1

5   **Postsynaptic plasticity of cholinergic synapses underlies the**

6   **induction and expression of appetitive memories in *Drosophila***

11   Carlotta Pribbenow[1], Yi-chun Chen[1,*], Michael-Marcel Heim[1,*], Desiree Laber[1,*], Silas

12   Reubold[1,*], Eric Reynolds[1,*], Isabella Balles[1], Raquel Suárez Grimalt[1,2], Carolin

13   Rauch[1], Jörg Rösner[4], Gregor Lichtner[1,5], Sridhar R. Jagannathan[1], Tania

14   Fernández-d.V. Alquicira[1], David Owald[1-3,+]

[1]Institute of Neurophysiology, Charité – Universitätsmedizin Berlin, corporate

member of Freie Universität Berlin and Humboldt-Universität zu Berlin, and

Berlin Institute of Health, Charitéplatz 1, 10117 Berlin, Germany

[2]Einstein Center for Neurosciences Berlin, Charitéplatz 1, 10117 Berlin

[3]NeuroCure, Charité – Universitätsmedizin Berlin, corporate member of Freie

Universität Berlin and Humboldt-Universität zu Berlin, and Berlin Institute of

Health, Charitéplatz 1, 10117 Berlin, Germany

[4]NWFZ, Charité – Universitätsmedizin Berlin, corporate member of Freie

Universität Berlin and Humboldt-Universität zu Berlin, and Berlin Institute of

Health, Charitéplatz 1, 10117 Berlin, Germany

[5]Universitätsmedizin Greifswald, Department of Anesthesia, Critical Care,

Emergency and Pain Medicine, Ferdinand-Sauerbruch-Straße, 17475

Greifswald, Germany

* equal contribution, in alphabetical order

+correspondence: david.owald@charite.de

In vertebrates, several forms of memory-relevant synaptic plasticity involve postsynaptic rearrangements of glutamate receptors. In contrast, previous work indicates that *Drosophila* and other invertebrates store memories using presynaptic plasticity of cholinergic synapses. Here, we provide evidence for postsynaptic plasticity at cholinergic output synapses from the *Drosophila* mushroom bodies (MBs). We find that the nicotinic acetylcholine receptor (nAChR) subunit α5 is required within specific MB output neurons (MBONs) for appetitive memory induction, but is dispensable for aversive memories. In addition, nAChR α2 subunits mediate memory expression and likely functions downstream of α5 and the postsynaptic scaffold protein Dlg. We show that postsynaptic plasticity traces can be induced independently of the presynapse, and that *in vivo* dynamics of α2 nAChR subunits are changed both in the context of associative and non-associative memory formation, underlying different plasticity rules. Therefore, regardless of neurotransmitter identity, key principles of postsynaptic plasticity support memory storage across phyla.

Keywords: learning and memory; postsynaptic plasticity; cholinergic synapses; *Drosophila;* mushroom bodies

Introduction

Changing the strength of defined chemical synaptic connections within associative networks is widely believed to be the basis of memory storage[1–3]. However, it is unclear and frequently debated to what degree underlying neurophysiological and molecular mechanisms are evolutionarily conserved. One main difference between vertebrates and invertebrates is that memory-storing synapses in vertebrates use glutamate as their primary transmitter, while those in invertebrates (at least for *Drosophila melanogaster* and *Sepia officinalis*) use acetylcholine[4–6]. Furthermore, it is widely established that invertebrate nervous systems utilize presynaptic plasticity for storing memories, while the degree to which postsynaptic plasticity can be used is largely unclear. In contrast, it is well established that storing information in vertebrates can depend on both pre- and postsynaptic mechanisms, including postsynaptic rearrangements of neurotransmitter receptors.

Detailed knowledge of the anatomical wiring and functional signaling logic of

the *Drosophila* learning and memory centers, the mushroom bodies (MBs)[5,7–21], allows

one to address whether, despite the use of different neurotransmitter systems, memory

storage modes are functionally comparable or evolutionarily conserved. The weights

of Kenyon cells (KCs) to MB output neuron (MBON) synapses are modulated by

dopaminergic neurons (DANs), which anatomically divide the MBs into at least 15

functional compartments, where information is stored on appetitive and aversive

associations, in addition to non-associative learning such as the relative familiarity of

an odor[5,7-12,15,22,23].

Studies so far have identified several traits pointing towards presynaptic

storage mechanisms within the KCs during memory formation[24–27]. Indeed, studies that

have blocked neurotransmitter release from KCs during learning[28–31] have brought

postsynaptic contributions to synaptic plasticity into question.

In vertebrates, typically, postsynaptic long-term changes[2,32,33] are mediated via

NMDA-sensitive glutamate receptors (NMDAR) that induce ('induction'[3]) an

expression phase ('expression'[3]) through changed glutamatergic AMPA receptor

(AMPAR) dynamics in dependence of postsynaptic scaffolds like PSD-95[34].

Invertebrate nAChRs in principle could take over similar functions to their glutamatergic

counterparts in vertebrates, despite their differing molecular characteristics[35]. Indeed,

nAChRs are pentamers that can be composed of homomeric assemblies of α subunits

or heteromeric combinations of different α and β subunits. The composition of subunits

determines the physiological properties of the nAChRs[35–38], and synaptic weights

could, in theory, be adjusted through the exchange of receptor subunits or entire

complexes.

Here, we capitalize on the genetic accessibility to individual output neurons of

the MBs to directly test whether postsynaptic receptors play a role in memory storage.

Derived from combined neurophysiological, behavioral, light microscopic and

molecular approaches, our data are supportive of a sequential role for nAChR subunits

in appetitive memory storage at the level of MBONs. Using artificial training protocols,

we demonstrate that postsynaptic calcium transients can change in response to

concurrent activation of dopaminergic neurons and application of acetylcholine,

circumventing KC output. Blocking KC output during appetitive, but not aversive,

learning abolishes memory performance. Moreover, specific knock-down of the α5

nAChR subunit, but none of the other six α subunits, in the M4/6 MBONs (also known

as MBON-γ5β'2a, MBON-β'2mp, MBON- β2β'2a and MBON-β'2mp bilateral) – an

output junction involved in coding appetitive and aversive memories[7,11,17] – impairs

immediate appetitive memories. Knock-down of α2 or α5, however, interferes with 3-

hour appetitive memories, as does knock-down of the scaffold Discs large (Dlg). We

report differential distribution of α subunits throughout the MB and demonstrate that

signal recovery of GFP-tagged subunits (as measured through fluorescence signal

recovered after photobleaching) is changed through plasticity protocols. In addition,

postsynaptically expressed non-associative familiarity learning also depends on α5

and α2 signaling as well as α2 dynamics. We propose a temporal receptor model and

speculate that, in *Drosophila*, nAChR subunits α5 and α2 take roles similar to NMDA

and AMPA receptors in vertebrates for memory induction and expression, indicating

that the general principle for postsynaptic plasticity independent of the

neurotransmitter system used, could be conserved throughout evolution.

Results

Neurotransmitter release from Kenyon cells is required for appetitive learning

One key argument in favor of exclusively presynaptic memory storage

mechanisms in *Drosophila* is based on experiments suggesting that blocking KC or

KC subset output selectively during learning leads to unaltered or mildly changed

memory performance[28–31]. If the postsynapse need not see the neurotransmitter during

training, it would likely be dispensable for memory induction. We revisited such

experiments and blocked KC output during T-maze training, exposing the animals

either to sugar-odor or shock-odor pairings (Fig. 1 a-b, Supplementary Fig. 1a-b).

We expressed the temperature-sensitive Dynamin mutant UAS-*Shibire^{TS}* (Shi)

at the level of KCs, trained animals at the restrictive temperature, and tested for

memory performance at permissive temperature 30 minutes later. These

manipulations allowed us to interfere with the synaptic vesicle exo-endocycle

specifically during conditioning, while reinstating neurotransmission afterwards.

Consistent with previous reports[28,30,31], a slight drop in aversive memory performance

(Fig. 1a) was not statistically different from their genetic controls, and also observable

in the permissive temperature controls (see Supplementary Fig. 1a). In contrast,

memories were completely abolished following block of KC output during appetitive
training (Fig. 1b, Supplementary Fig. 1b).

We next asked whether the requirement for neurotransmission during
appetitive learning was specific to the KC output synapse. To do so, we took an
analogous approach, this time blocking neurotransmission from downstream M4/6
(MBON-γ5β'2a, MBON-β'2mp, MBON- β2β'2a and MBON-β'2mp bilateral) MBONs
during appetitive training. We focused on the M4/6 set of MBONs as blocking these
during memory retrieval crucially interferes with appetitive memory expression, while,
on a physiological level, memory-related plasticity is observable[5,7,13]. When blocking
M4/6 during appetitive training, but not retrieval, memory scores were similar to those
of control groups (Fig. 1c), suggesting that the sites of plasticity are likely to be the KC
to MBON synapse in general, with one major site specifically being the connections
between KCs and M4/6 MBONs.

Thus, our experiments suggest that neurotransmitter release from KCs during
training is required for the formation of appetitive memories, but is less crucial for the
formation of aversive memories.

The α5 nAChR subunit is required for induction, and α2 for expression of
appetitive memories

Requirement for presynaptic neurotransmitter release alone does not
necessarily mean that postsynaptic plasticity is involved in appetitive memory
formation. To address a putative role for postsynaptic sensitivity in memory formation,
we next interfered with the postsynaptic receptor composition. Given that KCs are
cholinergic, we screened for memory requirement of all nicotinic α-subunits at the level
of the M4/6 MBONs (Fig. 2) using previously published[4, 51] genetically-targeted RNAi.
We concentrated on the nAChR α subunits, as they are crucial components for all
possible heteromeric or homomeric receptor pentamers[36].

When flies were tested for immediate appetitive memory, only knock-down of
the α5 subunit produced performance that was statistically different from the controls
(Fig. 2a, Supplementary Fig. 2a,i). Testing 3-hour appetitive memory performance
revealed a significant memory impairment in flies with α5, α1 and α2 knock-down (Fig.

2b, Supplementary Fig. 2b,j-l). While α5 subunits can form homomeric channels[38], α1

and α2 can partake in heteromeric channels together[37]. We therefore concentrated on

the α5 and α2 nAChR subunits in subsequent analyses.

To exclude developmental contributions to the observed memory defects, we

repeated the immediate and 3-hour appetitive memory experiments for α5 as well as

the 3-hour appetitive memory experiments for α2 knock-down animals, while

suppressing RNAi expression using the temperature-sensitive Gal4 repressor Gal80[ts]

during development, up until 3-5 days before memory testing. Memory impairments

were confirmed in all cases (Fig. 2e-g), but not detected in temperature controls

(Supplementary Fig. 2e-g).

We also tested aversive immediate and 3-hour memory using the same genetic

settings (Fig. 2c,d, Supplementary Fig. 2c,d). None of the knock-downs differed

significantly from controls, with the exception of α7 at the 3-hour time point. As,

comparable to vertebrate systems, α7 also plays a significant role at presynaptic

neurites[39], we did not follow up on this observation in this study.

As M4/6 output is also required for appropriate aversive memory expression[7,11],

α2 and α5 knock-down not impacting aversive memory performance suggested that

the observed appetitive memory impairments were not simply a consequence of lost

postsynaptic sensitivity to acetylcholine. To further corroborate this, we turned to a

brain explant preparation and applied acetylcholine focally to the dendrites of M4/6

neurons that expressed the calcium indicator GCamp6f, for both control and knock-

down settings, in the presence of the blocker of voltage-gated sodium channels

TTX[4,40]. Dendritic calcium transients were comparable between all groups

(Supplementary Fig. 3e). We also observed presynaptic sensitivity in all genotypes

(*not shown*) after applying acetylcholine to the presynaptic MBON boutons, making

presynaptic deficits following α2 or α5 knock-down unlikely.

Therefore, we conclude that, at the level of M4/6 neurons, immediate and 3-

hour appetitive memories are affected by knock-down of the α5 subunit, whereas 3-

hour memories also require the presence of α1- and α2-bearing receptors in addition.

The observed temporal profile of requirement for memory of α1- and α2-bearing

receptors relative to those incorporating the α5 subunit, could potentially point to a

temporal sequence of receptor function during initial memory formation and
subsequent memory expression.

The postsynaptic scaffold Dlg is required for 3-hour appetitive memory

At mammalian glutamatergic synapses postsynaptic receptor-mediated
changes in synaptic weight rely on receptor stabilization or destabilization that can be
mediated via scaffolding molecules. One such scaffold, PSD-95, that is mostly involved
in AMPA receptor motility, is conserved at *Drosophila* synapses. The orthologue
Dlg[41,42] is expressed throughout the brain, with mushroom body compartment-specific
enrichment noted previously[43] (also compare Fig. 4a,b). We investigated appetitive
and aversive memory performance following M4/6-specific knock-down of Dlg (Fig. 2a-
d, Supplementary Fig. 2a-d). Performance scores comparable to controls were found
for both immediate appetitive and aversive memories (Fig. 2a,c, Supplementary Fig.
2). Dlg knock-down, however, specifically abolished 3-hour appetitive memory
performance (Fig. 2b,d, Supplementary Fig. S2b), while Gal80[ts] experiments excluded
a developmental defect (Fig. 2h, Supplementary Fig. 2h). The temporal profile of Dlg
requirement therefore closely matched that of $\alpha2$ nAChR subunits.

Bypassing the presynapse: induction of persistent associative plasticity in the
postsynaptic compartment

Recent ultrastructural data has revealed direct synaptic connections between
dopaminergic neurons and MBONs[23,44], giving rise to a motif that could allow for
postsynaptic plasticity induction (see schematic in Fig. 3a). In order to directly test
whether postsynaptic plasticity could take place at the level of MBONs, we next
conducted neurophysiological proof-of-principle experiments.

To minimize plasticity induced by acute sensory experiences or general
network activity, we used an explant brain preparation bathed in TTX from flies
expressing the red light-activatable channelrhodopsin CsChrimson in a subset of
dopaminergic neurons (PAM neurons; R58E02-LexA) and the calcium indicator
GCaMP6f in M4/6 MBONs.

While dopamine release was controlled by red light flashes, neurotransmitter
release from KCs was mimicked by focal pressure ejection of acetylcholine to the
dendrites of the M6 (MBON-γ5β'2a) MBON (M6 was chosen for technical reasons, as
these neurons are most accessible for the used imaging technique). We verified that
KC presynapses do not respond to acetylcholine application[4], using both calcium
imaging and imaging of synaptic vesicle exocytosis at the level of KC axons (Fig. 3c
and Supplementary Fig. 3f,g). The observed absence of KC activation, with
acetylcholine being applied from an external source (Fig. 3a), minimized noise
attributable to possible presynaptic contributions.

Our protocols consisted of training phases where we differentiated between
temporal pairing of acetylcholine and optogenetic activation of dopaminergic neurons
('paired', Fig. 3b,h), dopamine only ('red-light only', Fig. 3b,e,g), or 'acetylcholine only'
(Fig. 3b,d,f). Acetylcholine application preceded (pre) and followed each training step
(post) to establish baseline responses and to assess synaptic weights following
training ('testing'). We found that test responses were significantly elevated following
the paired condition (Fig. 3h). This potentiation was not observed when testing after
acetylcholine only or dopamine only training (Fig. 3f,g). Importantly, we also did not
observe any changes when pairing acetylcholine application with red light in non-
CsChrimson-expressing controls (Supplementary Fig. 3c,d).

Because we are using global acetylcholine application instead of sparse
activation of single synapses, these experiments likely do not reflect *in vivo*
physiological settings[7]. However, our proof of principle experiments demonstrate that
postsynaptic sensitivity can change independently of the presynapse. Indeed, the
observed postsynaptic potentiation was not observed, when knocking-down α2 in M4/6
(Supplementary Fig. 3h), consistent with postsynaptic plasticity being linked to the
requirement of nicotinic receptors in memory storage.

Non-uniform distribution of nAChR α-subunits throughout MB compartments

Our data so far are suggestive of α2-containing nicotinic receptors being
involved in appetitive memory storage. To test whether receptor levels were
interdependent, we made use of a newly established CRISPR-based genomic
collection of GFP-tagged endogenous nAChR subunits (Woitkuhn, Pribbenow,
Matkovic, Sigrist and Owald, *unpublished*) covering all α subunits (with the exception

of α3) under control of their endogenous promoter, allowing for analyses of receptor
distribution and dynamics in a dense neuropile *in situ*.

We first characterized receptor subunit signals throughout the 15 MB
compartments, several of which have been shown to be involved in the encoding of
specific memories. We found a non-uniform distribution (Fig. 4a,b, Supplementary Fig.
4) that was unique for each subunit, indicating considerable heterogeneity of receptor
composition.

α5, which is required for immediate and 3-hour appetitive memories, was
abundant throughout the γ lobe, including γ5 (innervated by M6) and slightly less at the
level of β'2 (innervated by M4 and in parts by M6). α2 subunits, required for 3-hour
appetitive memories, showed similarly high relative abundance in β'2 and γ5 (Fig.
4a,b). Of note, these subunits were also detected in other MB output compartments,
such as α'3, which harbors MBONs involved in non-associative familiarity learning[15].

We next evaluated, whether the fluorescent signals of the α2 and α5 subunits
(with α5 potentially functioning upstream of α2, Fig. 2) observed in the β'2 and γ5
compartments were derived from receptors within the dendritic processes of M4/6. To
do so, we performed cell-specific knock-down experiments using VT1211-Gal4 and
quantified the relative fluorescent signal of the knock-down compartment relative to
the neighboring unmanipulated compartments (Fig. 4c-f).

Knock-down of the α5 nAChR subunit reduced the relative α5^{GFP} signal
specifically and significantly in γ5 and β'2 (Fig. 4e,f). α5 abundance was, however,
unaltered when knocking-down α2 or Dlg, which is in line with α5 functioning as a
possible trigger for plasticity processes.

Likewise, confirming that the observed signal was derived from M4/6 MBON
dendrites, α2 knock-down reduced relative α2^{GFP} levels significantly throughout the β'2
and γ5 compartments (Fig. 4c,d). Strikingly, we also observed reduced α2 nAChR
subunit levels following α5 subunit knock-down in the β'2 compartment or Dlg knock-
down in the β'2 and γ5 compartments (Fig. 4c,d), which would be in line with a Dlg-
dependent sequential requirement of receptor subunits during memory formation (also
compare behavioral data in Fig. 2).

Our data therefore are consistent with a role of α5 nAChR subunits and Dlg

functioning upstream of α2 subunit-positive receptors, at least within the β'2

compartment.

nAChR subunits shape synaptic MB output properties

We next focused on implications of α2 subunit knock-down on postsynaptic

function of M4/6 MBONs. Axonal calcium transients had previously been shown to be

decreased following knock-down of α subunits[4]. However, both increased or

decreased postsynaptic drive could lead to changed dendritic integration properties

underlying reduced signal propagation[45].

We expressed GCaMP6f in M4/6 MBONs (Fig. 5a), and exposed the flies

repeatedly to alternating puffs of the odors octanol (OCT) and MCH (Supplementary

Fig. 5a,b). We focused our experiments on the β'2 compartment (Fig. 5), as this is

innervated by M4 MBONs that show input-specific plasticity following appetitive

learning[7]. Initial dendritic odor responses were not statistically different between α2

subunit knock-down and controls (Fig. 5b,c), while initial odor-evoked dendritic calcium

transients were significantly elevated following knock-down of α5 (Fig. 5b,c).

Importantly, while control animals showed a relative facilitation in odor-specific calcium

transients after several exposures to OCT, we did not detect this in α2 knock-down

animals (Fig. 5d,e). Odor responses following α5 knock-down, however, clearly

depressed after multiple odor exposures (Fig. 5f), indicating that lack of α5 can lead to

pre-potentiated synaptic transmission, while α2 nAChR subunit knock-down interfered

with odor-evoked baseline transmission properties to a lesser extent. We did not

observe any changes in calcium signals at the level of the corresponding KC axons,

further supporting that the observed plasticity was of postsynaptic origin

(Supplementary Fig. 5h,i).

The α2 *nAChR subunit is required for the formation of appetitive memory*

***traces* in vivo**

Of note, the observed facilitation in M4/6 of controls was not apparent when

using repeated application of MCH, indicating that parameters underlying the observed

non-associative plasticity are odor-dependent (Supplementary Fig. 5e). However, the

only slight effects on base-line MCH-induced responses observed for α2 knock-down

animals, allowed us to next investigate the role of α2 during *in vivo* associative

appetitive learning.

We performed *in vivo* training under the microscope experiments using an

absolute paradigm, pairing odor (MCH) exposure with *ad libitum* sugar feeding during

training. Comparing odor responses at the level of M4 before and after training

revealed a marked depression for control animals in line with previous observations[7,17]

(Fig. 5g). Importantly, after MBON-specific knock-down of α2, this training-induced

plasticity was not observed (Fig. 5h).

Together, our data point towards a mechanism, where nicotinic receptor

subunits shape synaptic properties (Fig. 5), with α2 as a postsynaptic substrate

underlying appetitive training-induced plasticity processes.

In vivo *imaging of postsynaptic receptor plasticity reveals altered α2 dynamics*

Structural changes at the level of the receptor composition are hallmarks of

postsynaptic plasticity expression in vertebrates. Typically, rearrangements can be

measured by altered dynamics (or motility) of receptors that can reflect incorporation

or removal of receptor complexes. We next sought to test whether dynamic receptor

behavior could serve as a structural correlate of cholinergic postsynaptic memory trace

expression. To do so, we turned to *in vivo* imaging experiments of the endogenously

tagged α2 subunit (Fig. 6, Supplementary Fig. 6). Flies were either tethered under the

two (Fig. 6) or the single photon confocal (Fig. 7) microscope and subjected to training

protocols. Endogenous *in situ* receptor dynamics at the level of the β'2 compartment

of the MB were estimated as fluorescence recovery after photobleaching (FRAP),

allowing us to also pick up small increases in signal (Fig. 6a-c).

We conducted artificial appetitive training protocols by exposing the tethered

animals to odor with or without simultaneous focal injection of dopamine to the β'2

compartment of the MB (Fig. 6a-c). We performed two different sets of protocols

comparing recovery of α2^{GFP} (Fig. 6b, Supplementary Fig. 6b).

Following photobleaching, individual flies were exposed to consecutive focal

injections of dopamine, odor, and odor paired with focal dopamine injection. MCH

induced significantly increased fluorescence recovery when compared to dopamine

injections only or odor paired with dopamine (Fig. 6d,e). Dopamine, therefore, does

not induce plasticity on its own, and further, it suppresses odor-induced recovery when

applied simultaneously with an odor. To rule out that recovery depended on the type

of odor used, we also conducted similar experiments using OCT, this time testing all

conditions in separate flies. After photobleaching, flies were either exposed to a focal

injection of dopamine, an odor stimulus, or odor paired with focal dopamine injection.

We only observed significant recovery in the odor only condition, whereas dopamine

and odor paired with dopamine induced no significant recovery (Supplementary Fig.

6a-d).

Thus, our data indicate that pairing odor presentation with dopamine

application stalls $\alpha 2^{GFP}$ dynamics, potentially by either stabilizing the already present

amount of receptor or hindering new incorporation of $\alpha 2$-containing receptors.

Interestingly the opposite, increased receptor dynamics, is observed after odor

exposure without reinforcer. Note that the absence of acute stimuli (constant air stream

only), did not induce signal recovery, demonstrating that it is the presence of the odor

that changes baseline receptor behavior (Supplementary Fig. 6e). Thus, stalling $\alpha 2$

dynamics can be correlated to depression of M4/6 MBON synapses[7,13] following

appetitive training (Fig. 5g,h).

Familiarity learning alters postsynaptic receptor dynamics

We next asked whether postsynaptic plasticity expressed through $\alpha 5$ and $\alpha 2$

subunit interplay could underlie other forms of learning represented in the MBs. We

turned to the $\alpha'3$ compartment at the tip of the vertical MB lobe that has previously

been shown to mediate odor familiarity learning. This form of learning allows the animal

to adapt its behavioral responses to new odors and, importantly, permits for assaying

direct odor-related plasticity. Importantly, this compartment follows different plasticity

rules, because the odor serves as both the conditioned (activating KCs) and

unconditioned stimulus (activating corresponding dopaminergic neurons)[15]. While

allowing us to test whether the so far uncovered principles could also be utilized in a

different context, it also provides a less complex test bed to further investigate whether
α5 functions upstream of α2 dynamics.

Confirming previous observations[15], a repeated odor application paradigm (Fig.
7a) led to the depression of postsynaptic calcium transients at the level of the α'3
MBONs (Fig. 7b). Importantly, we did not detect a corresponding depression on the
presynaptic side when imaging arbors of a sparse α'β' KC driver line within α'3 (Fig.
7b), further indicating that memories were predominantly stored postsynaptically in this
compartment. We next performed *in vivo* FRAP experiments following familiarity
learning paradigms. After odor training, we observed clear recovery rates of α2^{GFP}
signals compared to the control group, however not of α5^{GFP} or Dlg^{GFP} (Fig. 7 c-f,
Supplementary Fig. 7). Therefore, increased α2 subunit dynamics are triggered
through training events and, at the level of the α'3 compartment, accompany
postsynaptic depression of the MBONs.

To invariantly test whether the observed recovery was attributable to α2
expressed in α'3 MBONs, we trained animals while knocking-down α2 specifically in
α'3 MBONs. In accordance with the observed signal recovery deriving from MBONs,
no recovery was observed after α2 knock-down (Fig. 7g). Importantly, we did not
observe α2^{GFP} recovery when performing specific α5 knock-down in α'3 MBONs (Fig.
7g), indicating a role of α5 upstream of α2 also in this compartment.

α5 subunits govern induction and α2 subunits expression of non-associative
familiarity learning

Finally, we tested whether interfering with α5 and α2 nAChR subunits at the
level of α'3 MBONs would also impact familiarity learning behavior (Fig. 8).
Flies were covered in dust and subjected to repeated odor exposures[15]. As expected[15],
control flies readily groomed to remove the dust, however typically stopped this action
when detecting the novel odor (Fig. 8a-c). Over subsequent trials, control flies learned
that this odor was familiar and stopped reacting to the stimulus, continuing grooming
(Fig. 8a-c, Supplementary Fig. 8). Expressing RNAi to the α2 subunit at the level of the
α'3 MBONs clearly impacted learning: flies learned with decreased efficacy and only
after several trials (Fig. 8a-d). Strikingly, α5 RNAi-expressing flies failed to stop

grooming even to the first stimulus. Indeed, they acted as if they had already learned
that an odor was familiar (Fig. 8a-d).

Together, our data are in line with a model where α5 can induce memory
formation, while lack of α5 leads to fully potentiated synapses. Subsequent expression
of memory traces requires α2-containing receptors. Importantly, recovery
accompanies synaptic depression at the level of the α'3 MBONs, while being
suppressed by paired training in the β'2 compartment. Moreover, α2 appears to be
involved in both depression and facilitation of synapses. Thus, synapses could
bidirectionally utilize plasticity of the same receptor subunit for storing different types
of information (Fig. 9).

Discussion

Synaptic weight changes are widely recognized as substrates for memory
storage throughout the animal kingdom. How synapses adapt in order to change their
efficacy during learning has been a focus of attention over the last decades. While it is
undisputed that both pre- and postsynaptic mechanisms of memory storage exist in
vertebrates, invertebrate memory-related synaptic plasticity has been largely localized
to the presynaptic compartment, although some debates exist[1]. The core of the debate
boils down to a key question: do vertebrates and invertebrates use similar mechanisms
to store memories or are there fundamental differences? A first clear difference
appears to be the use of different neurotransmitter systems, glutamate and
acetylcholine respectively, in the vertebrate and *Drosophila* learning centers[4].

Postsynaptic plasticity in appetitive memory storage

Here, we use the genetic tractability of the *Drosophila* system to directly
address postsynaptic plasticity during memory storage in invertebrates. Large
amounts of evidence from *Drosophila* so far suggests a presynaptic mode of memory
storage[24–28,46]. Moreover, it was demonstrated that block of KCs during learning does
not interfere with memory performance[28,30,31], although some studies blocking KC
subsets did find impairments[29,47] in the context of short-term appetitive memory. Lack

of necessity to detect neurotransmitter by the postsynapse in the course of memory

storage was interpreted as evidence for a presynaptic storage mechanism. If the

receptor does not 'see' the signal, it is dispensable for 'interpreting' it. Here, we

revisited such experiments and found, in accordance with previous studies, only mild,

if any, requirement for aversive memory storage. We, however, fully abolished

appetitive memories (Fig. 1) by blocking KC output during acquisition, providing a

model for postsynaptic plasticity (Fig. 9) that is induced and expressed through distinct

nAChR subunits (Fig. 2-8).

Our study hints towards different pre- and postsynaptic storage mechanisms

underlying aversive and appetitive memories. It also argues against the assumption

that appetitive and aversive memories will necessarily use the same molecular

machinery to store information. Findings in the past, predominantly based on

investigating aversive memories, have been generalized to learning per se. Indeed,

postsynaptic contributions have been ruled out for a synaptic junction required for

storage of aversive but not appetitive memories, which is fully consistent with our

findings[22]. Interestingly, arguing for a division of appetitive and aversive storage sites,

subpopulations of KCs have been implicated in aversive and appetitive memory

respectively[48]. Lastly, we do not wish to exclude a potential involvement of

postsynaptic plasticity in aversive memory formation per se. On the contrary, it is

conceivable that aversive memories also could have an appetitive component (release

from punishment). However, our experiments suggest that these influences are not as

crucial as for appetitive memories, at least in the context of single trial differential

associative learning.

Postsynaptic plasticity at the KC to MBON synapse

Recent anatomical studies[23,44] have reported both dopaminergic innervation of

presynaptic KC compartments as well as somewhat unexpectedly direct synapses

between presynaptic dopaminergic terminals and MBONs. We devised an experiment

where we substituted KC input to the postsynaptic MBON compartment through

artificial acetylcholine injection, while rendering dopaminergic neurons switchable

through optogenetics. A protocol that trained and subsequently tested the synaptic

junction between KCs and MBONs, demonstrates that plasticity (represented by a

change in calcium responses to acetylcholine injection) was inducible by pairing

dopaminergic with postsynaptic MBON activation that lasted beyond the training stage
and was observable by mere 'recall-like' activation of the system (Fig. 3).

Our proof-of-principle experiments uncovered the ability to potentiate after
pairing M6 MBON activation and stimulating a broad population of dopaminergic
neurons that convey information on sugar, water or the relative valence of aversive
stimuli[5], while we find postsynaptic plasticity to be required for appetitive memory
performance (Fig. 3). However, previous studies looking into 'natural' appetitive sugar
conditioning uncovered a relative depression in M4 (another MBON of the M4/6
cluster) dendrites, when comparing the responses of the paired (CS+) and unpaired
odor (CS-) one hour after appetitive conditioning[5,7,13]. Moreover, we here show that *in*
vivo appetitive absolute training depresses subsequent responses to the trained odor
(Fig. 5). It is important to note that, here (Fig. 3), for our *in vitro* experiments, we
perform global activation of the postsynaptic compartment and not the natural typical
coverage of 5 % of input synapses per odor[49] (that allow for differential conditioning).
Induced changes are therefore likely not comparable to the natural settings, where
sparse sets of KCs and dopaminergic neurons are active within a tight temporal
window. Moreover, we here abolish network contributions (by suppressing active
signal propagation), to be able to concentrate on synaptic mechanisms during plasticity
induction. Thus, our artificial training (Fig. 3) through global dendritic activation likely
does not mirror precise physiological conditions allowing for plasticity of a sparse set
of synapses to convey odor-specificity to a memory, and should therefore only be
viewed as a proof of principle for postsynaptic plasticity induction per se. However, the
relatively small amplitude of potentiation observed actually fits previous[7] *in vivo*
observations. Importantly, because we are using TTX and local training of KC to M6
synapses in our experiments, we are furthermore missing additional disinhibition that
in vivo is mediated via the GABAergic MVP2 MBON[13,19]. Of note, similar protocols[50]
that involved broad activation of KCs (and thus did not circumvent the presynaptic
compartment) have demonstrated comparable plasticity induction at this synaptic
junction.

Local acetylcholine application to the MB can also activate calcium transients
in dopaminergic presynaptic terminals[51]. Therefore, our protocol could in principle
include some dopaminergic contributions already at baseline level. However, control
experiments using the paired training protocol in the absence of CsChrimson
expression in dopaminergic neurons, do not show any signs of plasticity (Fig. 3 and
Supplementary Fig. 3). Moreover, it has been previously demonstrated that, to actually

release dopamine from the presynaptic terminal, a coincident signal via carbon

monoxide is required[52,53]. Therefore, an unwanted activation of dopaminergic neurons

in our experiments is unlikely.

It should also be noted that M4, which shows depression (Fig. 5), and M6 have

common but also distinct physiological roles, for instance during aversive memory

extinction[13]. Besides that, different temporal requirements for M4 and M6 memory

expression have been reported[11]. It is therefore possible that physiological changes in

the context of appetitive learning led to different plasticity profiles in M4 and M6

neurons respectively, or that initial potentiation over time can be reverted to

depression. As noted above, MBON drive is bidirectionally modifiable and has the

propensity to both potentiate and depress[7,11,17,26].

We did not observe plasticity after paired training following α2 knockdown *ex*

vivo (Supplementary Fig. 3). In line with that, our *in vivo* experiments directly establish

a connection between learning-triggered synaptic plasticity and the α2 subunit (Fig. 5).

Appetitive training induced synaptic depression at the level of the β'2-MBON dendrites

of control animals. In animals where we concurrently knock-down α2 in M4/6 MBONs,

this synaptic depression, however, was not observable.

Nicotinic receptors could follow temporal sequence

Lasting plasticity traces as observed here (Fig. 3) appear to fit the core criteria

for long-term potentiation of vertebrate glutamatergic postsynapses[3,54]. Plasticity can

be divided into an 'induction'[3] period mediated via NMDA receptors and a subsequent

'expression'[3] period that requires altered AMPA receptor dynamics[3]. Our findings here

lead to a model, where the nicotinic α5 subunit is required for the induction of appetitive

memories at *Drosophila* MBONs (Fig. 2, 9). We propose that α5 nAChR subunits (that

can form homomeric channels[38,39]) could take on a similar role to NMDARs. α5 would

gate the potentiation or depression of synaptic strength influencing the incorporation

or exchange of additional receptor subunits or complexes. In line with this, we show

that knock-down of α5 subunits interferes with familiarity learning in the α'3

compartment of the MBs: flies no longer form familiarity memories, they react to a novel

odor the same way as to a familiar one, 'as if they had learned that this new odor was

familiar before' (Fig. 7, 8). Moreover, we do not observe α5 subunit dynamics (Fig. 7),

whereas knock-down of α5 leads to decreased levels of α2 subunits (Fig. 4), and α2 dynamics are no longer observable when knocking-down α5 in the MBONs of the α'3 compartment (Fig. 7). Thus, we can draw first analogies to glutamatergic systems governing plasticity in vertebrates. Whether more core criteria are met for the comparison of invertebrate and vertebrate plasticity systems, further depends on whether the here observed receptor dynamics will actually translate to exo-/endocytosis of postsynaptic receptors or lateral diffusion of receptor subunits along the MBON dendrites. Our established system should provide the means to investigate this further in the future.

Interestingly, high levels of dendritic activation after α5 knock-down are translated to reduced axonal calcium transients[4], effectively leading to decreased signal transduction within the MBON. Of note, MBONs do not appear to exhibit prominent spines on their dendrites[23] (but see section below: *Are cholinergic and glutamatergic synapses interchangeable?*). Therefore, increased dendritic activation could lead to a change in membrane resistance and result in synaptic interference.

α2 subunit-positive receptors mediate memory expression

We also find that later forms of appetitive memory expression require both the α2 and α1 receptor subunits (Fig. 2). A recent study[37] has demonstrated that, when expressed heterologously, these subunits can co-assemble to form heterodimers with β subunits, which, depending on the precise composition of these channels, can harbor different properties, potentially reminiscent to AMPAR[55]. However, MB distribution profiles of α1 and α2 subunits do not match completely, for instance at the level of the γ5 or α'2 compartments (Fig. 4), indicating that they could also partake in different or independent receptor configurations. It would be worthwhile to compare receptor localization with single cell sequencing results[8].

Importantly, α2 subunit knock-down at M4/6 MBONs does not affect immediate appetitive memories, but later stages of memory (Fig. 2). We show in a complex neuropile, that *in vivo*, on a time scale of 10 to 20 minutes after learning, exposure to odor induces changed α2 receptor subunit dynamics (signal recovery after photobleaching) that are suppressed by simultaneous dopamine exposure ('learning', Fig. 6). Therefore, changes in α2 dynamics could match the temporal profiles of

memory expression and consolidation processes taking place in an early phase
following training (the first 20 minutes) to establish longer-term memories (3 hours). It
should be noted that, whether dopamine pairing would also lead to a reduction of α2
levels over time, potentially explaining synaptic depression, cannot be assessed with
our experimental settings at this time, but would appear as a valid possibility.

We show that familiarity learning can take place when knocking down α2
nAChR subunits in α'3 MBONs in principle (Fig. 8), however, at clearly decreased
efficacy and only after several trials. We speculate that the observation of memories
still being expressed *per se* in this context, could be explained by redundancies with
α1 or other subunits (but see heterogeneous localization and enrichment in different
MB compartments, Fig. 4). Redundancies could also explain why we partially observe
functional phenotypes after knock-down of individual subunits, but only moderate
structural changes. We also would like to point out that subunits we did not identify as
absolutely required for memory expression (Fig. 2) in this study could nonetheless
partake in distinct phases of plasticity processes.

Similar to AMPAR, localization of α2, but not α5 subunits, depends on Dlg, the
orthologue to vertebrate PSD-95 and PSD-93. Interestingly, the latter has been
implicated in structural integrity of cholinergic nicotinic receptor arrangements[59] in
vertebrates.

α2 dynamics as general plasticity mode?

In the context of both familiarity learning and appetitive conditioning, odor
exposure induces increased α2 subunit dynamics (Fig. 6, 7) accompanying
postsynaptic depression[7,15] (Fig. 7), while not or mildly affecting α5 subunits (for
familiarity learning). Therefore, the same basic mechanisms, odor-induced α2 receptor
dynamics, seem to express two opposed plastic outcomes in the context of associative
and non-associative memories and contribute to different learning rules across MB
compartments[22,56]. We speculate that α2 dynamics induced by odor in the M4/6
dendrites could be reminiscent of dark currents in the vertebrate visual system[57]
allowing for rapid adaptation with low levels of synaptic noise. Receptor exchange at
the level of M4/6 dendrites would actually take place when no associations are formed
and stalled when dopaminergic neurons (triggered by sugar) are simultaneously active

with KCs (triggered by odor). Indeed, repeated OCT stimulation led to a facilitation of
calcium transients (potentially corresponding to an increase of receptor incorporation,
Fig. 5-6), while depression (in this case likely to be mediated by removal of receptors,
but see above) is triggered by paired training (Fig. 5). In contrast, at the level of the α'3
compartments, odor activates both MBONs and dopaminergic neurons. Here, the
plasticity rule would be reversed. Synaptic depression is accompanied by actively
changing the receptor composite. We speculate that such plasticity could function
reminiscent of mechanisms observed for climbing fiber-induced depression of parallel
fiber to Purkinje cell synapses[58]. However, whether increased dynamics can be
translated to more incorporation or removal of α2-type receptors, or depending on the
plasticity rule both, will require high resolution imaging experiments in the future.
686
687
688 ***Are cholinergic and glutamatergic synapses interchangeable?***
689
Our study fuels the question of how unique properties of individual
neurotransmitter systems at synapses are. While dopamine signaling is remarkably
conserved between invertebrates and vertebrates, cholinergic and glutamatergic
systems appear, now more than before (with this study), somewhat interchangeable.
While vertebrates (but also evolutionarily distant *C. elegans*) for instance use
acetylcholine at the neuromuscular junction and store memories predominantly at
glutamatergic synapses, it is the other way around in *Drosophila* and other
invertebrates, such as *Sepia*[4–6,60,61]. Now we show that, at cholinergic synapses, α5
and α2 subunits, at least to a certain extent, behave in a potentially comparable way
to NMDARs and AMPARs at glutamatergic synapses during postsynaptic plasticity
which underlies memory storage. In this context, we offer several lines of evidence that
invertebrates utilize postsynaptic plasticity during memory storage and not as
previously assumed rely on presynaptic mechanisms only.
703
704   We therefore propose that, across phyla, postsynaptic plasticity, with the
705 propensity to store memories and adapt network function plastically, can take place
706 regardless of neurotransmitter identity.
707
708   One key difference between the dendritic arbors of the MBONs analyzed in this
709 study compared to dendrites of glutamatergic neurons in vertebrates, is a lack of
710 anatomical spines (Supplementary Fig. 9a). Without spines, how can input-specificity

be preserved at MBON postsynaptic densities? KC output to MBON input analysis of the recently published fly hemibrain connectome[62] (neuprint.org) suggests that, at the ultrastructural level, MBON postsynaptic densities are separated spatially (Supplementary Fig. 9a). Compartmentalization could therefore be mediated by, for instance, biochemical separation of PSDs. Importantly, input-specific plasticity has been shown to be inducible in non-spiny neurons in vertebrates, with diffusion barriers established, e.g., through calcium buffers, between postsynaptic densities[63].

Interestingly, our MBON input analysis further revealed that postsynaptic plasticity mechanisms could actually add a layer to promote input-specificity. Indeed, we find that single presynaptic KC release sites that innervate MBON dendrites can also target other MBONs and/or other postsynaptic targets simultaneously (Supplementary Fig. 9b-f). Plasticity confined to single postsynaptic densities innervated by a KC terminal could therefore change the weight of transmission for one target (e.g., MBONs involved in memory storage, such as M4[7]), while not changing the weight of the connection to other targets (for example MBONs not involved in a specific action or other targets of non-MBON identity, Supplementary Fig. 9g). It should be noted that this architecture does not exclude presynaptic plasticity mechanisms[70] (for instance following aversive conditioning). Indeed, we would speculate that synaptic connections can be subdivided into distinct compartments on both the pre- and the postsynaptic side, potentially through transsynaptic molecules[61,64], allowing for fine-tuned and target-dependent changes of parameters within either side of a synapse.

A molecular plasticity sequence?

Together, we propose a model (Fig. 9) in which α5-subunit containing receptors could mediate the early phase of postsynaptic memory storage and we speculate this could lead to elevated postsynaptic calcium flux (not addressed in this study). Concurrent events see changed dynamics of the α2 receptor. Nicotinic receptor subunits hereby could interact with adaptor proteins to bind to Dlg, reminiscent to what is known for AMPAR[35]. Importantly, we identify elevated α2 subunit dynamics in the context of associative and non-associative memory expression. Increased α2 subunit dynamics in both cases are triggered by odor application. At the level of M4/6, suppressed dynamics would correspond to synaptic depression, while at the level of α'3 MBONs increased dynamics may result in postsynaptic depression. Therefore,

747 different learning rules could govern the incorporation or exchange or mobilization of
748 receptors in or out of synapses. The precise molecular and biophysical parameters
749 underlying these plasticity rules are currently unknown and will need to be addressed
750 in the future. One option could include potential exchange of α2 subunits for a receptor
751 complex with higher calcium permeability.
752
Our findings are consistent with the current mushroom body skew model[5],
where the summed MBON output will determine an animal's choice. However, we add
an additional layer, already at the MBON input site. Changes do not happen, as
previously believed, at the presynaptic compartment only, but potentially at both
synaptic compartments. Thus, the power to store (potentially conflicting) information
separately at either the pre- or postsynaptic site, equips the system with additional
flexibility. How precisely pre- to postsynaptic and post- to presynaptic signaling is
regulated will need to be addressed in the future, but will likely involve transsynaptic
signaling routes[61,64]. Importantly, the identified modes of postsynaptic plasticity will
open avenues for investigations looking into pre- versus postsynaptic contributions
during reversal learning, reconsolidation and extinction learning[13,33].
764
765

Methods

Fly genetics

Flies were raised on standard food under standard laboratory conditions unless
stated otherwise (25°C, 65 %, 12-hour light-dark cycle)[7,40]. Driver lines used were
MB011B (Split-Gal4)[10], MB112C[10] (Split-Gal4), MB461B[10] (Split-Gal4), MB027B[15]
(Split-Gal4), R13F02-Gal4[10], OK107-Gal4[4], VT1211-GAL4[7], and R58E02-LexA[26]. We
used the following UAS-nAChR^{RNAi} flies[4,51]: Bloomington stock numbers 28688, 27493,
27671, 31985, 25943, 27251 and 25835. Additionally, we used[41,42] Dlg^{S97}-RNAi as well
as UAS-Dlg^{GFP}, tubP-GAL80^{ts} [48], UAS-Shi^{ts1} [48], 247-dsRed[7], LexAop-CsChrimson,
UAS-GCamp6f[4,13,26], and UAS-SynaptoPhluorin[65]. Gal80^{ts} flies were raised at 18-20°C
and were placed at 32°C 3-5 days before the experiment. Note that complex genotypes
did not always permit usage of MB011B for genetical access to M4/6 neurons
throughout the manuscript. In that case, in order to reduce genetic complexity, we used
VT1211-Gal4.

Behavior

T-maze memory

780　
781　
3-9-day old mixed-sex populations were trained and tested together as
previously described[7]. Odors used were 3-Octanol (OCT, Aldrich) and 4-
Methylcyclohexanol (MCH, Aldrich) diluted in mineral oil (approximately 1:100 for
aversive, 1:1000 for appetitive memory, absolute concentrations were minimally
adjusted to prevent odor bias). For aversive protocols, flies were exposed to the CS+
for 1 minute with 12 1.5 seconds long 120 V electric shocks (interstimulus interval: 3.5
seconds) followed by 45 seconds of air, 1 minute of CS- exposure and another 30
seconds of air. Flies were given 2 minutes to choose between the CS+ and CS- in a
T-Maze during retrieval in the dark. For appetitive conditioning flies were starved for
20 - 24 hours before the experiment. Flies were exposed to the CS- for 2 minutes. After
30 seconds, flies were exposed to the CS+ paired with sugar for 2 minutes followed by
another 30 seconds of air. Performance indices were calculated as described
previously[7]. Time of retrieval is stated in the figures. For Shi[ts] experiments, flies were
kept at 32°C 30 minutes prior to and during training and brought to room temperature
directly afterwards. Room temperature was approximately 23°C. For Fig. 2 and
Supplementary Fig. 2 behavioral data sets from separate experiments were pooled.
Note that 'screening hit' data displayed in Fig. 2a,b and Supplementary Fig. 2a,b were
replotted to allow for comparison of genotypes with the corresponding genetic controls
in Supplementary Fig. 2i-m.

Familiarity learning

801　
Familiarity training was essentially performed as described before[15] with slight
adjustments. Flies were covered in yellow dust (Reactive Yellow 86, Fisher Scientific)[15]
and placed in a cylindrical custom designed chamber. To ensure a constant air stream
we placed the chamber between an air and a vacuum pump (800ml/min). Air
permeable cotton wool was used to close the open ends of the chamber. The air supply
was either connected to pure mineral oil or MCH diluted in mineral oil at a concentration
of 1:50. For switching between odor and mineral oil, a clamp was manually opened
and closed. Video recording was performed at 26 frames per second. For recordings
and analyses we used Python (v3.6) in Anaconda Jupyter Notebook environment.

811　

Imaging

Confocal single photon imaging and receptor quantification

Fixed explant brain imaging. Brains were dissected on ice, fixed in 4% paraformaldehyde (Sigma) for 20 minutes and placed in PBST (0.1% Triton) for 30 minutes followed by washing with PBS for 20 minutes twice. Vectashield was used as mounting medium. Flies were 2-8 day-old females raised at room temperature.

Recording endogenous fluorescence. Imaging was performed using a confocal single photon inverse microscope (Leica SP5/STED) equipped with a 64x oil objective. Laser power and gain were adjusted between experiments, making normalization of the signals necessary. Values for the heatmap in Fig. 4 were normalized to the mean MB fluorescence value to ensure comparability. Voxel size was (height x width x depth) 123 nm x 123 nm x 500 nm. ROIs were drawn manually in ImageJ using the 247-dsRed channel for orientation (Fig. 4a). Heat maps were created in Microsoft Excel. For quantifications following knock-down, the γ5 compartment was normalized to γ4 ((γ5- γ4)/ γ4), and the β'2 to the β'1 compartment ((β'2- β'1)/ β'2) of the same animal. Each 'n' corresponds to one hemisphere.

In vivo two photon imaging of receptor dynamics

Fluorescence recovery after photobleaching (FRAP) experiments were performed *in vivo*. 2-8 day-old flies were anesthetized on ice and mounted in a custom-made chamber. The head capsule was opened under room temperature sugar-free HL3-like saline, and legs were immobilized with wax[7]. Sugar-free HL3-like saline containing 30 units of Papain (Roche) was applied to the head capsule for 8 minutes to digest the brain's glial sheath and facilitate removal. Images were acquired using a multi photon microscope (Nikon) with a 25× water-immersion objective, controlled by Nikon NIS Elements software. Diluted odors (MCH or OCT in mineral oil 1:1000) were delivered on a clean air carrier stream using a 6-channel delivery system (CON electronics). The flies were subjected to experimental conditions including either no odor (air), odor only, odor paired with local dopamine (10 mM) injection via a micropipette, or local dopamine injection only (see Fig. 6b and Supplementary Fig. 6b for experimental protocol schematics). Photobleaching was accomplished using focused, high intensity laser exposure for ~1 minute. Analysis of fluorescence recovery was performed using FIJI. ROIs were manually selected and the percent recovery fluorescence was calculated by subtraction of the post-bleaching baseline

fluorescence and division by the pre-bleaching baseline fluorescence. The percent
recovery values were log-transformed and a linear mixed effects model without
intercept was fit using the interaction between condition and time as fixed effect - to
determine condition-specific differences of the recovery kinetics - and time as random
effect (R package lme4). A linear mixed effect model was used to appropriately model
repeated measures within animals. Significance of recovery of individual conditions
was assessed using the regression coefficients of the condition-time interaction of the
linear mixed model. Differences of recovery between pairs of conditions were tested
using pairwise comparisons of estimated marginal means of the linear mixed model (R
package emmeans). Correction for multiple pairwise comparisons was performed
using Tukey's method.

In vivo *confocal single photon imaging of receptor dynamics and calcium*
transients

3-4 days after eclosure, female flies were prepared as described above and
imaged. Imaging was performed using a SP5 single-photon confocal microscope
(Leica microsystems). Recording frame rate was 3 Hz. For bleaching high laser power
was used focusing on the α'3 compartments for 15-25 seconds. The baseline was
recorded after bleaching, immediately before fixed inter-stimulus interval-training[15].
OCT was presented ten times for a second with a six second pause in between. Odor
delivery (CON electronics) was controlled by the Leica acquisition software. After
training, the same brain plane was recorded for 10 seconds with a pixel size of 200 nm
in time intervals of 0, 5, 10, 15, 20, 30, 60 minutes after training. For control
experiments air only was delivered to the chamber. Images of the same time interval
recordings were averaged and processed in ImageJ. Gaussian blur (σ = 0.5) was
applied for smoothing and ROIs were selected manually.

In vivo *two photon calcium imaging*

To measure odor responses, female 3-6-day old flies expressing UAS-
GCaMP6f and UAS-RNAi to α2 or α5 at the level of M4/6 were tethered under the
multiphoton microscope (Femtonics), essentially as described before[7,66]. 5 alternating
1 second OCT and MCH puffs were applied with 30 seconds in between each
presentation. Fluorescent signals were recorded from dendrites in the β'2 MB
compartment using MESc software (Femtonics) at a frame rate of roughly 31 Hz. ROIs
incorporating the dendritic arbors were manually drawn. Data was processed using a
Savitzky-Golay filter. Further analysis was performed using Matlab. For absolute

training, following protocol was applied: after initial testing for odor responses, flies
were exposed to odor puffs (MCH) twice with a 30-second gap between the
applications. Corresponding odor responses were averaged. Training consisted of
odor application while the fly fed on a sucrose droplet provided by a custom-made
feeding arm[67]. After a two-minute break, two odor puffs with a gap of 30 seconds were
applied. Again, corresponding odor responses were averaged. Areas under the curve
(AUCs) were calculated using Matlab and the first 4 seconds following odor onset were
analyzed in order to cover entire responses. The AUCs pre- and post-training were
normalized to the mean pre-training values of a group respectively. The averaged test
responses pre-training were compared to the average post-training responses using a
paired t-test or Wilcoxon matched-pairs signed rank test.

Explant brain widefield imaging, neurotransmitter application and optogenetics

Postsynaptic plasticity induction: Brains of 3-10 day-old mixed sex flies were
dissected on ice. Flies expressed CsChrimson[tdTomato] under control of R58E02-LexA
and UAS-GCaMP6f (and α2 RNAi) under control of MB011B. The head capsule and
sheath were removed in carbogenated solution (103 mM NaCl, 3 mM KCl, 5 mM N-
Tris, 10 mM trehalose, 10 mM glucose, 7 mM sucrose, 26 mM NaHCO$_3$, 1 mM
NaH$_2$PO4, 1.5 mM CaCl$_2$, 4 mM MgCl$_2$, 295 mOsm, pH 7.3) with forceps. The brain
was subsequently perfused with carbogenated solution containing TTX (2 µM; 20 ml /
10 min flow speed) and imaged using an Olympus MX51WI wide field microscope with
a 40x Olympus LUMPLFLN objective and an Andor iXON Ultra camera controlled by
Solis software. An Olympus U25ND25 light filter was placed in the beam path to
minimize baseline CsChrimson activation. A custom designed glass microcapillary was
loaded with uncarbongenated solution containing 0.1 mM acetylcholine and
maneuvered to the M6 dendrites. The injection pressure of a P25-1-900 picospritzer
was calibrated between 3-8 psi (depending on initial calcium responses). Each local
acetylcholine application spanned 15 ms with a 4 s inter-injection interval.

Three pulses of acetylcholine followed by a 2-3 minute break were recorded
after which the optogenetic response was assessed by applying 2 seconds red light
pulses with an inter-red light-interval of 4 seconds. 3 acetylcholine pulses were
recorded followed by either 5 acetylcholine injections, 5 red light pulses or both paired.
For paired training both stimuli began simultaneously and the acetylcholine injection
lasted for 15 ms (and gave rise to a calcium transients typically lasting > 1 seconds,
please see example in Fig. 3), while the paired red light pulse lasts for 2 seconds,
allowing for maximal temporal overlap. This process was repeated 5x, with a 4 second

break between trials. Following the training trial, 3 final acetylcholine test injections
were applied after 1 minute. For analysis, the first of the 3 acetylcholine injections was
always discarded because of initial dilution of the capillary tip and the remaining 2 peak
intensities were averaged.

All peaks within an experiment were quantified relative to the fluorescence
baseline that we calculated for pre and post training acetylcholine responses.
Baselines were set independently for each pre- and post-training recording using the
polynomial interpolation function in NOSA[68]. For investigating α2 knock-down, only the
paired condition was tested.

For controls not expressing CsChrimson, we used VT1211-Gal4 driving UAS-
GCaMP6f, instead of MB011B, for technical reasons. This was combined with either
expression of R58E02-LexA or LexAop-CsChrimson[tdTomato]. Only paired training was
investigated in this context.

Excitability of Kenyon cell axons: To test whether KC axons were excited by
focal acetylcholine injections at the level of M4/6 dendrite innervation, either UAS-
SynaptoPhluorin or UAS-GCaMP6f were expressed under the control of OK107-Gal4.
Following the acetylcholine injection experiment, the capillary was exchanged with a
capillary containing the same solution with additional 300-400 mM KCl, to evaluate
tissue health (not shown for GCaMP6f imaging). To pick up potentially small changes,
we increased the injection pressure to 8-14 Psi and the injection time to 150-225 ms
(GCaMP6f, 8s inter-injection interval and 3 consecutive injections) and 300-525 ms
(SynaptoPhluorin, with an 8s inter-injection interval and 3 consecutive injections).

Images were analyzed using NOSA[40,68] and GraphPad Prism.

Statistics

Statistical analyses were performed as stated in the previous methods sections
and figure legends. Data were always tested for normality using a Shapiro-Wilk test. If
normally distributed, data were analyzed using ANOVA followed by post-hoc test or a
(paired) t-test. If not normal, we used a Kruskal Wallis followed by post-hoc test, or a
Wilcoxon matched-pairs signed rank test.

Tagged receptor subunits

946   All subunits were tagged using CRISPR technology and motifs previously
947   described[69]. Further details will be published separately and can be requested from
948   the corresponding author.

Connectome analysis

In Supplementary Fig. 9, we analyzed the partial connectome of the female
adult   fly   brain   (hemibrain   v1.2.1)[62]   using   the   neuprint-python   package
(https://github.com/connectome-neuprint/neuprintpython). To investigate the synaptic
relationship between KC presynapses and MBON postsynaptic sites, we first identified
KCs   with   the   status   'Traced'   in   the   connectome.
Second, for each MBON of interest, we identified the relevant KCs connected to it
(which is a subset of the count in the previous step). Third, for each KC identified, we
selected each presynaptic terminal (x,y,z locations of synapses) connected to the
MBON of interest, and for each of these presynapses, we identified all synaptic
partners residing on the postsynaptic side. Fourth, for the MBON of interest, we
counted the number of postsynaptic connections per individual presynapses (that also
contain the specific MBON, Supplementary Fig. 9 d,e,f). Finally, we identified the
composition of all neurons identified as co-postsynaptic partners of KC to MBON
synapses.

Data availability

967   The datasets generated during and/or analysed during the current study are available
968   from the corresponding author on reasonable request.

Code availability

972   The code used in this study is available at
973   https://github.com/jagannathancolabs/2022_postsynapticplasticity

Acknowledgements

We thank Anatoli Ender, Johannes Felsenberg, Davide Raccuglia, Lisa
Scheunemann, Stephan Sigrist, Uli Thomas and Scott Waddell for comments on the
manuscript, Stephan Sigrist and Uli Thomas for reagents, the Janelia and Vienna fly
projects, and the Bloomington Stock Center and VDRC for fly lines as well as Daisuke
Hattori and Yoshi Aso for help with the familiarity experiments. Multiphoton and single
photon confocal imaging was partially performed using microscopes of the AMBIO and
NWFZ core facilities of the Charité. **Funding:** Funded by the Deutsche
Forschungsgemeinschaft (DFG, German Research Foundation) under Germany´s
Excellence Strategy – EXC-2049 – 390688087, the Emmy Noether Programme, TP
A27 of SFB958 (184695641), and TP A07 of SFB1315 (327654276) to DO. S.R.J is
supported by the Walter Benjamin Programme of the DFG. **Author contribution**:
Conceptualization, C.P., M-M.H., E.R., S.R., D.L., Y-C.C., D.O., Investigation, C.P., M-
M.H., E.R. D.L., R.S.-G., C.R., S.R., Y-C.C., I.B., T.F., G.L., S.R.J.; Resources, D.O.;
Writing, D.O., C.P.; Instrumentation: JR; Comments, M-M.H., E.R. D.L., R.S.-G., S.R.,
Y-C.C., S.R.J..

Competing interests: Authors declare no competing interests.

Supplemental Information:
Supplementary Figure 1. Permissive temperature controls accompanying Fig. 1.

Supplementary Figure 2. Genetic controls and alternate data display accompanying
Fig. 2.

1002

1003 Supplementary Figure 3. Control experiments for Fig. 3.

1004

1005 Supplementary Figure 4. Detailed distribution of α subunits in the MB accompanying
1006 Fig. 4.

1007

1008 Supplementary Figure 5. Additional data accompanying Fig. 5.

1009

1010 Supplementary Fig. 6. Receptor subunit recovery, accompanying Fig. 6.

1011

Supplementary Fig. 7. Dlg^{GFP} FRAP, accompanying Fig. 7.

1013

Supplementary Fig. 8. Additional ethograms, accompanying Fig. 8.

1015

Supplementary Fig. 9. Input analysis of KC to MBON synapses, referred to in legend
of Fig. 9.

1018

References

1. Glanzman, D. L. Common Mechanisms of Synaptic Plasticity in Vertebrates and Invertebrates. Curr Biol 20, R31–R36 (2010).

2. Korte, M. & Schmitz, D. Cellular and System Biology of Memory: Timing, Molecules, and Beyond. Physiol Rev 96, 647–93 (2016).

3. Nicoll, R. A. A Brief History of Long-Term Potentiation. Neuron 93, 281–290 (2017).

4. Barnstedt, O. et al. Memory-Relevant Mushroom Body Output Synapses Are Cholinergic. Neuron 89, 1237–47 (2016).

5. Owald, D. & Waddell, S. Olfactory learning skews mushroom body output pathways to steer behavioral choice in Drosophila. Curr Opin Neurobiol 35, 178–84 (2015).

6. Shomrat, T. et al. Alternative sites of synaptic plasticity in two homologous "fan-out fan-in" learning and memory networks. Curr Biology Cb 21, 1773–82 (2011).

7. Owald, D. et al. Activity of Defined Mushroom Body Output Neurons Underlies Learned Olfactory Behavior in Drosophila. Neuron 86, 417–427 (2015).

8. Aso, Y. et al. Nitric oxide acts as a cotransmitter in a subset of dopaminergic neurons to diversify memory dynamics. Elife 8, e49257 (2019).

9. Aso, Y. et al. Mushroom body output neurons encode valence and guide memory-based action selection in Drosophila. Elife 3, e04580 (2014).

10. Aso, Y. et al. The neuronal architecture of the mushroom body provides a logic for associative learning. Elife 3, e04577 (2014).

11. Bouzaiane, E., Trannoy, S., Scheunemann, L., Plaçais, P.-Y. & Preat, T. Two independent mushroom body output circuits retrieve the six discrete components of Drosophila aversive memory. Cell Reports 11, 1280–92 (2015).

12. Cohn, R., Morantte, I. & Ruta, V. Coordinated and Compartmentalized Neuromodulation Shapes Sensory Processing in Drosophila. Cell 163, 1742–55 (2015).

13. Felsenberg, J. et al. Integration of Parallel Opposing Memories Underlies Memory Extinction. Cell 175, 709-722.e15 (2018).

1060   14. Felsenberg, J., Barnstedt, O., Cognigni, P., Lin, S. & Waddell, S. Re-evaluation of
1061   learned information in Drosophila. Nature 544, 240–244 (2017).
1062
1063   15. Hattori, D. et al. Representations of Novelty and Familiarity in a Mushroom Body
1064   Compartment. Cell 169, 956-969.e17 (2017).
1065
1066   16. Ichinose, T. et al. Reward signal in a recurrent circuit drives appetitive long-term
1067   memory formation. Elife 4, e10719 (2015).
1068
1069   17. Lewis, L. P. C. et al. A Higher Brain Circuit for Immediate Integration of Conflicting
1070   Sensory Information in Drosophila. Curr Biology Cb 25, 2203–14 (2015).
1071
1072   18. Pai, T.-P. et al. Drosophila ORB protein in two mushroom body output neurons is
1073   necessary for long-term memory formation. P Natl Acad Sci Usa 110, 7898–903
1074   (2013).
1075
1076   19. Perisse, E. et al. Aversive Learning and Appetitive Motivation Toggle Feed-
1077   Forward Inhibition in the Drosophila Mushroom Body. Neuron 90, 1086–99 (2016).
1078
1079   20. Séjourné, J. et al. Mushroom body efferent neurons responsible for aversive
1080   olfactory memory retrieval in Drosophila. Nat Neurosci 14, 903–10 (2011).
1081
1082   21. Plaçais, P.-Y., Trannoy, S., Friedrich, A. B., Tanimoto, H. & Preat, T. Two pairs of
1083   mushroom body efferent neurons are required for appetitive long-term memory
1084   retrieval in Drosophila. Cell Reports 5, 769–80 (2013).
1085
1086   22. Hige, T., Aso, Y., Modi, M. N., Rubin, G. M. & Turner, G. C. Heterosynaptic
1087   Plasticity Underlies Aversive Olfactory Learning in Drosophila. Neuron 88, 985–98
1088   (2015).
1089   23. Takemura, S. et al. A connectome of a learning and memory center in the adult
1090   Drosophila brain. Elife 6, e26975 (2017).
1091
1092   24. Bilz, F., Geurten, B. R. H., Hancock, C. E., Widmann, A. & Fiala, A. Visualization
1093   of a Distributed Synaptic Memory Code in the Drosophila Brain. Neuron 106, 963-
1094   976.e4 (2020).
1095
1096   25. Boto, T., Louis, T., Jindachomthong, K., Jalink, K. & Tomchik, S. M. Dopaminergic
1097   modulation of cAMP drives nonlinear plasticity across the Drosophila mushroom body
1098   lobes. Curr Biology Cb 24, 822–31 (2014).
1099
1100   26. Handler, A. et al. Distinct Dopamine Receptor Pathways Underlie the Temporal
1101   Sensitivity of Associative Learning. Cell 178, 60-75.e19 (2019).
1102
1103   27. Ehmann, N., Owald, D. & Kittel, R. J. Drosophila active zones: from molecules to
1104   behaviour. Neurosci Res 127, 14–24 (2017).
1105
1106   28. Dubnau, J., Grady, L., Kitamoto, T. & Tully, T. Disruption of neurotransmission in
1107   Drosophila mushroom body blocks retrieval but not acquisition of memory. Nature 411,
1108   476–480 (2001).
1109
1110   29. Krashes, M. J., Keene, A. C., Leung, B., Armstrong, J. D. & Waddell, S. Sequential
1111   Use of Mushroom Body Neuron Subsets during Drosophila Odor Memory Processing.
1112   Neuron 53, 103–115 (2007).
1113

1114　30. McGuire, S. E., Le, P. T. & Davis, R. L. The Role of Drosophila Mushroom Body
1115　Signaling in Olfactory Memory. Science 293, 1330–1333 (2001).
1116
1117　31. Schwaerzel, M., Heisenberg, M. & Zars, T. Extinction Antagonizes Olfactory
1118　Memory at the Subcellular Level. Neuron 35, 951–960 (2002).
1119
1120　32. Kandel, E. R., Dudai, Y. & Mayford, M. R. The molecular and systems biology of
1121　memory. Cell 157, 163–86 (2014).
1122
1123　33. Lüscher, C. & Malenka, R. C. Drug-evoked synaptic plasticity in addiction: from
1124　molecular changes to circuit remodeling. Neuron 69, 650–63 (2011).
1125
1126　34. Won, S., Levy, J. M., Nicoll, R. A. & Roche, K. W. MAGUKs: multifaceted synaptic
1127　organizers. Curr Opin Neurobiol 43, 94–101 (2017).
1128
1129　35. Thompson, A. J., Lester, H. A. & Lummis, S. C. R. The structural basis of function
1130　in Cys-loop receptors. Q Rev Biophys 43, 449–99 (2010).
1131
1132　36. Dent, J. A. Advances in Experimental Medicine and Biology. Adv Exp Med Biol
1133　683, 11–23 (2010).
1134
1135　37. Ihara, M. et al. Cofactor-enabled functional expression of fruit fly, honeybee, and
1136　bumblebee nicotinic receptors reveals picomolar neonicotinoid actions. Proc National
1137　Acad Sci 117, 16283–16291 (2020).
1138
1139　38. Lansdell, S. J., Collins, T., Goodchild, J. & Millar, N. S. The Drosophila nicotinic
1140　acetylcholine receptor subunits Dα5 and Dα7 form functional homomeric and
1141　heteromeric ion channels. Bmc Neurosci 13, 73 (2012).
1142
1143　39. Eadaim, A., Hahm, E.-T., Justice, E. D. & Tsunoda, S. Cholinergic Synaptic
1144　Homeostasis Is Tuned by an NFAT-Mediated α7 nAChR-Kv4/Shal Coupled Regulatory
1145　System. Cell Reports 32, 108119 (2020).
1146
1147　40. Raccuglia, D. et al. Network-Specific Synchronization of Electrical Slow-Wave
1148　Oscillations Regulates Sleep Drive in Drosophila. Curr Biology Cb 29, 3611-3621.e3
1149　(2019).
1150
1151　41. Bachmann, A. et al. Cell type-specific recruitment of Drosophila Lin-7 to distinct
1152　MAGUK-based protein complexes defines novel roles for Sdt and Dlg-S97. J Cell Sci
1153　117, 1899–1909 (2004).
1154
1155　42. Soukup, S.-F., Pocha, S. M., Yuan, M. & Knust, E. DLin-7 Is Required in
1156　Postsynaptic Lamina Neurons to Prevent Light-Induced Photoreceptor Degeneration
1157　in Drosophila. Curr Biol 23, 1349–1354 (2013).
1158
1159　43. Crittenden, J. R., Skoulakis, E. M., Han, K. A., Kalderon, D. & Davis, R. L. Tripartite
1160　mushroom body architecture revealed by antigenic markers. Learn Mem Cold Spring
1161　Harb N Y 5, 38–51 (1998).
1162
1163　44. Eichler, K. et al. The complete connectome of a learning and memory centre in an
1164　insect brain. Nature 548, 175–182 (2017).
1165
1166　45. Stuart, G. J. & Spruston, N. Dendritic integration: 60 years of progress. Nat
1167　Neurosci 18, 1713–1721 (2015).
1168

1169 46. Tully, T., Preat, T., Boynton, S. C. & Vecchio, M. D. Genetic dissection of
1170 consolidated memory in Drosophila. Cell 79, 35–47 (1994).
1171
1172 47. Trannoy, S., Redt-Clouet, C., Dura, J.-M. & Preat, T. Parallel Processing of
1173 Appetitive Short- and Long-Term Memories In Drosophila. Curr Biol 21, 1647–1653
1174 (2011).
1175
1176 48. Perisse, E. et al. Different kenyon cell populations drive learned approach and
1177 avoidance in Drosophila. Neuron 79, 945–56 (2013).
1178
1179 49. Honegger, K. S., Campbell, R. A. A. & Turner, G. C. Cellular-resolution population
1180 imaging reveals robust sparse coding in the Drosophila mushroom body. J Neurosci
1181 Official J Soc Neurosci 31, 11772–85 (2011).
1182
1183 50. Zhao, X., Lenek, D., Dag, U., Dickson, B. & Keleman, K. Persistent activity in a
1184 recurrent circuit underlies courtship memory in Drosophila. Elife 7, e31425 (2018).
1185
1186 51. Cervantes-Sandoval, I., Phan, A., Chakraborty, M. & Davis, R. L. Reciprocal
1187 synapses between mushroom body and dopamine neurons form a positive feedback
1188 loop required for learning. Elife 6, e23789 (2017).
1189
1190 52. Ueno, K. et al. Carbon Monoxide, a Retrograde Messenger Generated in
1191 Postsynaptic Mushroom Body Neurons, Evokes Noncanonical Dopamine Release. J
1192 Neurosci Official J Soc Neurosci 40, 3533–3548 (2020).
1193
1194 53. Ueno, K. et al. Coincident postsynaptic activity gates presynaptic dopamine
1195 release to induce plasticity in Drosophila mushroom bodies. Elife 6, e21076 (2017).
1196
1197 54. Bliss, T. V. & Lomo, T. Long-lasting potentiation of synaptic transmission in the
1198 dentate area of the anaesthetized rabbit following stimulation of the perforant path. J
1199 Physiology 232, 331–56 (1973).
1200
1201 55. Greger, I. H., Watson, J. F. & Cull-Candy, S. G. Structural and Functional
1202 Architecture of AMPA-Type Glutamate Receptors and Their Auxiliary Proteins. Neuron
1203 94, 713–730 (2017).
1204
1205 56. Aso, Y. & Rubin, G. M. Dopaminergic neurons write and update memories with
1206 cell-type-specific rules. Elife 5, e16135 (2016).
1207
1208 57. Hagins, W. A., Penn, R. D. & Yoshikami, S. Dark Current and Photocurrent in
1209 Retinal Rods. Biophys J 10, 380–412 (1970).
1210
1211 58. Ito, M. Cerebellar Long-Term Depression: Characterization, Signal Transduction,
1212 and Functional Roles. Physiol Rev 81, 1143–1195 (2001).
1213
1214 59. Neff, R. A., Gomez-Varela, D., Fernandes, C. C. & Berg, D. K. Postsynaptic
1215 scaffolds for nicotinic receptors on neurons. Acta Pharmacol Sin 30, 694–701 (2009).
1216
1217 60. Ackermann, F., Waites, C. L. & Garner, C. C. Presynaptic active zones in
1218 invertebrates and vertebrates. Embo Rep 16, 923–38 (2015).
1219
1220 61. Owald, D. & Sigrist, S. J. Assembling the presynaptic active zone. Curr Opin
1221 Neurobiol 19, 311–8 (2009).
1222

1223 62. Scheffer, L. K. et al. A connectome and analysis of the adult Drosophila
1224 centralbrain. Elife 9, e57443 (2020).
1225
1226 63. Goldberg, H., Tamas, G., Aronov, D. & Yuste, R. Calcium Microdomains in Aspiny
1227 Dendrites. Neuron 40, 807-821 (2003).
1228
1229 64. Tang, A.-H. et al. A transsynaptic nanocolumn aligns neurotransmitter release to
1230 receptors. Nature 536, 210-214 (2016).
1231
1232 65. Lin, A. C., Bygrave, A. M., Calignon, A. de, Lee, T. & Miesenböck, G. Sparse,
1233 decorrelated odor coding in the mushroom body enhances learned odor discrimination.
1234 Nat Neurosci 17, 559–568 (2014).
1235
1236 66. Böhme, M. A. et al. Rapid active zone remodeling consolidates presynaptic
1237 potentiation. Nat Commun 10, 1085 (2019).
1238
1239 67. Lin, S. et al. Neural correlates of water reward in thirsty Drosophila. Nat Neurosci
1240 17, 1536–1542 (2014).
1241
1242 68. Oltmanns, S. et al. NOSA, an Analytical Toolbox for Multicellular Optical
1243 Electrophysiology. Front Neurosci-switz 14, 712 (2020).
1244
1245 69. Raghu, S. V., Joesch, M., Sigrist, S. J., Borst, A. & Reiff, D. F. Synaptic
1246 Organization of Lobula Plate Tangential Cells in Drosophila: Dα7 Cholinergic
1247 Receptors. J Neurogenet 23, 200–209 (2009).
1248
1249 70. Stahl, A. *et al.* Associative learning drives longitudinally graded presynaptic
1250 plasticity of neurotransmitter release along axonal compartments. *Elife* **11**, e76712
1251 (2022).

1252
1253
1254
1255

1256 **Figures**

1257

1258 **Figure 1**

1259

1260

Figure 1: KC neurotransmitter release is required for the acquisition of appetitive
memories.

a-c) Flies expressing temperature-sensitive Shibire (Shi) within KCs or MBONs are trained
at restrictive temperature (32°C), and subsequently placed at permissive temperature (23°C)
throughout the consolidation and retrieval phase. Memory performance was tested 30 minutes
after training at permissive temperature. Shi blocks neurotransmitter release at 32°C.

a) Block of neurotransmitter release from KCs (driver line R13F02-Gal4) during training
does not impact 30 min aversive memory performance. Bar graphs: mean ± SEM; n = 7 – 8;
one-way ANOVA followed by Tukey's test (p > 0.05).

b) Block of neurotransmitter release from KCs (driver line R13F02-Gal4) during training
impairs 30 min appetitive memory performance. Bar graphs: mean ± SEM; n = 10 – 16; Kruskal-
Wallis followed by Dunn's test (p < 0.05), * = p < 0.05.

c) Block of neurotransmitter release from M4/6 MBONs (driver line MB011B) during
training does not impact 30 min appetitive memory performance. Bar graphs: mean ± SEM; n
= 14 – 24; one-way ANOVA followed by Tuckey's test (p > 0.05).

Also see Supplementary Fig. 1 for further information.

1277

1278

Figure 2

1280

1281

Figure 2: Specific nAChR α subunits are needed for specific memories in M4/6 neurons.

a)   Immediate appetitive memories are impaired following RNAi knock-down of the α5 nAChR subunit in M4/6 MBONs (driver line MB011B). Bar graphs: mean ± SEM; n = 8 – 13, for controls: n = 20; one-way ANOVA followed by Dunnett's test (p < 0.05), * = p < 0.05.

Note: data depicted correspond to initial screen, please see Supplementary Fig. 2i for alternate
display including all genetic controls.

b) RNAi knock-down of the α1, α2, α5 nAChR subunits or Dlg in M4/6 MBONs (driver line
MB011B) impair 3-hour appetitive memories. Bar graphs: mean ± SEM; n = 12 – 26, for
controls: n = 38; Kruskal-Wallis followed by Dunn's test ($p < 0.05$), * = $p < 0.05$.

Note: data depicted correspond to initial screen, please see Supplementary Fig. 2j-m for
alternate display including all genetic controls.

c) Immediate aversive learning is not impaired by RNAi knock-down of any subunit in
M4/6 MBONs (driver line MB011B). Bar graphs: mean ± SEM; n = 6 – 8, for controls: n = 12;
Kruskal-Wallis followed by Dunn's test ($p > 0.05$).

d) 3-hour aversive memory is not affected by knock-down of α subunits with the exception
of α7 (driver line MB011B). Bar graphs: mean ± SEM; n = 21 – 32, for controls: n = 61. Kruskal-
Wallis followed by Dunn's test ($p < 0.05$), * = $p < 0.05$.

e) RNAi knock-down of the α5 subunit in M4/6 MBONs (driver line VT1211-Gal4) is
suppressed during development using Gal80^{ts}. 3-5 days before the experiment RNAi-knock-
down was induced. Immediate memory is significantly impaired. Bar graphs: mean ± SEM;
n = 6 – 7; Kruskal-Wallis followed by Dunn's test ($p < 0.05$), * = $p < 0.05$.

f) RNAi knock-down of the α2 subunit in M4/6 MBONs (driver line VT1211-Gal4) is
suppressed during development using Gal80^{ts}. 3-5 days before the experiment RNAi knock-
down was induced. 3-hour memories are significantly impaired. Bar graphs: mean ± SEM; n =
16 – 17; one-way ANOVA followed by Tukey's test ($p < 0.05$), * = $p < 0.05$.

g) RNAi knock-down of the α5 subunit in M4/6 MBONs (driver line VT1211-Gal4) is
suppressed during development using Gal80^{ts}. 3-5 days before the experiment RNAi knock-
down was induced. 3-hour memories are significantly impaired. Bar graphs: mean ± SEM; n =
25 – 27; one-way ANOVA followed by Tukey's test ($p < 0.05$), * = $p < 0.05$.

h) RNAi knock-down of Dlg in M4/6 MBONs (driver line VT1211-Gal4) is suppressed
during development using Gal80^{ts}. 3-5 days before the experiment RNAi knock-down was
induced. 3-hour memories are significantly impaired. Bar graphs: mean ± SEM; n = 8 – 11; one-
way ANOVA followed by Tukey's test ($p < 0.05$), * = $p < 0.05$.

Also see Supplementary Fig. 2 for further information.

1316
1317

1318 **Figure 3**

1319
1320
Figure 3: Induction of postsynaptic plasticity bypassing the presynapses.

a) Connectivity scheme of MB output synapses. Cholinergic KCs and dopaminergic
neurons are presynaptic to M6 MBONs. Only connections relevant for this protocol are shown
for simplicity. Red light pulses trigger release of dopamine (DA) from dopaminergic neurons
(R58E02-LexA > lexAop-CsChrimson[tdTomato]), while KC input is circumvented and mimicked by
focal acetylcholine (ACh; 0.1 mM) injections to M6 dendrites in an explant brain preparation.
Postsynaptic responses at the level of M6 are measured using GCaMP6f (MB011B > UAS-
GCaMP6f). TTX in the bath suppresses feedback signaling and overall network activity within
the circuit.

b) Training scheme (top). Baseline responses to ACh application were initially established
(pre). Subsequent training protocols consist of either pairing ACh application with simultaneous
activation of dopaminergic neurons (purple connection lines), activation of dopaminergic

neurons ('red light only, red connection lines), or ACh only (blue, connection lines). This is
followed by a test trial (post) through ACh application.

c) Averaged traces of axonal KC calcium changes induced by focal ACh injections. No
apparent transients are observable; n = 7. Line is mean ± SEM.

d) Sample trace of dendritic M6 calcium changes induced by focal ACh injections.

e) Sample trace of dendritic M6 calcium changes induced by red-light pulses.

f-h) Above: sample calcium traces in response to ACh injections recorded from M6
dendrites pre (black traces) and post (colored traces) training. Below: peak quantification.

f) Changes in acetylcholine-evoked calcium transients; comparison of mean peaks pre-
and post 'ACh only' training. Before-after plots and bar graphs (mean); n = 13, ratio paired t-
test.

g) Changes in acetylcholine-evoked calcium transients; comparison of mean peaks pre-
and post 'red light only' training. Before-after plots and bar graphs (mean); n = 10, ratio paired
t-test.

h) Changes in acetylcholine-evoked calcium transients; comparison of mean peaks pre-
and post 'paired' training. Before-after plots and bar graphs (mean); n = 18, ratio paired t-test,
* = $p < 0.05$.

Also see Supplementary Fig. 3 for further information

Figure 4

Figure 4: nAChR α subunit localization throughout the MB: MBON-specific RNAi alters subunit distribution.

a) Representative images of the GFP-tagged nAChR subunits α2 and α5 as well as Dlg. Individual images displayed here are taken from different animals. For other subunits see Supplementary Fig. 4. Scale-Bar: 20 µm. Left: merge of α subunit signal (green) with MB compartments marked with 247-dsRed (magenta). Compartments are indicated by dashed-lines. Top row: γ compartments; middle row: α, β', β and γ compartments, bottom row: α', α and β compartments.

b) Quantification of all GFP-tagged α receptors (except for the α3 subunit). GFP signals for the indicated MB compartments are relative to the mean intensity of the GFP signal of the complete MB; n = 7 – 18.

c) Knock-down of Dlg or α2 in M4/6 neurons (driver line VT1211-Gal4) significantly reduces the α2^{GFP} fluorescence within the γ5 compartment (relative to unmanipulated γ4). Bar graph: normalized mean ± SEM; n = 8 – 20; Kruskal-Wallis followed by Dunn's test (p < 0.05), * = p < 0.05.

d) Knock-down of either the α2 or the α5 nAChR subunit or Dlg in M4/6 neurons (driver line VT1211-Gal4) decreases the relative fluorescence signal of α2^{GFP} in the β'2 compartment (relative to unmanipulated β'1). Bar graph: normalized mean ± SEM; n = 9 – 20; one-way ANOVA followed by Dunnett's test (p < 0.05), * = p < 0.05.

e) Knock-down of α5 in M4/6 neurons (driver line VT1211-Gal4) decreases the α5^{GFP} signal in the γ5 compartment (relative to unmanipulated γ4). Bar graph: normalized mean ± SEM; n = 9 –28; Kruskal-Wallis followed by Dunn's test (p < 0.05), * = p < 0.05.

f) α5^{GFP} fluorescence is significantly decreased in the β'2 compartment (relative to unmanipulated β'1) after knock-down of α5 in M4/6 neurons (driver line VT1211-Gal4). Bar graph: normalized mean ± SEM; n = 10 – 28; Kruskal-Wallis followed by Dunn's test (p < 0.05), * = p < 0.05.

Also see Supplementary Fig. 4 for further information.

Figure 5

Figure 5: α2 is required for learning-associated plasticity *in vivo*.

a) Scheme indicating the dendritic imaging area **(b-f)** at the level of the β'2 compartment.

b) Averaged traces of GCaMP6f responses to OCT from control (black), α2 subunit RNAi (blue) and α5 subunit RNAi (red; driven in M4/6 respectively; VT1211-Gal4 as driver line) flies. Solid traces are mean, shaded areas SEM; n = 10 – 12. Odor exposure is indicated by bar.

c) Area under curve quantifications of averaged odor responses show significantly elevated odor responses to OCT following α5 knock-down in M4/6 neurons (driver line VT1211-Gal4). Mean ± SEM; n = 10 – 12; Kruskal-Wallis followed by Dunn's test (p < 0.05); * = p < 0.05.

d) Control flies show a significant increase between the first and the fifth response to OCT. Before-after plots; n = 10; paired t-test; * = p < 0.05.

e) α2 RNAi flies show no difference between the first and fifth odor response to OCT. nAChR subunit RNAi is driven in M4/6 neurons (driver line VT1211-Gal4). Before-after plots; n = 12; Wilcoxon matched-pairs signed rank test.

f)   α5 RNAi flies show a significant decrease in calcium transients over the course of
consecutive odor exposures. nAChR subunit RNAi is driven in M4/6 neurons (driver line
VT1211-Gal4). Before-after plots; n = 12; paired t-test; * = $p < 0.05$.

g)   Control flies show a significant decrease in GCaMP6f responses to MCH (trained odor)
following absolute training (axonal imaging). Before-after plots and bar graphs (mean); n = 6;
Wilcoxon matched-pairs signed rank test; * = $p < 0.05$.

h)   α2 RNAi flies show no significant decrease in GCaMP6f response to MCH (trained
odor) following absolute training (driver line VT1211-Gal4, axonal imaging). Before-after plots
and bar graphs (mean); n = 8; Wilcoxon matched-pairs signed rank test.

Also see Supplementary Fig. 5 for further information.

Figure 6

Figure 6: α2 nAChR subunits dynamically rearrange.

a) Scheme of site of dopamine injection during fluorescence recovery after photobleaching (FRAP) experiments at the level of the KC-MBON synapses of the β'2 compartment.

b) FRAP experimental protocol. After bleaching the baseline picture was taken followed by dopamine injection, odor presentation and odor presentation simultaneously with dopamine injection in the same fly. Fluorescence recovery was monitored at the 10- and 20-minute time points.

c) Example image of α2GFP expression; white dashed box shows the β'2 output zone; dopamine injection pipette (with Texas Red) is labelled in magenta. Scale bar: 10 μM.

d) Regression of fluorescence recovery after photobleaching following MCH exposure (blue line), MCH exposure simultaneously with dopamine (DA) injection (purple line), dopamine injection alone (red line).

e) Regression coefficient. Bar graphs: regression coefficients of recovery kinetics ± standard error of regression; n = 9 – 10, linear mixed effects model followed by pairwise comparison from estimated marginal trends. * = p < 0.05.

Also see Supplementary Fig. 6 for further experiments.

Figure 7

1432
1433

Figure 7: Non-associacive plasticity alters postsynaptic α2 subunit receptor dynamics.

a) Training scheme indicating odor application, bleaching and imaging time points. MCH
was given 10 times for 1 second with a pause of 6 seconds in-between. Images were taken

after training in absence of odor immediately afterwards and 5, 10, 15, 20, 30 and 60 minutes
later.

b) Top: calcium peaks in response to odor stimuli of presynaptic KCs (MB369B as driver
line) and adjacent postsynaptic MBONs (driver line: MB027B). Individual calcium responses to
trials 1 and 10 for MBONs (orange lines) and KCs (blue lines). Bottom: Averaged calcium
responses to odor stimuli of presynaptic KCs and postsynaptic MBONs of trials 2 - 4 and 8 - 10
respectively. Responses decrease at the level of MBONs but not at the level of KCs over ten
trials. Box plots are median and 75 % quartiles; n = 15; Kruskal-Wallis followed by Dunn's test
($p < 0.05$), * = $p < 0.05$.

c) Scheme of α'3 compartment analyzed and representative α5^{GFP} fluorescent image
(smoothed). Scale bar: 10 µm.

d) Example images of α2^{GFP} FRAP experiment at the level of the α'3 compartment at
specific time points before and after training. Top row, after training; bottom row: control
settings. Scale bar: 2 µm.

e) FRAP of α2^{GFP} nAChR subunit in the α'3 compartment after odor presentation. α2^{GFP}
shows significant recovery following odor training compared to the controls. Recovery rate is
normalized to the baseline recorded after selective bleaching of the α'3 MB compartment. Box
plots are median and 75 % quartiles; n = 4 – 6; multiple t - tests with Sidak-Bonferroni correction,
* = $p < 0.05$.

f) FRAP of α5^{GFP} subunit in the α'3 compartment after odor presentation. α5^{GFP} did not
show significant recovery compared to the controls. Recovery rate is normalized to the baseline
recorded after selective bleaching of α'3 MB compartment. Box plots are median and 75 %
quartiles; n = 5 – 7, multiple t - tests with Sidak-Bonferroni correction.

g) FRAP of α2^{GFP} nAChR subunit in the α'3 compartment after odor presentation and
knockdown of either the α2 or α5 subunit in the α'3 MBON (driver line MB027B). α2^{GFP} shows
no recovery 30 min after odor training in α2 or α5 knockdown animals compared to the controls.
Recovery rate is normalized to the baseline recorded after selective bleaching of the α'3 MB
compartment. Box plots are minimum value to maximum value; n = 4 – 5; Kruskal-Wallis
followed by Dunn's test ($p < 0.05$), * = $p < 0.05$.
1466

1467 Also see Supplementary Fig. 7 for further information.

Figure 8

Figure 8: α2 and α5 nAChR subunits are required for non-associative familiarity learning

at the level of α'3 MBONs.

a)　　　Knock-down of α nAChR subunits at the level of α'3 MBONs alters odor familiarity learning and the probability to stop grooming. α2 RNAi knock-down delays familiarity learning effects to novel odors. α5 RNAi knock-down flies do not show a novelty response at all.

b)   Grooming behavior response of dusted flies following the repeated presentations of a
novel odor (MCH). Ethogram of grooming behavior (blue) during ten intervals of odor
exposures. Horizontal lines in each trial correspond to a single experimental fly within a trial
group. Not grooming (grey) flies can further be categorized between pausing and wandering
(see Supplementary Figure 8). n = 15.

c)   Knock-down of α2 subunit in α'3 MBONs (driver line MB027B) impairs odor familiarity
learning significantly by showing a higher probability to terminate grooming responses during
the learning period. The learning period is defined as the odor exposure rounds following the
first exposure). Bottom graph: non-linear representation of grooming flies over ten training trials.
Note that α5 behavioral responses are best described by linear representation. Box plots are
median and 75 % quartiles; n = 9, one-way ANOVA followed by Dunnett's test ($p < 0.05$) * = p
< 0.05.

d)   Knock-down of α5 subunits in α'3 MBONs (driver line MB027B) leads to an increased
probability to start grooming earlier. 'Grooming' is defined here as constantly grooming during
2 and 3 seconds after odor delivery. Bottom graph, non-linear representation of grooming flies
over ten training trials. Box plots are median and 75 % quartiles; n = 9, Kruskal-Wallis followed
by Dunn's test ($p < 0.05$), * = p < 0.05.

Also see Supplementary Fig. 8 for further information.
1494

1495 **Figure 9**

1496
1497 **Figure 9: Model of postsynaptic plasticity sequence across compartments**

1498
Our data are consistent with a model in which α5-subunit containing receptors (red) mediate
the early phase of postsynaptic memory storage, potentially by leading to elevated calcium flux
(not addressed in this study) at individual postsynaptic densities (see discussion and
Supplementary Fig. 9 showing separated PSDs and analyses concerning input specificity).
Concurrent events see changed dynamics of the α2 receptor (blue). Nicotinic receptor subunits
hereby potentially interact with adaptor proteins to bind to Dlg. Importantly, we identify elevated
α2 subunit dynamics in the context of associative (M4; KC (green) and dopaminergic neuron
(orange) activation needed for memory formation) and non-associative (α'3 MBONs; odor
activates both KCs (green) and DANs (purple)) memory expression. Increased α2 subunit
dynamics in both cases are triggered by odor application. At the level of M4, suppressed
dynamics (concurrent activation of KC (green) and DAN (orange)), would correspond to
postsynaptic depression, while at the level of α'3 MBONs increased dynamics could result in
postsynaptic depression. Therefore, different learning rules might govern the incorporation,
exchange or stabilization of receptors in or out of synapses. Please see discussion for further
details.

1514

Supplementary Information

Supplementary Figure 1: Permissive temperature controls accompanying Fig. 1.

a)   Permissive temperature control for experiments shown in Fig. 1a. 30 min aversive memory performance when training at 23°C (driver line R13F02-Gal4). Bar graphs: mean ± SEM; n = 7 – 9; Kruskal-Wallis followed by Dunn's test (p = 0.08).

b)   Permissive temperature control for experiments shown in Fig. 1b. 30 min appetitive memory performance when training at 23°C (driver line R13F02-Gal4). Bar graphs: mean ± SEM; n = 6 – 8; Kruskal-Wallis followed by Dunn's test (p = 0.7).

1538
1539
1540
Supplementary Figure 2: Genetic controls and alternate data display of data presented
in Fig. 2.

a) Immediate appetitive memory is not impaired in genetic control groups. Bar graphs:
mean ± SEM; n = 7 – 11, for controls: n = 16, one-way ANOVA followed by Tukey's test (p >
0.05).

b) 3-hour appetitive memory is not impaired in genetic control groups. Bar graphs: mean
± SEM; n = 7 – 33, for controls: n = 41, one-way ANOVA followed by Tukey's test (p > 0.05).

c) Immediate aversive learning is not impaired in genetic control groups. Bar graphs:
mean ± SEM; n = 6 – 10, for controls: n = 13, Kruskal-Wallis followed by Dunn's test (p > 0.05).

d) 3-hour aversive memory is not impaired in genetic control groups. Bar graphs: mean ±
SEM; n = 6 – 10, for controls: n = 15, Kruskal-Wallis followed by Dunn's test (p > 0.05).

e) RNAi knock-down of the α5 subunit in M4/6 MBONs (driver line VT1211-Gal4) is
suppressed using Gal80^{ts}. Immediate memories is not impaired. Bar graphs: mean ± SEM; n =
9; one-way ANOVA followed by Tukey's test.

f) RNAi knock-down of the α2 subunit in M4/6 MBONs (driver line VT1211-Gal4) is
suppressed using Gal80ts. Mid term memory is not impaired. Bar graphs: mean ± SEM; n = 9
– 11, one-way ANOVA followed by Tukey's test.

g) RNAi knock-down of the α5 subunit in M4/6 MBONs (driver line VT1211-Gal4) is
suppressed using Gal80ts. Mid term memory is not impaired. Bar graphs: mean ± SEM; n = 10
– 11, one-way ANOVA followed by Tukey's test.

h) RNAi knock-down of Dlg in M4/6 MBONs (driver line VT1211-Gal4) is suppressed
using Gal80ts. Mid term memory is not impaired. Bar graphs: mean ± SEM;
n = 8 –10, one-way ANOVA followed by Tukey's test.

(**i-m**) Alternate display (see Methods for details) for appetitive memory experiments shown in
Fig. 2a,b:

i) Immediate appetitive memories are impaired following RNAi knock-down of the α5
nAChR subunit in M4/6 MBONs (driver line MB011B). Bar graphs: mean ± SEM; n = 9 – 21,
one-way ANOVA followed by Tukey's test (p < 0.05), * = p < 0.05.

j) 3-hour appetitive learning is impaired by RNAi knock-down of the α1 nAChR subunit in
M4/6 MBONs (driver line MB011B). Bar graphs: mean ± SEM; n = 25 − 51, Kruskal-Wallis
followed by Dunn's test (p < 0.05), * = p < 0.05.

k) RNAi knock-down of the α2 nAChR subunit in M4/6 MBONs (driver line MB011B)
impairs 3-hour appetitive memories. Bar graphs: mean ± SEM; n = 27 − 51, Kruskal-Wallis
followed by Dunn's test (p < 0.05), * = p < 0.05.

l) 3-hour appetitive learning is impaired by RNAi knock-down of the α5 nAChR subunit in
M4/6 MBONs (driver line MB011B). Bar graphs: mean ± SEM; n = 27 − 51, one-way ANOVA
followed by Tukey's test (p < 0.05), * = p < 0.05.

m) RNAi knock-down of Dlg in M4/6 MBONs (driver line MB011B) impairs 3-hour
appetitive memories. Bar graphs: mean ± SEM, n = 13 − 31; one-way ANOVA followed by
Tukey's test (p < 0.05), * = p < 0.05.

1581
1582 Note that data in panels a-b and i-m were replotted to allow comparison between all genotypes.

1583

Supplementary Figure 3: Control experiments and non-normalized data display for Fig.
3.

a) Connectivity scheme of MB output synapses. Also shown in Fig. 3.

b) Training scheme.

c) Control experiments for Fig. 3 f-h. Paired training protocol in the absence of LexAoP-
CsChrimson. Before-after plots and bar graphs (mean); n = 9, ratio paired t-test.

d) Control experiments for Fig. 3 f-h. Paired training protocol in the absence of R58E02-
LexA. Before-after plots and bar graphs (mean); n = 8, Wilcoxon matched-pairs signed rank
test.

e) GCaMP6f responses following focal ACh injections to the γ5 compartment of the MB
are not significantly altered following α2 or α5 knock-down in the M4/6 MBONs (driver line
VT1211-Gal4). The bath contains TTX to suppress spontaneous neural activity. Bar graphs:
mean ± SEM; n = 12 – 23, one-way ANOVA followed by Dunnett's test ($p > 0.05$).

f) Averaged SynaptopHluorin responses following focal ACh injections (0.1 mM) to the
γ5 compartment of the MB using the OK107-Gal4 driver (KCs), n = 8. Shaded area depicts
SEM.

g) Averaged SynaptpHluorin responses following focal KCl injections (300-400 mM) to the
γ5 compartment of the MB using the OK107-Gal4 driver (KCs), n = 4. Shaded area depicts
SEM.

1611
1612 **h)** Left: sample calcium traces in response to ACh injections recorded from M6 dendrites
1613 pre (black traces) and post (colored traces) training. Right: peak quantification.
1614 Following RNAi knock-down of the α2 subunit in M4/6 MBONs no significant potentiation was
1615 detected after paired acetylcholine application and dopaminergic neuron activation. Before-
1616 after plots and bar graphs (mean); n = 15, ratio paired t-test.

1617

1618
1619
1620 **Supplementary Figure 4: Detailed distribution of α subunits in the MB accompanying**
1621 **Fig. 4.**

1622

1623 Example image planes of GFP expression at the level of the MB compartments for all α nAChR
1624 subunits (except for α3) and Dlg. Pictures are taken from different animals. Scale-Bar: 20 μm.
1625 Left: merged image of α subunit signal (green) with MB compartments marked by 247-dsRed
1626 (magenta). Top two rows: γ compartments; middle two rows: γ, β' and β compartments, bottom
1627 three rows: α'/β' and α/β, with bottom row: α' and α compartments.

1628
1629
1630 **Supplementary Figure 5: Additional data and display of non-normalized data**
1631 **accompanying Fig. 5.**
1632

a) Scheme indicating the imaging area at the level of the β'2 compartment.

b) Odor exposure protocol. 5 OCT stimuli were alternatingly administered with 5 MCH
stimuli. 1 second odor puffs were separated by 30 seconds breaks. Odor responses are
measured using GCaMP6f as in Fig. 5.

c) Averaged traces of GCaMP6f (calcium) responses to MCH from control (black), α2
(blue) and α5 RNAi (red; driven in M4/6 respectively, driver line VT1211-Gal4) flies. Solid traces
are mean, shaded areas SEM; n = 10 - 12. Odor exposure is indicated by bar.

d) Area under curve quantification of averaged odor responses show significantly
elevated odor responses to MCH following α5 knock-down in M4/6 neurons (driver line VT1211-
Gal4). Mean ± SEM; one-way ANOVA followed by Dunnett's test (p < 0.08), * = p < 0.05; n =
10 – 12.

e) Control flies show no significant increase between the first and the fifth response to
MCH. n = 10; paired t-test.

f) α2 RNAi flies show no difference between the first and fifth odor response to MCH.
nAChR subunit RNAi is driven in M4/6 neurons (driver line VT1211-Gal4). n = 12; paired t- test.

g) α5 RNAi flies show a significant decrease in calcium transients over the course of
consecutive odor exposures. nAChR subunit RNAi is driven in M4/6 neurons (driver line
VT1211-Gal4). n = 12; Wilcoxon matched-pairs signed rank test; * = p < 0.05.

h) KCs (driver line R13F02-Gal4) show no significant change in calcium transients over
the consecutive OCT odor exposures. Before-after plots. n = 10, paired t- test.

i) KCs (driver line R13F02-Gal4) show no significant change in calcium transients over
the consecutive MCH odor exposures. Before-after plots. n = 10, paired t- test.

1655

1656
1657
1658 **Supplementary Figure 6: Receptor subunit recovery, accompanying Fig. 6.**
1659
a)   Scheme of site of dopamine injection during fluorescence recovery after
photobleaching (FRAP) experiments at the level of the KC-MBON synapses of the β'2
compartment.

b)   FRAP experimental protocol. After bleaching, a baseline picture was taken followed
by odor presentation or odor presentation simultaneously with dopamine injection or dopamine
injection by itself. Fluorescence recovery was monitored at the 10 and 20 minute time points.

c)   Example image of α2^{GFP} expression (the same image is shown here as in Fig. 6); white
dashed box shows the β'2 output zone; dopamine injection pipette (with Texas Red) is labelled
in magenta. Scale bar: 10 μm.

d)   Regression of fluorescence recovery after photobleaching following OCT exposure
(blue line), or OCT exposure simultaneously with dopamine (DA) injection (purple line), or
dopamine injection alone (red line). n = 3 –5, t-test, Satterthwaite's method for approximating
the degrees of freedom. *: regression coefficient $p < 0.05$ in the linear mixed effect model. Only
'OCT only' exposure shows recovery that significantly differs from 0.
1674
1675 **e)**   Control settings for Fig. 6 and the experiments shown in this figure. After bleaching,
1676 α2^{GFP} flies were exposed to air only. n = 9, t-test, Satterthwaite's method for approximating the

1677   degrees of freedom. The regression coefficient is not significantly different from 0.

a   Dlg^{GFP}

1686
1687   **Supplementary Figure 7: Dlg^{GFP} FRAP, accompanying Fig. 7**

1688
1689   **a)**   FRAP of Dlg^{GFP} in α'3 MBONs. Dlg^{GFP} did not show significant recovery.
1690   Recovery rate is normalized to the baseline recorded after selective bleaching of the α'3
1691   MB compartment. n = 5 - 7; multiple t-tests.

Supplementary Figure 8: Additional ethograms, accompanying Fig. 8.

a, b) Ethograms of the behavioral responses of flies shown in Fig. 8 with additional behavioral categories of pausing and wandering (when not grooming). Ethograms show pausing (red, **a**) which is defined as not moving and not grooming or wandering (dark red, **b**) which is defined as moving around in the chamber.

Supplementary Figure 9: Input analysis of KC to MBON synapses, accompanying Fig. 9.

a) Reconstructed example synapses from electron microscopic (EM) volume (neuprint.org)[62]: two different KCs connect to the same MBON on the postsynaptic side.

b)   EM image (neuprint.org) showing a KC presynapse simultaneously innervating two
other KCs and the M4 MBON. Scale bar: 500 nm.

c)   EM image (neuprint.org) showing a KC presynapse simultaneously innervating the
M4 MBON and two sites of another MBON.

d)   Analysis of number of postsynaptic partners for each KC presynapse identified
providing input to M4. The histogram shows the distribution of KC synapses to M4 relative to
how many postsynaptic partners the KC presynapses have.

e)   Analysis of number of postsynaptic partners for each KC presynapse identified
providing input to M6. The histogram shows the distribution of KC synapses to M6 relative to
how many postsynaptic partners the KC presynapses have.

f)   Analysis of number of postsynaptic partners for each KC presynapse identified
providing input to α'3 MBONs (pooled). The histogram shows the distribution of KC synapses
to α'3 MBONs relative to how many postsynaptic partners the KC presynapses have.

g)   Percentage of types of neurons that share a KC presynapse with a given MBON.

1722

1723