Four Individually Identified Paired Dopamine Neurons Signal Taste Punishment in Larval *Drosophila*

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25   **Abstract**

Dopaminergic neurons (DANs) carry out multiple tasks in the brain, including the transmission of information related to rewards and punishments across various animal species. They are responsible for evaluating sensory input, storing resulting associations as memory, and continuously updating them based on their relevance and reliability. Accurate

30   comprehension of the dopaminergic system's operation necessitates an understanding of the specific functions mediated by individual DANs. To this end, our research employs *Drosophila* larvae, which possess approximately 12,000 neurons in their brains, of which only around 1% (approximately 120) are DANs.

The presynaptic projections to the mushroom body (MB) - a brain region pivotal for

35   associative olfactory learning in insects - are limited to only eight larval dopaminergic neurons. These DANs are further subdivided into two clusters: the primary protocerebral anterior medial cluster (pPAM) comprises four cells, and the dorsolateral 1 cluster (DL1) comprises the remaining four cells. Our findings confirm previous research that demonstrates that the pPAM DANs innervating the MB's medial lobe encode for a gustatory

40   sugar reward signal. Furthermore, we have identified four DANs in the DL1 cluster - DAN-c1, DAN-d1, DAN-f1, and DAN-g1 - each of which innervates distinct compartments of the MB peduncle, lateral appendix, and vertical lobe. Optogenetic activation of DAN-f1 and DAN-g1 alone suffices to substitute for punishment. Furthermore, optogenetic inhibition, calcium imaging results and electron microscopy-based reconstruction of all sensory input circuits to

45   the four DL1 DANs demonstrate that each DAN encodes a different aspect of punishment, with DAN-g1 being of central importance for the salt dependent teaching signal.

To summarize, our investigation has revealed the existence of a cellular division of labor among larval DANs concerning the transmission of dopaminergic reward (pPAM cluster) and punishment signals (DL1 cluster). Individual DANs in each cluster encode for distinct but

50   partially overlapping aspects of the teaching signal. The striking resemblance in the

organizing principle of larval DANs with that of its adult counterpart and the mammalian basal ganglion suggests that there may be a limited number of efficient neural circuit solutions available to address more complex cognitive challenges in nature.

Introduction

55   To cope with a constantly changing environment, animals benefit from learning behavioral

rules that need constant adaptation and updating. To test this ability, researchers often use

Pavlovian or classical conditioning experiments (Pavlov 1927), which examine an organism's

capacity to form associations between sensory cues, called conditioned stimuli (CS), and

rewards or punishments, referred to as aversive and appetitive unconditioned stimuli (US) or

60   teaching signals (Gerber et al 2009, Glanzman 1995, Heisenberg 2003, Menzel 2022,

Waddell 2013, Weber et al 2023b, Widmann et al 2018). Dopaminergic neurons (DANs) and

other modulatory neurons mediate teaching signals that allow animals to classify a given CS

into positive or negative valence based on past experience (Cognigni et al 2018, Menzel

2001, Schultz 2015, Thum & Gerber 2019, Watabe-Uchida et al 2017). For example, when a

65   CS is paired with simultaneous activation of a specific set of DANs in the PAM cluster, larval

or adult *Drosophila* subsequently perceive the odor as attractive (Burke et al 2012,

Eschbach et al 2020, Liu et al 2012, Rohwedder et al 2016, Schroll et al 2006) – a similar

effect was seen after optogenetic activation of DANs located in the ventral tegmental area of

the mouse brain to induce conditioned place preference (Tsai et al 2009). In contrast,

70   activation of larval DL1 DANs (Eschbach et al 2020, Schroll et al 2006, Weiglein et al 2021),

adult PPL1 DANs (Aso et al 2012, Aso & Rubin 2016, Aso et al 2010, Claridge-Chang et al

2009), or DANs in the posterior tail of the striatum of mice were shown to reinforce

avoidance learning (Menegas et al 2018). These examples illustrate that the dopaminergic

system constitutes a highly conserved and fundamental neural principle of the brain, which

75   has been retained throughout the process of evolution and executes comparable functions

across diverse animal species, including even humans (Frick et al 2022).

Our research focuses on the *Drosophila* larva, an organism endowed with a relatively simple

central nervous system, composed of approximately 12,000 neurons (Dumstrei et al 2003,

Winding et al 2023). Classical conditioning in larvae can be analyzed by standardized

80   behavioral assays (Almeida-Carvalho et al 2017, Apostolopoulou et al 2013, Scherer et al

2003, Weber et al 2023a), in conjunction with a wealth of genetic methods for targeting single molecules and neurons (Dietzl et al 2007, Li et al 2014, Perkins et al 2015, Port et al 2020). In addition, the connectome of the brain was recently reconstructed, which enables the extraction and analysis of related wiring motifs, as well as entire learning and memory

85   circuits (Winding et al 2023). Based on these advances, it was possible to establish a more fundamental cellular and synaptic understanding of the larval memory circuit, the mushroom body (MB) network. The MB is a higher-order parallel fiber system present in numerous invertebrate brains, including hemimetabolous as well as holometabolous insects and their larval stages (Strausfeld et al 1998, Strausfeld et al 2020). Analogous to the vertebrate

90   cerebrum, the MB in insects is responsible for forming and maintaining associations (Heisenberg 2003, Menzel 2001, Menzel 2022, Thum & Gerber 2019, Tomer et al 2010, Waddell 2013). In the first developmental stage (L1), the larval MB comprises approximately 110 intrinsic Kenyon cells (KC) per hemisphere, which sparsely encode for conditioned stimulus (CS) information, and is dominated by olfactory input (Eichler et al 2017, Masuda-

95   Nakagawa et al 2009).

The transmission of olfactory information from the peripheral sensory organ is achieved via only 21 olfactory receptor neurons (ORNs) per body side, which in turn send signals across 35 uni- and multiglomerular projection neurons, ultimately reaching the MB KCs (Berck et al 2016, Fishilevich et al 2005, Kreher et al 2005, Masuda-Nakagawa et al 2009, Ramaekers et

100   al 2005). KCs receive modulatory input from 17 neurons, consisting of eight dopaminergic neurons, four octopaminergic neurons, and five neurons with unknown signaling molecule identity (Eichler et al 2017, Rohwedder et al 2016, Saumweber et al 2018, Selcho et al 2009). The DANs are divided into two clusters based on the location of their soma and related neuronal lineage, namely the primary protocerebral anterior medial cluster (pPAM)

105   and the dorsolateral 1 cluster (DL1). Each DAN innervates a specific and non-overlapping site in the MB, which, together with one to three output neurons, defines a distinct compartment (Eichler et al 2017). The larval MB comprises a total of 11 compartments, eight

of which are defined by DAN input (Eichler et al 2017, Saumweber et al 2018). The four DANs of the pPAM cluster (DAN-h1, DAN-i1, DAN-j1, and DAN-k1) innervate the four related compartments (h, i, j, and k) of the medial lobe and provide teaching signals for sugar rewards (Eichler et al 2017, Rohwedder et al 2016, Schleyer et al 2020). The four DANs of the DL1 cluster (DAN-c1, DAN-d1, DAN-f1, and DAN-g1) innervate the lower peduncle (c), the lateral appendix (d), the intermediate (f), and the lower vertical lobe (g) (Eichler et al 2017, Saumweber et al 2018). Previous studies have shown that the DL1 cluster is capable of mediating aversive teaching signals (Eschbach et al 2020, Schroll et al 2006, Selcho et al 2009, Weiglein et al 2021). Moreover, larvae can use a variety of natural stimuli as aversive teaching signals such as heat, electric shock, vibration, bitter substances (like quinine or caffeine), and high salt concentrations (low salt concentrations, however, are rewarding) (Aceves-Pina & Quinn 1979, Apostolopoulou et al 2016, Apostolopoulou et al 2014, Eschbach et al 2011, Gerber & Hendel 2006, Khurana et al 2009, Khurana et al 2012, Niewalda et al 2008, Pauls et al 2010, von Essen et al 2011). However, it is currently unknown if and how these natural stimuli are represented by the DL1 cluster.

We only know that optogenetic stimulation of nociceptive sensory neurons and their downstream basin neurons has been observed to activate three out of the four DL1 DANs, namely DAN-d1, DAN-f1, and DAN-g1 (Eschbach et al 2020). Furthermore, the simultaneous presentation of odor with optogenetic activation of these three DANs has been found to elicit aversion in larvae (Eschbach et al 2020, Weiglein et al 2021). These findings suggest that the activation of DL1 DANs may induce a punishing effect during training, based on somatosensory perception. But can these four individual DANs also process other aversive sensory modalities such as chemosensory information? Utilizing electron microscopy-based reconstruction, all first and second order input neurons of the MB, consisting of a complete set of 102 neuron pairs, were examined (Eschbach et al 2020, Winding et al 2023). The analysis identified numerous afferent sensory inputs via 20 pairs of feedforward neurons (FFNs), which can now be integrated with the published wiring diagram

135   of all chemosensory input neurons (Miroschnikow et al 2018). This enables a more comprehensive understanding of the distinct modes of action of the four DL1 DANs in relation to the processing of different sensory modalities.

Consequently, we have investigated the role of the four DL1 DANs in the larval brain regarding their involvement in encoding a salt-dependent aversive teaching signal at the

140   single-cell level. Utilizing behavioral assays, fluorescence microscopy, calcium imaging experiments, and connectome analyses, we have established a more comprehensive description of the coding space of the DANs at the single-cell level. Our investigation unveils a cellular specialization among the four DL1 DANs, where individual DANs code for different but partially overlapping aspects of the aversive teaching signal

145 **Results**

The larval dopaminergic system is functionally bipartite to convey positive and negative teaching signals

There are eight DANs in the larval brain that innervate the mushroom body that can be anatomically subdivided into four pPAM cluster and four DL1 cluster neurons (Figure 1A-C

150 and Figure 1- figure supplement 1 A-C). Three of the four pPAM cluster DANs can be labeled with the driver line R58E02-Gal4 (Figure 1- figure supplement 1A-E) (Rohwedder et al 2016). Pairing an odor presentation with simultaneous optogenetic blue light activation of R58E02 positive DANs using a mutant microbial-type rhodopsin ChR2^{XXL}, which depolarizes neurons artificially (Dawydow et al 2014), establishes an appetitive olfactory memory (Figure

155 1- figure supplement 1F and G) (Rohwedder et al 2016). Please be aware that the activation of blue light itself does not cause any disruptive behavioral side effects, as it does not independently induce memory formation. In line with this finding, it was shown that ablation or inhibition of R58E02 positive DANs strongly reduces appetitive olfactory memory (Rohwedder et al 2016) but has no functional significance for aversive olfactory memory

160 (Figure 1- figure supplement 1H and I) (Rohwedder et al 2016). Therefore, it can be concluded that pPAM cluster DANs are not involved in the encoding of aversive teaching signals. How are aversive teaching signals encoded?

The driver line TH-Gal4 can be used to label the four DANs of the DL1 cluster (Figure 1A-E), as well as most other DANs, except for a small subset of DANs, such as pPAM DANs and

165 neurons in the subesophageal zone (Figure 1D and E) (Selcho et al 2009). By coupling odor presentation with simultaneous optogenetic activation of TH-Gal4 positive DANs through ChR2^{XXL}, an aversive olfactory memory can be established (Schroll et al 2006), which was observed after both one (Figure 1F) and three (Figure 1G) odor-DAN activation pairings. Consistent with these findings, aversive associative memory formed after one or three

170 training cycles of odor high salt pairings were significantly reduced when DANs activity was

inhibited optogenetically via GtACR2 (Figure 1H and I), a light-gated channel from the algae *Guillardia theta* (Mohammad et al 2017). The observed partial reduction may be attributable to biological, technical, or systemic factors, or a combination of these (a more detailed description can be found in the corresponding part of the discussion).In these experiments

175 salt has to be added during the test to initiate the recall of the aversive memory (indicated as red test plates, Figure 1F-I) (Gerber & Hendel 2006, Schleyer et al 2015, Schleyer et al 2011). Previous studies have shown that the larva only recalls aversive odor-high salt memory when tested in the presence of the teaching signal. The prevailing explanation suggests that the larva engages in memory-based search during the test, but only if the

180 memory content can be used in a beneficial way (Gerber & Hendel 2006, Schleyer et al 2015, Schleyer et al 2011).

Based on our findings and previously published data on both larval and adult stages, we propose a functional division of labor among DANs projecting to the MB. Specifically, pPAM neurons are likely to convey a dopaminergic appetitive teaching signal, whereas DL1

185 neurons may transmit a dopaminergic aversive teaching signal. Nevertheless, the findings related to DL1 DANs necessitate additional validation as the TH-Gal4 driver line used in the experiment labels approximately 120 cells, thus limiting the ability to draw cell-specific conclusions.

190 *Anatomical single-cell analysis of DL1 DANs*

In an earlier investigation, we conducted a screening of several thousand Gal4 lines in order to construct a high-resolution atlas of the mushroom body at the cellular level via light microscopy (Saumweber et al 2018). A subsequent reconstruction of the wiring diagram of the entire MB at the electron microscopic level encompassed not only individual neurons but

195 all synaptic connections (Eichler et al 2017). Utilizing this anatomical work, we identified a total of 102 Gal4 lines featuring MB extrinsic neuron expression patterns. These lines were

subjected to intersectional strategies in order to restrict expression patterns to single

neurons (Eschbach et al 2020, Saumweber et al 2018). For the present study, we employed

eight split-Gal4 lines chosen from this set, as they contained only one or two DANs of the

200   DL1 cluster (Figure 2).

DAN-c1 was specifically labeled by SS02160 (Figure 2B), DAN-d1 by MB328B and MB143B

(Figure 2C and Figure 2- figure supplement 1B), DAN-f1 by SS02180 and MB145B (Figure

2D and Figure 2- figure supplement 1C) and DAN-g1 by SS01716 (Figure 2E). In addition,

we used the split-Gal4 lines MB065B (Figure 2F), SS01702 (Figure 2G) and MB054B

205   (Figure H-K) that specifically labeled DAN-c1/DAN-f1, DAN-c1/MBIN-e1 or DAN-f1/DAN-g1,

respectively. For our evaluation we crossed each split-Gal4 line with the reporter strain UAS-

mCD8::GFP;mb247-LexA,lexAop-mRFP (Burke et al 2012) that allowed us to monitor the

expression pattern (green) in the background of a MB specific reference staining (magenta).

Our results confirm the reported specificity even despite the use of different reporter lines

210   and lacking antibody staining (Eschbach et al 2020), as we analyzed only endogenous GFP

expression using a more cautious and faster whole mount staining protocol (see Material

and Methods). In addition, we also verified that the DANs labeled via MB054B are indeed

dopaminergic as the GFP expression overlaps with an anti-TH antibody staining (Figure 2-

figure supplement 2). Having established anatomical validity, we subsequently employed

215   these lines to explore the physiological response of each DAN within the DL1 cluster to

sugar and salt stimuli. For the purpose of analyzing DAN-d1 and DAN-f1, we limited our

investigation to the split-Gal4 lines MB328B and SS02180 as these lines displayed more

robust expression levels.

220   *Physiological single-cell analysis of DL1 DANs*

In order to investigate how high salt teaching signals are represented at the level of

individual DL1 DANs, we conducted calcium imaging experiments using a microfluidic chip

on intact and immobilized larvae, as previously described (Si et al 2019). We expressed
GCaMP6m under the control of four different split-Gal4 lines specific to DL1 DANs (SS02160
225   (DAN-c1), MB328B (DAN-d1), SS02180 (DAN-f1), and SS01716 (DAN-g1)), as well as the
pPAM specific R58E02-Gal4, and exposed the larvae to solutions containing 100 mM salt, 1
M salt, or 500 mM fructose. Our results showed that only DAN-c1 responded to 100 mM salt
(Figure 3A). While DAN-c1, DAN-d1, and DAN-g1 responded to 1 M salt (Figure 3A, B, and
D). Conversely, pPAM DANs showed a reduced signal in response to 1 M salt (Figure 3E).
230   Both DAN-c1 and pPAM DANs showed a calcium increase in response to 500 mM fructose
(Figure 3A and E). Although DAN-f1 did not respond to either salt concentration, a reduction
in the calcium signal was seen for 500 mM fructose (Figure 3C). Overall, these results
provide support for the functional division of the larval dopaminergic system at the
physiological level, where most DL1 neurons are activated by high salt concentrations, with
235   one exception inhibited by fructose. In contrast, pPAM cluster neurons display the opposite
response, being inhibited by high salt concentrations and strongly activated by fructose.

Behavioral single-cell analysis of DL1 DANs

Next, we analyzed the role of single DL1 DANs in aversive olfactory learning (Figure 2-
240   figure supplement 1D-G, Figure 4- figure supplement 1-4). We employed split-Gal4 lines
SS02160 (DAN-c1), MB328B (DAN-d1), SS02180 (DAN-f1), and SS01716 (DAN-g1) to
express the apoptosis proteins Hid and Reaper, resulting in the ablation of the respective
individual DL1 DANs. The ablated animals were subjected to the aversive odor-high salt
memory assay, using one (Figure 4- figure supplement 1A-D) and three training cycles
245   (Figure 4- figure supplement 1E-H). We utilized two different training protocols to
differentiate between the effects on short-term memory (STM) and anesthesia-resistant
memory (ARM), as previous studies have demonstrated that larvae exhibit both after one
training cycle, whereas only ARM is present after three training cycles (Widmann et al 2016).

However, we found that the ablation of individual DL1 DANs had no effect on aversive olfactory memories in any of the eight experiments conducted (Figure 4- figure supplement 1A-H). For DAN-d1 and DAN-f1 we verified these results by using the additional split-Gal4 lines MB143B and MB145B (Figure 2- figure supplement 1D-G). In addition, ablation of DAN-d1, DAN-f1 and DAN-g1 individually did not have any impact on odor-quinine and odor-fructose memories established after one cycle training (Figure 4- figure supplement 2 and 3). Moreover, MB input neurons that are not dopaminergic (OAN-a1, a2, MBIN-b1, b2, and OAN-e1) were also found to have no relevant effect (Figure 4- figure supplement 4). These results suggest that although DL1 DANs respond to high salt concentrations (Figure 3), they are not individually necessary in ablation experiments for mediating the aversive teaching signal. The data suggest that the aversive teaching signal may be distributed across multiple neurons, indicating possible functional redundancy or, alternatively, developmental compensation that could attenuate the effects of DL1 cell ablation. However, it is possible that there are developmental compensatory effects for DL1 cluster DANs, allowing the remaining three neurons (or other cells) to take over the function of the ablated one. This hypothesis was analyzed later in optogenetic inhibition experiments.

The combination of DAN-f1 and DAN-g1 is required to establish aversive odor-high salt memories

In order to investigate a possible redundancy among DANs, we proceeded by using split-Gal4 lines MB054B (DAN-f1/DAN-g1), MB065B (DAN-c1/DAN-f1), and SS01702 (DAN-c1/MBIN-e1) to express apoptosis proteins Hid and Reaper. This approach selectively ablated either two DL1 DANs or one DL1 DAN and another MB input neuron with unknown neurotransmitter identity. It should be noted that no additional split-Gal4 lines are presently available for further DL1 DAN combinations or the complete DL1 cluster. We then subjected the larvae to the aversive odor-high salt memory assay, using one (Figure 4A-C) and three

275　(Figure 4D-F) training cycles. Only larvae with ablated DAN-f1/DAN-g1 exhibited significantly

　　reduced aversive olfactory memory after one cycle training (Figure 4A). No memory

　　reduction was observed when ablating DAN-c1/DAN-f1 or DAN-c1/MBIN-e1 (Figure 4B and

　　C). Ablation of all three sets of neurons, including MB054B, had no effect on aversive

　　olfactory memory after three training cycles (Figure 4D-F). To ascertain the independence of

280　the DAN-f1/DAN-g1 neuron-mediated impairment of aversive olfactory memory after one

　　cycle training from the specific odors used, hexyl acetate and benzaldehyde were employed

　　instead of amyl acetate and benzaldehyde. The results of these experiments confirmed the

　　previous findings, as ablation of DAN-f1/DAN-g1 neurons significantly reduced aversive

　　odor-high salt memory after one cycle training (Figure 4G), but not after three cycles (Figure

285　4J). Innate olfactory and gustatory behavior was not impaired by DAN-f1/DAN-g1 ablation

　　(Figure 4- figure supplement 5).

　　Subsequently, we investigated the selectivity of DAN-f1/DAN-g1 neurons concerning the

　　nature of the aversive teaching stimulus. To this end, we utilized quinine as an aversive

　　teaching signal, a stimulus that has been demonstrated to be learnable by the larva, as

290　evidenced by previous studies (Apostolopoulou et al 2014, El-Keredy et al 2012). Removal

　　of the DAN-f1/DAN-g1 neurons using the split-Gal4 line MB054B had no effect on aversive

　　odor-quinine memory independent of the number of training cycles (Figure 4H and K).

　　Furthermore, ablation of DAN-f1/DAN-g1 did not alter appetitive odor-fructose memories

　　after one or three training cycles (Figure 4I and L). Therefore, the DL1 DAN-f1/DAN-g1

295　combination seems to be specifically required for aversive odor-high salt memory. This is

　　further supported by anatomical data that verified the effectiveness of Hid and Reaper using

　　the MB054B split-Gal4 line (Figure 4- figure supplement 6).

Individual DL1 DANs can instruct an aversive memory recalled on a salt plate

300   In order to determine if the activation of individual DL1 DANs signals aspects of the natural

high salt punishment, we conducted a one training trial learning experiment using ChR2^{XXL} to

artificially activate DAN-c1, DAN-d1, DAN-f1, DAN-g1, and the combination of DAN-f1/DAN-

g1. We used split-Gal4 lines SS02160, MB328B, SS02180, SS01716, and MB054B,

respectively (Figure 5). Simultaneously activating DAN-f1/DAN-g1 via ChR2^{XXL} during an

305   odor presentation establishes an aversive olfactory memory (Figure 5E and F). These

results are comparable to those observed for TH-Gal4 (as shown in Figure 1F and G).

Additionally, this memory is observed after one (Figure 5E) and three training cycles (Figure

5F). Furthermore, single-cell analysis indicates that individual activation of DAN-f1 and DAN-

g1, but not of DAN-d1 and DAN-c1 (please note that experimental animals are different from

310   zero and significantly different to one control), can induce an aversive odor memory (Figure

5A-D).

An individual DL1 DAN is acutely necessary to establish an aversive odor-high salt memory

Next, we examined the extent to which the individual DL1 DANs are acutely required for the

315   transmission of a functional aversive teaching signal. We have already shown that the

combination of DAN-f1/DAN-g1 is required for aversive odor-high salt memory (Figure 4A

and G). However, in these experiments DANs were removed during the entire development,

which may have led to compensatory mechanisms or side effects.

To overcome this limitation, we left the individual DL1 DANs intact during embryonic and

320   larval development and used GtACR2 to selectively inhibit DAN activity only during training,

thus isolating the teaching signals from other DAN functions. Optogenetic inhibition of

individual DL1 DANs showed no effect on aversive odor-high salt memories (Figure 6A-C),

except for DAN-g1 that led to a small but significant reduction (Figure 6D). Optogenetic

inhibition of DAN-f1/DAN-g1 resulted in a similar reduced aversive odor-high salt memory.

325   This was the case after one training cycle (Figure 6E) and three training cycles (Figure 6F).

Thus, the learning deficit in this experiment was more pronounced than after ablation of these two DL1 DANs throughout development (Figure 4D).

A synaptic reconstruction of all chemosensory inputs for each individual DL1 and pPAM
DAN.

Certainly, a comprehensive analysis from the sensory neurons responsible for salt perception, via the interneurons involved in salt information processing, up to the dopaminergic DL1 teaching neurons would be highly desirable to understand why and how the different DANs respond to salt. Regrettably, such an analysis is currently not feasible due to our limited understanding of salt perception and which sensory cells encode this crucial information (Liu et al 2003). Therefore, this question cannot be answered by the larval connectome. However, it is possible to analyze how qualitative differences between individual DL1 DANs (Figure 3-6) are represented in the connectome, since not all DANs code equally for the punishment teaching signal. Therefore, we compared the input wiring diagrams of the individual DANs of the pPAM and DL1 clusters by mapping the synaptic connections from sensory neurons - to interneurons - to individual DANs, using the full larval brain EM volume (Figure 7, Figure 7- figure supplement 1) (Eschbach et al 2020, Miroschnikow et al 2018, Winding et al 2023).

We found that DAN-c1, DAN-d1, DAN-f1, and DAN-g1 (black circle) clearly differ in their downstream connections. They do however receive input from 1, 6, 6 or 11 sensory-to-DAN interneurons (orange circle), respectively (Figure 7A), which in turn means that these DL1 DANs are connected to 3, 42, 35, and 9 sensory neurons (numbers in colored circuits below). Please note that we restricted our analysis on direct sensory-to-DANs interneurons (single hop analysis) and did not expand the analysis onto two hop or three hop connections to keep the evaluation restricted to the shortest paths from sensory neurons to DANs. There is no direct input from sensory neurons onto DL1 DANs. Sensory input thus appears to be

increasingly instructing DAN-c1, DAN-d1, DAN-f1, and DAN-g1. The analysis also shows that all four DL1 DANs receive information from all gustatory senses via these interneurons (coded by the yellow color for sensory composition). In addition, DAN-d1, as well as DAN-f1, receive information from olfactory and thermosensory neurons (Figure 7D). In general, less processed sensory input seems to be more important for DAN-f1 and DAN-g1 than for DAN-c1 and DAN-d1, as they get 17% and 11% of their entire input from these sensory-to-DAN interneurons in contrast to 1% and 5%, respectively. The four DL1 DANs show a conserved connection motif, as they get additional input from other upstream neurons (US) that are not directly linked to sensory neurons (grey circles; between 17% and 37%), as well as from MB KCs (purple circles; between 36% and 49%) and little MBON feedback (dark-red circle; between 1% and 2%). Examination of the different input types of DL1 DANs presented here, shows that DAN-f1 and DAN-g1 receive the highest proportion of sensory input relative to other DANs. Even by looking at the sensory-to-DAN interneurons individually, one can again see the familiar pattern of differences and partial overlap (Fig. 7D). DAN-d1, DAN-f1 und DAN-g1 all get input from interneuron pair 1 in contrast to DAN-c1. However again, DAN-f1 and DAN-g1 are most strongly connected to this neuronal pair (indicated by the thickness of the lines). DAN-f1 and DAN-g1 are therefore also the most similar in terms of their gustatory input (due to their connection via interneuron pair 1), despite all the clearly recognizable differences.

For the pPAM DANs we found the same general connectivity principles as for DL1 DANs regarding input from MBONs, KCs, and non-sensory upstream neurons (Fig 7B). Please note that the DAN-h1 neuron is not present at the early L1 EM brain volume and thus cannot be included in this evaluation (Figure 7B and E). In contrast, however, it is striking that pPAM DANs seem to get input from more interneurons connected to the sensory system (up to 16 for DAN-j1) signaling additional internal somatosensory and CO_2 information. A closer inspection of the sensory-to-DAN interneurons showed that 27 of the 35 neurons are specifically innervating only DL1 or pPAM DANs (Figure 7C). Only 8 out of 35 interneurons

connect to both clusters. Based on a calculated hub score sensory-to-DAN interneurons pair

380   1 (FB2IN-12) and pair 5 (FFN-21) were likely most significant for instructing DL1 DANs with

sensory information. Whereas pairs 10 (FFN-17) and 11 (FFN-12) were – based on

connectivity – most important in instructing DL1 and pPAM DANs. For the three pPAM DANs

sensory-to-DAN interneuron pairs 16, 17, 18, 19 and 20 resulted in the highest hub scores

(FFN-37, FFN-4, FFN-5, FFN-18 and FB-4).

385   The difference in cellular connectivity of DL1 and pPAM DANs is also reflected in different

local input brain hubs: axo-dendritic synapses of sensory-to-DAN interneurons with DL1

DANs formed a characteristic hook-shape in the superior protocerebrum; in contrast, pPAM

DANs get synaptic input from sensory-to-DAN interneurons in the anterior inferior

protocerebrum (Figure 7F).

Discussion

A functional assignment for individual DANs of the DL1 cluster in their ability to encode a salt punishment teaching signal.

The discovery of DL1 neurons as aversive teaching signal mediators in larval *Drosophila*

395 (Eschbach et al 2020, Schroll et al 2006, Selcho et al 2009, Weiglein et al 2021) complements previous findings that demonstrate the sufficiency of four DANs from the pPAM cluster as an appetitive reinforcement signal in these animals (Rohwedder et al 2016). This dichotomy demonstrates an organizational principle shared by adult *Drosophila* and larvae (Aso et al 2012, Aso & Rubin 2016, Aso et al 2010, Burke et al 2012, Claridge-Chang

400 et al 2009, Das et al 2014, Huetteroth et al 2015, Liu et al 2012, Schleyer et al 2020, Schroll et al 2006, Selcho et al 2009, Weiglein et al 2021, Yamagata et al 2015), despite a significant reduction in cell numbers. The principle appears to apply to mammals as well (Groessl et al 2018, Lammel et al 2012, Menegas et al 2018, Schultz 2015). But how do individual DL1 neurons contribute to the overall teaching signal? Is the signal evenly

405 distributed across multiple cells, supporting the mass action hypothesis, or do individual DANs specifically encode for the salt stimulus (Rohwedder et al 2016)? Our results show that the four DL1 DANs perform this function in a mixture of both forms. There is no unique labeled line because both ablation of each individual DAN (Figure 2- figure supplement 1D-G, Figure 4- figure supplement 1) and the acute optogenetic inhibition (Figure 6) do not

410 abolish the odor-high salt memory. The teaching signal is encoded in a redundant manner within the DL1 cluster, yet its distribution among the four DANs is non-uniform and relies heavily on the functionalities of DAN-g1 and DAN-f1 (Table 1, Figure 4-6). Each DAN possesses a distinct identity, which is defined by its specific, albeit partially overlapping, neuronal input circuitry and physiological response to salt (Table 1, Figure 3 and 7).

415 DAN-c1 responds to both positive and negative gustatory stimuli (Figure 3A). However, the activity of the cell is not sufficient or necessary for aversive memory (Figure 4- figure

supplement 1A and E, Figure 5A and Figure 6A). These results suggest that DAN-c1 activity does not encode a taste-specific teaching signal, which is supported by EM analysis, as this particular cell is only weakly connected to three sensory neurons via a single sensor-to-DAN interneuron (Figure 7A and D). However, this interpretation might depend on the type and duration of DAN stimulation, as it was recently shown that - as in our case - optogenetic activation does not induce aversive memory, but thermogenetic 30 minute stimulation in a non-reciprocal one trail paradigm leads to aversive memory (Qi et al 2024). In addition, it could also be that the function of DAN-c1 is more to integrate state-dependent network changes, similar to the role of some DANs in innate and state-dependent preference behavior of adults (Boto et al 2019, Cazale-Debat et al 2024, Cohn et al 2015, Lewis et al 2015, Mohammad et al 2024, Siju et al 2020). DAN-d1 activity appears to have a negligible impact on the formation of odor-high salt memories in the paradigms we tested. Even though the neuron responds to high salt concentrations (Figure 3B) and receives input from 42 sensory neurons via six sensor-to-DAN interneurons (Figure 7A and D), its activity did not induce an aversive memory (Figure 5B) and was not found to be essential for odor-high salt memory (Figure 2- figure supplement 1D and E, Figure 4- figure supplement 1B and F, Figure 5B). Nonetheless, it is possible that DAN-d1 could instruct other salt-dependent memory types, established after non-associative training, second-order conditioning, or reversal learning (Kacsoh et al 2015, Konig et al 2019, Mancini et al 2019, Paranjpe et al 2012, Tabone & de Belle 2011, Yamada et al 2023). Other learning experiments in larvae (Eschbach et al 2020, Weiglein et al 2021) demonstrate that DAN-d1 activation can indeed guide the formation of aversive odor memory. However, these studies employ three training repetitions rather than a single repetition, as in our approach. Additionally, they utilize absolute conditioning with a single odor, in contrast to our protocol of differential conditioning involving two odors. Furthermore, their testing does not incorporate salt as an aversive gustatory stimulus. The observed differences in results suggest that DAN-d1 activation may only establish aversive memory if the stimulation is applied multiple times and is more distinctly presented during training. . In combination, DAN-f1 activity and DAN-g1 seem to

445   have a crucial function in high salt punishment. DAN-f1 is inhibited by sugar (Figure 3C) and

obtains input from 35 sensory neurons through six sensor-to-DAN interneurons (Figure 7A

and D). Despite ablation and acute suppression of activity having no impact on the retention

of a memory related to a high salt stimulus (Figure 2- figure supplement 1C, E and G, Figure

4- figure supplement 1C and G, Figure 6C), DAN-f1 activity can instruct the formation of an

450   aversive memory (Figure 5C). However, there is the possibility that the impact of the ablation

experiments could be more pronounced than what is observed here, given the potential for

developmental compensation. This has been anatomically demonstrated in the adult

dopamine mutants by serotonergic cells (Niens et al 2017). The identified function for DAN-

f1 activity is consistent with previous research, although their experimental designs differ

455   from our own in the test situation (Eschbach et al 2020, Weiglein et al 2021). Consequently,

it is plausible that DAN-f1 activity encodes for a teaching signal that carries additional

information beyond high salt. It is evident that DAN-g1 exerts the most potent effect.

Following exposure to high salt, this cell becomes active (Figure 3D), is crucial for the

formation of an odor-high salt memory (Figure 6D) and can even establish an aversive

460   memory that can be recalled on a high salt containing test plate (Figure 5D). Additionally, it

receives extensive sensory input from nine neurons through 11 distinct interneurons (Figure

7A and D). Therefore, DAN-g1 is the only cell of the DL1 cluster that satisfies all

prerequisites to independently encode the high salt teaching signal. In summary, the DL1

cluster appears to be arranged in a reasonable functional configuration. The instructive

465   signal is encoded by a mere four cells, with the requisite of partial redundancy to preclude

complete loss of function in the event of single-cell failure. Nevertheless, the cells manifest

variation, thereby covering different and cumulative characteristics of stimulation.

However, these cellular-level interpretations are inevitably oversimplified. The MB054B split

line occasionally exhibits a weak indication of a third cell (Figure 2J), introducing a degree of

470   variability in the experiments. Similarly, DAN-f1 demonstrates no direct response to salt

(Figure 3C), suggesting it is not essential for salt learning, yet it triggers a potent aversive

memory (Figure 5C). Nevertheless, it is worth mentioning that the physiological and EM analyses were conducted using first instar larvae, whereas the light microscopy experiments and behavioral tests were performed with third instar larvae. Hence, it is possible that developmental-specific alterations may have occurred, which were not accounted for in this study. Regrettably, our understanding of how salt is perceived in larvae remains limited (Liu et al 2003). Assuming peripheral organ perception as primary, the left and right interneuron 1 (also called FB2IN-12), connected to DAN-d1, DAN-f1, and DAN-g1, is presumed to be centrally important as it encodes for the main peripheral gustatory information (encoded by yellow color, Figure 7C, first and second line). However, these DANs display no response to low salt concentrations and varying responses to high salt concentrations and fructose. In contrast, DAN-c1 is the only cell that responds to high and low concentrations of salt (Figure 3A) but is not connected to interneuron pair 1 or other sensory-to DAN interneurons (except interneuron 20L; Figure 7C). Further experiments are therefore necessary to refine our initial cellular-level description and comprehend the functional spectrum of individual DL1 DANs more comprehensively.

Furthermore, other factors must also be taken into account. Blocking almost all TH-positive cells does not result in the complete disappearance of the salt memory. Instead, it is merely reduced by half (Figure 1H and I). There could be several reasons for this outcome, or a combination thereof. For example, it is plausible that the UAS-GtACR2 effector does not completely suppress the activity of dopaminergic neurons. Additionally, the memory may encompass different types (Widmann et al 2016), not all of which are associated with dopamine function. It is also noteworthy that TH-Gal4 does not encompass all dopaminergic neurons - even a neuron from the DL1 cluster is missing (Selcho et al 2009). Considering that we are using high salt concentrations in this experiment, it is also conceivable that no taste-driven memories are formed solely due to the systemic effects of salt (e.g., increased osmotic pressure). These effects naturally complicate the interpretation of the data at the single-cell level.

The functioning of the DL1 cluster DANs as a multimodal node for punishment.

500   Larvae, of course, perceive not only high salt concentrations as punishing, but also other sensory stimuli such as bitter substances (quinines and caffeine), electric shock, temperature, mechanosensory input (vibration), and light (Aceves-Pina & Quinn 1979, Apostolopoulou et al 2016, Apostolopoulou et al 2014, Eschbach et al 2011, Gerber & Hendel 2006, Khurana et al 2012, Pauls et al 2010, von Essen et al 2011). However, the

505   extent to which the DL1 cluster is involved in coding for most of these aversive teaching signals remains unclear. Currently, comparisons at the level of DANs can only be made for sensory inputs that are dependent on high salt or mechanosensation. Interestingly, similar to high salt concentrations, somatosensory information can also result in the establishment of aversive olfactory memory in larvae (Eschbach et al 2020). The transmission of these

510   somatosensory teaching signals is orchestrated by mechanosensory neurons from the chordotonal organs, class IV multidendritic nociceptive neurons distributed along the body, and multisensory basin neurons (Hwang et al 2007, Jovanic et al 2016, Ohyama et al 2015, Tracey et al 2003). The connectome revealed that these mechanosensory neurons also link to the DL1 cluster through three to four interneurons and elicit the activation of DAN-d1,

515   DAN-f1, and DAN-g1 (Eschbach et al 2020). Upon comparing the interneurons that relay high salt and mechanosensory information to the DL1 cluster, six pairs of interneurons are of critical significance, respectively. These pairs include FFN-20, FB2N-12, FB2N-19, FFN-23, FFN-29, and FB2N-15 for mechanosensory information (Eschbach et al 2020), and FB2N-12, FB2N-18, FB2IN-11, FB2IN-6, FFN-21, and FFN-24 for high salt concentrations (Figure

520   7; Figure 7- figure supplement 1A, rightmost column). Interestingly, only one pair of interneurons - FB2N12 - is shared between the two types of information and the other five types are specific to the sensory modality. The wiring patterns of DAN input neurons exhibit a notable consistency, with a limited number of neurons dedicated to a specific sensory modality, and single neurons shared across different modalities. This results in a unique

525   combination of cellular codes that are partly redundant and partly specific for the four DL1

DANs. While certain DANs may hold greater significance for a particular information, such as DAN-d1 for somatosensory information, the DAN-g1 cell appears to play a general central role for encoding an aversive teaching signal. DAN-g1 responds to all aversive stimuli tested thus far and its activity has proven informative for all aversive memories analyzed to date. In order to gain a deeper comprehension of the processing of diverse sensory modalities in the larval brain, it is crucial to broaden the analytical framework presented in our work to encompass visual stimuli, various bitter substances, and temperature stimuli.

However, the overall depiction presented above of the information flow and processing of DL1 DANs is an oversimplification. Indeed, DL1 and pPAM DANs are among the most complex and highly recurrent neurons in the brain (Winding et al 2023). This exceptional level of recurrent connectivity enables DANs to provide high-dimensional feedback, which enables them to encode a diverse range of features and engage in parallel computations. Such computations can guide a working memory, contain feedback from neurons that integrate both learned and innate values, and receive long-range feedback from descending neurons that encode motor commands; all in addition to polysynaptic feedforward inputs from the entire set of sensory modalities.

A cellular comparison of DAN function across *Drosophila* metamorphosis

A recent study conducted by Truman and colleagues explored the integration, or lack thereof, of individual input and output neurons from the larval MB into the adult network following metamorphosis (Truman et al 2023). The study demonstrated that while all four DANs of the pPAM cluster perished, three out of the four cells from the DL1 cluster persisted as part of the adult MB circuitry. Only DAN-f1 altered its innervation in the adult brain, relocating from the larval MB to the adult superior medial protocerebrum. In contrast, DAN-c1, DAN-d1, and DAN-g1 maintained their function as MB DANs in the adult system, identified as PPL1-γ1pedc, PPL1-γ2α'1, and PPL1-γ1, respectively. Due to the lack of

appropriate genetic tools, data on the function of PPL1-γ1 is scarce (Aso et al 2014). Therefore, a functional comparison between larval and adult DANs is only possible for the DAN-c1/PPL1-γ1pedc and DAN-d1/PPL1-γ2α'1 combination. Embryonic-born DAN-c1/PPL1-γ1pedc, DAN-d1/PPL1-γ2α'1, and DAN-g1/PPL1-γ1 exhibit sustained activity and respond to various sensory stimuli, including electric shock, temperature, or bitter taste, that can instruct short lasting aversive memories (Aso et al 2012, Aso & Rubin 2016, Aso et al 2010, Claridge-Chang et al 2009, Das et al 2014, Galili et al 2014, Huang et al 2024, Kirkhart & Scott 2015, Tomchik 2013, Villar et al 2022, Vrontou et al 2021). Despite the massive reorganization of the mushroom body, the basic function of DAN-c1/PPL1-γ1pedc and DAN-d1/PPL1-γ2α'1, mediating punishment teaching signals that instruct short-term memories, remains the same. Evidence indicates that the underlying molecular signaling pathways maintain their identity and kinetics throughout development, with larval and adult interstimulus interval curves for DAN-d1/PPL1-γ2α'1 being almost identical (Aso & Rubin 2016, Huang et al 2024, Weiglein et al 2021). During adulthood, PPL1-DANs are responsible for numerous other functions, including innate olfactory responses, suppression of appetite memory in response to nutritional status, pre-exposure learning, memory reconsolidation, and forgetting (Berry et al 2015, Berry et al 2018, Felsenberg et al 2017, Jacob et al 2021, Krashes et al 2009, McCurdy et al 2021, Placais et al 2012, Siju et al 2020, Tian et al 2016, Vrontou et al 2021). Additionally, they are involved in various behaviors like locomotion and sleep (Berry et al 2015).

An open question is whether larval DL1 DANs encode these same functions to a similar extent. Our study establishes a fundamental framework that can be leveraged to investigate this issue. Considering the numerical simplicity of the larval nervous system, the available genetic tools, and the complete connectome, it may be possible to attain an understanding of the larval dopaminergic system at the level of single cells and single synapses. It is plausible that nature has a limited number of effective neural circuit solutions for complex cognitive and behavioral problems, such as the remarkable ability of brains to learn from a

small number of examples. Therefore, comprehending the larval dopaminergic system may

580   provide insight into the shared circuit motifs that exist across the animal kingdom.

Materials and Methods

<u>Fly Strains</u>

Flies were raised and maintained on *Drosophila* standard food at 25°C, 60-80% relative

585 humidity and a 14/7-hour light/dark cycle. For anatomical analysis, UAS-mCD8::GFP (Bloomington stock center no. 32194) (Selcho et al 2009), UAS-mCD8::GFP;mb247-LexA,lexAop-mRFP/TM3,Sb (Burke et al 2012) and UAS-mCD8::GFP;nSyb-LexA,lexAop-mRFP/TM6B (Burke et al 2012, Riabinina et al 2015) were used to analyze the morphology of DANs. For behavioral experiments, UAS-hid,rpr (Abbott & Lengyel 1991, White et al

590 1996), UAS-GtACR2 (Mohammad et al 2017) (Bloomington stock center no. 92984) and UAS-ChR2^{XXL} (Dawydow et al 2014) (Bloomington stock center no. 58374) were used to ablate or optogenetically silence or activate neurons by blue light (470 nm). UAS-GCaMP6m (Chen et al 2013) (Bloomington stock center no. 42748) flies were used for Ca^{2+}-imaging experiments. The Gal4 strains TH-Gal4 (Friggi-Grelin et al 2003) and R58E02-Gal4

595 (Rohwedder et al 2016) (Bloomington stock center no. 41347) were used to manipulate different sets of DANs. Split-Gal4 lines MB054B, MB065B, MB328B, MB143B, MB145B, SS02160, SS02180, SS01716, SS01958, SS21716, SS24765, SS01702 (Eschbach et al 2020, Pfeiffer et al 2010, Saumweber et al 2018) were used for anatomical, physiological and behavioral experiments. w^{1118} flies (Bloomington stock center no. 3605) were used to

600 obtain heterozygous driver and effector control groups. See supplementary table 1 for more information.

<u>Anatomical Analysis</u>

Immunostaining

605 To confirm expression patterns of R58E02-Gal4, TH-Gal4 and MB054B split-Gal4, flies were crossed to UAS-mCD8::GFP. To assess the functionality of the ablation by effector UAS-

hid,rpr, split-Gal4 MB054B was crossed with the latter. Furthermore, MB054B>+ and +>UAS-hid,rpr were anatomically examined for validation. All flies were raised at 25°C for seven days before dissection. Third instar larvae (wandering stage) were put on ice and

610   dissected in PBS (Phosphate Buffered Saline; Sigma Aldrich, cat. no. P4417). Brains were fixed in 4% formaldehyde solution (in PBS, Thermo Scientific, cat. no. 047392.9M) for 20 minutes at room temperature. After eight rinses in PBT (PBS with 3% Triton X-100, Sigma Aldrich, cat. no. X100), brains were blocked with 5% normal goat serum (NGS, Sigma Aldrich, cat. no. G9023) in PBT for 1 hour at room temperature and incubated for 48 hours

615   with primary antibodies at 4°C. Before application of the secondary antibodies for at least 24 hours at 4°C, brains were washed seven times with PBT. After that, larval brains were again washed eight times with PBT and mounted on poly-L-lysin-coated coverslips (Janelia FlyLight DPX mounting receipt: https://www.janelia.org/project-team/flylight/protocols), dehydrated through a series of increasing concentrations of ethanol and cleared three times

620   for 5 minutes in xylene (Sigma Aldrich, cat. no. 247642). At the end, larval brains were mounted in DPX mounting medium (dibutyl phthalate in xylene, Sigma Aldrich, cat. no. 06522) and left to rest in the dark for at least 24 hours before imaging. Primary antibodies were: rat anti-N-Cadherin (1:50; Hybridoma, cat. no. DN-Ex #8-s), rabbit anti-GFP (1:1000; Life Technologies, cat. no. A6455) ,mouse 4F3 anti-DLG (1:200; Hybridoma, cat. no. 4F3

625   anti-discs large) and mouse anti-TH (1:50, Immunostar, cat. no. 22941). Secondary antibodies were: goat anti-rat Alexa Fluor 647 (1:500 or 1:250, for anti-N-Cadherin, Life Technologies, cat. no. A21247), goat anti-rabbit Alexa Fluor 488 (1:500, for anti-GFP, Life Technologies, cat. no. A11008), goat anti-mouse Alexa Fluor 568 (1:500, for anti-DLG, Life Technologies, cat. no. A10037) and goat anti-mouse Cy3 (1:200, for anti-TH, life

630   technologies, cat. no. A10521).

Native fluorescence

For native fluorescence, we used a previously described protocol for adult flies (Pitman et al 2011) and adapted it to larvae. All split-Gal4 lines were crossed to UAS-mCD8::GFP;mb247-

LexA,lexAop-mRFP/TM3,Sb, while split-Gal4 line MB054B was additionally crossed to UAS-mCD8::GFP;nSyb-LexA,lexAop-mRFP/TM6B and raised at 25°C for seven days. Third instar larvae (wandering stage) were put on ice and dissected in PBS. Brains were fixed under vacuum in 4% formaldehyde solution (in PBS) for 25 minutes at room temperature. After five washes in PBT (PBS with 3% Triton X-100), larval brains were rinsed twice directly and once for 10 minutes under vacuum in PBS at room temperature. Samples were mounted on poly-L-lysin-coated coverslips and embedded in Vectashield (Vector Laboratories, cat. no. H-1000-10). Slides were stored in the dark and cold for at least 24 hours before imaging.

Confocal microscopy

Confocal microscopy was conducted on a Zeiss LSM800 confocal laser scanning microscope with ZEN 2.3 software. All images were projected and adjusted with ImageJ (Fiji is just ImageJ, Version 1.53c, Java 1.8.0_172 (64-bit)). Final Figures were arranged and labeled in Affinity Publisher 2.1.1.

Functional Imaging and microfluidics

For larval imaging experiments, adult flies were transferred to larvae collection cages (Genesee Scientific) containing grape juice agar plates and 180 mg of fresh yeast paste per cage. Flies were allowed to lay eggs on the agar plate for 1–2 days before the plate was removed for collection of larvae. Calcium imaging experiments were performed in L1 larvae (2 days AEL). We used a previously described method for microfluidic delivery of chemicals in aqueous form with simultaneous imaging of calcium activity in intact larvae (Si et al 2019). All experiments used an 8-channel microfluidic chip equipped with a vacuum port to stabilize

the animal's head. The same tastants were used as in the behavioral experiments (NaCl: VWR Chemicals, cat. no. 27810.364; D-Fructose: Sigma Aldrich cat. no. 47740). Stimuli

660 consisted of 5-10 s pulses interleaved with 15 s water washout periods. An L1 larva was washed with deionized water and loaded into the microfluidic device using a 1 mL syringe filled with Triton X-100 (0.1% [v/v]) solution. The animal was pushed to the end of the loading channel with its dorsal side facing the objective. GCaMP6m signal was recorded using an inverted Nikon Ti-E spinning disk confocal microscope and a 60X water immersion

665 objective (NA 1.2). A CCD microscope camera (Andor iXon EMCCD) captured frames at 30 Hz. Recordings from at least 5–7 larvae were collected for each condition. For each larva the average response over 5s around the peak value during stimulus presentation was compared to chance level analyzed via normal t-test. Responses during stimulation were normalized based on the baseline response 10s before the stimulation. P-values for each

670 experiment are given in the respective figure panel. Each graph shows the mean calcium signal plotted as the relative response strength ΔF/F and the related standard error of the mean on the y axis. The time in seconds is given below each graph on the x axis. The grey box indicates the duration of the stimulus application. The sample size of each group (N=5-7) is given above each row. n.s. $p > 0.05$; * $p < 0.05$.

Behavioral Experiments

Associative olfactory learning

Standard experiments were done as described before (Apostolopoulou et al 2013, Gerber et al 2013, Hendel et al 2005, Michels et al 2017, Widmann et al 2018). Learning experiments

680 were conducted on assay plates (85 mm diameter, Sarstedt, cat. no. 82.1472) filled with a thin layer of either 2.5% (w/v) pure agarose solution (Sigma Aldrich, cat. no. A9539) or 2.5% (w/v) agarose plus either 1.5 M sodium chloride solution (VWR Chemicals, cat. no. 27810.364), 2 M D-Fructose solution (Sigma Aldrich cat. no. 47740) or 10 mM quinine

solution (quinine hemisulfate; Sigma Aldrich cat. no. Q1250). Before closing and labeling the

685   lids, solutions were let to cool down at room temperature to avoid condensation. Plates were

stored at 18°C and used within five days. As olfactory stimuli, we used two different odor

combinations: either amyl acetate (AM, Sigma Aldrich, cat. no. 46022) diluted 1:250 in

paraffin oil (Sigma Aldrich, cat. no. 76235) and undiluted benzaldehyde (BA, Sigma Aldrich,

cat. no. 12010) or benzaldehyde and hexyl acetate (HA, Sigma Aldrich, cat. no. 108154)

690   both diluted 1:100 in paraffin oil. For both odor combinations, 10 µL odor were loaded into

custom-made Teflon containers (4.5 mm diameter) with perforated lids (Scherer et al 2003).

A minimum of 30 early third instar larvae were collected and exposed to a first odor (amyl

acetate or hexyl acetate) while crawling on pure agarose for five minutes, followed by five

minutes exposition to a second odor (benzaldehyde) on agarose medium with either 1.5 M

695   sodium chloride, 10 mM quinine as negative reinforcer or 2 M D-Fructose as positive

reinforcer (Odor1/Odor2+). For three cycle training, two more repetitions of training trials

were performed. A second group of larvae received reciprocal training (Odor1+/Odor2).

Please note that occurrence of odor and reinforcer during training have been randomized to

exclude sequential or positional effects. After one or three training cycles, larvae were

700   transferred onto test plates containing either pure agarose (reward learning) or agarose plus

salt or quinine (punishment learning) on which odor 1 and odor 2 were represented on

opposite sides. After five minutes, larvae were counted on each side of the test plate

(#Odor1, #Odor2) or in a 10 mm neutral zone in the middle of the plate (#Neutral). A

Preference index (PREF) for each group of larvae was calculated as follows:

$$PREF_{Odor1/Odor2+} = \frac{\#Odor2 - \#Odor1}{\#Total}$$

$$PREF_{Odor1+/Odor2} = \frac{\#Odor1 - \#Odor2}{\#Total}$$

To measure specifically the effect of associative learning, we then calculated the associative performance index (PI) as follows:

$$PI = \frac{PREF_{Odor1/Odor2+} + PREF_{Odor1+/Odor2}}{2}$$

710   Negative Performance Indices represent aversive associated learning, whereas positive values represent appetitive learning. Division by 2 ensures scores are bound within (-1; 1).

Odor preference test

Pure agarose (2.5% w/v) plates were cast as previously described (Apostolopoulou et al 2013, Gerber et al 2013, Michels et al 2017, Scherer et al 2003, Widmann et al 2018).
715   Furthermore, the odors were prepared according to the dilutions used in the learning experiments. An odor container (either AM 1:250, BA undiluted, HA 1:100, or BA 1:100) was placed on one side of the agarose plate. An empty container was placed on the opposite side to exclude visual or other side effects. Approximately 30 third instar feeding stage
720   larvae were collected and placed on the plate for 5 min. After this time, larvae were counted, subdivided into larvae on the odor side (#Odor), larvae on the empty container side (#Empty) and larvae in a 10 mm neutral zone in the middle of the plate, as well as larvae that were on the lid of the Petri dish (#Neutral). Preferences were calculated as follows:

$$PREF_{Odor/Empty} = \frac{\#Odor - \#Empty}{\#Total}$$

Positive values indicate a preference for the respective odor, while negative values indicate
725   odor avoidance. The preference tests were performed in different orientations (odor container on plate up/down/left/right) to exclude directional bias.

Gustatory preference test

High salt-dependent choice behavior experiments were performed using standard methods
730　(Hendel et al 2005, Huser et al 2017, Huser et al 2012, Niewalda et al 2008, Selcho et al
2009, Widmann et al 2016). A 2.5% (w/v) agarose solution was boiled as previously
described and a thick layer was poured into Petri dishes. After cooling, half of the agarose in
the Petri dish was removed and filled by 2.5% (w/v) agarose solution with 1.5 M sodium
chloride. For the choice experiment, about 30 third instar feeding stage larvae were put in
735　the middle of the Petri dish and let them crawl for five minutes. Larvae were then counted as
being located on the sodium chloride side (#NaCl), pure agarose side (#Agarose), or a
neutral area of about 10 mm width in the middle of the plate (#Neutral), as well as larvae that
were on the lid of the Petri dish (#Neutral). The gustatory preference indices for sodium
chloride were calculated as follows:

$$PREF_{NaCl} = \frac{\#NaCl - \#Agarose}{\#Total}$$

740　Negatives values for preference indices indicate an aversion to sodium chloride. The
gustatory preference tests were performed in different orientations (NaCl solution on plate
up/down/left/right) to exclude directional bias.

Optogenetic substitution experiment

745　To substitute an actual salt punishment by remotely activating DANs, we used UAS-
ChR2^{XXL}. Effector lines were crossed to w^{1118} to obtain appropriate genetic controls. Larvae
were maintained in vials with *Drosophila* standard food on 25°C and wrapped in aluminum
foil to ensure development in constant darkness. A group of 30 feeding-stage third-instar
larvae were placed onto plates containing 2.5% agarose and exposed to either AM or BA.
750　During the presentation of the first odor, the larvae were exposed to blue light (470 nm, 220

lux) for 5 minutes. The second odor BA (undiluted) was presented in darkness for five minutes on a pure agarose plate. As described for odor-high salt learning, training was performed reciprocally and the sequence of training trials was alternated across repetitions of the experiment. Furthermore, a control group, identical to the experimental group, underwent no blue light stimulation during the learning paradigm. Data were then scored as above. Please note, tests were done in the presence of 1.5 M sodium chloride to test for the specific quality of the memory. The activation of blue light itself does not cause any disruptive behavioral side effects, as it does not induce an appetitive or aversive memory on its own.

Optogenetic inhibition of neuronal activity

To acutely block synaptic output, we used UAS-GtACR2. Flies were maintained on standard food supplemented with 0.5 mM all-*trans*-retinal (ATR, Sigma Aldrich, cat. no. R2500) at 25°C as described before (Meloni et al 2020). Vials were wrapped in aluminum foil to ensure larval development in constant darkness. Groups of about 30 third instar feeding stage larvae received a reciprocal two-odor training as described before, exposed to blue light (470 nm, 1100 lux) during the entire training phase. The sequence of training trials was alternated across repetitions of the experiment. The test was performed for 5 minutes in darkness on an agarose plate mixed with 1.5 M sodium chloride. Control experiments were performed with the same genotype but with standard food lacking 0.5 mM all-*trans*-retinal. Please note that the blue light exposure itself does not cause any disruptive behavioral side effects, as it does not impair odor-high salt learning of control animals. Data were then calculated and scored as mentioned above.

Evaluation of DAN input wiring diagrams

Pie charts of sensory profiles were calculated using the percentage of total synaptic input of interneurons as fraction (thereby ignoring other inputs to show distribution of sensory origins). Percentages then give the percentage of total sensory synaptic input to interneurons. The calculation of the hub score was done in the following way: Fraction of

780   total synaptic input from all sensory neurons to defined interneurons (see IDs) was multiplied by the total fraction of input of the DANs from this interneuron. For anatomical reconstructions and visualizations included here, we made use of earlier published data (Eichler et al 2017, Eschbach et al 2020, Miroschnikow et al 2018, Winding et al 2023).

785   Statistical Analysis

Behavioral results were analyzed in GraphPad Prism 8.4.3. To test for normal distribution, Shapiro-Wilk test was applied. Groups that did not violate the assumption of normal distribution, were analyzed with an unpaired t-test (comparison of two groups) or a one-way ANOVA followed by a Tukey's post hoc test (comparisons between groups larger than two).

790   For nonparametric statistics, Mann-Whitney test (comparison between two groups) or Kruskal-Wallis followed by Dunn's multiple comparisons tests were applied. Furthermore, one sample t-test for parametric or Wilcoxon signed-rank test for nonparametric data sets were performed to compare means or medians against chance level. Results are visualized in box-plots, indicating the median as middle line, 25%/75% quantiles as box boundaries and

795   minimum/maximum performance indices as whiskers. Each data point is represented as black dot and sample sizes are noted within graphics. Asterisks and "n.s." indicate $p < 0.05$ and $p > 0.05$, respectively. The source data and results of all statistical tests are documented in the Supplemental Information file.

Supplemental Information

Supplemental Information includes Supplemental Experimental Procedures and additional figures and can be found with this article online.

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Author contributions

Conceptualization, D.W., K.V., A.M., M.P. and A.S.T; Methodology, D.W., K.V. and A.S.T;
815 Investigation, D.W., K.V., A.M., M.P. and A.S.T; Writing, D.W., K.V., A.M., M.P. and A.S.T; Supervision, D.W. and A.S.T.

Figures

820   **Figure 1: The larval dopaminergic system is subdivided in two functionally distinct**

clusters. (A) The larval dopaminergic neurons (DANs) can be anatomically subdivided into

the primary protocerebral anterior medial (pPAM) and dorsolateral 1 (DL1) cluster based on

their cell body position. **(B)** The DL1 cluster (cell bodies in green and purple) consists of four

DANs providing input to the c, d, g, and f compartment of the vertical lobe, peduncle and

825 lateral appendix of the MB (in grey). **(C)** Schematic representation of the larval MB using single letter acronyms based on (Saumweber et al 2018) to indicate compartment innervation by MB input and output neurons. DL1 DANs specifically innervate the compartments c, d, f and g in the vertical lobe, lateral appendix and peduncle, whereas pPAM DANs innervate the shaft (h) and medial lobe (I, j, k) of the MB. **(D,E)** Four DL1 DANs

830 (DAN-c1, DAN-d1, DAN-g1, and DAN-f1) are included in the expression pattern of the TH-GAL4 driver line. However, expression of a UAS-mCD8::GFP reporter via TH-GAL4 labels many more neurons (in green, anti-GFP) throughout the entire CNS (in red and blue, anti-N-cadherin and anti-discs large), in total about 100 neurons. **(F)** To test whether optogenetic activation of the DL1 DANs is sufficient to substitute for a punishment, we used the TH-

835 GAL4 driver in combination with UAS-ChR^{XXL}. Experimental and control larvae were trained by simultaneously presenting and odor and blue light and thus artificial activation of DL1 DANs, whereas a second odor was presented in darkness. Only larvae of the experimental genotype ($p > 0.05$, N=15), but not of the two genetic controls (both $p < 0.05$, for each group N=15), retrieve an aversive associative olfactory memory. The same result was seen after

840 one or three training trials (**F** and **G**, respectively). **(H, I)** To test for the acute function of the DL1 DANs in aversive associative olfactory memory, we expressed GtACR2 via the TH-GAL4 driver. Acute optogenetic inhibition of synaptic output from DL1 and other DANs reduced odor high salt memory. Experimental larvae raised on supplemented food (0.5 mM all-*trans*-retinal, ATR) and trained in blue light performed on a lower level than control

845 animals kept on standard food ($p < 0.05$, for each group N=15). A memory impairment was seen after one and three training trials (**G** and **H**, respectively). All behavioral data is shown as box-plots. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The precise sample size is given below each box-plot. n.s. $p > 0.05$; * $p < 0.05$. Scale bars:

850 in **C** 50 μm and in **D** 25 μm. The source data and results of all statistical tests are documented in the Supplemental Information file.

Figure 2: Anatomical single cell analysis of DL1 DAN specific split-GAL4 driver lines.

(A) The larval MB is organized into 11 compartments: CX calyx; IP and LP intermediate and lower peduncle; LA lateral appendix; UVL, IVL, and LVL upper, intermediate, and lower vertical lobe; SHA, UT, IT, LT shaft as well as upper, intermediate, and lower toe of the medial lobe. Single-letter synonyms of compartment names are given as "a–k". These letters are used to indicate compartment innervation by the MB input and output neurons

(Saumweber et al 2018). DL1 cluster DANs are DAN-c1, DAN-d1, DAN-g1, and DAN-f1 that innervate the respective four different compartments of the MB. **(B-J)** Individual split-Gal4 driver lines were crossed with the reporter strain UAS-mCD8::GFP;mb247-lexA,lexAop-mRFP. Third instar larval brains were dissected, fixed and mounted to visualize the fluorescent reporter signal labeling the MB (mb247-lexA,lexAop-mRFP shown in magenta) and the respective DAN pattern (GFP shown in green). (B-E) SS02160 (DAN-c1), MB328B (DAN-d1), SS02180 (DAN-f1), SS01716 (DAN-g1) each specifically label a single DL1 DAN (cell bodies are highlighted by white arrowheads). **(F–K)** Two neurons can be seen in MB065B, SS01702, and MB054B split-Gal4 that express in DAN-c1/DAN-f1, DAN-c1/MBIN-e1, and DAN-f1/DAN-g1. Please note that MB065B shows strong expression DAN-f1 but weaker staining in DAN-c1. **(H–K)** MB054B showed reliable strong expression in DAN-f1 and DAN-g1. In some brains a third weak cell body was visible right next to the other two DANs (**I**; GFP channel inverted and shown in black; **J**; cell body highlighted with gray arrowhead). Due to the low expression level we were not able to identify this cell given that only the g and f compartment of the MB were innervated **(H)**. **(K)** Analysis of the entire brain via native fluorescence expression of GFP (green) and n-Syb (magenta) did not reveal additional cells for MB054B split-Gal4. Scale bars: **(B-J)** 20 µm, **(K)** 50µm.

Figure 3: Calcium responses of DANs to gustatory stimulation. Four different split-Gal4 lines and the R58E02 driver line were crossed with UAS-GCaMP6m to express a calcium reporter in DANs. The responses of each DAN towards 100 mM NaCl, 1 M NaCl, and 500 mM fructose was tested in intact larvae using a microfluidic chip-based setup. **(A)** Calcium responses in DAN-c1 (N= 5) were induced by gustatory stimulation with 100 mM NaCl (orange), 1 M NaCl (red) and 500 mM fructose (green, for all $p < 0.05$). **(B)** DAN-d1 (N=5) calcium responses were only seen after 1 M NaCl stimulation (red, $p < 0.05$), but not after 100 mM NaCl (orange, $p > 0.05$) and 500 mM fructose stimulation (green, $p > 0.05$). **(C)** Stimulation with 100 mM (orange, $p > 0.05$) and 1 M NaCl (red, $p > 0.05$) did not induce

calcium responses in DAN-f1 (N=6). However, stimulation with 500 mM fructose reduced the obtained calcium signal (green, $p < 0.05$). **(D)** 1 M NaCl (red, $p < 0.05$), but not 100 mM NaCl (orange, $p > 0.05$) and 500 mM fructose (green, $p > 0.05$), induced a calcium response in DAN-g1 (N=7). **(E)** pPAM DANs calcium responses (N=5) were only seen after 500 mM fructose stimulation (green, $p < 0.05$). Stimulation with low (orange) and high salt concentrations (red) did not increase calcium signals, however both reduced pPAM activity ($p < 0.05$). Each graph shows the mean calcium signal plotted as the relative response strength $\Delta F/F$ and the related standard error of the mean on the y axis. The time in seconds is given below each graph on the x axis. The grey box indicates the duration of the stimulus application. The sample size of each group (N=5-7) is given above each row. n.s. $p > 0.05$; * $p < 0.05$. The source data and results of all statistical tests are documented in the Supplemental Information file.

Figure 4: Ablation of DAN-f1 and DAN-g1 together impairs aversive olfactory memory.

In all panels associative performance indices are shown for tests immediately after classical

905   conditioning. In the upper panels **(A-C)** larvae are trained once by pairing an olfactory

stimulus with high salt punishment, whereas in the lower panels **(D-F)** three training cycles

were applied. Schematic overviews for both conditioning protocols are shown on the left.

The three different DL1 DAN specific split-Gal4 driver MB054B, MB065B, and SS01702 that

each label two neurons were crossed to the effector UAS-hid,rpr to induce apoptosis **(A-F)**.

910   **(A)** With MB054B used as driver strain to ablate the DL1 DAN combination DAN-f1/DAN-g1,

the aversive associative performance index of the experimental group was decreased

compared to both controls ($p < 0.05$, N=15 for each group). **(B-F)** In all other experiments

ablation of different DL1 DAN combinations did not reveal a phenotype. (in all experiments at

least one or even both control groups are compared to the experimental group $p > 0.05$, for

915   each group N=15-25). Please note, that this also includes MB054B crossed with UAS-hid,rpr

tested after three training trials **(D)**. **(G)** To verify the memory phenotype of MB054B crossed

with UAS-hid,rpr tested after one trail conditioning, we repeated the experiment using the

odor pair hexyl acetate (HA) and benzaldehyde (BA). Again, experimental larvae tested after

one trial learning showed a robust decrease in odor-high salt memory when compared to

920   both genetic control groups ($p < 0.05$, for each group N=15). The memory phenotype was

not seen after three training trials (**J**, $p > 0.05$, for each group N=15). **(H, K)** With MB054B

used as driver strain to ablate the DL1 DAN combination DAN-f1/DAN-g1, aversive odor-

quinine memory was not impaired after one or three cycle conditioning ($p > 0.05$ when

comparing experimental and control groups, for each group N=15). **(I, L)** Similarly, appetitive

925   odor-fructose learning after one and three cycle conditioning was not impaired when ablating

DAN-f1/DAN-g1 ($p > 0.05$ when comparing experimental and control groups, for each group

N=15). All behavioral data is shown as box-plots. Please note that the same data set for the

effector control was plotted in subfigures E and F as respective behavioral experiments have

been conducted in parallel. Differences between groups are highlighted by horizontal lines

930   between them. Performance indices different from random distribution are indicated below

each box-plot. The precise sample size of each group is given below each box-plot. n.s. p > 0.05; * p < 0.05. The source data and results of all statistical tests are documented in the Supplemental Information file.

Figure 5: Optogenetic DL1 DAN activity can substitute for salt punishment. In all panels associative performance indices are shown for tests immediately after classical conditioning. In panels **(A-E)** larvae are trained once by pairing an olfactory stimulus with artificial blue light activation, whereas in panel **(F)** three training cycles were applied.

940 Schematic overviews for both conditioning protocols are shown to the left of **(A)** and **(F)**. **(A-D)** To test whether optogenetic activation of the individual DL DANs DAN-c1, DAN-d1, DAN-f1, and DAN-g1 is sufficient to substitute for a punishment, we used the split-Gal4 lines SS02160, MB328B, SS02180, and SS01716 in combination with UAS-ChR2^{XXL}. **(E, F)** For simultaneous optogenetic activation of DAN-f1/DAN-g1 we used MB054B. Larvae of the

945 experimental genotypes for DAN-c1, DAN-f1, DAN-g1 and the DAN-f1/DAN-g1 combination (for all p < 0.05, for each group N=15), but not for DAN-d1 (N=15) and all genetic controls (for all p > 0.05, for each group N=15), showed an aversive associative memory. The results imply that in the tested conditions, the punishment signal can be mediated by the artificial activation of all individual DL1 DANs, with the exception of DAN-d1. All behavioral data is

950 shown as box-plots. Please note that the same data set for the effector control was plotted in

subfigures B-E as respective behavioral experiments have been conducted in parallel. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The sample size of each group (N=15) is given below each box-plot. n.s. p > 0.05; * p < 0.05. The source

955   data and results of all statistical tests are documented in the Supplemental Information file.

Figure 6: Optogenetic inhibition of DL1 DAN activity impairs aversive olfactory memory. In all panels associative performance indices are shown for tests immediately after

960 classical odor high salt conditioning. In panels **(A-E)** larvae are trained once by pairing an olfactory stimulus with an aversive high salt stimulation, whereas in panel **(F)** three training cycles were applied. Schematic overviews for both conditioning protocols are shown to the left of **(A)** and **(F)**. **(A-D)** To test whether optogenetic inhibition of the individual DL DANs DAN-c1, DAN-d1, DAN-f1, and DAN-g1 during training impairs punishment signaling, we

965 used the split-Gal4 lines SS02160, MB328B, SS02180, and SS01716 in combination with UAS-GtACR2 and blue light stimulation during the entire training phase. **(E, F)** For simultaneous optogenetic inhibition of DAN-f1/DAN-g1 we used MB054B. **(A-C)** Larvae with inhibited DAN-c1, DAN-d1, or DAN-f1 function during training showed no impairment of odor-high salt memory comparable to controls that were kept on standard food without

970 supplemented all-*trans*-retinal (ATR, 0.5 mM) and received the same protocol (for all p > 0.05, N=15 for each group). **(D-F)** In contrast inhibition of DAN-g1 alone, or the combination of DAN-f1/DAN-g1 after single trial and three trial conditioning impaired odor-high salt

memory compared to controls (for all $p < 0.05$, for each group N=15). This shows that DAN-g1 function is of central importance for signaling a salt punishment teaching signal. All

975   behavioral data is shown as box-plots. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The precise sample size of each group is given below each box-plot. n.s. $p > 0.05$; * $p < 0.05$. The source data and results of all statistical tests are documented in the Supplemental Information file.

Figure 7: Interneurons and hub analysis of sensory to DAN pathways. (A, B) Schematic graph representation of individual DL1 **(A)** and pPAM **(B)** DANs. The outer ring at the bottom of each scheme represents the sensory composition of neurons targeting sensory-to-DANs interneurons (orange circle). The type of sensory information is encoded by the respective color (ACa: anterior central sensory compartment, AVa: ventral anterior sensory compartment, ACal: lateral anterior central sensory compartment, ACp: posterior anterior central sensory compartment, ACpl: posterior-lateral anterior central sensory compartment, VM: ventromedial sensory compartment, TD CO_2: tracheal dendritic neurons responding to CO_2, ORNs: olfactory receptor neurons). DAN input neurons are shown that get no direct sensory input (grey circles). Individual DANs are shown in the middle of the scheme as black circles. They are connected to MB KC (purple circles), which in turn connect to mushroom

body output neuron (MBONs; dark red circles). Arrows indicate the direction of the synaptic

995   connection and its strength (coded by arrow thickness). Numbers in circles indicate number

of neurons. The percentages indicate the proportion relative to the total input that the cell

receives from the specific neuronal partners. For example, DAN-f1 receives 17% of its input

from six different sensory-to-DAN interneurons (yellow circle), 17% from 19 other, non-

sensory interneurons, 41% from 64 MB KCs, and 2% from two MBONs. **(C)** Dot plot showing

1000   the importance of interneurons acting as sensory to DAN hub. Dot size was calculated using

the fraction of total input an interneuron receives from sensory neurons multiplied by the

fraction of total input this interneuron gives to an DAN. Colored backgrounds of dots are

highlighted in orange for the connections with a hub size of 0.001 or above. **(D, E)**

Schematic of graph representation. The outer ring represents the sensory composition of

1005   neurons targeting upstream neurons of DANs. The type of sensory information is encoded

by the respective color. Synaptic threshold for upstream interneurons of DANs = 3 and of

upstream sensory neurons = 1. Line thickness to interneurons and targets represents the

percentage of synaptic input. White or orange circles connected to the outer ring represent

the interneuron layer. The inner ring represents individual target neurons, DL1 and pPAM

1010   DANs. The identity of each DAN and interneuron is given by the label in its related circle. **(F)**

EM reconstruction of DL1 and pPAM DANs (grey) highlighting their presynaptic sites in red

(for DL1 DANs) and green (for pPAM DANs). At the top a horizontal view of the brain is

shown. At the bottom a frontal view of the brain is shown.

 Table 1: A summary of the characteristics of the individual DL1 DANs.

		DAN-c1	DAN-d1	DAN-f1	DAN-g1
Behavior	DAN ablation	No	No	No	No
	DAN substitution	No	No	Yes	Yes
	DAN inhibition	No	No	No	Yes
Physiology		NaCl↑ Fructose↑	NaCl↑	Fructose↓	NaCl↑
Connectomics		# of sensory-to-DAN interneurons			

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Supplemental Data

Figure 1 – Figure Supplement 1: The dopaminergic pPAM cluster encodes for a teaching signal in larvae. (A, B) The pPAM cluster (cell bodies in blue) consists of four DANs providing input to the h, i, j, and k compartments of the medial lobe of the mushroom body (MB; in grey) – the larval memory center. **(C)** Schematic representation of the larval MB using single letter acronyms based on (Saumweber et al 2018) to indicate compartment

innervation by MB input and output neurons. DL1 DANs specifically innervate the compartments c, d, f and g in the vertical lobe, lateral appendix and peduncle, whereas pPAM DANs innervate the shaft (h) and medial lobe (I, j, k) of the MB. **(D, E)** Three pPAM DANs (DAN-h1, DAN-i1, and DAN-j1) are specifically labeled via the R58E02 driver line. Expression of a UAS-GFP reporter via R58E02 labels only these three DANs (in green, anti-GFP) throughout the entire CNS (in red and blue, anti-N-cadherin and anti-discs large) besides one additional cell that does not innervate the MB (asterisk in **E**). **(F)** To test whether optogenetic activation of the pPAM DANs is sufficient to substitute for a reward, we used the R58E02 driver in combination with UAS-ChR^{XXL} to express a modified form of Channelrhodopsin2. Experimental and effector control larvae were trained via presenting and odor stimulus together with blue light stimulation and thus artificial activation of pPAM DANs, whereas a second odor was presented in darkness. Furthermore, a genetic control group received the same training protocol without blue light stimulation. Only larvae of the experimental genotype ($p > 0.05$), but not of the two genetic controls (both $p < 0.05$, for each group N=15), retrieve an appetitive associative olfactory memory. This shows that activation of the pPAM DANs is sufficient to mediate a reward teaching signal. The same effect was seen applying one or three training trials (**F and G**, respectively). **(H, I)** To test for the acute function of the pPAM DANs in aversive associative olfactory memory, we expressed GtACR2, an anion-conducting, light-gated channel via the R58E02 driver. Acute optogenetic inhibition of synaptic output from pPAM DANs during training did not change odor high-salt memory. Experimental larvae kept on supplemented food with 0.5 mM all-*trans*-retinal (ATR) or genetic controls, maintained on standard food performed at the same level ($p > 0.05$, for each group N=15). Intact odor high-salt memory was seen after one or three training trials (**H** and **I**, respectively). All behavioral data is shown as box-plots. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The precise sample size of each group is given below each box-plot. n.s. $p > 0.05$; * $p < 0.05$. Scale bars: **(D)** 50 µm, **(E)** 25 µm. The

 source data and results of all statistical tests are documented in the Supplemental

Information file.

Figure 2 – Figure Supplement 1: Ablation of individual DL DANs does not impair aversive olfactory memory reinforced by high salt. (A) The larval MB is organized into 11 compartments. Letters from a-k are used to indicate compartment innervation by the MB input and output neurons. **(B, C)** Individual split-Gal4 driver lines were crossed with the reporter strain UAS-mCD8::GFP;mb247-lexA,lexAop-mRFP. Third instar larval brains were dissected, fixed and mounted to visualize the fluorescent reporter signal labeling the MB (mb247-lexA,lexAop-mRFP shown in magenta) and the respective DAN pattern (GFP shown

in green). **(B)** MB143B labels specifically DAN-d1. **(C)** MB145B labels DAN-f1 and very weakly several additional cells. **(D, E)** Aversive associative performance indices are shown for tests immediately after classical odor-high salt conditioning using one training trial. A schematic overview of the conditioning protocol is shown at the left. Neither DAN-d1 nor DAN-f1 function is required. Even when tested using two alternative split-Gal4 lines MB143B and MB145B. **(F-G)** The same result can also be seen after three cycle odor-high salt conditioning. A schematic overview of the conditioning protocol is shown at the left. In summary, in all four behavioral experiments the aversive associative performance indices of the experimental groups (N=15-20) were not decreased compared to the two (in all cases $p > 0.05$, N=15-25) or at least one genetic control group (in **F**, $p < 0.05$ comparing UAS control and experimental larvae). All behavioral data is shown as box-plots. Please note that the same data sets for the effector control were plotted in subfigures D and E (one cycle) and F and G (three cycle), as respective behavioral experiments have been conducted in parallel. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The precise sample size of each group is given below each box-plot. n.s. $p > 0.05$; * $p < 0.05$. The source data and results of all statistical tests are documented in the Supplemental Information file.

1375 **Figure 2 – Figure Supplement 2: Split-Gal4 MB054B covers three dopaminergic neurons in the DL1 cluster. (A-F)** Split-Gal4 MB054B driver strain was crossed to UAS-mCD8::GFP effector strain. **(A)** A z projection of a larval central brain is shown using anti-GFP to label expression pattern of Split-Gal4 MB054B (cyan), anti-TH to label TH-positive cells (yellow) and anti-N-cadherin to visualize the neuropil (magenta). The DL1 cluster of the

1380 left hemisphere is highlighted. **(B,C)** Single channel visualization of all TH-positive cells **(B)** and GFP expressing cells **(C)** of the crossed Split-Gal4 MB054B driver line to effector UAS-mCD8::GFP within the larval central brain. **(D)** A detailed z projection of the DL1 cluster is shown using anti-GFP to label expression pattern of Split-Gal4 MB054B (cyan) and anti-TH to label TH-positive cells (yellow). Co-labeling of anti-TH and anti-GFP is highlighted by

1385 arrowheads. **(E,F)** Single channel visualization of all TH-positive cells **(E)** and GFP expressing cells **(F)** of the Split-Gal4 MB054B driver line crossed to effector UAS-mCD8::GFP within the DL1 cluster. Scale bars: **(A-C)** 50 µm, **(D-F)** 10 µm.

1390 **Figure 4 – Figure Supplement 1: Ablation of individual DANs does not impair aversive olfactory memory.** In all panels aversive associative performance indices are shown for tests immediately after odor-high salt classical conditioning. In the upper panels **(A-D)** larvae

are trained once, whereas in the lower panels **(E-H)** three training cycles were applied. Schematic overviews for both conditioning protocols are shown on the left. The four different DL1 DAN specific split-Gal4 driver strains SS02160, MB328B, SS02180, and SS01716 were crossed to the effector UAS-hid,rpr to induce apoptosis **(A-H)**. Cell-specific ablation of distinct DL1 DANs did not impair odor-high salt memories after either one or three cycle training. In no case were associative performance indices upon DL1 DAN ablation lower than in both genetic controls (**A-H**, in all experiments at least one or even both control groups are compared to the experimental group p > 0.05, for each group N= 15-20 for one cycle training, and N= 15-25 for three cycle training). All behavioral data is shown as box-plots. As the respective behavioral experiments in subfigures A-D and E-H were conducted in parallel, the same data sets for the corresponding effector control were plotted. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The exact sample size of each group is given below each box-plot. n.s. p > 0.05; * p < 0.05. The source data and results of all statistical tests are documented in Supplemental Information file.

1410 **Figure 4 - Figure Supplement 2: Ablation of individual DL DANs does not impair aversive olfactory memory reinforced by bitter quinine.** In all panels aversive associative performance indices are shown for tests immediately after classical odor-quinine conditioning using one training trial. A schematic overview of the conditioning protocol is shown to the left of **(A)**. **(A-C)** The three different DL1 DAN split-Gal4 driver MB328B,

1415 SS02180, and SS01716 that specifically label DAN-d1, DAN-f1, and DAN-g1, respectively, are crossed with the effector UAS-hid,rpr to induce apoptosis. In all three experiments the aversive associative performance indices of the experimental groups (N= 15) were not decreased compared to the two control groups (in all cases $p > 0.05$, for each group N=15). All behavioral data is shown as box-plots. Please note that the same respective data set for

1420 the effector control was plotted in all subfigures, as these behavioral experiments were conducted in parallel. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The exact sample size of each group is given below each box-plot. n.s. $p > 0.05$; * $p < 0.05$. The source data and results of all statistical tests are documented in

1425 Supplemental Information file.

 Figure 4 – Figure Supplement 3: Ablation of individual DL DANs does not impair appetitive olfactory memory. In all panels appetitive associative performance indices are shown for tests immediately after classical odor-fructose conditioning using one training trial. A schematic overview of the conditioning protocol is shown to the left of **(A)**. **(A-C)** The three different DL1 DAN split-Gal4 driver MB328B, SS02180, and SS01716 that specifically label

1435 DAN-d1, DAN-f1, and DAN-g1, respectively, are crossed with the effector UAS-hid,rpr to induce apoptosis. In all three experiments the appetitive associative performance indices of the experimental groups (N=15) were not decreased compared to the two related control groups (in all cases p > 0.05, for each group N=15). All behavioral data is shown as box-plots. Differences between groups are highlighted by horizontal lines between them.

1440 Performance indices different from random distribution are indicated below each box-plot. The precise sample size of each group is given below each box-plot. n.s. p > 0.05; * p < 0.05. The source data and results of all statistical tests are documented in the Supplemental Information file.

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Figure 4 – Figure Supplement 4: Functional analysis of mushroom body input neurons that are not dopaminergic. In panels **(A-C)** larvae are trained once by pairing an olfactory stimulus with high salt punishment, whereas in panel **(D-F)** three training cycles were applied. Schematic overviews for both conditioning protocols are shown to the left of

1450 **(A)** and **(D).** The three different mushroom body input split-Gal4 driver SS24765, SS21716, and SS01958 that specifically label OAN-a1,a2, MBIN-b1,b2, and OAN-e1 are crossed with the effector UAS-hid,rpr to induce apoptosis. Ablation of any of the three types of input neurons did not impair odor-high salt memory regardless of whether larvae were trained once or three times (**A-C** and **D-F**). In all cases the experimental groups (N=15) performed

1455 on the same level as both (**A-C** and **F**, in all cases p > 0.05, N=15-20) or at least one genetic

control (**D, E**, Gal4 control p > 0.05 and UAS-hid,rpr control p < 0.05, N=15-25). All behavioral data is shown as box-plots. As the respective behavioral experiments in subfigures A-C for one cycle and D and F for three cycles were conducted in parallel, the same data sets for the corresponding effector control were plotted. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The precise sample size of each group is given below each box-plot. n.s. p > 0.05; * p < 0.05. **(G-I)** Individual split-Gal4 driver lines were crossed with the reporter strain UAS-mCD8::GFP;mb247-lexA,lexAop-mRFP. Third instar larval brains were dissected, fixed and mounted to visualize the fluorescent reporter signal labeling the MB (mb247-lexA,lexAop-mRFP shown in magenta) and the respective neuronal pattern (GFP shown in green). SS24765 (OAN-a1,a2), SS21716 (MBIN-b1,b2), and SS01958 (OAN-e1) each specifically label a specific input neuron pair **(G, H)** or a single input neuron **(I)**. Scale bars: 20 μm. The source data and results of all statistical tests are documented in the Supplemental Information file.

Figure 4 – Figure Supplement 5: Sensory acuity tests. In panels **(A-E)** naïve larval preferences towards different chemosensory stimuli are shown. Olfactory preference of MB054B split Gal4 crossed to UAS-hid,rpr to ablate DAN-f1/DAN-g1 are shown in **(A-D)**. For all the tested odor stimuli, amyl acetate diluted 1:250 in paraffin oil, pure benzaldehyde, hexyl acetate diluted 1:100 in paraffin oil, and benzaldehyde diluted 1:100 in paraffin oil, no difference between the experimental group (N=15) and the two genetic control groups was seen (for all cases p > 0.05, for each group N=15). **(E)** Gustatory high salt preference of MB054B split-Gal4 crossed to UAS-hid,rpr to ablate DAN-f1/DAN-g1 (N=15) was not reduced but rather increased in their avoidance of 1.5 M high salt concentration compared to both genetic controls (both p < 0.05, for each group N=15). All behavioral data is shown as box-plots. Differences between groups are highlighted by horizontal lines between them. Performance indices different from random distribution are indicated below each box-plot. The precise sample size of each group is given below each box-plot. n.s. p > 0.05; * p < 0.05. The source data and results of all statistical tests are documented in the Supplemental Information file.

Figure 4 – Figure Supplement 6: Anatomical validation of effector UAS-hid,rpr

1490 **functionality to ablate neurons of MB054B Split-Gal4 in the DL1 cluster.** To confirm the

functionality of the effector UAS-hid,rpr, anatomical examination of the single effector line

UAS-hid,rpr **(A-C''')**, the Split-Gal4 driver line MB054B **(D-F''')**, as well as the cross of both

(G-I''') were examined. **(A, D, G)** A z projection of a larval central brain of the respective

strain or cross is shown using anti-TH to label TH positive cells (yellow) and anti-N-cadherin

1495 to visualize the neuropil (magenta). **(B, E, H)** Single channel visualization of all TH-positive

cells (black) of the respective larval brains shown in **(A, D, G)**. The DL1 cluster is highlighted

by violet dashed lines. **(C)** Detailed z projection of the left hemispheric DL1 cluster of effector

line UAS-hid,rpr is shown as single channel (anti-TH, black). Cell bodies of the DL1 cluster

neurons are highlighted by violet asterisk. **(C'-C''')** Partial projection of the z-stack shown in

1500 **(C)** to distinguish single cell bodies. Anatomical analysis revealed seven TH-positive cell

bodies in the DL1 cluster of UAS-hid,rpr, highlighted by violet arrowheads. **(F)** Detailed z

projection of the left hemispheric DL1 cluster of Split-Gal4 line MB054B is shown as single channel (anti-TH, black). Cell bodies of the DL1 cluster neurons are highlighted by violet asterisk. **(F'-F''')** Partial projection of the z-stack shown in **(F)** to distinguish single cell

1505 bodies. Anatomical analysis revealed seven TH-positive cell bodies in the DL1 cluster of MB054B, highlighted by violet arrowheads. **(I)** Detailed z projection of the left hemispheric DL1 cluster of the crossed Split-Gal4 driver strain MB054B to UAS-hid,rpr is shown as single channel (anti-TH, black). Cell bodies of the DL1 cluster neurons are highlighted by violet asterisk. **(I'-I''')** Partial projection of the z-stack shown in **(I)** to distinguish single cell bodies.

1510 Crossing Split-Gal4 MB054B to effector line UAS-hid, rpr led to ablation of the respective MB054B cell bodies (see also Figure 2H-K), leaving only four TH-positive cell bodies (highlighted by violet arrowheads) in the DL1 cluster. Scale bars: 50 µm **(A,B, D,E, G,H)** and 20 µm **(C-C''', F-F''', I-I''')**.

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Figure 7 – Figure Supplement 1: A sensory neuron to DAN map based on EM connectivity. (A) Adjacency matrix of all neurons used in this study (no synaptic threshold)

showing sensory compartments (magenta) to interneuron (yellow) to DL1 (red) and pPAM DANs (green). All synapses between shown neuron types were included. Top: Nine different sensory origins synapse onto upstream neurons (interneurons) of the DANs. Interneuron ID and neuron name is listed on the left and right of the matrix. From top to bottom the connectivity of 35 interneurons are shown that establish the complete single hop circuit between sensory neurons and DANs. Please note that most of the interneurons form pairs (15 paired and 5 unpaired interneurons). **(B)** Schematic of all sensory neurons that innervate the DL1 cluster (left) and the pPAM cluster (right). The top shows a frontal view, at the bottom a view from above along the longitudinal axis is given. The sensory neurons upstream of DANs are color coded based sensory origin, which is explained at the bottom of the figure. **(C)** Comparison of the similarity of DL1 and pPAM sensory input. Most of the sensory input is either specific for the DL1 cluster (57 neurons, magenta) or the pPAM cluster (61 neuron, green). Only a small number of sensory neurons (7 neurons, black) instructs both clusters. **(D)** Frontal and sagittal views are shown from top to bottom that highlight the distinct sets of interneurons connecting DL1 DANs (4 neurons, red) or pPAM DANs (16 neurons, green) with the sensory neurons.

 Supplemental Table 1:

Experimental models: organisms/strains			
Drosophila: w¹¹¹⁸	Bloomington Drosophila Stock Center	RRID: BDSC: 3605	N/A
***Drosophila:* UAS-mCD8::GFP**	Bloomington Drosophila Stock Center	RRID: BDSC: 32194	Pfeiffer et al. 2010
***Drosophila:* UAS-mCD8::GFP; nSyb-LexA, IOP-mRFP/TM6B**	Wolf Hütteroth	N/A	combined from Burke et al. 2012 and Riabinina et al. 2015
***Drosophila:* UAS-mCD8::GFP; mb247-LexA, IOP-mRFP/TM3,Sb**	Wolf Hütteroth	N/A	Burke et al. 2012
***Drosophila:* UAS-hid,rpr**	Sprecher lab	N/A	Abbott and Lengyel 1991; White et al. 1996
***Drosophila:* UAS-GtACR2**	Bloomington Drosophila Stock Center	RRID: BDSC: 92984	Mohammad et al. 2017
***Drosophila:* UAS-ChR2^{XXL}**	Bloomington Drosophila Stock Center	RRID: BDSC: 58374	Dawydow et al. 2014
***Drosophila:* UAS-GCaMP6m**	Bloomington Drosophila Stock Center	RRID: BDSC: 42748	Chen et al. 2013
***Drosophila:* TH-Gal4**	Bloomington Drosophila Stock Center	RRID: BDSC: 95268	Friggi-Grelin et al. 2003
***Drosophila:* R58E02**	Bloomington Drosophila Stock Center	RRID: BDSC: 41347	Jenett et al. 2012
***Drosophila:* MB054B**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* MB065B**	Bloomington Drosophila Stock Center	RRID: BDSC: 68281	Eschbach et al. 2020
***Drosophila:* MB143B**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* MB145B**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* MB328B**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* SS01702**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* SS01716**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* SS01958**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* SS02160**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* SS02180**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* SS21716**	Zlatic lab	N/A	Eschbach et al. 2020
***Drosophila:* SS24765**	Zlatic lab	N/A	Eschbach et al. 2020

Chemicals, antibodies, and recombinant proteins			
Phosphate Buffered Saline	Sigma-Aldrich	N/A	Cat. no. P4417
Triton X-100	Sigma-Aldrich	CAS: 9002-93-1	Cat. no. X100
4% Formaldehyde	Thermo Scientific	CAS: 50-00-0	Cat. no. 047392.9M
Normal goat serum	Sigma-Aldrich	N/A	Cat. no. G9023
Xylene	Sigma-Aldrich	CAS:1330-20-7	Cat. no. 247642
dibutyl phthalate in xylene	Sigma-Aldrich	CAS: 84-74-2	Cat. no. 06522
NaCl	VWR Chemicals	CAS: 7647-14-5	Cat. no. 27810.364
D-Fructose	Sigma-Aldrich	CAS: 57-48-7	Cat. no. 47740
Agarose	Sigma-Aldrich	CAS: 9012-36-6	Cat. no. A9539
Quinine	Sigma-Aldrich	CAS: 207671-44-1	Cat. no. Q1250
Amyl acetate	Sigma-Aldrich	CAS: 628-63-7	Cat. no. 46022
Paraffin oil	Sigma-Aldrich	CAS: 8012-95-1	Cat. no. 76235
Hexyl acetate	Sigma-Aldrich	CAS: 142-92-7	Cat. no. 108154
Benzaldehyde	Sigma-Aldrich	CAS: 100-52-7	Cat. no. 12010
all-*trans*-retinal	Sigma-Aldrich	CAS: 116-31-4	Cat. no. R2500
rat anti-N-Cadherin	Hybridoma	RRID: AB_528121	Cat. no. DN-Ex #8-s
rabbit anti-GFP	Life Technologies	RRID: AB_221570	Cat. no. A6455
mouse 4F3 anti-DLG	Hybridoma	RRID: AB_528203	Cat. no. 4F3 anti-discs large
goat anti-rat Alexa Fluor 647	Life Technologies	RRID: AB_141778	Cat. no. A21247
goat anti-rabbit Alexa Flour 488	Life Technologies	RRID: AB_143165	Cat. no. A11008
goat anti-mouse Alexa Fluor 568	Life Technologies	RRID: AB_2535804	Cat. no. A21235
mouse anti-TH	Immunostar	RRID: AB_572268	Cat. no. 22941
goat anti-mouse Cy3	Life Technologies	RRID: AB_10373848	Cat. no. A10521
Software and algorithms			
Fiji version 1.53c (64-bit)	NIH		https://fiji.sc/
GraphPad Prism 8.4.3	GraphPad Software, La Jolla, CA		https://www.graphpad.com/scientific-software/prism/
Adobe Photoshop	Adobe Systems, San Jose, CA		https://www.adobe.com/de/products/photoshop.html
Affinity Publisher 2.1.1	Serif (Europe) Ltd., Nottingham, UK		https://affinity.serif.com/de/publisher/
ZEN 2.3 software	Carl Zeiss Microsocopy Germany GmbH		https://www.zeiss.de/mikroskopie/produkte/mikroskopsoftware/zen-lite/zen-lite-download.html