#)). We conclude that the Pair1 neurons are
108 present from larval to adult stages, and that the Pair1 neurons are enriched for postsynaptic partners in the T1
109 neuromere.

111 **Pair1 neurons maintain the same molecular profile from larval to adult stages**

112 If Pair1 neurons persist from larva to adult, they may express the same transcription factor (TF) profile at
113 both stages. We screened a small collection of TF markers for expression in the larval and adult Pair1
114 neurons, and in all cases we found identical expression ([Figure 2](#)). Larval and adult Pair1 neurons expressed
115 Hunchback (Hb), Sex combs reduced (Scr), and Bicoid (Bcd); but did not express Visual system homeobox
116 (Vsx1) or Nab ([Figure 2](#)). These results support the conclusion that Pair1 persists from larva to adult,
117 maintaining both molecular and morphological features, and raises the interesting possibility that the three
118 TFs (Hb, Bcd, Scr) may provide a molecular code that directs both larval and adult Pair1 morphology and/or
119 connectivity (see Discussion).

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121 **Pair1 activation arrests forward locomotion in adults**

122 We previously showed that larval MDN persists in adults and can induce backward locomotion at both stages
123 despite the obvious difference in motor output – limbless crawling vs. six-legged walking (Carreira-Rosario et
124 al., 2018). This raised the question of whether the adult Pair1 neuron also maintains its larval function, i.e. to
125 pause forward locomotion. To test this hypothesis, we used Pair1-Gal4 to express the red light-gated cation
126 channel CsChrimson (Chrimson) to activate Pair1 neurons in the adult. Experimental flies were fed all-*trans*
127 retinal (ATR; required for Chrimson function) whereas control flies were fed vehicle only.

128 Control flies exposed to red light did not pause or arrest forward locomotion, did not show an
129 increased probability of pausing, and did not have a decrease in distance traveled during the stimulus interval.
130 In contrast, experimental flies expressing Chrimson in Pair1 neurons showed a near complete arrest of
131 forward locomotion, an increased probability of pausing, and a reduced distance traveled during the stimulus
132 interval (Figure 3A-C; Figure 3 - Supplement 1). These effects were reversed after turning off the red light,
133 with the exception of a slightly reduced distance travelled, likely due to a lingering physiological effect of the
134 30 sec Pair1 activation (Figure 3A,C). Pair1 activation resulted in an increase in immobile flies (Figure 3E)
135 and a corresponding decrease in whole body translocation (defined as “large movements”, Figure 3F).
136 Importantly, Pair1 activation did not prevent small body part movements such as those involved in grooming
137 (defined as “small movements”, Figure 3G). Note that Pair1-Gal4 off-target expression is common but
138 variable from fly to fly, whereas its expression in Pair1 neurons is fully penetrant; because the Chrimson-
139 induced behavior is also fully penetrant, we conclude that the arrest in forward locomotion is due to
140 Chrimson activation of the Pair1 neurons. We conclude that Pair1 activation prevents a single behavior –
141 forward locomotion – but does not produce general paralysis or interfere with non-translocating limb
142 movements.

143 **MDN and Pair1 are synaptic partners during adulthood**

144 Given that MDN and Pair1 are synaptic partners in the larvae (Figure 1 - Supplement 1), MDN and Pair1
145 persist into adulthood (Figure 1 and 2), and MDN and Pair1 both regulate the same behavior in larvae and
146 adults (Figure 3) (Carreira-Rosario et al., 2018), we hypothesized that MDN and Pair1 may also be synaptic
147 partners during adulthood. To test this hypothesis, we used the MDN-LexA and Pair1-Gal4 to label MDN
148 and Pair1 neurons individually in the same animal (Figure 4A,B). We observed MDN and Pair1 neurites in
149 close proximity to each other (Figure 4C-E). To determine if MDN and Pair1 are synaptic partners in this
150 region of neuropil, we utilized t-GRASP (targeted GFP reconstitution across synaptic partners), an activity-
151 independent method to label synaptic contact sites (Shearin et al., 2018). Control flies only expressing pre-t-
152 GRASP in MDN did not have detectable t-GRASP signal (Figure 4F). However, flies expressing pre-t-
153 GRASP in MDN and post-t-GRASP in Pair1 had t-GRASP signal, indicating that MDN and Pair1 form
154 synapses (Figure 4G). We conclude that MDN and Pair1 are synaptic partners during adulthood.

155 **Discussion**

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159 Together with our earlier work (Carreira-Rosario et al., 2018), our results here show that a core decision-
160 making circuit is preserved from larval stages into the adult. This decision-making circuit contains MDN and
161 its monosynaptically-coupled Pair1 neuron, allowing the fly to switch between antagonistic behaviors:
162 forward versus backward locomotion. Our work raises several interesting questions: Do many other larval
163 neural circuits persist and have similar function in adults? Are the cues that establish MDN-Pair1 connectivity
164 in the larvae also used to re-establish MDN-Pair1 connectivity in the adult?

165 How much of the larval MDN-Pair1 circuit is maintained into the adult? The larval circuit contains
166 the MDN partners Pair1, ThDN, and A18b, and the Pair1 partner A27h (Figure 5). In addition to MDN, we
167 show here that Pair1 is maintained. There is no Gal4 line or markers for the ThDN neuron, and the only
168 A18b line has extensive off-target expression, so the fate of these two neurons is unknown. In contrast, the
169 A27h interneuron, which regulates forward crawling in the larvae, undergoes apoptosis during pupal stages
170 (data not shown) and thus does not regulate forward walking in the adult. This is not surprising as the A27h
171 neurons are located in the abdominal segments, which do not have a role in adult walking.

172 How much of the adult MDN-Pair1 circuit is present in the larvae? Recent work mapping the adult
173 MDN circuit has identified over 30 VNC neurons downstream of MDN, including the LBL40 and LUL130
174 neurons required for hindleg backward stepping (Figure 5; Feng et al., 2020). Recent work has also identified
175 adult neurons important for forward walking (Bidaye et al., 2020), but their relationship to adult MDN is
176 unknown. In the future, it will be interesting to see if any of these adult neurons are present in the larvae,
177 particularly those regulating forward and backward walking, and determine if they are also MDN or Pair1
178 target neurons.

179 Elegant recent work has shown that initiation of forward walking requires the forelegs, innervated by
180 motor neurons in the prothoracic segment, whereas initiation of backward walking requires the hindlegs,
181 innervated by motor neuron in the metathoracic segment (Feng et al., 2020). This is consistent with our
182 finding that the adult Pair1 neuron innervates the prothoracic neuropil, a site well-positioned to arrest motor
183 activity driving foreleg stepping and initiation of forward locomotion. Similarly, adult MDN synaptic partners
184 primarily innervate the metathoracic neuromere (Feng et al., 2020), a good location for inducing hindleg
185 stepping and initiation of backward walking. A similar spatial segregation is likely to occur in the larva, where
186 forward crawling is induced by A27h in posterior segments, and backward crawling is induced in anterior
187 segments (Fushiki et al., 2016; Tastekin et al., 2018).

188 In larvae, Pair1 activation only causes a pause in forward locomotion – after pausing, the animal
189 returns to baseline speed regardless of the red-light stimulus duration (Carreira-Rosario et al., 2018; Tastekin
190 et al., 2018). In contrast, Pair1 activation in adults results in a persistent arrest in forward locomotion for the
191 duration of the red light stimulus, although non-translocating limb movements are not affected. We speculate
192 these differences may be due to differences in Pair1 downstream synaptic partners, with more redundancy in
193 the larval circuit. Understanding the function of the neurons downstream of adult Pair1 in the T1 neuromere
194 is likely to provide insights into these behavioral differences.

195 Here we identify a transcription factor combination (Hb, Bcd, Scr) that persists in Pair1 neurons
196 from larvae to adults. It is intriguing to speculate that these transcription factors may be required for cell
197 surface molecule expression used to establish the MDN-Pair1 synaptic specificity in the embryo as well as to
198 re-establish MDN-Pair1 synaptic specificity following pupal remodeling. Perhaps these transcription factors
199 drive expression of the same cell surface molecules at both stages, or even continuously to maintain
200 functional connectivity.

201 Individual neurons that have similar functions in larva and adults have been identified, including
202 select motor neurons, sensory neurons (Consoulas et al., 2002, 2000; Levine, 1984; Truman, 1992; Weeks,
203 2003), and Kenyon cells of the mushroom body (Eichler et al., 2017; Li et al., 2020). However, it remains
204 unknown whether any of their synaptic partners also persist and retain the same pattern of connectivity. Our
205 work is the first, to our knowledge, to show that a pair of synaptically connected interneurons – the core of a
206 decision-making circuit – can persist from larva to adult and perform similar functions at both stages.
207 Remarkably, both MDN and Pair1 undergo dramatic pruning and regeneration events during metamorphosis,
208 only to re-form synapses with each other following neuronal remodeling. This suggests that synapse
209 specificity cues are maintained from the late embryo, where MDN-Pair1 connectivity is first established, into

210 pupal stages, where MDN-Pair1 connectivity is re-established. The importance of the MDN-Pair1 decision-
211 making circuit is highlighted by its persistence from embryo to adult, despite adapting to different sensory
212 input and motor output at each stage. Perhaps other descending or ascending interneurons will also persist
213 into adults, switching inputs and outputs as needed. Indeed, the idea that a core decision-making circuit that
214 is stable across developmental stages is supported by recent elegant TEM reconstruction of neural circuits at
215 different stages of *C. elegans* development (Witvliet et al., 2020). Here the authors conclude that "Across
216 maturation, the decision-making circuitry is maintained whereas sensory and motor pathways are substantially
217 remodeled." These results, together with ours, raise the possibility that preservation of decision-making
218 interneuron circuit motifs may be functional modules that can be used adaptively with different sensorimotor
219 inputs and outputs. The presence of this circuit motif in both flies and worms suggests that it may be an
220 ancient evolutionary mechanism for assembling sensorimotor circuits.

221

222 **Materials and Methods**

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224 Fly husbandry

225 All flies were reared in a 25°C room at 50% relative humidity with a 12 hr light/dark cycle. All comparisons
226 between groups were based on studies with flies grown, handled and tested together.

227

228 Fly Stocks

- 229 1) *R75C02-Gal4* (Pair1 line; BDSC #39886)
- 230 2) *UAS-myr::GFP* (BDSC #32198)
- 231 3) *UAS-CsChrimson::mVenus* (Vivek Jayaraman, Janelia Research Campus)
- 232 4) *VT044845-lexA* (adult MDN line; a gift from B. Dickson, Janelia Research Campus)
- 233 5) *UAS-mCD8::RFP, LexAop-mCD8::GFP;;* (BDSC #32229)
- 234 6) *LexAop-pre-t-GRASP, UAS-post-t-GRASP* (BDSC #79039)
- 235 7) *Hs-KD, 3xUAS-FLP; 13xLexAop(KDRT.Stop)myr:smGdP-Flag/ CyO-YFP ;*
236 *13xLexAop(KDRT.Stop)myr:smGdP-V5, 13xLexAop(KDRT.Stop)myr:smGdP-HA, nSyb-(FRT.Stop)-*
237 *LexA::p65/R75C02-Gal4* (line to permanently label Pair1; Doe Lab; modified from (Ren et al., 2016))

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239 Gal4 driver "immortalization"

240 Immortalization flies (see genotype #7, above) were allowed to lay eggs for four hours. Newly hatched larvae
241 were placed in a food vial, and at 96 hours ALH the food vial was partially submerged in a 37°C water bath
242 for 5 minutes, allowing the hs-KD to act as a recombinase to remove the KDRT Stop cassette, resulting in
243 nSyb-LexA driving HA expression permanently in the neurons expressing Pair1-Gal4 at the time of heat
244 shock (96h ALH). After the heat shock, larvae in the food vial recovered at 18°C for 5 minutes, and then
245 grown to adulthood at 25°C.

246

247 Immunostaining and imaging

248 Standard confocal microscopy and immunocytochemistry methods were performed as previously described
249 (Carreira-Rosario et al., 2018). Primary antibodies used recognize: GFP (rabbit, 1:500, ThermoFisher,
250 Waltham, MA; chicken, 1:1500, Abcam12970, Eugene, OR), HA (rat, 1:100, Sigma, St. Louis, MO),
251 Hunchback (mouse, 1:400, AbcamF18-1G10.2), Sex combs reduced (mouse, 1:10, Developmental Studies
252 Hybridoma Bank, Iowa City, IA), Bicoid (rat, 1:100, John Reintz, University of Chicago, Illinois), Vsx1
253 (guinea pig, 1:500, Claude Desplan, NYU, New York), Nab (guinea pig, 1:500, Stefan Thor, University of
254 Queensland, Brisbane, Australia) and t-GRASP signal (rabbit GFP G10362, 1:300, Invitrogen). Secondary
255 antibodies were from Jackson ImmunoResearch (Donkey, 1:400, West Grove, PA). Confocal image stacks
256 were acquired on a Zeiss 800 microscope. All images were processed in Fiji (<https://imagej.net/Fiji>) and
257 Adobe Illustrator (Adobe, San Jose, CA). Images were processed as described previously (Carreira-Rosario et
258 al., 2018). The primary neurites of Pair1 were traced using the Simple Neurite Tracer in Fiji.

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Adult Behavioral Experiment

Adult behavior was assayed using two arenas, a closed loop arena (Figure 3) and an open field arena (Figure 3 – Supplement 1). For the closed loop arena, adult female flies 1 day after eclosion were transferred to standard cornmeal fly food supplemented with 100 mL 0.5 mM ATR or 100% ethanol for 4 days (changed every 2 days). Animals, with intact wings, were starved for 4 hrs and then placed in arenas and their behavior was recorded as described previously (Carreira-Rosario et al., 2018). Flies were exposed to low transmitted light, red light, and low transmitted light again for 30 sec each. This was done three times for each animal. To calculate different parameters, the recorded videos were tracked and analyzed using the CalTech Fly Tracker (Fontaine et al., 2009) and JABA (Kabra et al., 2013). The speed, distance and behavior reported were specific to the first trial. The reported speeds are the average speed of each second. The pausing probability was calculated as previously described (Carreira-Rosario et al., 2018). “Pre” defines the 30 secs prior to red light exposure, “light” defines the 30 secs of red light exposure and “post” defines the 30 secs after red light exposure. Immobile movements were defined as the fly not translocating and not moving other body parts. Small movements were defined as the fly not translocating but moving body parts (i.e. grooming, moving wings). Large movements were defined as the fly translocating its body. All behavior measures were normalized by dividing them by the group average “pre” values.

For the open field arena, adult flies were fed ATR and vehicle as described above. 3 animals were placed in a circular arena with a diameter of 14.5 cm and height of 0.5 cm. After 5 min for environmental acclimation, animal behavior was recorded at 25 FPS using a Basler aCA2040-25gm GigE camera under infrared light for 4 sec followed by 4 secs under red light and another 4 sec under infrared light, as described previously (Risse et al., 2013). The was repeated 3 times, and tracked and analyzed as described above.

Statistics

All statistical analysis (t-test, one-way and two-way ANOVA with Bonferroni’s multiple comparison tests) were performed with Prism 9 (GraphPad software, San Diego, CA). Numerical data in graphs show individual measurements (animals), means (represented by red bars) or means \pm S.E.M. (dashed lined), when appropriate. The number of replicates (n) is indicated for each data set in the corresponding legend.

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Competing Interests

None.

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303 **Figure legends**

304

305 **Figure 1. The Pair1 neuron persists from larval to adult stages.**

306 (A-B) Pair1 neurons (cell body: yellow asterisk; neurites: yellow arrowhead) in the larval CNS (gray outline) at
307 28h ALH (A) and 96h ALH (B). Here and in subsequent panels are maximum intensity projections of
308 confocal sections containing the Pair1 neurons; anterior, up; dorsal view. Significant 'off-target' expression
309 marked with white arrowheads. Scale bar, 50 μm . (A'-B') Enlargement of the brain regions boxed in A,B.
310 Scale bar, 20 μm . (A''-B'') Tracing to show Pair1 neuron morphology. Genotype: +; *UAS-myr::GFP*; *R75C02-*
311 *Gal4*.

312 (C-D) Pair1 neurons (cell body: yellow asterisk; neurites: yellow arrowhead) in the pupal CNS (gray outline) at
313 24h APF (C) and 96h APF (D). Significant 'off-target' expression marked with white arrowheads. Scale bar,
314 50 μm . (C'-D') Enlargement of the brain regions boxed in C, D; cell body: yellow asterisk, neurites: yellow
315 arrowhead. Scale bar, 10 μm . (C'') Tracing to show Pair1 neuron morphology. (D'') Focal plane showing
316 Pair1 cell bodies (region boxed in D', cell body marked with yellow asterisks). Scale bar, 10 μm . (D''') Tracing
317 to show Pair1 neuron morphology. Note that Pair1 can be followed to T1 in the 3D confocal stack but is
318 difficult to represent here due to fasciculation of Pair1 with off-target neurons. Genotype: +; *UAS-myr::GFP*;
319 *R75C02-Gal4*.

320 (E) Pair1 neurons (cell body: yellow asterisk; neurites: yellow arrowhead) in the 4d adult CNS (gray outline)
321 Significant 'off-target' expression marked with white arrowheads. Scale bar, 50 μm . (E') Enlargement of the
322 brain region boxed in E. Scale bar, 10 μm . (E'') Focal plane showing Pair1 cell bodies (region boxed in E',
323 cell body marked with yellow asterisks). Scale bar, 10 μm . (E''') Tracing to show Pair1 neuron morphology.
324 Genotype: +; *UAS-myr::GFP*; *R75C02-Gal4*.

325 (F) Pair1 neurons (cell body: yellow asterisk; neurites: yellow arrowhead) permanently labeled at 96h ALH
326 and visualized in the 4d old adult. See methods for details. Significant 'off-target' expression marked with
327 white arrowheads. Scale bar, 50 μm . (F') Enlargement of the brain region boxed in F; Pair1 cell body: yellow
328 asterisk; Pair1 neurites: yellow arrowhead. Scale bar, 10 μm . (F'') Focal plane showing Pair1 cell bodies
329 (region boxed in F', cell body marked with yellow asterisks). Scale bar, 10 μm . (F''') Tracing to show Pair1
330 neuron morphology. Genotype: *Hs-KD,3xUAS-FLP*; *13xLexAop(KDRT.Stop)myr::smGdP-Flag* +;
331 *13xLexAop(KDRT.Stop)myr::smGdP-V5*, *13xLexAop(KDRT.Stop)myr::smGdP-HA*, *nSyb(FRT.Stop)LexA::p65*
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333 **Figure 1 – Supplement 1. MDN axon and Pair1 dendrite target the same neuropil in the larval brain.**

334 The TEM volume of the newly hatched larva "Seymore" showing the axon (green) and dendrite (blue)
335 domains of a single MDN and Pair1 neuron defined by the location of pre- and postsynapses. Left: MDN
336 axon and dendrite domains. Middle: Pair1 axon and dendrite domains. Right: MDN axon and Pair1 dendrite
337 are closely entwined the same region of neuropil (white bracket).
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339 **Figure 2. The Pair1 neuron expresses the same molecular markers at larval and adult stages.**

340 (A) Schematic of the larval brain showing region of Pair1 neurons (red box) enlarged in panels below.
341 Anterior up, dorsal view.

342 (B-G) Larval Pair1 neurons (left column), indicated markers (middle column), and merge (right column) at
343 28h ALH. In some cases the second Pair1 neuron is out of the focal plane, but both Pair1 neurons always
344 have the same gene expression profile. Markers detect the following transcription factors: Hb, Hunchback;
345 Scr, Sex combs reduced; Bcd, Bicoid; Vsx1, Visual system homeobox 1; and Nab. Scale bar, 5 μm . (G)
346 Summary: marker expression matches that in adults. Genotype: +; *UAS-myr::GFP*; *R75C02-Gal4*.

347 (H) Schematic of the adult brain showing region of Pair1 neurons (red box) enlarged in panels below.
348 Anterior up, dorsal view.

349 (I-N) Adult Pair1 neurons (left column), indicated markers (middle column), and merge (right column) in 4d
350 old adult. Scale bar, 5 μ m. (N) Summary: marker expression matches that in larvae. Genotype: +; *UAS-*
351 *myr::GFP*; *R75C02-Gal4*.

352

353 **Figure 3. Pair1 activation for 30s arrests forward locomotion but does not cause paralysis in adults**

354 (A) Speed (mm/sec) of adult flies expressing Chrimson in Pair1 neurons following neuronal activation
355 (+ATR, blue) or no activation (vehicle control, black) in a closed loop arena. Speed was recorded for the 30s
356 prior to activation, the 30s light-induced activation (red stipple), and 30s after activation. Mean \pm S.E.M, n =
357 10. Genotype for this and all subsequent panels: *UAS-CsChrimson::mVenus*; +; *R75C02-Gal4*.

358 (B) Probability of forward locomotion pausing upon light-induced Pair1 activation (ATR treatment, blue)
359 compared to vehicle control (black). Statistics: t-test, $p < 0.001$; n = 10.

360 (C) Total distance traveled pre-light stimulus (“pre”), during the light stimulus (“light”) and post-light
361 stimulus (“post”) (terminology used here and in subsequent panels) of flies fed ATR (Pair1 activation, blue)
362 compared to controls (fed vehicle, no Pair1 activation, black). Statistics: two-way ANOVA: drug treatment,
363 $F(1, 18) = 111.3$, $p < 0.0001$; time, $F(1.867, 33.61) = 47.03$, $p < 0.0001$; interaction $F(2, 26) = 38.24$, $p <$
364 0.001 ; Bonferroni’s multiple comparisons between drug treatments within each timepoint: pre, $p > 0.9999$;
365 light, $p < 0.0001$; post, $p = 0.0001$; n = 10.

366 (D) Percent time doing large movements (whole body translocation, light grey), small movements (body part
367 movement but no translocation, dark grey) or no movements (immobile, black) of flies fed vehicle (left side)
368 or ATR (right side) during each time phase (pre, light, post).

369 (E) Normalized duration of time spent immobile during each timepoint (pre, light, post) for flies fed ATR
370 (Pair1 activation, blue) compared to controls fed vehicle (black). Statistics: two-way ANOVA: drug treatment,
371 $F(1, 18) = 112.8$, $p < 0.0001$; time, $F(1.930, 34.74) = 25.55$, $p < 0.0001$; interaction, $F(2, 36) = 27.81$, $p <$
372 0.0001 ; Bonferroni’s multiple comparisons between drug treatments within each timepoint: pre, $p > 0.9999$;
373 light, $p < 0.0001$; post, $p = 0.0022$; n = 10.

374 (F) Normalized duration of time spent doing small movements during each timepoint (pre, light, post) for
375 flies fed ATR (Pair1 activation, blue) compared to controls fed vehicle (black). Statistics: two-way ANOVA:
376 drug treatment, $F(1, 18) = 5.111$, $p = 0.036$; time, $F(1.923, 34.62) = 10.82$, $p = 0.0003$; interaction, $F(2, 36) =$
377 4.225 , $p = 0.0225$; Bonferroni’s multiple comparisons between drug treatments within each timepoint: pre, p
378 > 0.9999 ; light, $p = 0.0022$; post, $p > 0.9999$; n = 10.

379 (G) Normalized duration of time spent doing large movements during each time phase (pre, light, post) for
380 flies fed ATR (Pair1 activation, blue) compared to controls fed vehicle (black). Statistics: two-way ANOVA:
381 drug treatment, $F(1, 18) = 53.56$, $p < 0.0001$; time, $F(1.869, 33.64) = 53.44$, $p < 0.0001$; interaction, $F(2, 36)$
382 $= 52.20$, $p < 0.0001$; Bonferroni’s multiple comparisons between drug treatments within each timepoint: pre,
383 $p > 0.9999$; light, $p < 0.0001$; post, $p = 0.0074$; n = 10.

384

385 **Figure 3 – Supplement 1. Pair1 activation for 4s arrests forward locomotion but does not cause**
386 **paralysis in adults in an open field arena**

387 (A) Speed (mm/sec) of adult flies in an open field arena. Flies were fed food supplemented with ATR (blue)
388 or ethanol (vehicle, black). Red square represents the presentation of the light stimulus.

389 (B) Probability of pausing upon light activation of Pair1 (ATR treatment, blue) compared to controls (vehicle
390 treatment, black) (t-test, $p < 0.001$; n = 12).

391

392 **Figure 4. MDN and Pair1 are synaptic partners in adults**

393 (A, B) MDN neurons (A) and Pair1 neurons (B) in the adult central brain. Neurons are in white, nc82
394 counterstain in magenta for whole brain orientation; cell bodies marked by yellow arrowheads. Here and in
395 subsequent panels shows maximum intensity projection of volume; anterior, up; dorsal view. Scale bar, 50
396 μm . Genotype: *UAS-mCD8::RFP, LexAop-mCD8::GFP; VT044845-LexA; R75C02-Gal4*
397 (C-E) MDN neurites (C), Pair1 neurites (D), and merge (E) in the subesophageal ganglion (red box in
398 schematic). Scale bar, 10 μm . Genotype: *UAS-mCD8::RFP, LexAop-mCD8::GFP; VT044845-LexA; R75C02-*
399 *Gal4*.
400 (F) No detectable t-GRASP signal was observed in the subesophageal ganglion without expression of the pre-
401 t-GRASP fragment in MDN. Scale bar, 10 μm . Genotype: ; *LexAop-pre-t-GRASP, UAS-post-t-*
402 *GRASP/R75C02-Gal4*.
403 (G) t-GRASP signals between MDN and Pair1 were observed in the subesophageal ganglion. Scale bar, 10
404 μm . Genotype: ; *VT044845-LexA; LexAop-pre-t-GRASP, UAS-post-t-GRASP/R75C02-Gal4*.

405
406 **Figure 5. Model describing the MDN-Pair1 circuit in larval and adult stages**

407 In the larva, activation of MDN by upstream sensory neurons activates backward crawling and inhibits
408 forward crawling. MDN neurons synapse onto A18b and Pair1 interneurons. A18b subsequently regulates
409 backward crawling by synapsing onto motor neurons. Pair1 synapses onto and inhibits the pre-motor neuron
410 A27h, which generates forward locomotion when activated. In the adult, activation of MDN by upstream
411 sensory neurons activated backward walking and inhibits forward walking. MDN neurons synapse onto
412 LBL40, LUL120 and Pair1. Activation of LBL40 and LUL130 generates backward walking. Activation of
413 Pair1 generates a pausing behavior, likely through the inhibition of neurons generating forward locomotion.
414 In both larvae and adult, MDN and Pair1 neurons (blue) persist and function as a core circuit to regulate
415 locomotion.

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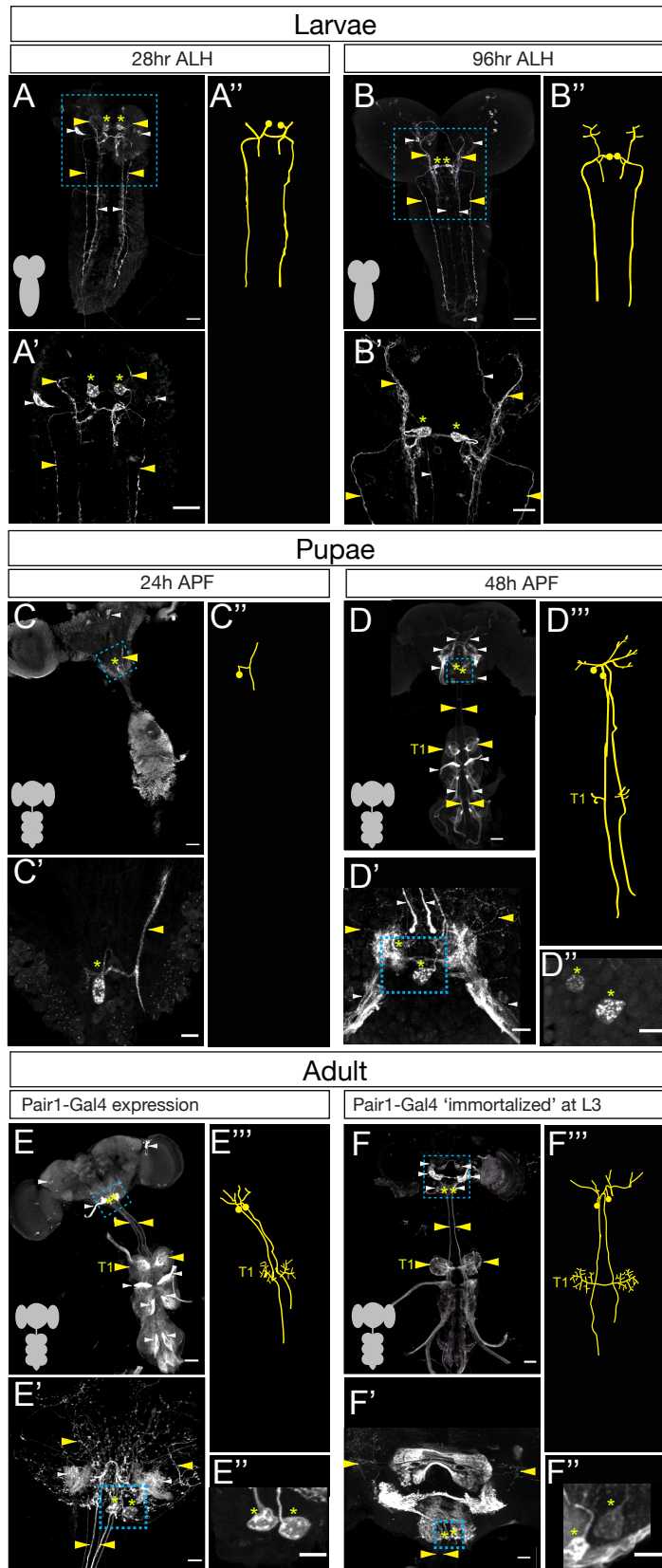


Figure 1

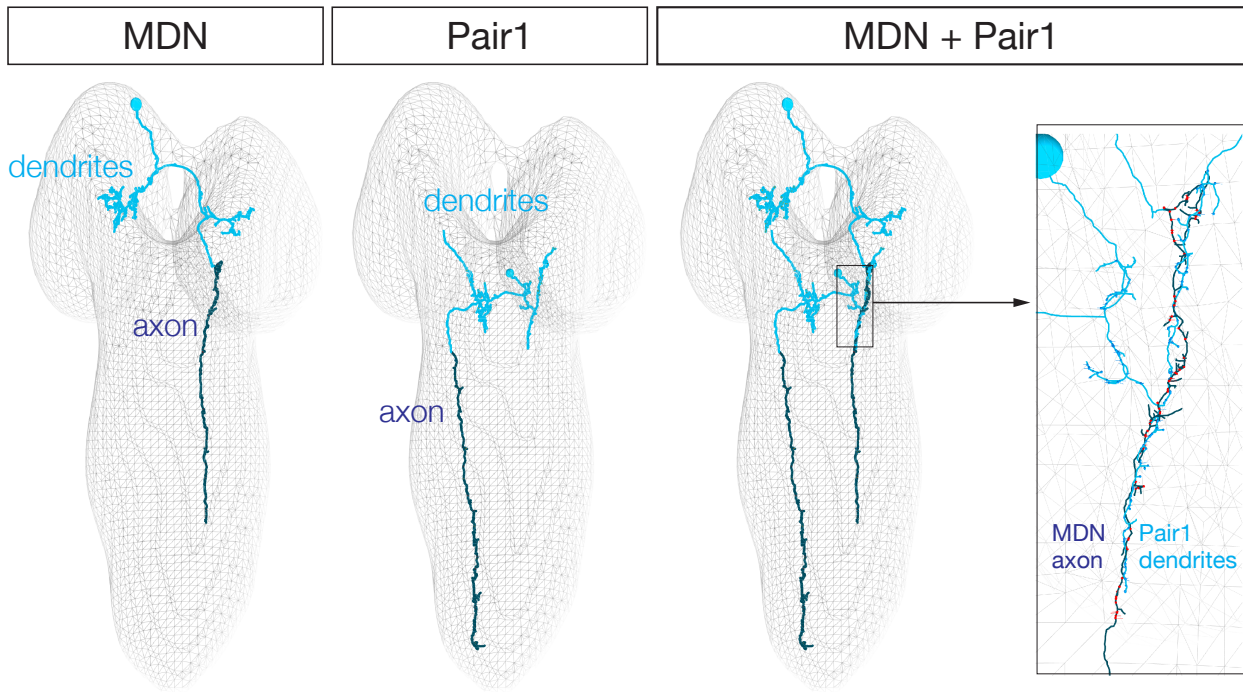


Figure 1 - supplement 1

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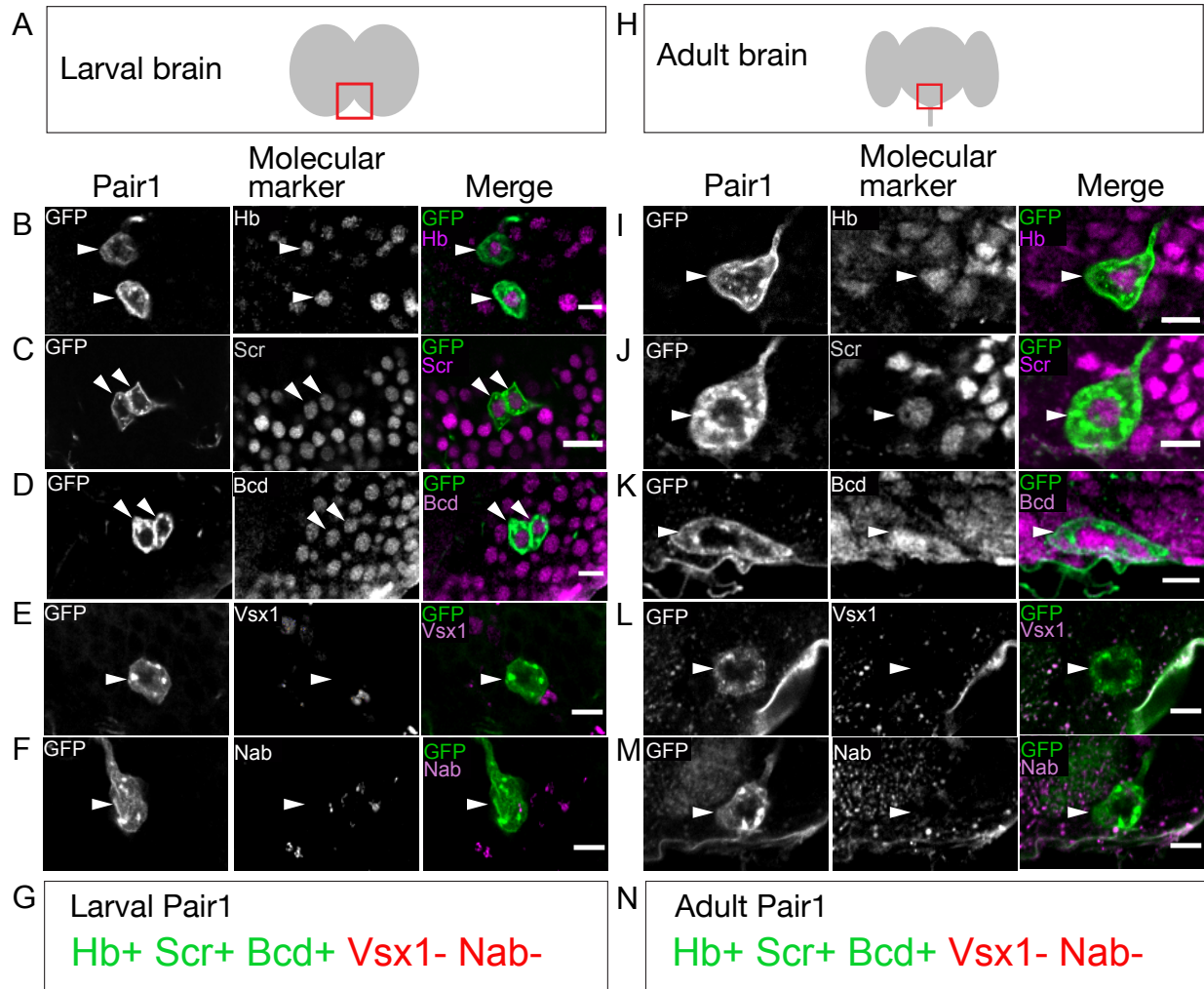


Figure 2

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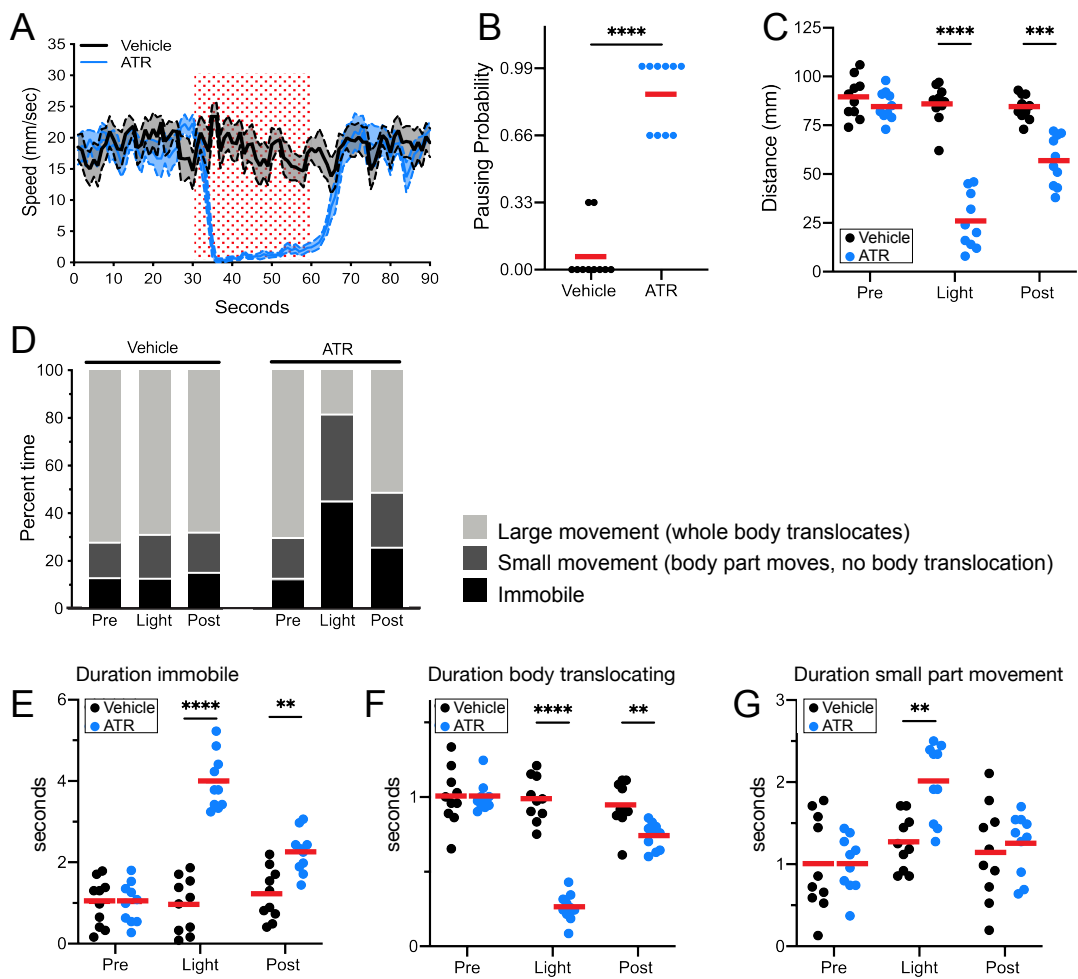


Figure 3

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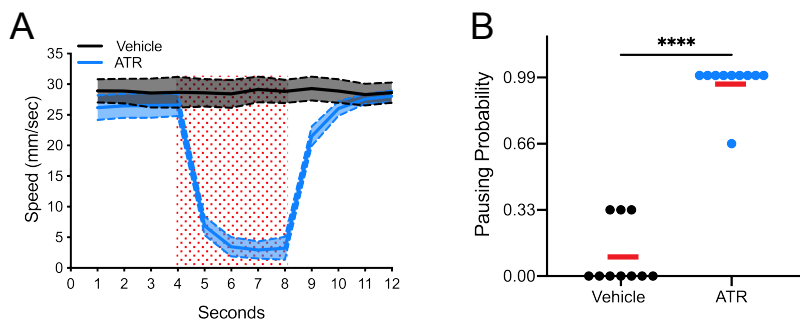


Figure 3 - Supplement 1

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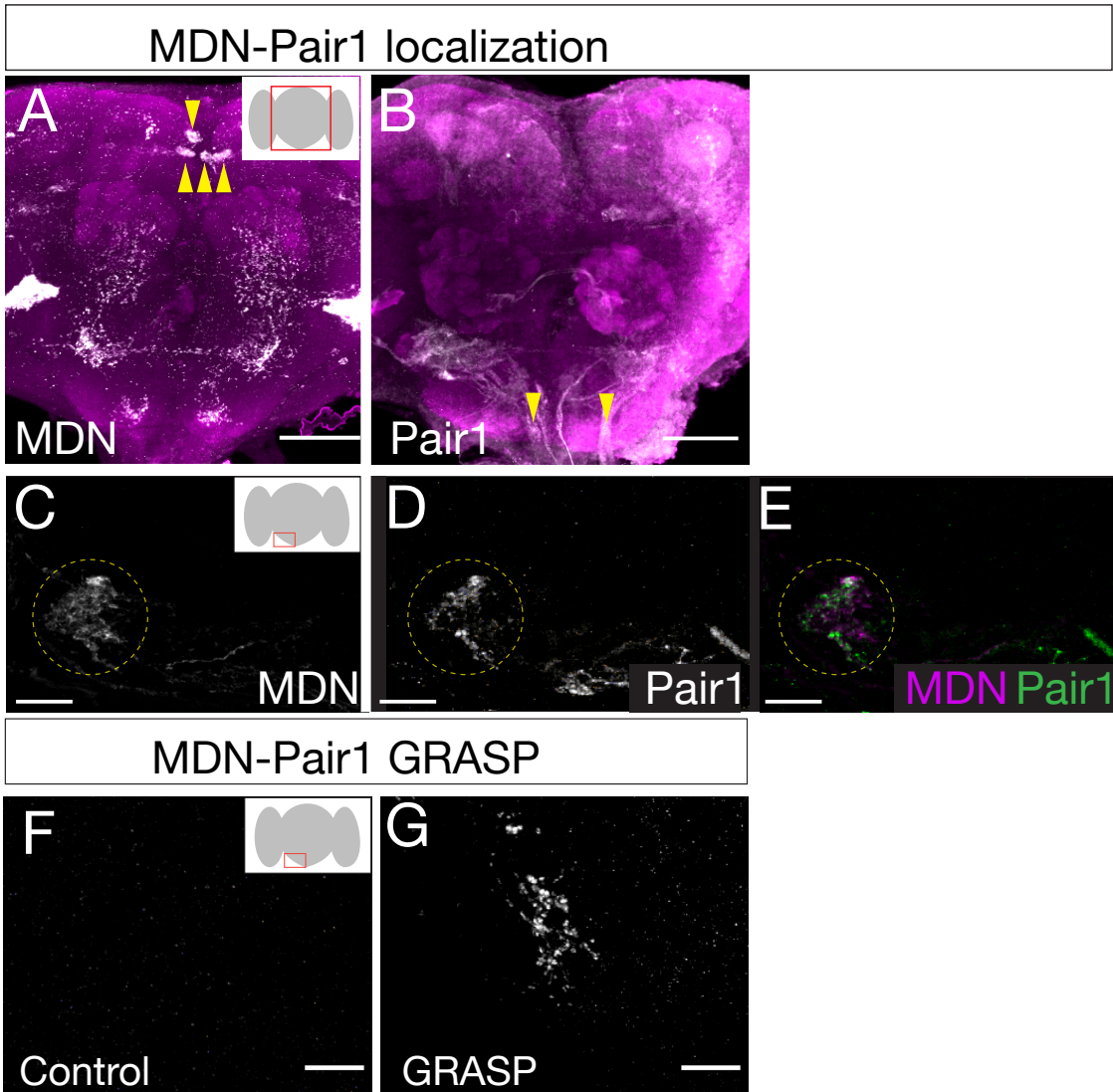


Figure 4

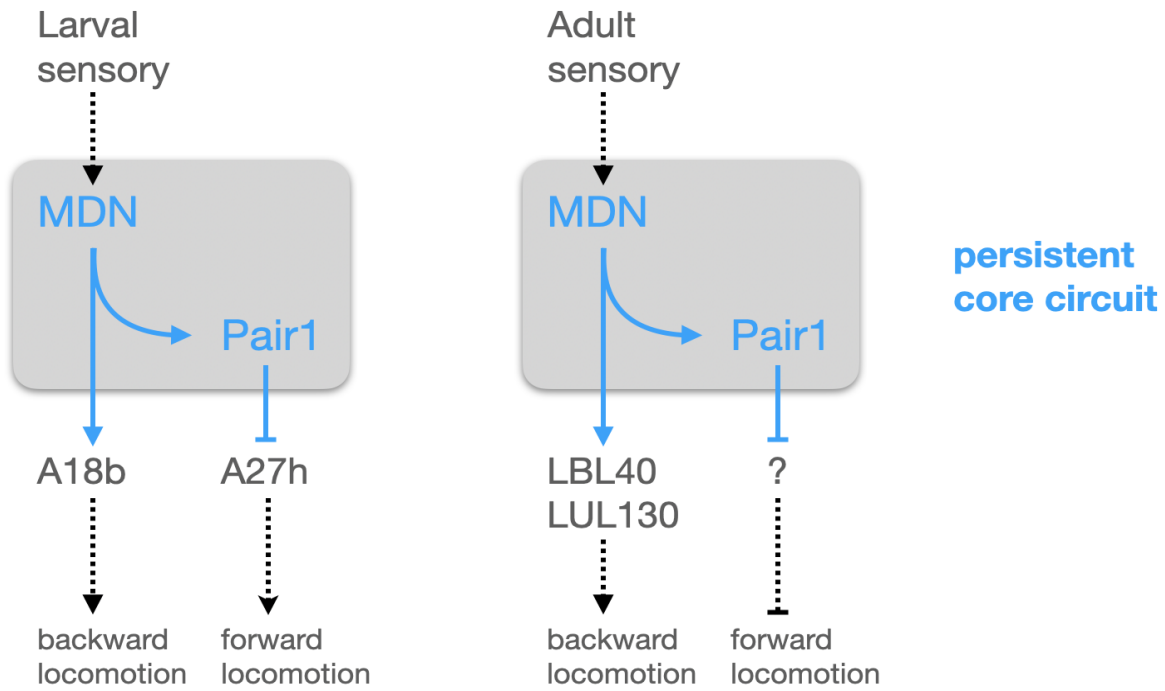


Figure 5

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