Prediction error drives associative olfactory learning and conditioned behavior in a spiking model of *Drosophila larva* 2 Anna-Maria Jürgensen^a, Panagiotis Sakagiannis^a, Michael Schlever^{b,c}, Bertram Gerber^{b,d,e}, and Martin Paul Nawrot^a ^aComputational Systems Neuroscience, Institute of Zoology, University of Cologne, Cologne, Germany ^bLeibniz Institute for Neurobiology (LIN), Department of Genetics, Magdeburg, Germany ^cInstitute for the Advancement of Higher Education, Faculty of Science, Hokkaido q University, Japan 10 ^dInstitute for Biology, Otto-von-Guericke University, Magdeburg, Germany 11 ^eCenter for Brain and Behavioral Sciences (CBBS), Otto-von-Guericke University, 12 Magdeburg, Germany 13

Abstract

14

Predicting reinforcement from the presence of environmental clues is an essential component 15 of guiding goal-directed behavior. In insect brains, the mushroom body is central to learning 16 the necessary associations between sensory signals and reinforcement. We propose a biologically 17 realistic spiking network model of the Drosophila larva olfactory pathway for the association of 18 odors and reinforcement to bias behavior towards approach or avoidance. We demonstrate that 19 prediction error coding through the integration of currently present and expected reinforcement 20 in dopaminergic neurons can serve as a driving force in learning that can, combined with a 21 synaptic homeostasis mechanism, account for experimentally observed features of acquisition 22 and loss of associations in the larva that depend on the intensity of odor and reinforcement and 23 temporal features of their pairing. To allow direct comparisons of our simulations with behav-24 ioral data [1], we model learning-induced plasticity over the complete time course of behavioral 25 experiments and simulate the locomotion of individual larvae towards or away from odor sources 26 in a virtual environment. 27

28 Introduction

Goal-directed behavior in dynamic situations benefits from the ability to predict future conditions in the environment from the occurrence of sensory clues. In insects, the mushroom body (MB) is the central brain structure for multi-sensory integration, involved in memory formation and recall [2, 3]. It is at the core of learning and retaining valuable associations between sensory inputs and reinforcement in the synapses between the MB intrinsic and its output neurons [4–7].

One of the underlying mechanisms is associative learning, a process that gradually establishes a relationship between two previously unrelated elements. In classical conditioning, the conditioned sensory stimulus (CS) obtains behavioral relevance through its concurrence with the reinforcing unconditioned stimulus (US), an acquisition process depending dynamically on their spatiotemporal proximity. The temporal evolution of this process has been formalized in the Rescorla-Wagner (RW) model (eqn. 1) [8].

$$\Delta V = \alpha \cdot (\lambda_{\rm US} - V(t)),$$

$$V(t + \Delta t) = V(t) + \Delta V.$$
(1)

Here, a CS obtains predictive power of concurrent or successive US [8], that depends on the 40 strength of the already acquired association between the CS and US V(t), allowing for anticipatory 41 behavior to the expected US [9, 10]. The acquisition of this association terminates when the US is 42 fully predicted. Until then, the change in associative strength ΔV is proportional to the difference 43 between the maximum associative strength (or asymptote) $\lambda_{\rm US}$ and the current associative strength 44 V(t) (eqn. 1). The maximum associative strength is a property of the US, determined mainly by 45 the intensity of the reinforcement. While the current associative strength V(t) is defined by the 46 shared learning history of CS and US [8]. The concept of prediction error (PE) [11] is a derivative 47 of the Rescorla-Wagner model [8]. The error signal equals the difference between the current $\lambda_{\rm US}$ 48 and the predicted value of US V(t). Over the course of the memory acquisition/training phase, the 49 pace of learning, which can be formalized as the slope of the acquisition curve, decreases as the PE 50 is reduced, minimizing the driving force for changes of the association [8, 11]. This difference is 51 multiplied with a learning rate parameter (α), here combined for the CS and the US (eqn. 1). 52

This continuous optimization of predictions, guided by the PE, could allow animals to efficiently adapt their goal-directed behavior in dynamic environments. Among the most relevant associations to be learned are those that enable the prediction of reward or punishment. Dopaminergic neurons (DANs) have long been known to encode information about reward and punishment. These types of neurons respond to the presence of rewards and punishments in the environment, both in vertebrates [12–17], as well as invertebrates [18–21]. The electrical stimulation or optogenetic activation of DANs induces approach or avoidance both in vertebrates [22–25] and invertebrates [20, 26–32]. In adult [5, 6, 33, 34] and larval [20, 35, 36] *Drosophila* this approach or avoidance learning is facilitated by

the modulation of MB output synapses by DAN activity. Ultimately DANs do not only signal the presence of rewards or punishments but have also been suggested to encode PE in various vertebrate species [16, 37–40] and might have a similar function in insects [19, 20, 32, 34, 41–44].

We utilize our spiking model of the Drosophila larva MB in one brain hemisphere that forms 64 associations of odors with reinforcement to further test the hypothesis that PE coding within this 65 circuit takes place in DANs that receive input from the output neurons of the MB or their down-66 stream partners [20, 45-47], that might provide feedback to the DANs. Beyond the scope of similar 67 models [20, 48–51] (see Discussion, section: Comparison with other MB models), we demonstrate 68 that this mechanism can reproduce the experimentally observed findings on the acquisition of asso-69 ciations of odors with reinforcement in a time-resolved manner [1]. To facilitate direct qualitative 70 and quantitative comparisons with animal behavioral data, we couple this model with a realistic 71 locomotory model of the larva [52] that captures the effects of learned associations on chemotactic 72 behavior in individual animals. 73

74 Results

⁷⁵ Connectome-based circuit model of the larval olfactory pathway

The network architecture of our model (fig. 1A) is based on the anatomy of the olfactory pathway 76 in one Drosophila larva brain hemisphere [20, 29, 53, 54] (for more details see Methods, section: 77 Network model). Peripheral processing is carried out by 21 olfactory receptor neurons (ORNs), each 78 expressing a different olfactory receptor type [53, 55, 56]. ORNs form one-to-one excitatory synaptic 79 connections with 21 projection neurons (PNs) and 21 local interneurons (LNs) in the antennal lobe 80 [53]. Each LN connects with all PNs via inhibitory GABAergic synapses, establishing a motif for 81 lateral inhibition within the antennal lobe. The 72 mature larval Kenyon cells (KCs) [54] are the 82 excitatory principal cells of the MB. Each KC receives excitatory input from 2-6 randomly selected 83 PNs [54]. The KCs are subjected to feedback inhibition, provided via the GABAergic anterior paired 84 lateral (APL) neuron, which receives input from all KCs [29]. Only mature KCs, characterized by 85 a fully developed dendrite, are included in this model, yielding a complete convergent synaptic KC 86 >APL connectivity. The output region of the MB is organized in compartments, in which the 87 KC axons converge with the dendrites of one or few MB output neurons (MBONs) [20, 54]. Our 88 model assumes two MBONs from two different compartments that are representative of two different 89 categories of output neurons of the MB that mediate either approach or avoidance [4–6, 33, 35, 36, 90 57–59] with a single MBON each. Both MBONs receive excitatory input from all of the KCs to 91 fully capture the information that is normally represented by the complete set of MBONs. Each 92 compartment is also innervated by a single DAN, signaling either reward or punishment and targeting 93 the KC>MBON synapses to facilitate learning (for a discussion of all simplifications compared to 94 the animal brain, see Methods, section: Network model). 95

⁹⁶ Learning through KC>MBON plasticity

We assume that the KC>MBON synapses undergo plasticity, based on strong experimental evidence 97 in larval [20, 35, 36] and adult flies [5, 6, 34]. This plasticity requires the convergence of the sensory 98 pathways in the form of KC activation and of the reinforcing pathway, mediated by neuromodu-99 latory DAN signaling at the synaptic site. We employ a two-factor learning rule (eqn. 2) at each 100 KC>MBON synapse (fig. 1 A,B). The first factor is expressed in the pre-synaptic KC activation by 101 an odor, tagging the synapse eligible for modification. This is modeled via an exponentially decay-102 ing eligibility trace $e_i(t)$, which is set to a 1 whenever the respective KC elicits a spike (fig. 1B). 103 The decay time constant determines the window of opportunity for synaptic change. The pres-104 ence of reinforcement (reward or punishment) constitutes the second factor and is signaled by the 105 reward-mediating DAN₊ or punishment-mediating DAN₋, respectively. Spiking of the DAN provides 106 a neuromodulatory reinforcement signal R(t) to the synaptic site. If a DAN spike coincides with 107 positive eligibility at the synapse, the respective synaptic weight is reduced. At each synapse i, the 108 reduction of synaptic weight Δw_i depends on the learning rate a (table S1) and is proportional to 109 the amplitude of the eligibility trace $e_i(t)$ (fig. 1 B): 110

$$\Delta w_i = -a \cdot e_i(t) \cdot R(t) \le 0. \tag{2}$$

We introduce a synaptic homeostasis mechanism (eqn. 3) that modulates the effects of plasticity 111 at each KC>MBON synapse to account for the experimentally observed loss of a learned associ-112 ation when reinforcement is omitted [41, 60, 61] and to ensure continued input to both MBONs. 113 With each MBON spike, the current weight w_i of each respective KC>MBON synapse is increased, 114 proportionally to the extent to which the weight differs from its original value w_{init} (table S1) and 115 multiplied with a homeostatic regulation factor h (table S1). This mechanism serves as an imple-116 mentation for the loss of the association when the reinforcement is omitted. While reinforcement 117 is present, the learning curve will either continue to rise or remain at the asymptote if already 118 saturated. The interaction of the two mechanisms of learning and unlearning at the level of the 119 individual KC>MBON synapses allows to include the loss of learned associations, when continued 120 reinforcement is omitted (see Discussion, section: A mechanistic implementation of the RW model) 121 and also ensures continued input to the MBONs, despite the reduction of input weights over the 122 course of the learning process (eqn. 2). The homeostatic factor h hereby serves as an implementation 123 of a time constant of this exponential process. The interaction of synaptic plasticity and homeostatic 124 regulation defines the magnitude of the weight at the next simulation timestep $t + \Delta t$ as 125

$$w_i(t + \Delta t) = w_i(t) + \Delta w_i + (w_{init} - w_i(t)) \cdot h.$$
(3)

It has been shown in behavioral experiments that specific MBONs encode a behavioral tendency to either approach or avoid a currently perceived stimulus, depending on the acquired stimulus valence [4–6, 36, 57, 58]. In the naive state of our model, all KC>MBON synapses have the same

¹²⁹ initial weights w_{init} (table S1), and hence the spiking activity of both MBONs is highly similar. ¹³⁰ Learning alters the KC>MBON synaptic weights and thus skews the previously balanced MBON ¹³¹ output. This acquired imbalance between MBON outputs biases behavior towards the approach or ¹³² avoidance of the conditioned odor. To quantify the effect of learning, we compute the behavioral ¹³³ bias BB (eqn. 4) from the firing rates of both MBONs over T = 1s as follows:

$$BB = \frac{MBON_{+} - MBON_{-}}{T}.$$
(4)

¹³⁴ Implementation of prediction error coding in the KC-MBON-DAN motif

In the larva, many DANs and other modulatory neurons receive excitatory or inhibitory input 135 from different MBONs, either in a direct manner or via one or two interneurons [20]. Based on 136 this observation, we constructed our hypothetical feedback motif (for similar models see discus-137 sion section: Comparison with other MB models). In the model, DANs are activated by external 138 reward/punishment signals and also receive excitatory and indirect inhibitory feedback from both 139 MBONs (fig. 1A). As the initial balance between the two MBON outputs shifts over the course of 140 the training process, the amount of excitatory and inhibitory feedback that DANs receive continues 141 to diverge, allowing the DANs to access the model's learning history. Ultimately DAN activation 142 signals the difference between the current external activation and the expected activation based 143 on prior learning, implemented as the difference between excitatory and inhibitory MBON>DAN 144 feedback. Including this feedback leads to learning curves that saturate when the reward is fully 145 predicted, and the prediction error approaches zero (fig. 2 A,D). This effect disappears, when the 146 feedback circuit is disabled (fig. 2 A). In this case the behavioral bias quickly reaches the maximum 147 value of the measure when the MBON₂ elicits no more spikes and can not encode further learning. 148 Increasing reward intensity, learning rate or odor intensity (see Methods, section: Experimental 149 protocols) foster a faster acquisition of the association and increases the maximum strength of the 150 association at the same time (fig. 2 A). 151

Increasing the reward intensity after a 2.5 min (black curve), or 5 min (gray curve) of appetitive training, results in a steeper slope of the learning curve and also increases the maximum during training trials of 2.5 min duration with increased reward intensity (fig. 2 B). Higher intensity of the reward results in an average DAN spike rate of 39.14Hz(std = 1.27(standard deviation)) compared to 33.11Hz(std = 1.34).

Additionally, we tested for loss of the acquired association as the reduction in behavioral memory expression, over the course of prolonged exposure to the CS without the US, following initial memory acquisition [8, 62]. We test this in our model experiments by presenting the odor, previously paired with reward, for an extended period of time, in the absence of reinforcement. During the test phase and without the presence of reward to trigger synaptic KC>MBON_ weight reduction, the extinction mechanism is no longer outweighed by learning and drives each individual weight back towards w_{init} (fig. 2 C, upper panel). We also demonstrate the interaction of the learning rule with this mechanism in figure S1, where the learning rate remains constant but the magnitude of the homeostatic regulation was manipulated to show that both mechanisms need to be in balance.

¹⁶⁶ Learned preference and behavior generalize to similar odors

We trained our model by pairing a reward with a single odor for 4min. After the training procedure, 167 we tested the behavioral bias either for the same or a different odor, following the experimental 168 approach used in the larva [63]. Mimicking the experimental data, we show that the odor preference 169 is highest if training odor and testing odor are identical in the case of training with 3-octanol. When 170 amylacetate is used during training, 3-octanol preference is increased (fig. 3A). Since 3-octanol 171 activates a subset of the ORNs activated by amylacetate (fig. 1 D), some of which with higher rates 172 than in the case of amylacetate, we also tested for generalization using a set of ORN activation 173 patterns with a controlled degree of overlap (see Methods, section: Sensory input, fig. 1D) and 174 show that with decreasing similarity, the generalization effect to a new odor is diminished (fig. 3 A). 175 Figure 3 B shows the network response to 30sec stimulations with amylacetate and 3-octanol in a 176 single exemplary model instance. On the level of the ORNs, 3-octanol merely activated a subset 177 of the amylacetate-activated neurons. The uniqueness of the odor identities is enhanced in the KC 178 population [64]. 179

¹⁸⁰ The model reproduces temporal features in trace conditioning experiments

Including an odor-evoked eligibility trace at the KC>MBON synapses allows the model to maintain 181 the sensory odor representation for a time window, during which reinforcement will trigger synaptic 182 change (fig. 1B). The time window between odor and reward onset (0, 10, 20, 30, 40, 50, 60, 120s)183 was varied for trace conditioning experiments with a 30s presentation of odor and reward that was 184 repeated three times. A small inter-stimulus-interval (ISI) of 10 to 30s leads to an increase in 185 behavioral bias compared to the complete overlap of odor and reinforcement (fig. 3 C), using the 186 extended window of opportunity for synaptic change triggered by each KC spike. Long ISIs do not 187 lead to learning as the eligibility trace declines back to zero during this time (fig. 3 C). These findings 188 match observations from experiments in larvae [29, 65, 66] with the caveat that the trace in the real 189 larva brain seems to extend for a slightly longer period of time, compared with our experiments. 190

¹⁹¹ The model reproduces paired and unpaired associative conditioning ex-¹⁹² periments

¹⁹³ To test if learning, driven by prediction error, can account for learned larval behavior, we replicated

¹⁹⁴ single-trial conditioning experiments performed with larvae [1] in simulation. In these experiments,

animals were trained with the odor amylacetate in a single trial of varying duration (1 - 8 min).

¹⁹⁶ To this end, larvae were placed on a Petri dish coated with an agar-sugar substrate and the odor in

two small containers for diffusion in the air (paired training). Either before or following this train-197 ing protocol, larvae underwent a single trial without sugar and odor. Afterward, the animals were 198 transferred to a new dish with two odor containers placed on different sides (one of them contained 199 amylacetate, and the other one was empty). This paired training was compared with an unpaired 200 protocol with separate (randomized order) presentations of amylacetate and sugar. Following the 201 paired training protocol (odor and reward are presented concurrently), the animals showed a ten-202 dency to approach the previously rewarded odor, as measured by the difference in the number of 203 animals on each side at the end of a 3min test phase, divided by the total number of animals. Fol-204 lowing the experimental literature, we will refer to this measure as the preference index ([1] eqn. 15). 205 The animals' preference is relatively consistent across training trials of different duration. Prolonged 206 paired training did not lead to an increase in preference (fig. 5 A). These experiments did not in-207 clude a test for odor preference before training, but untrained larval odor preference of odors used in 208 learning experiments has been demonstrated elsewhere [67-69]. This paired training was compared 209 with an unpaired protocol with separate (randomized order) presentations of amylacetate and sugar. 210 Here the extent to which animals preferred amylacetate over no odor varied with the duration of 211 the training trial. The longer the duration of the training, the more the preference index decreased 212 from an initially high value but saturated around 2.5min (fig. 5 A). 213

We aimed to replicate these behavioral experiments on two levels. Firstly, we focused on the 214 direct model output that reflects the strength of the acquired association between amylacetate 215 and reward (behavioral bias, eqn. 4) and later also simulated behavior based on these biases. We 216 simulated both the paired and unpaired training protocol (fig. 4 B). While the unpaired training 217 yielded almost no behavioral bias, the models that underwent the paired training show an increased 218 behavioral bias, that depended on the duration of the training and saturated for longer training 219 duration (fig. 4B). The simulation results reported in figure 4B were obtained using odor-naive 220 models that exhibited no odor preference, prior to training. To account for the experimental finding 221 that real larvae often do have an odor preference even without any training [67–69], we readjusted 222 our experiments to include a pre-training period of 10 minutes to start the conditioning experiments 223 with the amylacetate-reward association already established. This adaptation of the protocol leads to 224 results (fig. 4 C) that match the results obtained in real behavioral experiments (fig. 5 A). The paired 225 condition in figure 4 C shows that once the behavioral bias is saturated (fig. 2 A), continued pairing 226 maintains the association, without further increasing it. Unpaired training on the other hand, causes 227 the behavioral bias to decrease and saturate at a lower level. For a discussion of different potential 228 causes of a reinforcement expectation prior to training, please refer to the discussion (Comparison of 229 modeling results to experimental findings). Figure 2A demonstrates that disabling MBON>DAN 230 feedback leads to a learning curve that does not saturate but instead increases with a steep slope 231 until it reaches the maximum value for the behavioral bias eqn. 4) with a MBON rate of 0. To verify 232 if this PE feedback mechanism is responsible for the difference between maintenance and loss of the 233

association in figure 4 C, we repeated the same experiment with disabled MBON>DAN feedback.

The behavioral bias overall is much higher, compared to the intact network (fig. 4 B). The maximum is reached before the test phase of even the shortest 1min training experiment, with no MBONspikes elicited.

Secondly, since the effect of training in lab experiments is quantified behaviorally via spatially 238 defined, group-level metrics (preference index and performance index (eqn. 15, eqn. 16), [1]), we 239 performed behavioral simulations of the testing phase with groups of virtual larvae for both the 240 paired and unpaired condition [1], allowing a straightforward comparison with the animal experi-241 ments (fig. $5 \, \text{A}$). To this end, we utilized a realistic model for the simulation of larval locomotion 242 and chemotactic behavior [52] that uses the behavioral bias at the output of the MB model as 243 a constant gain factor to modulate the locomotory behavior of individual larvae towards or away 244 from a spatially placed odor source in a virtual arena (see Methods, section: Realistic modeling of 245 larval locomotion). The resulting preference indices, acquired across groups of independently sim-246 ulated larvae (fig. 5 C), can directly be compared to the experimentally obtained preference indices 247 (fig. 5 A). We also compare performance indices from our simulated experiments (fig. 5 D) with those 248 from the lab experiments (fig. 5 B) and find that the model can replicate these when accounting for 249 the odor preference at the beginning of the experiment. 250

²⁵¹ Discussion

Seeking rewards and avoiding punishments by predicting change in the environment is a major 252 motivator of animal behavior. Sensory clues can acquire the necessary predictive power to guide 253 behavior through classical conditioning, an associative learning process potentially driven by re-254 ward/punishment PE [8, 11], as observed in vertebrates [16, 37-40]. To test the biological plausi-255 bility of the proposed PE coding motif in the larval MB and test its capacity to explain behavioral 256 data we implemented a spiking network model of the olfactory pathway, coupled with a simulation 257 of locomotory behavior [52]. We demonstrate that our model of PE coding results in saturating 258 group-level and individual learning curves, where the slope and maximum of the learning curve are 259 determined by the intensity of both the reward and the odor signal. Learning is also influenced by 260 the timing of odor and reinforcement and can be extinguished if reinforcement is omitted during the 261 presentation of the sensory clue. After verifying that this circuit motif enables learning as predicted 262 by the PE theory, we show that it can also explain time-resolved larval behavior in conditioning 263 experiments. 264

A mechanistic implementation of the RW model

A number of predictions can be derived from the phenomenological RW model [8] and tested in our mechanistic model thereof. We found that regardless of odor/reward intensity or the model's learning rate, the strength of the odor-reward association (quantified as the behavioral bias) saturates over time (fig. 2 A), as the strength of the already acquired association V(t) approaches the maxi-

mum value supported by the given reinforcement input ($\lambda_{\rm US}$) (eqn. 1). Consequently, our model's 270 acquisition curve saturates at a higher value when the intensity of the reinforcement is increased 271 (fig. 2 A,B), as predicted by the RW model, in which a stronger US should result in a higher value 272 of $\lambda_{\rm US}$ [8]. In our model, a higher reinforcement intensity relates to a higher input rate into the 273 respective DAN (see Methods, section: Sensory input) which translates into more frequent DAN 274 spikes within a given window of 1 second, used to compute the behavioral bias (eqn. 4). This defines 275 the asymptote of the learning curve. According to the RW model, increasing either the intensity of 276 the odor or the learning rate α [8] should lead to faster acquisition of the association. In our model, 277 the learning rate directly influences the increment of each respective synaptic weight Δw^{i} , while an 278 increase of the odor intensity allows for a more frequent execution of the weight update routine, by 279 influencing the eligibility trace (eqn. 2). 280

The RW model predicts that the omission of reward should result in the loss of the learned 281 association (eqn. 1, [8]). From the equation itself, we can not infer if this loss is due to extinction 282 or forgetting. Extinction, characterized by the possibility of recovery of the association, after its 283 temporary loss [70], has been demonstrated in adult [71, 72] but not larval Drosophila. To retain 284 the association for recovery, extinction relies on the formation of parallel memory traces for the 285 acquisition and the loss of the association [41, 60]. The mechanism implemented in our model is 286 overwriting the association, since the homeostatic mechanism drives the synaptic weights toward 287 their initial value, thereby deleting the learned association with no chance of recovery, but only in 288 the presence of olfactory input, eliciting MBON spikes. The resulting behavior during the extinction 289 phase of the experiments presents itself in a similar way, while the underlying mechanism is different. 290

²⁹¹ Comparison of modeling results to experimental findings

A variety of experiments have demonstrated group-level acquisition curves that saturate over mul-292 tiple training trials or with increasing duration of a single trial in olfactory conditioning [1, 51, 293 73-75]. To replicate larval behavior in reward learning experiments [1] with varying duration of the 294 learning phase (fig. 5 A,B) we trained our model with an odor and reward in a paired vs. unpaired 295 fashion (fig. 4B). Real larvae show a strong odor preference even after a very short training and 296 no significant increase in their preference when trained in a paired manner for longer periods of 297 time [1, 67]. Instead, the animals trained in an unpaired protocol start out with a similarly high 298 odor preference, which then decreases over time [1, 67]. This behavior is very counter-intuitive since 299 the coincidence of odor and reward should yield an association of those two stimuli and thus an 300 increased behaviorally expressed preference for the CS [8]. To resolve this contradiction, we include 301 the observation that animals might not be naive to the training odor prior to the beginning of the 302 experiments in the model. In that case, the animals would enter into the experiment with an already 303 established reward prediction that would be violated during unpaired training. Three scenarios lend 304 themselves as plausible causes of this effect: Firstly, accidental conditioning over the course of their 305 lifespan during which they are raised on a food substrate while being exposed to air that carries 306

many different odorants. Alternatively, or in fact, additionally, the animals might exhibit an innate 307 preference for many odors [76-78]. Finally, the presence of the reward during reward-only phases 308 might lead to an association of the experimental context with that reward (previously discussed 309 by Saumweber et al. [67]). The resulting reward expectation (solely based on the always present 310 context), unmet during the odor-only phases could lead to a prediction error signal. All three candi-311 date explanations would yield a similar projection for the unpaired experimental protocol: A reward 312 expectation acquired prior to the actual experiment would cause a violation of that expectation 313 during odor-only trials of the unpaired experiments. In all three cases, the animal's preferences 314 might also generalize to a broader array of odors, leading to an overall preference for some odors, as 315 observed experimentally. To test this hypothesis we pre-trained our model before simulating condi-316 tioning experiments (fig. 4 C) and observed that this allows us to reproduce the animal experiments 317 (fig. 5 A,B). Including odor preference at the beginning of the experiment ensures the model not only 318 behaves in accordance with the RW model [8], but also fits the animal experimental results [1]. A 319 possible alternative explanation could be a sensory habituation process to the odor that might cause 320 odor preferences to decrease over time, resulting in the observed patterns for unpaired learning. In 321 the paired condition this effect might be abolished by the continued presentation of odor and reward 322 together [79]. 323

Thus far we have tested our model in experiments where the CS and US presentation were fully 324 overlapping (paired conditions). We now consider different onset times, with the onset of the CS 325 always preceding the onset of the US (fig. 3,C). For these experiments we used a shorter duration of 326 30s for both CS and US presentation, repeated over three acquisition trials to mimic experimental 327 conditions in larval experiments [29, 66] that used optogenetic activation of DANs as a proxy for 328 sugar reward. Similar to their experiments we show that the behavioral bias clearly depends on the 329 temporal delay between CS and US (fig. 3,C). Complete temporal overlap of CS and US (ISI=0) 330 does not seem to expend the full potential of learning the association, instead partial overlap yields 331 stronger associations due to the extended window of opportunity for synaptic change triggered by 332 the odor's eligibility trace. In our model, the eligibility trace e(t) represents a molecular process that 333 maintains the odor signal locally in the KC>MBON synapses (eqn. 2). Zeng et al. [80] demonstrate 334 that feedback from the serotonergic dorsal paired medial neuron onto the KCs directly influences 335 the length of the KC eligibility trace, making it a candidate mechanism for associative learning with 336 a delayed US. Appetitive and aversive trace conditioning experiments have been conducted with 337 larvae [29, 65, 66] and adult flies and other insects [74, 81–83]. In all of these experiments where 338 the CS is presented before the US demonstrate that longer inter-stimulus intervals abolish learning 339 of the CS-US association when no KC odor representation persists during the reinforcement period. 340 In the cases of shorter intervals, the experimental data is not entirely conclusive. Either the odor 341 preference was higher for partial or no overlap, compared with complete overlap [29, 83] or highest 342 for complete overlap [51, 66, 74]. 343

³⁴⁴ We also looked at the extent of reinforcement generalization to novel odors. Experiments have

shown that associations between an odor and reinforcement generalize, to a varying extent, to other 345 odors, as shown in experiments [63, 84]. Previous modeling experiments have also shown that 346 reinforcement generalization depends on odor similarity in adult insects [48, 85–87]. In our larval 347 model, we also demonstrate both generalization to other odors, as well as a loss in strength, compared 348 to the training odor (fig. $3 \, \text{A}$). We also show that the extent of the generalization depends on the 349 similarity of the training and test odor, as measured by the overlap of the input patterns (fig. 1D). 350 The larval pathway with its relatively small coding space [53, 55] might be especially prone to such 351 poor discriminative abilities. 352

³⁵³ 0.1 Model predictions for behavioral experiments

Our approach targets two hypotheses: Firstly, symmetrical inhibitory and excitatory feedback from 354 MBONs to DANs should yield a circuit capable of saturating learning curves as predicted by the 355 RW model [8], due to PE [11] driving the learning process, which has also been suggested by 356 previous models [20, 48–51]. Secondly, saturating learning curves, driven by PE should translate 357 into (simulated) animal behavior, when comparing different training duration and intensities of 358 reinforcement. We were able to test these hypotheses in model experiments, on the level of MB 359 readout (behavioral bias, eqn. 4, fig. 2, 4)) and through the comparison of animal and simulated 360 behavior of artificial larvae (fig. 5). While the simulation results fit nicely with the real larval behavior 361 in an experiment with a varied training duration ([1], fig. 5), ultimately, the role of MBON>DAN 362 feedback needs to be tested in behavioral animal experiments, directly manipulating this feedback. 363 Some specific predictions that could be tested in such experiments are: 364

- Learning curves of individual animals should saturate over time when KC>MBON feedback is intact.
- When the MBON>DAN feedback is removed after some training, the learning curve should increase with a steeper slope and might not saturate.
- Increasing or decreasing the intensity of the odor or the reinforcement should lead to saturation on a higher or lower level, respectively.
- The removal of the KC>MBON feedback should weaken or abolish the saturation of the learning curve over time.

Based on our modeling results, we support the idea that the error computation between the prediction and reality of reinforcement is done in the DANs and relies on MBON>DAN feedback. Our hypotheses for experiments are based on this assumption. Nevertheless, some saturation, that is not based on PE, might still occur, even if MBON input to DANs is removed. The entire MB circuitry consists of many more elements than our model implementation and would presumably have additional mechanisms to ensure homeostatic balance and continued MBON input, potentially leading to some weaker form of saturation in the learning pro.

³⁸⁰ Comparison with other MB models

Models of the learning in the MB, based on plasticity at the MB output synapses, without PE coding, 381 have been around for some time, both for Drosophila [85, 87-89] and other insects [86, 90]. In all 382 of these models, plasticity is mediated by the activity of modulatory neurons (e.g., dopaminergic), 383 coinciding with either KC [86, 87] or coordinated KC and MBON activity [85, 88, 89]. These models 384 can perform associative learning of a stimulus, paired with reinforcement [85–89], as well as more 385 complicated forms of learning such as second order conditioning [89] and matching to sample [88] or 386 reinforcement generalization tasks, the extent of which depends on the stimulus similarity [85, 86]. 387 Additionally, some models were successfully tested in patterning tasks [85, 86], where combinations 388 of stimuli are reinforced, while their individual components are not or vice versa. Models in which 389 synaptic plasticity is driven not solely by the activity of modulatory neurons, but by a prediction 390 error signal lend themselves to studying the evolution of learning over time (either over several 391 trials, or in a continuous manner), and its dependency on the learning history. We hypothesize that 392 such mechanisms for PE coding in the MB involve the modulatory DANs [19, 20, 32, 34, 41–44] 393 and are based on MBON feedback to the DANs, serving as a manifestation of previous learning. 394 Recently a number of modeling approaches have targeted the idea of PE coding in DANs in the adult 395 Drosophila [48–51] as well as in the larval MB [20]. In these models, some form of MBON>DAN 396 feedback is implemented, allowing these models to fulfill some of the predictions of the PE theory [8, 397 11]. One of the most fundamental predictions is the saturation of the learning curve across time, as 398 the prediction error decreases, demonstrated in a trial-based manner in some of those models [48–51] 399 as well as the loss of an acquired association [20, 48-50]. Some of the previously published models 400 include mechanisms for either permanent loss of the association in memory or extinction (parallel 401 associations in memory). Within the MB circuitry, the formation of a parallel extinction memory 402 involves an additional DAN of opposite valence [20, 48-50], whereas complete loss is implemented 403 as a process of changing the KC>MBON weights in the opposite direction of the learning process 404 [51, 89], as done in our model. Additionally, some of these models capture temporal dynamics 405 of learning experiments to some extent by utilizing eligibility traces in the KC>MBON synapses 406 [20, 50, 51], to our understanding, none have tested these predictions in continuous experiments 407 with spiking dynamics. Therefore, beyond the scope of these contributions, we implemented PE 408 coding mechanistically in a fully spiking network equipped with synaptic eligibility that we train 409 and test in continuous experiments to allow for the assessment of dynamic change in the model's 410 odor preference. In combination with a time-continuous behavioral simulation [52] during memory 411 retention tests, this allowed for straightforward comparison with larval experiments. 412

Prediction error coding is not the only mechanism discussed in the literature to explain such phenomena in learning. Gkanias et al. tested a PE-based learning rule against a different dopaminebased learning rule that dos not require the presence of the CS as a reference point for expected reinforcement [87] in a more complex circuit model consisting of a number of interconnected microcircuits. They show that both methods can produce a saturating learning curve across trials. Their

alternative learning rule, embedded in a multi-compartment structure of the MB can also explain
extinction, blocking, and second order conditioning, by relying on interactions between different
MBONs and DANs that encode different memory processes.

421 Outlook

Some experimentally observed effects in insect learning can not be captured by the RW model [8] 422 and are thus not targeted by our model implementation. Among them are CS and US pre-exposure 423 effects [91–94] that might be explained by changes either in attention to the CS or habituation to 424 the CS or the US, caused by prolonged exposure prior to training, rather than changes in associative 425 strength (for a review see [95]). Also interesting, but not directly predicted by the RW model [8] 426 is the experimental observation of second order conditioning in adult *Drosophila* [96–99], where a 427 second CS2 is paired with the CS, after this CS has acquired an association with the US. Through the 428 CS2-CS pairing without the US, the CS2 acquires predictive power of the US. Different mechanisms 429 have been proposed to be involved in causing this effect [98, 100]. Among them is an excitatory 430 synaptic KC>DAN connection, strengthened during first order conditioning, that would allow the 431 KC odor representation to activate the DAN as a substitute for reinforcement during the CS2-CS 432 pairing. Exploring this phenomenon using network models could yield valuable insights into the 433 Drosophila circuit, as well as aid in our general understanding of PE coding. Insect experiments 434 have provided mixed evidence for other phenomena that can be predicted from the RW model, such 435 as blocking [101-104] and hints at conditioned inhibition [105-107] that would be interesting to 436 investigate. Furthermore, expanding the model to include different MB output compartments would 437 offer a perspective to explore parallel associations regarding the same stimulus [41]. This could 438 enable temporary loss of the learned association, while simultaneously retaining parallel memory 439 for recovery (extinction vs. forgetting). Ultimately more possible directions arise from the major 440 benefit of using a spiking model, which offers the potential to conduct experiments at high temporal 441 resolution, instead of in a trial-based manner [20, 48-50]. In a future closed-loop approach that 442 connects our continual learning MB model with the locomotory model in the full temporal resolution, 443 we intend to simulate a behaving agent to investigate the temporal dynamics of adaptive behavior 444 in analogy to the tracking experiments of real larva [73, 108–111]. 445

$_{446}$ Methods

447 Network model

All neurons are modeled as leaky integrate-and-fire neurons with conductance-based synapses. They elicit a spike, whenever the threshold V_T is crossed(parameters provided in table S1). Each neuronal membrane potential v_i is reset to the resting potential V_r whenever a spike occurs, followed by an absolute refractory period of 2 ms, during which the neuron does not integrate any inputs. Any

neuron from a given population $(v^{\rm O}, v^{\rm P}, v^{\rm L}, v^{\rm K}, v^{\rm A}, v^{\rm M}, v^{\rm D})$ is governed by the respective equation 452 for ORNs, PNs, LNs, KCs, APL, MBONs and DANs (eqs. (5) to (11), fig. 1A). Depending on 453 the neuron type, in addition to a leak conductance g_L , the equations consist of excitatory g_e and 454 inhibitory synaptic input g_i . In the case of the DANs, one excitatory $g_e^{M \mp D}(E_E - v_i^D)$ and inhibitory 455 $g_i^{M\pm D}(E_I - v_i^D)$ input represent the two types of MBON feedback for the reward and punishment 456 encoding DAN, respectively. An additional spike-triggered adaptation conductance was implemented 457 for ORNs, KCs, MBONs, and DANs (eqn. 12, [64]), in accordance with our current knowledge of the 458 adaptive nature of ORNs in the larva [112] and the adult fly [113, 114]. Adaptation in KCs has so 459 far only been demonstrated in other insects [115, 116]. In the model of these neurons, the adaptation 460 conductance g_{Ia} is increased with every spike and decays over time with τ_{Ia} . The mechanism of 461 synaptic plasticity is described in the results section (Learning through KC>MBON plasticity). 462

$$C_m \frac{d}{dt} v_i^O = g_L^O (E_L^O - v_i^O) + g_e^{InputO} (E_E - v_i^O) - g_{Ia} (E_{Ia} - v_i^O)$$
(5)

$$C_m \frac{d}{dt} v_i^L = g_L^L (E_L^L - v_i^L) + g_e^{OL} (E_E - v_i^L)$$
(6)

$$C_m \frac{d}{dt} v_i^P = g_L^P (E_L^P - v_i^P) + g_e^{OP} (E_E - v_i^P) - g_i^{LP} (E_I - v_i^P)$$
(7)

$$C_m \frac{d}{dt} v_i^K = g_L^K (E_L^K - v_i^K) - g_i^{APLK} (E_I - v_i^K) + g_e^{PK} (E_E - v_i^K) - g_{Ia} (E_{Ia} - v_i^K)$$
(8)

$$C_m \frac{d}{dt} v_i^A = g_L^A (E_L^A - v_i^A) + g_e^{KAPL} (E_E - v_i^A)$$
(9)

$$C_m \frac{d}{dt} v_i^M = g_L^M (E_L^M - v_i^M) + g_e^{KM} (E_E - v_i^M)$$
(10)

$$C_m \frac{d}{dt} v_i^D = g_L^D (E_L^D - v_i^D) - g_i^{M_{\pm}D} (E_I - v_i^D) + g_e^{M_{\mp}D} (E_E - v_i^D) + g_e^{InputD} (E_E - v_i^D)$$
(11)

$$\frac{d}{dt}g_{Ia} = -\frac{g_{Ia}}{\tau_{Ia}}.$$
(12)

463 All code for the model implementation is accessible via

464 https://github.com/nawrotlab/PEcodingDosophilaMB

We based our circuit model on the larval connectome both in terms of connectivity as well as numbers of neurons in each population [20, 53, 54] and introduced simplifications to support the mechanic investigation of the MBON>DAN feedback circuit and its role in PE coding and excluded a number of connections that have been demonstrated in the larva. Due to limited availability of

anatomical, functional, and behavioral data most of our circuit implementation is based on the first 469 instar larva [20, 53, 54], while the information on the APL connectivity within the circuit originates 470 from studies on the third instar larva [29]. Behavioral experiments used for comparison with our 471 simulation results were also performed with third instar larvae [1, 29, 66]. We demonstrate that our 472 model based on the less developed circuit in the first instar larva is sufficient to reproduce animal 473 behavior as observed in the older animals. From the anatomy of the first instar larva we excluded 474 DAN>KC [54] and DAN>MBON synapses [54] that may play an additional role in learning-induced 475 plasticity at KC>MBON synapses [54], the details of which are not fully known. Instead, we induce 476 plasticity purely via the simulation of a neuromodulatory effect of the DANs onto the KC>MBON 477 synapses (54). We also neglect recurrent interactions among KC themselves 54. Many of these 478 interactions affect KC that encode different sensory modalities, which are not included in our purely 479 olfactory model. Furthermore, we simplified the connectivity between LNs and PNs [53] and between 480 PNs and KCs to 2-6 PN inputs per KC, which excludes the set of KCs in the larva that receives 481 exclusive input from only one PN [54]. This modification supported model robustness with respect 482 to odor encoding within the small set of 72 KCs. Finally, from the population of ≈ 25 larval MBONs 483 we only modeled two and correspondingly adapted KC>MBON synapses to provide both MBONs 484 with input from all KCs. 485

486 Sparse odor representation

We implemented four mechanisms supporting population- and temporal sparseness in the MB odor 487 representation [64]. Population sparseness is defined as the activation of only a small subset of 488 neurons by any given input [117]. In this circuit population sparseness is enhanced through lateral 489 inhibition (via LNs), inhibitory APL feedback, and the divergent connectivity from PNs to a larger 490 number of KCs [64]. Temporal sparseness indicates that an individual neuron responds with only a 491 few spikes to a specific stimulus configuration [118-120], which supports encoding dynamic changes 492 in the sensory environment [121, 122]. In our model temporal sparseness is facilitated by spike 493 frequency adaptation, an adaptive process to prolonged stimulus exposure, in ORNs and KCs and 494 by inhibitory feedback via the APL[64]. 495

496 Sensory input

In the olfactory pathway of larval *Drosophila* any odor activates up to $\approx 1/3$ of ORNs, depending on its concentration [112, 123]. We implemented receptor input with stochastic point processes to ORNs via synapses to mimic the noise in a transduction process at the receptors. Each of the 21 receptor inputs is modeled according to a gamma process (shape parameter k= 3). The spontaneous firing rate of larval ORNs has been measured in the range of 0.2 - 7.9 Hz, depending strongly on odor and receptor type [123, 124]. ORNs in our model exhibit an average spontaneous firing rate of 8.92Hz (std=0.2). We constructed realistic olfactory input across the ORN population for amylacetate and

3-octanol by estimating ORN spike frequency from the calcium signals measured in the receptor 504 neurons [112] (dilution of 10^{-4} [112]), ensuring the spike rates would not exceed the rates reported 505 by [123]. They showed that using an even stronger odor concentration (dilution 10^{-2}) ORN never 506 exceeded a frequency of 200Hz. Due to the lower concentration used for amylacetate and 3-octanol 507 (fig. 1D) [112] in our experiments and because Kreher et al, 2005 measured only the first 0.5s 508 after odor onset when the effects of spike frequency in ORNs are the weakest (leading to higher 509 spike rates) we decided to use a maximum of 150Hz in odor activated ORNs. After generating the 510 gamma process realizations we clipped multiple spikes occurring in each time step of the simulation 511 discarding all but the first spike in each time step. Similar to the odor input, the presence of either 512 reward or punishment in the experimental context was implemented as input to the DAN₊/DAN₋. 513 Regular gamma spike trains (k = 10) were generated and clipped for the odor input. 514

To assess the effects of odor similarity on generalization we in addition created four artificial odors (A,B,C,D) (fig. 1 D) and quantified the pair-wise distances in ORN coding space using the cosine distance (eqn. 13), where vectors a and b each represent the input spike rate of two odors.

$$D_{cos} = 1 - \frac{\sum_{i=1}^{n} a_i \cdot b_i}{\sqrt{\sum_{i=1}^{n} a_i^2} \cdot \sqrt{\sum_{i=1}^{n} b_i^2}}.$$
(13)

The cosine distance between odors A and B equals 0.21, 0.77 between odors A and C, and 0.99 between odors A and D. The comparison of amylacetate and 3-octanol yields a distance of 0.16.

520 Experimental protocols

The experiments reported here belong to one of three categories. The first was performed to provide insight into the model and the effects of specific circuit functions on synaptic plasticity, and prediction error coding. To this end, we used amylacetate as the primary odor input. We varied the intensity of the reward via the frequency of gamma spike train, provided as input into the DAN₊ (either 500Hz or 550Hz, resulting in an average output spike rate of 33.11/39.14Hz), and the learning rate α (0.6nS or 0.8nS). Additionally, MBON>DAN feedback was either enabled or disabled (fig. 1 A).

Experiments belonging to the second category were designed to replicate larva lab experiments 527 to allow for a direct comparison with our model results. With these comparisons, we aim to validate 528 the model and show to what extent our assumptions about the circuit functions allow us to recreate 529 experimental data (fig. 5). Replicating lab experiments also provide more insights into the circuit 530 mechanisms and offers alternative interpretations of the phenomena observed in data from animal 531 experiments. Our implementations of the lab experiments were set up following the general procedure 532 described in the Maggot Learning Manual [125]. Regardless of the specific protocol used in different 533 experiments, larvae are placed into Petri dishes in groups of 30 animals. They are allowed to move 534 around freely on the substrate that contains reinforcing substances, such as sugar or bitter tastants. 535 During the entire time, they are subjected to specific odorants, emitted from two small containers 536 in the dish to create permanent and uniformly distributed odor exposure within the dish. In the 537

analogy of the experimental setting, in our simulated experiments, each model instance is trained individually through the concurrent presentation of olfactory stimulation and reward. One-minute intervals with only baseline ORN stimulation were included between training trials to simulate the time needed in the lab experiments for transferring larvae between Petri dishes. Unless otherwise specified and test phases refer to 3 min, during which only odors are presented. All simulations were implemented in the network simulator Brian2 [126].

544 Realistic modeling of larval locomotion

Behavior during the testing phase of the olfactory learning experiment is simulated via the freely 545 available python-based simulation platform Larvaworld (https://github.com/nawrotlab/larvaworld, 546 [52]). A group of 30 virtual larvae is placed with random initial orientation around the center of 547 a 100 mm diameter Petri dish and left to freely behave for 3 minutes. The previously conditioned 548 odor is placed at one side of the dish, 10 mm from the arena's boundary. Each larva features a 549 bi-segmental body, supervised by a layered control architecture [52]. The basic layer of the control 550 architecture is a locomotory model, capable of realistic autonomous exploration behavior. It consists 551 of two coupled oscillators, one of which represents the crawling apparatus that generates forward 552 velocity oscillations, resembling consecutive peristaltic strides [52]. The other oscillator generates 553 alternating left and right lateral bending, manifested as oscillations of angular velocity [127]. The 554 crawling and the bending oscillators are coupled via phase-locked suppression of lateral bending to 555 capture the bend dependency on the stride-cycle phase during crawling (weathervaning). Finally, 556 intermittent crawling is achieved by a superimposed intermittency module that generates alternating 557 epochs of crawling and stationary pauses, with more headcasts for orientation during the latter [52]. 558 Modulation of behavior due to sensory stimulation is introduced at the second, reactive layer of 559 the control architecture. An odor signal can transiently alter both, the amplitude and frequency 560 of the lateral bending oscillator, which biases free exploration towards approach or avoidance along 561 an olfactory chemical gradient. This modulation of behavior is directly influenced via top-down 562 signaling from the third, adaptive layer of the control architecture. In our approach, the spiking 563 MB model populates the adaptive layer and its learning-dependent output, defined as the behavioral 564 bias BB (i.e. the difference in MBON firing rates, eqn. 4), provides the top-down signal [36]. We 565 formalize the gain of behavioral modulation as 566

$$G = g \cdot BB. \tag{14}$$

which is directly proportional to the behavioral bias and the additional proportionality factor g = 0.5.

A set of 10 * 30 trained MB model instances is used to generate 10 groups of 30 simulated larvae. The preference index and the performance index [1] for these simulations are illustrated in figure 5. Preference indices (Pref) are computed individually for the paired and the unpaired experiments

⁵⁷² [1], based on the number of animals on each side (odor vs. empty) of the Petri dish at the end of ⁵⁷³ the test phase.

$$Pref = \frac{count_{odor} - count_{no \ odor}}{count_{odor} + count_{no \ odor}}.$$
(15)

The Performance indices (PI) are computed from the preference indices of the paired and unpaired experiments [1].

$$PI = \frac{Pref_{paired} - Pref_{unpaired}}{2}.$$
 (16)

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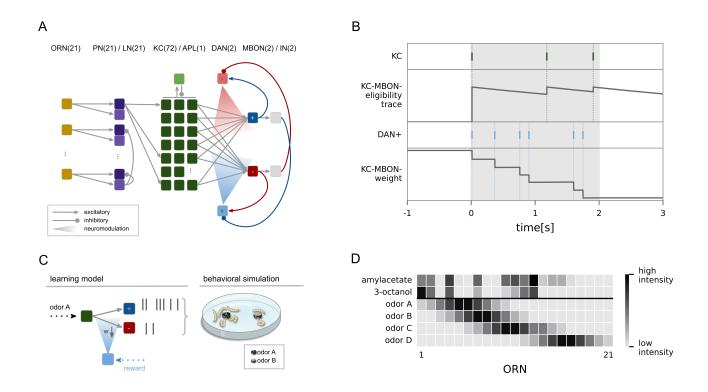


Figure 1: Network mechanisms. (A) Network model of the *Drosophila* larva olfactory pathway including all neurons and connections implemented. One-to-one feed-forward connections between 21 olfactory receptor neurons (ORN) and 21 projection neurons (PN)/local interneurons (LN) and from 2-6 PN to each of the 72 Kenvon cells (KC). Lateral inhibition from each LN innervates all PNs and recurrent feedback inhibition from the anterior paired lateral (APL) neuron is provided onto all KCs. The MB output region is organized in two distinct compartments. The upper compartment holds the approach encoding $MBON_{+}$ and is innervated by the punishment mediating DAN_{-} , the lower compartment holds the avoidance mediating MBON_ and is innervated by the reward mediating DAN₊. Each DAN can exert a neuromodulatory effect on the plastic KC>MBON synapses within its compartment. MBONs provide excitatory and inhibitory (via gray interneurons) feedback to the DANs. (B) Sketch of synaptic weight change at a single KC>MBON synapse with respect to the synaptic eligibility trace elicited by KC spikes and the occurrence of reward-triggered spikes in DAN_{+} . Amylacetate is paired with a reward for 2s (gray shaded area). (C) To generate simulated larval behavior in the petri dish during the test phase of the learning experiments, we utilized our locomotory model [52], based on the behavioral bias (eqn. 4) acquired by the MB model during the training phase. The behavioral bias is used directly as input to the locomotory model. (D) All odors (see Methods, section: Sensory input) were used in the experiments. Naturalistic odor patterns for amylacetate and 3-octanol as well as four artificial patterns (odorA,odorB,odorC,odorD) with varying distances (see Methods, section: Sensory input) from odorA. Each odor activates a different set of input neurons with a different spike rate, as indicated by the color bar.

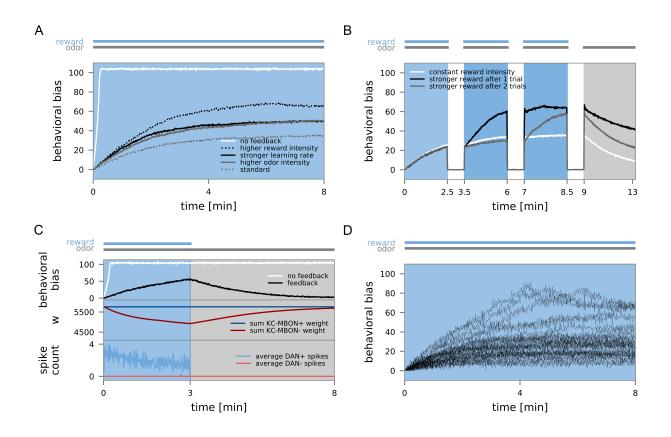


Figure 2: Learning with prediction errors. (A) N = 30 model instances were trained with the odor amylacetate (CS) and reward (US, blue background). MBON>DAN feedback, the reward/odor intensity, and the learning rate were manipulated in separate experiments. The odor preference (behavioral bias, eqn. 4) was measured continuously in windows of 1 sec and averaged over all model instances. (B) N = 30 model instances were trained during three trials with amylacetate and reward (blue background). Reward intensity was either constant across the three training trials (white curve), or enhanced during the third (gray) or the second and third trials (black). The training was followed by a 3 min test phase with odor only (gray background). (C)N = 30 model instances were trained with amylacetate and reward (blue background). (D) Individual acquisition curves for N = 30 model instances (standard experiment fig. 2 A).

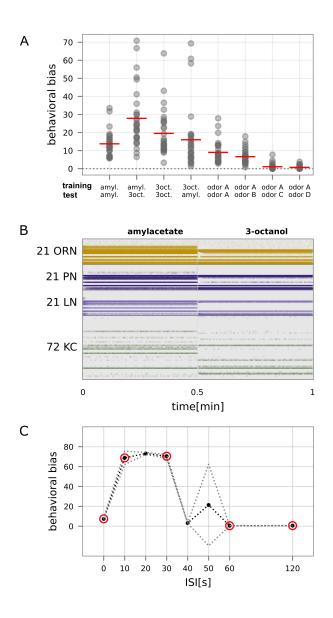


Figure 3: **Reward generalization and trace conditioning.** (A) The behavioral bias generalizes to odors that differ from the training odor after a 4min training (3min test phase). We conducted simulation experiments with different combinations of training and testing odor, each for 10 groups (gray circles represent the mean of a single group) of N = 30 larvae, and red lines indicate the mean between groups. The behavioral bias is highest when the training and the testing odor are the same. (B) Spiking activity in the network during the presentation of amylacetate (left) and 3-octanol (right) in a single naive model instance. (C) Simulated trace conditioning experiments with odor (amylacetate) and reward. Inter-stimulus interval (ISI) indicates the time between odor and reward onset. The black line displays the mean, gray lines the std over N = 10 groups of 30 model instances each. Conditions circled in red correspond to the conditions also used in animal experiments [29, 66]

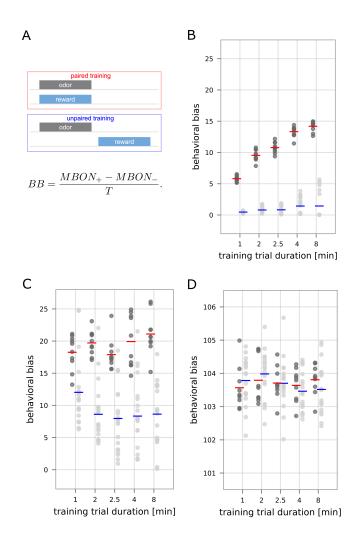


Figure 4: Paired and unpaired learning in the MB model. (A) Schematic overview of the paired vs. unpaired training protocol. (B) The model's behavioral bias for training with amylacetate and reward for N = 10 paired (dark gray, mean in red) and N = 10 unpaired (light gray, mean in blue) experiments with groups of 30 modeled larvae each. In the unpaired condition, half of the groups were trained with the odor preceding the reward, for the other half, the reward preceded the odor. (C) Model behavioral bias for amylacetate for N = 30 paired and N = 30 unpaired experiments with randomized order of odor and reward. Prior to the conditioning experiment the model instances underwent a 10min pre-training period, during which odor and reward were paired. (D) Model behavioral bias for amylacetate for N = 30 paired and N = 30 unpaired experiments with randomized order of odor and reward. Prior to the conditioning experiments with randomized order of odor and reward. The MBON>DAN feedback was disabled. Prior to the conditioning experiment the model instances underwent a 10min pre-training and N = 30 unpaired experiments with randomized order of odor and reward. The MBON>DAN feedback was disabled. Prior to the conditioning experiment the model instances underwent a 10min pre-training.

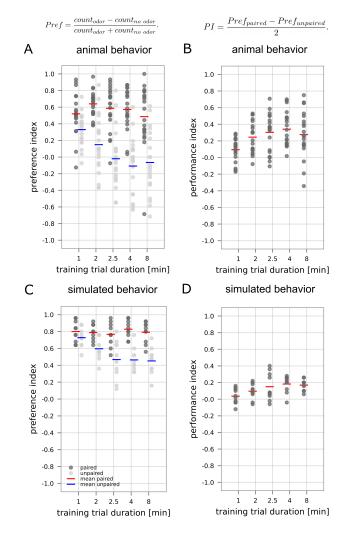


Figure 5: Replicating behavioral experiments with paired and unpaired training. (A) Experimental preference indices for anylacetate for 20 groups of 30 real animals each for paired and unpaired experiments with randomized order of odor and reward [1]. (B) Experimental performance indices for anylacetate computed between preference in paired and unpaired real animal experiments [1]. (C) The simulated behavior is based on the protocol in A. Simulated preference indices for anylacetate for N = 10 paired and N = 10 unpaired experiments with varied order of odor and reward. (D) Simulated performance indices for anylacetate computed between preference in paired and unpaired and unpaired simulation experiments.

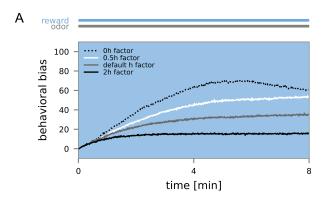


Figure S1: The effect of the homeostatic mechanism on the learning curve. (A) N = 30 model instances were trained with the odor amylacetate (CS) and reward (US, blue background). The odor preference (behavioral bias) was measured continuously in windows of 1 sec and averaged over all model instances. The learning rate was the same in all three experiments, while the magnitude of the homeostatic regulation h (eqn. 3,table S1) was either at its default value, at 0, or at half or twice the magnitude of the default value.

Neuron Parameters

Capacitance ORN	C_m^O	$100 \mathrm{pF}$
Capacitance PN	C_m^P	$30 \mathrm{pF}$
Capacitance LN	C_m^L	$50 \mathrm{pF}$
Capacitance KC	C_m^K	$30 \mathrm{pF}$
Capacitance APL	C_m^A	$200 \mathrm{pF}$
Capacitance MBON	C_m^M	$100 \mathrm{pF}$
Capacitance DAN	C_m^D	$100 \mathrm{pF}$
Leak Conductance ORN	g_L^O	$5\mathrm{nS}$
Leak Conductance PN	g_L^P	$2.5 \mathrm{nS}$
Leak Conductance LN	g_L^L	$2.5 \mathrm{nS}$
Leak Conductance KC	g_L^K	$5\mathrm{nS}$
Leak Conductance APL	g^A_L	$5\mathrm{nS}$
Leak Conductance MBON	g_L^M	$5\mathrm{nS}$
Leak Conductance DAN	g_L^D	$5\mathrm{nS}$
Leak Potential ORN	E_L^O	-60mV
Leak Potential PN	E_L^P	$-59 \mathrm{mV}$
Leak Potential LN	E_L^L	$-59 \mathrm{mV}$
Leak Potential KC	E_L^K	$-55 \mathrm{mV}$
Leak Potential APL	E_L^A	-60mV
Leak Potential MBON	E_L^M	-60mV
Leak Potential DAN	E_L^D	-60mV
Threshold Potential ORN	V_T^O	-35mV
Threshold Potential PN	V_T^P	-30mV
Threshold Potential LN	V_T^L	-30mV
Threshold Potential KC	V_T^K	-35mV
Threshold Potential APL	V_T^A	-30mV
Threshold Potential MBON	V_T^M	-30mV
Threshold Potential DAN	V_T^D	-30mV
Resting Potential ORN	V_r^O	-60mV
Resting Potential PN	V_r^P	$-59 \mathrm{mV}$
Resting Potential LN	V_r^L	$-59 \mathrm{mV}$
Resting Potential KC	V_r^K	$-55 \mathrm{mV}$
Resting Potential APL	V_r^A	-60mV
Resting Potential MBON	V_r^M	-60mV
Resting Potential DAN	V_r^D	-60mV
Refractory Time	$ au_{ref}$	2ms

Excitatory Potential	E_E	$0 \mathrm{mV}$
Inhibitory Potential	E_I	-75mV
Excitatory Time Constant	$ au_e$	5ms
Inhibitory Time Constant	$ au_i$	$10 \mathrm{ms}$
Plasticity Parameters		
Eligibility Trace Time Constant	$ au_{eligibility}$	5s
Learning Rate	α	$0.3 \mathrm{nS}$
Synaptic Weights		
Weight Input-ORN	wORNinputORN	3nS
Weight ORN-PN	wORNPN	10nS
Weight ORN-LN	wORNLN	4nS
Weight LN-PN	wLNPN	1 nS
Weight PN-KC	wPNKC	1 nS
Weight KC-APL	wKCAPL	20 nS
Weight APL-KC	wAPLKC	50 nS
Weight KC-MBON	wKCMBON	80nS
Weight Input-DAN	wDANinputDAN	$2.5 \mathrm{nS}$
Excitatory Weight MBON-DAN	wMBONDANex	4nS
Excitatory Weight MBON-local interneuron	$wMBONMBON_LN$	35nS
Inhibitory Weight local interneuron-DAN	wMBONDANin	70 nS
Normalization Factor KC-MBON	$normalization_{f}actor$	0.0001
Adaptation Parameters		
Adaptation Time Constant	$ au_{Ia}$	1000m
Adaptation Reversal Potential	E_{Ia}	-90mV
Increase of Spike Frequency Adaptation Conductance ORN	ORN_{SFA}	$0.1 \mathrm{nS}$
Increase of Spike Frequency Adaptation Conductance KC	KC_{SFA}	0.02 nS
Increase of Spike Frequency Adaptation Conductance MBON	$MBON_{SFA}$	$0.1 \mathrm{nS}$
Increase of Spike Frequency Adaptation Conductance DAN	DAN_{SFA}	$0.1 \mathrm{nS}$
Simulation Parameters		
Time Step	dt	$0.1 \mathrm{ms}$

Table S1: Network parameters.