Higher-order olfactory neurons in the lateral horn supports odor valence and odor identity coding in *Drosophila*

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Abstract

Understanding neuronal representations of odor-evoked activities and their progressive transformation from the sensory level to higher brain centers features one of the major aims in olfactory neuroscience. Here, we investigated how odor information is transformed and represented in higher-order neurons of the lateral horn, one of the higher olfactory centers implicated in determining innate behavior, using *Drosophila melanogaster*. We focused on a subset of third-order glutamatergic lateral horn neurons (LHNs) and characterized their odor coding properties in relation to their presynaptic partner neurons, the projection neurons (PNs) by two-photon functional imaging. We found that odors evoke reproducible, stereotypic and odor-specific response patterns in LHNs. Notably, odor-evoked responses in these neurons are valence-specific in a way that their response amplitude is positively correlated with innate odor preferences. We postulate that this valence-specific activity is the result of integrating inputs from multiple olfactory channels through second-order neurons. GRASP and micro-lesioning experiments provide evidence that glutamatergic LHNs obtain their major excitatory input from uniglomerular PNs, while they receive an odor-specific inhibition through inhibitory multiglomerular PNs. In summary, our study indicates that odor representations in glutamatergic LHNs encode hedonic valence and odor identity and primarily retain the odor coding properties of second-order neurons.

Introduction

Insects are the most successful taxon among the whole animal kingdom in terms of their distribution and ability to survive in a multitude of environmental conditions. Largely they rely on their olfactory sense to carry out their fundamental goal directed behaviors, such as food navigation, mating, ovipositing or escape from predators. The powerful ability to detect odor cues, to evaluate the information efficiently with a relatively small number of neurons and to transform the neuronal signal into an appropriate behavioral output, makes the insect olfactory system a premier model system for olfactory research. Numerous studies have investigated the neuronal representation of odors at successive neuronal layers from the periphery to higher brain levels using *Drosophila melanogaster* as a model organism (Bhandawat et al., 2007; Ng et al., 2002; Root et al., 2007; Schmuker et al., 2007; Schmuker & Schneider, 2007; Seki et al., 2017; Wilson

et al., 2004). Although much progress has been made in understanding odor coding at the antennal lobe (AL) level (Bhandawat et al., 2007; Galizia, 2014; Ng et al., 2002; Wilson et al., 2004), the coding strategies and processing mechanisms of higher brain centers still remain largely elusive. In this regard, the lateral horn (LH) has recently gained attention as a crucial signal processing center integrating both innate as well as learned behavioral information (Das Chakraborty & Sachse, 2021). Several studies during recent years have advanced our understanding of the anatomical and functional properties of higher-order lateral horn neurons (LHNs) regarding odor processing (Das Chakraborty & Sachse, 2021; Dolan et al., 2019; Frechter et al., 2019; Jeanne et al., 2018; Varela et al., 2019). The generation of several LH cell type specific lines, characterization of polarity and neurotransmitter identity of LHNs as well as the establishment of detailed EM connectomic data sets have led to a significant progress in the field to study the function of specific LHN classes (Dolan et al., 2019; Frechter et al., 2019). The LH is comprised of three categories of neurons, which include LH input neurons (LHINs, which are mainly olfactory projection neurons (PNs) along with mechanosensory, thermosensory and gustatory neurons), LH local neurons (LHLNs) and LH output neurons (LHONs) (Bates et al., 2020; Dolan et al., 2019; Frechter et al., 2019). In terms of PN-LHN connectivity, the olfactory PNs deriving from individual glomeruli of the AL form stereotyped and conserved connections with certain LHNs (Fişek & Wilson, 2014; Jeanne et al., 2018; Jefferis et al., 2007; Marin et al., 2002; Wong et al., 2002). Although all kind of connections are possible, PNs having similar odor tuning patterns are prone to target similar LHN types (Jeanne et al., 2018). Certain pairs of narrowly tuned glomeruli encoding ecologically relevant odors and eliciting specific kinds of behavior, such as courtship, aggregation or food seeking, converge onto the same LHN types and have been shown to be over-represented in the LH in terms of synaptic densities (Bates et al., 2020; Jeanne et al., 2018). Furthermore, a high amount of divergence has also been described to occur at the level of PN to LHN connectivity (Huoviala et al., 2018). Altogether these complex connectivity patterns in addition to direct pooling of feed-forward inputs from PNs innervating different glomeruli results in broader tuning patterns of LHNs compared to their presynaptic PNs (Bates et al., 2020; Frechter et al., 2019). In addition to the observed broadly tuned LHNs, narrowly tuned LHNs also exist, which receive input from a single type of PN and which are assumed to be further modulated by odorant selective inhibition through inhibitory neurons (Fişek & Wilson, 2014). Although several studies agree that odors are compartmentalized in the LH based on either their chemical

identity (Frechter et al., 2019), behavioral significance (Bates et al., 2020; Jeanne et al., 2018; Jefferis et al., 2007) or hedonic valence (Strutz et al., 2014; Tanaka et al., 2004), it still remains controversial how the odor information is transformed from the PN to the LHN level and which odor features are coded by subtypes of LHNs.

In the present study, we aimed to elucidate the odor coding and processing strategies of LHNs by investigating how a neuronal subset of particular neurotransmitter identity encodes different odor features in the LH and how this representation is correlated to their presynaptic partner neurons in the AL, the uni- and multiglomerular PNs (uPN, mPN). Using photoactivatable GFP, we first identified diverse clusters of LHNs based on their different neurotransmitter identities and further focused our detailed analysis exclusively on glutamatergic LHNs. Using *in vivo* two-photon functional imaging, we characterized several aspects of odor-evoked activity in these neurons, such as odor-specific response patterns, reproducibility of repeated stimulations as well as stereotypy across different individuals. We could successfully demonstrate that attractive and aversive odors are clearly segregated and that their response amplitudes are positively correlated with the innate behavioral preference to an odor. We also dissected how excitatory input from uPNs and odor-specific inhibition from mPNs contribute to the fine-tuning of odor-specific response patterns of LHNs. Altogether this study demonstrates a significant role of glutamatergic LHNs regarding olfactory processing and extends our knowledge about the transformation processes of neuronal information taking place from the periphery to higher brain levels, such as the LH.

Results

Identification of higher-order neurons in the LH

To unravel distinct neuronal circuits in the LH, we expressed photoactivatable GFP (*UAS-C3PA-GFP*) (Ruta et al., 2010) under different Gal4 driver lines with various neurotransmitter and neuromodulator identities (i.e. *Cha-Gal4*, *GAD1-Gal4*, *dVGlut-Gal4*, *TH-Gal4* and *tdc2-Gal4*) (**Figure 1A, B**). To precisely define the upper and lower limit of the LH, we labeled uPNs using *GH146-QF*, *QUAS-td tomato* in the background as a landmark in every experiment. Using two-photon excitation we photoactivated *C3PA-GFP* throughout the LH neuropil with laser pulses of

760 nm in the *in vivo* fly brain. Photoactivation of the LH using driver lines for cholinergic (*Chac-Gal4*) and GABAergic neurons (*GAD1-Gal4*) revealed clusters of LHNs with labeled somata along with cholinergic and GABAergic PNs, respectively. The labeled cholinergic and GABAergic LHN clusters were located ventrolaterally to the LH. Selective labeling of glutamatergic neurons in the LH by photoactivation using the *dVGlut-Gal4* driver line revealed several clusters of LHNs cell bodies positioned dorsomedial, ventrolateral and ventral to the LH with major neurite tracts entering the LH at distinctive locations. Glutamatergic LHNs are comprised of a large number of labeled somata (~ 40) and a few ventrolateral and ventral clusters possessing long neuronal tracts (**Figure 1B**). The labelled glutamatergic LHNs may include both LHLNs and LHONs (Frechter et al., 2019). Photoactivation of the LH using drivers for dopaminergic (*TH-Gal4*) and octopaminergic neurons (*tdc2-Gal4*) revealed the presence of both kinds of LHNs (i.e. LHLNs, LHONs) in the LH (**Figure 1A**). These neurons innervate the LH sparsely with a very long neuronal tract and their somata positioned far away from the LH. Since we observed a high abundance of glutamatergic neurons in the LH, which were non-overlapping with second-order neurons (i.e. PNs) deriving from the AL, we confined our further experiments on this subset of LHNs to elucidate their functional properties regarding odor coding.

Glutamatergic LHNs reveal reproducible, stereotypic and odor-specific responses that emerge from the PN level

To elucidate how odors are represented in the LH, we monitored odor-evoked functional responses of glutamatergic LHNs. To do so, we expressed the genetically encoded calcium sensor *UAS-GCaMP6f* (Chen et al., 2013) that has a high sensitivity and low basal fluorescence under the control of *dVGlut-Gal4*. In order to accurately and consistently identify a comparable focal plane for functional imaging across different animals, we expressed *QUAS-td tomato* in uPNs using *GHI46-QF* in the background as a landmark. We performed *in vivo* two-photon functional imaging from the glutamatergic LHNs at three different focal planes to comprehensively monitor odor responses throughout the LH (i.e. upper, middle and lower plane). We used a panel of 14 ecologically relevant odors at a concentration of 10^{-3} , which have been shown to be behaviorally either attractive or aversive (Knaden et al., 2012). Our imaging data show that each odor evoked a specific activity pattern across all three focal planes in the LH (**Figure 2A**). Since the LH appears

rather homogenous and lacks clear morphological landmarks, we used a grid approach for imaging data analysis (**Figure 2B**) (see Methods for details) and determined the $\Delta F/F$ to analyze the neuronal activity across odors for each animal. At first, we compared the reproducibility of the evoked responses of these glutamatergic LHNs to two sequential presentations of each odor. This analysis of neuronal activity for repeated odor stimulations clearly showed that the odor-evoked responses were highly reproducible both qualitatively and quantitatively across trials (**Figure 2C**), which is further depicted by a high correlation coefficient across repeated stimulations for all odors (**Figure 2D**). Hence, although the odor responses appear generally broad, each odor induced an unique response pattern in these glutamatergic LHNs (**Figure 2D**). Furthermore, a cluster analysis based on the linkage distance between the spatial patterns of each odor response reveals that the majority of the replicates for each odor are clustered together, confirming that individual odors are represented in an odor-specific manner by glutamatergic LHNs (**Figure 2F**). To investigate whether the observed odor response properties of glutamatergic LHNs derived from their presynaptic partner neurons, we determined the odor representations of uPNs in the LH to the same odor set. Previous studies based on the innervation pattern of PNs reported a compartmentalization in the LH either based on the hedonic valence of an odor, i.e. attractive versus aversive odors (Seki et al., 2017), the odor category, i.e. pheromones versus food odors (Jefferis et al., 2007) or with regard to the behavioral ‘odor scene’ (Bates et al., 2020; Jeanne et al., 2018). However, whether an odor-specific response map exists at the uPN level in the LH has so far not been unambiguously shown with a functional approach. We therefore expressed *UAS-GCaMP6f* under control of *GH146-Gal4* and monitored odor-evoked responses at the two-photon microscope from the same three focal planes as used for the LHN imaging experiments. Notably, when we compared the odor response properties of uPNs with those of LHNs we observed that the PNs possess a significantly reduced correlation coefficient as compared to LHNs, implying that odors are better segregated at the uPN level in the LH than at the third-order neuron level (**Figure 2E**). In addition, a cluster analysis based on the linkage distance between the spatial patterns of each odor response at the PN level confirms that uPNs also reveal odor-specific response patterns in the LH (**Figure 2G**). Altogether, these observations clearly demonstrate and signify that uPNs as well as glutamatergic LHNs exhibit an odor-specific response map in the LH.

Response amplitude of glutamatergic LHNs reflects odor valence

Next, we investigated whether hedonic valence is still preserved in the LH level in this particular subset of LHNs as it has been described for both types of PNs, i.e. uPNs and mPNs (Knaden et al., 2012; Strutz et al., 2014). We therefore quantified and pooled the correlation coefficients of all odor responses of glutamatergic LHNs to all attractive and aversive odors and compared those that shared the same valence (i.e. within valence) as well as for odors having an opposing valence (i.e. across valence) for all three focal planes measured (**Figure 3A**). Interestingly, we observed that the responses to odors with the same valence were significantly more similar, depicted by a higher correlation coefficient, compared to the odor responses across valence. In order to visualize the odor representations in a multidimensional space, we employed a principal component analysis (PCA) of the odor response profiles by taking into account two parameters, the spatial response patterns as well as the intensity of odor-evoked responses by each odor. As expected, the attractive and aversive odor responses were segregated significantly throughout the LH (* $p < 0.05$, one way ANOSIM) confirming that odors are indeed categorized according to their behavioral value by these third-order glutamatergic LHNs (**Figure 3B**). Notably, when only the response amplitude was considered, it was obvious that attractive odors in general evoked a stronger activity in glutamatergic LHNs compared to responses to behaviorally aversive odors across all three focal planes (**Figure 3C and D**). Next, we examined whether odor responses of second-order uPNs also reflect the hedonic valence of an odor within the LH as it has been shown for the AL level previously (Knaden et al., 2012). A PCA of the odor response profile of uPNs revealed that the attractive and aversive odors are clearly segregated (** $p < 0.005$, one way ANOSIM) at the PN level also in the LH (**Figure 3E**). In order to investigate if, similar to LHNs, the response magnitude in uPNs contributes to the observed clear segregation, we averaged the overall response activities to attractive and aversive odors across all three planes. Indeed, we observed a similar trend, meaning that uPNs responded stronger to attractive odors than to aversive ones in the LH (**Figure 3D and F**). This observation was novel and unexpected and tempted us to postulate that the flies' innate odor preference might be determined by the response amplitude of uPNs and glutamatergic LHNs. To address whether the odor-evoked response strength is correlated with behavioral preference, we performed T-maze assays to the same odor set as used for the functional imaging experiments. As shown in previous studies, flies displayed different levels of attraction and aversion to individual odors (Knaden et al., 2012; MacWilliam et al., 2018; Min et al., 2013;

Strutz et al., 2014). Plotting the olfactory preference indices of all tested odors with the response strengths of glutamatergic LHNs indeed revealed a significant correlation (**Figure 3G**) showing that odors eliciting stronger LHN activity induce stronger attraction, while odors that elicit lower LHN responses induce a stronger aversion. Interestingly, the response strength of uPNs did not correlate significantly with the olfactory preference indices (**Figure 3H**), although uPNs activity displayed a valence-specific representation (**Figure 3C**). This observation suggests that the activity in the subset of glutamatergic LHNs may contribute significantly to the observed behavioral odor preferences. However, this correlation does not confirm any causality between the response strength of glutamatergic LHNs and the behavioral output. To test a causal relationship, it would be necessary to silence or artificially activate the glutamatergic LHNs and to determine the functional consequences with regard to behavioral valence. However, since the line *dVGlut-Gal4* is broadly expressed throughout the whole brain, genetic manipulations would not be confined to the subset of glutamatergic LHNs and would therefore be difficult to evaluate. One might argue that the observed differential LHN responses emerge already at the sensory neuron level and are not the result of neuronal processing within the AL or LH. To examine this idea, we obtained odor-evoked response of olfactory sensory neurons (OSNs) to individual odors from the DoOR database (Münch & Galizia, 2016) and plotted the response strengths of glutamatergic LHNs either against the overall OSN activity (**Figure 3I**) or the number of ORs (olfactory receptors) activated by a specific odor (**Figure 3J**). Interestingly, we could not observe any correlation in either case, indicating that the differential LHN activity observed here is not deriving from the OSN level, but is the result of second-order neuronal processing. Notably, the comparison between uPNs and LHNs responses also did not show any correlation (**Figure 3K**), further suggesting that the differential activity observed in the glutamatergic LHNs is not a mere relay of the AL input through uPNs, but rather processed by either mPNs or other LHNs to generate the observed graded odor responses in higher-order glutamatergic LHNs.

Polarity and connectivity of glutamatergic LHNs with uPNs and mPNs

To determine the neuronal polarity of the glutamatergic LHNs, we expressed pre- and postsynaptic markers (*UAS-sypGC3*, *UAS-homerGC3*) under the control of *dVGlut-Gal4*. Immunohistochemistry followed by confocal scanning demonstrated that these neurons possess

both, presynaptic dendrites and postsynaptic axon terminals in the LH (**Figure 4A**, left and right panel), indicating that the cell cluster of glutamatergic LHNs probably comprises both LHLNs as well as LHONs (Dolan et al., 2019). The presence of presynaptic terminals suggests that these neurons provide input to other LHONs, LHLNs or even perform a feedback inhibition onto PN. To assess the neuronal connectivity of these neurons with excitatory and inhibitory PNs at the morphological level in detail, we employed the GRASP technique (Feinberg et al., 2008). Initially, we employed two split-GFP fragments, tagged to the extracellular domain of the CD4 transmembrane protein (hereafter referred to as *CD4-GRASP*). To determine the connectivity between uPNs and glutamatergic LHNs, we expressed one fragment of GFP in glutamatergic LHNs, using *dVGlut-Gal4* and the other fragment in uPNs, using *GH146-LexA* (**Figure 4B**, left panel). Next, to examine whether mPNs and glutamatergic LHNs are synaptic partner neurons, we employed a similar approach by using *dVGlut-LexA* and *MZ699-Gal4* (**Figure 4B**, right panel). In both cases we observed a clear neuronal connectivity to the glutamatergic LHNs. However, since *CD4-GRASP* is not synaptically targeted and can potentially lead to false positive signals at non-synaptic locations, we employed in addition a GFP 1-10 fragment, which is tagged to the synaptic protein neurexin, resulting in an enhancement of synaptic specificity (Fan et al., 2013; Shearin et al., 2018). Neurexin is present predominantly at the presynaptic site (Fan et al., 2013; Xing et al., 2018), but also occurs along with postsynaptic terminals (Taniguchi et al., 2007). When we performed GRASP by expressing neurexin in glutamatergic LHNs, we observed a clear GFP signal in the LH (**Figure 4C**, left panel) implying that glutamatergic LHNs provide presynaptic output onto uPNs and therefore convey a feedback inhibition to the AL. Such a feedback connection needs to be verified in future experiments. Next, in order to determine the synaptic connectivity between glutamatergic LHNs and mPNs, we expressed neurexin in mPNs and observed distinct GFP puncta in the LH implying that also mPNs form presynaptic and probably postsynaptic connections with glutamatergic LHNs (**Figure 4C**, right panel).

Glutamatergic LHNs receive major excitatory input from uPNs and an odor-specific inhibition from mPNs

In order to further elucidate the neural connectivity and to verify our GRASP experiments, we determined the functional connectivity of glutamatergic LHNs with uPNs and mPNs by transection

experiments using two-photon laser-mediated micro-dissection. To manipulate the input from uPNs, we micro-lesioned the iACT tract entering the LH and compared the odor-evoked responses to vinegar and benzaldehyde from the glutamatergic LHNs before and after laser ablation (**Figure 5**, schematic upper panel). After transecting the iACT tract we observed that the odor-induced responses to both, vinegar and benzaldehyde were completely abolished across all three focal planes of the LH (**Figure 5 A-A'', B-B''**). This finding implies that glutamatergic LHNs receive their major excitatory input from the AL through uPNs. To verify whether this complete elimination of the odor-evoked activity is not just a technical side effect of the laser ablation procedure, we also monitored odor responses from the glutamatergic LHNs in the other, untreated brain hemisphere before and after micro-lesioning, which proved not to be affected (**Figure 5-figure supplement 1**). In order to examine the functional connectivity between mPNs and glutamatergic LHNs, we also transected the mACT tract entering the LH and monitored odor responses from glutamatergic LHNs before and after laser transection. Notably, transecting the mACT led to a significant increase in the odor-evoked responses in the middle and upper plane of the glutamatergic LHNs (**Figure 5 C-C'', D-D''**). We further observed that the mACT transection affected the odor-evoked responses differentially in a way that the response increase was higher in the case of vinegar than for benzaldehyde. This observation indicates that the glutamatergic LHNs receive an odor selective inhibition from mPNs which is stronger for the food odor vinegar than for the repellent odor benzaldehyde.

mPN- mediated inhibition onto glutamatergic LHNs contributes to odor specificity

Next, we aimed to understand the impact of the observed odor-specific inhibition of mPNs with regard to the odor-specific responses of glutamatergic LHNs in the LH. To address this issue, we monitored odor-evoked responses in the glutamatergic LHNs to two repeated stimulations of vinegar and benzaldehyde (depicted as 1 and 2 in the schematic in **Figure 6A, A'**), followed by measuring LHN activity after micro-lesioning the mACT tract (depicted as 3 in **Figure 6A, A'**). Since, we observed a significant response increase following mPN-mediated inhibition only in the middle and upper plane (**Figure 5C', C'', D' and D''**), we confined our analysis to these two areas. In order to determine whether the laser ablation of mACT alters also the odor-evoked spatial pattern in addition to an increase in response amplitude, we compared the correlation coefficients

between the repeated responses before laser ablation to those after the transection. Our results revealed that removal of the inhibitory input from mPNs leads to a reduced correlation coefficient of the odor response patterns evoked by vinegar in LHNs (**Figure 6B, B'**). Interestingly, such an effect was not observed in the case of benzaldehyde although we observed a weak mPN-mediated inhibition for this odor before (**Figure 6B, B'**). Since the odor-evoked spatial patterns were modified after silencing the mPN input into the LH, we next wondered whether this also affected the odor specificity of the LHN responses. We therefore compared the correlation coefficients ‘across odors’ (i.e. between vinegar and benzaldehyde) before and after microlesion (**Figure 6C**). Strikingly, we observed that the odor-evoked response patterns of vinegar and benzaldehyde became increasingly similar after removal of the mPN-mediated inhibition depicted as an increased correlation coefficient after laser ablation. This finding suggests that the inhibitory input from mPNs onto glutamatergic LHNs is crucial to generate a distinct and odor-specific response pattern for individual odors at the LH level.

Discussion

Our study functionally characterizes a subset of glutamatergic higher-order neurons in the LH regarding odor coding and processing. We demonstrate that glutamatergic LHNs respond in a reproducible, stereotypic and odor-specific manner and these response properties emerge at the level of presynaptic uPNs (**Figure 6D**). Notably, the differential activity levels of glutamatergic LHNs to attractive and aversive odors are positively correlated to the olfactory behavioral preference indicating that these neurons are mainly tuned to attractive odors. The response features do not arise from the OSN level, but rather derive from local processing within the LH by integrating inputs from multiple olfactory channels through uPNs, which also show valence specific odor representation. Furthermore, laser transection experiments demonstrate that these higher-order neurons receive their major excitatory input from uPNs and an odor-specific inhibitory input from mPNs. Lastly, our data show that the observed mPN-mediated inhibition seems to be required for generating an odor-specific response map in the LH (**Figure 6D**).

Glutamatergic LHNs respond in a stereotypic and odor-specific manner

A growing body of evidence suggests the existence of an anatomical and functional stereotypy in early processing centers of the insect olfactory pathway (Hildebrand & Shepherd, 1997; Korsching, 2002). This stereotypy becomes obvious first at the sensory neuron (OSN) level, where OSNs expressing a certain OR target and converge on a stereotypic glomerulus resulting in a conserved spatial map in the AL between different individuals (Wang et al., 2003; Wilson et al., 2004). This anatomical stereotypy was shown to be retained at the postsynaptic PN level (Jefferis et al., 2007; Tanaka et al., 2004).

Several studies support the notion that an anatomical stereotypy might also be present at the level of the LH, in particular shown for the PN to LHN connectivity (Bates et al., 2020; Fişek & Wilson, 2014; Jeanne et al., 2018; Jefferis et al., 2007). Along this line, functional studies have demonstrated that LHNs respond in a reproducible and stereotyped manner to odors and this stereotypy is a general feature of the LH (Fişek & Wilson, 2014; Frechter et al., 2019). However, how an ensemble of LHNs integrates inputs from several olfactory channels, i.e. the presynaptic excitatory and inhibitory PNs, and whether each odor induces a specific and stereotyped response pattern in the LH was not clearly addressed before. In our study we demonstrate that each odor is represented by an odor-specific activity pattern in the LH, while the examined glutamatergic LHNs display broader tuning patterns than their presynaptic partner neurons. Although it has been assumed previously that odor specificity may not be encoded in higher-order brain centers (Grabe & Sachse, 2017; Strutz et al., 2014), our findings are in accordance with a recent study by Frechter et al. (2019) who demonstrate the existence of 33 different LH cell types exhibiting stereotypic odor response properties with increased tuning breadth than PNs. The observed broader tuning breadth of LHNs could be due to their property of integrating various inputs from multiple odor channels, e.g. one LHN receives on average excitatory input from ~ 5.2- 6.2 glomeruli (Frechter et al., 2019; Jeanne et al., 2018). Here, we observed that uPNs are more efficient in encoding odor identity than the glutamatergic LHNs in the LH (**Figure 6D**), whereas LHNs reveal an improved categorization of odors either based on behavioral significance, ‘odor scene’ or chemical group (Bates et al., 2020; Frechter et al., 2019; Jeanne et al., 2018). However, the distinct odor-specific response map by glutamatergic LHNs observed in our study suggests that the dimensionality of

odor features might not get reduced but still retains information about the odor specificity at the third-order processing stage.

Coding of hedonic odor valence

Our functional imaging recordings revealed that odor valence is encoded by glutamatergic LHNs leading to different activation strengths and patterns for attractive and aversive odors in the LH. This observation is well in line with previous studies that have revealed that odor-evoked responses in higher brain centers are generally categorized according to certain odor features as already mentioned above. For example, using functional imaging or patch-clamp recordings of second-order olfactory neurons revealed the existence of distinct attractive and aversive odor response domains in the LH formed by uPNs and mPNs (Seki et al., 2017; Strutz et al., 2014). Such a categorization according to hedonic valence is also visible in our study, when the odor-evoked responses of glutamatergic LHNs were plotted in a PCA, taking into account the spatial response patterns as well as the intensity of activity. Although no prominent spatial domain of attractive or aversive odors was evident in our recordings, we observed that attractive odors evoked a generally stronger activity when compared to aversive odors in this subset of LHNs. We noted a similar trend in second-order uPNs. However, their response strength was neither correlated with the olfactory preference determined in behavior nor the odor response properties of LHNs. The observed significant correlation between the amount of odor-evoked activity in glutamatergic LHNs to the behavioral valence of an odor leads us to postulate that the activity strengths of higher-order olfactory neurons to odor stimulation might determine the behavioral response – an assumption that needs to be tested in future studies.

Our study suggests that glutamatergic LHNs use different strategies to extract different features of odor information, 1) conserving the identity of an olfactory stimulus by forming an odor-specific activity map and, 2) encoding the valence of an odor by integrating information from multiple olfactory channels. It is an ongoing debate regarding how neurons in the LH evaluate an odor stimulus. It can be noted that irrespective of how an animal detects, encodes and categorizes an odor at different olfactory processing levels, the behavioral output is in any case a binary choice, reflected in either to approach (positive) or to leave (negative), to copulate (positive) or to reject

(negative), to oviposit (positive) or to find another suitable oviposition site (negative) (Galizia, 2014). Hence, based on the context or ecological relevance, an odor can just be evaluated either as ‘pleasant’ or ‘unpleasant’, which is well reflected by the response properties of glutamatergic LHNs regarding their valence-specificity and their correlation between response strength and behavioral odor preference.

Notably, such a correlation between response intensity and behavioral preference has also been observed in previous studies, where the amplitude of food odor-evoked activity in neuropeptide F (dNPF) neurons was found to strongly correlate with food odor attractiveness (Beshel & Zhong, 2013). Another study that combined functional imaging with tracking of innate behavioral responses revealed that the behavioral output could be accurately predicted by a model summing up the normalized glomerular responses, in which each glomerulus contributes a small but specific part to the resulting odor preference (Badel et al., 2016). At the level of the LH, LHNs then integrate the olfactory information from the glomerular responses conveyed via uPNs and mPNs. In one of our previous studies we demonstrated that mPNs respond differently to attractive and aversive odors, and mediate behavioral attraction (Strutz et al., 2014). In this context our study complements this previous finding by showing that also uPNs display distinct valence-specific responses in the LH to attractive and aversive odors. Information from these two PN pathways become integrated and processed in the LH resulting in valence-specific activities in glutamatergic LHNs, which may in turn determine the relative behavioral preference.

mPN-mediated inhibition generates odor-specific response patterns in the LH

We demonstrate that mPNs inhibit the glutamatergic LHNs in an odor-selective manner leading to an odor-specific response pattern. According to our observations, glutamatergic LHNs receive a stronger inhibition from mPNs in response to the odor vinegar than to benzaldehyde. In the absence of this inhibition, we noted that in addition to an increased response amplitude and altered odor representation, the activity patterns of different odors became more strongly correlated and hence more similar. We therefore conclude that the mPN-mediated selective inhibition on this glutamatergic subset of LHNs is necessary to maintain odor specificity. Along this line, a previous study has reported that mPNs provide an odor-selective input to vlpr neurons, another class of

third-order LHNs (Liang et al., 2013). According to these authors, this odor-specific modulation depends on the nature of the odor and results from the stereotyped connectivity of mPNs in the AL as well as in the LH. Although this study provides evidence that uPNs are not presynaptically inhibited by mPNs, another study established that mPNs indeed inhibit uPNs in the LH, facilitating odor discrimination (Parnas et al., 2013). In addition to uPNs and vlpr neurons, our study identifies another class of recipient neurons (glutamatergic LHNs) that receives mPN-mediated odorant-selective inhibition. The glutamatergic LHNs population in our study comprises glutamatergic LHONs as well as LHLNs (Dolan et al., 2019). Although we were not able to clearly differentiate the functional properties between different populations of LHONs and LHLNs, since we used a general marker for glutamatergic neurons, our study provides the first understanding of how odors are integrated, transformed and finally represented in the LH by an ensemble of glutamatergic LHNs.

Intriguingly, the neurotransmitter identity of this class of LHNs opens up another interesting aspect: Knowing that glutamate can act as an excitatory (Aungst et al., 2003; Das et al., 2011a) or inhibitory neurotransmitter (Liu & Wilson, 2013), as well as a coincident detector (Das et al., 2011b), depending upon the receptors present in the postsynaptic neurons, further experiments are needed to reveal the consequences of glutamatergic LHN input onto their postsynaptic partner neurons. Certainly, the presence of an impressive amount of vesicular glutamate in the LH points towards a significant role of glutamatergic LHNs with regard to odor coding and processing at this higher brain center.

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Author contributions

S.D.C and S.S designed research; S.D.C and H.C performed research; S.D.C, H.C and S.S analyzed data; B.S.H. provided intellectual and financial support, S.D.C, B.S.H. and S.S. wrote the paper.

Declaration of Interests

The authors declare no conflict of interest.

Materials and methods:

Fly stocks

Flies were raised on autoclaved cornmeal-yeast-sucrose-agar food in a 12-h light/dark cycle at 25°C incubator. The following lines have been used for functional imaging: *dVGlut-Gal4* (II) (Mahr & Aberle, 2006) (Bloomington 26160), *GHI46-Gal4* (II) (Stocker et al., 1997) (from Leslie Vosshall's lab.), *GHI46-QF*, *QUAS mtd Tomato* (BDSC 30037) and *UAS-GCaMP6f* (III) (Chen et al., 2013) (Bloomington 52869). For photoactivation experiments *Cha-Gal4* (II) (Bloomington 6798), *GAD1-Gal4* (II) (Ng et al., 2002) (Bloomington 51630), *dVGlut-Gal4*, *TH-Gal4* (III) (Zhang et al., 2007) (from Mani Ramaswami's lab), *tdc2-Gal4* (II) (Koon et al., 2010) (NCBS, India) and *UAS-C3PA* (Ruta et al., 2010) (gift from Sandeep Datta) flies have been used. Wild type *Canton-S* flies have been used for behavioral experiments.

Photoactivation

UAS-C3PA was expressed under *Cha-Gal4*, *GAD1-Gal4*, *dVGlut-GAL4*, *TH-Gal4* and *tdc2-Gal4* in the background of *GHI46-QF*, *QUAS mtd Tomato* for the photoactivation experiment. An initial pre-photoactivation scan of the whole LH area was taken at 925 nm with a 40× water immersion objective. The extent of the LH in z section was identified based on the innervation pattern of uPNs (visualized by *GHI46-QF*, *QUAS mtd Tomato*). A region of interest in different focal planes covering the entire LH area was photoactivated for ~1-2 min followed by 2-min rest using ~7mW

of 760 nm of laser. We allowed 10–15 min for photoactivated GFP to diffuse in more distal neural processes. The post-photoactivation scan was taken using the same set-up as used for the pre-photoactivation scan.

Functional imaging

All functional imaging experiments were performed on 4-7 day old mated female flies. Flies were dissected for optical imaging according to the protocol by Strutz et al. (2014). Flies were briefly immobilized on ice and then mounted onto a custom-made Plexiglas stage with a copper plate (Athene Grids, Plano). A needle before the head was placed to stabilize the proboscis and to align the head properly. Protomp II composite (3M ESPE) was used to fix the head with the copper plate. We bent the anterior part of the fly's head with fine gold wire, and a small plastic plate having a round window was placed on top. We sealed the head with that plate using two-component silicone (Kwik Sil) and leaving the center part open to make a cut. The cuticle between the eyes and the ocelli was gently cut under saline (130 mM NaCl, 5 mM KCl, 2 mM MgCl₂, 2 mM CaCl₂, 36 mM saccharose, 5 mM Hepes, 1 M NaOH, pH 7.3). The cuticle was either bent forward and fixed to the silicon or removed. All fat, trachea, and air sacs were removed carefully.

Functional imaging was performed using a 2-photon laser scanning microscope (2PCLSM, Zeiss LSM 710 meta NLO) equipped with an infrared Chameleon Ultra™ diode-pumped laser (Coherent, Santa Clara, CA, USA) and a 40× water immersion objective lens (W Plan-Apochromat 40×/1.0 DIC M27). The microscope and the laser were placed on a smart table UT2 (New Corporation, Irvine, CA, USA). The fluorophore of GCaMP6f was excited with 925 nm. For each individual measurement, a series of 40 frames (corresponding to 10 s) acquired at a resolution of 256 × 256 pixels was taken with a frequency of 4 Hz. During the entire 10 s of recording each odor was delivered after 2 s (8 frames) for 2 s (8 frames). To cover the entire LH, images were acquired from three different focal planes (i.e. upper, middle and lower), each 15µm apart in z direction. *GH146-QF*, *QUAS-td tomato* was always used in the background to precisely locate the planes for all imaging experiments. A set of 14 odors was used including phenylacetaldehyde (PHA), vinegar (VIN), 2,3 butanedione (BUD), acetoin acetate (AAC), ethyl acetate (ETA), propionic acid (PAC) and gamma-butyrolactone (G-BUT) as attractive odors, 1-octanol (OCT), benzaldehyde (BEA),

acetophenone (ACP), methyl salicylate (MSC), geosmin (GEO) and linalool (LIN) as aversive odors and cis-vaccenyl acetate (cVA) as a pheromone. All odors have been used at a concentration of 10^{-3} diluted in mineral oil. Vinegar was diluted in double distilled water. Flies were imaged for up to 1 hr, with a minimum inter-stimulus interval of one minute. For odor delivery we used a computer-controlled odor delivery system (described in (Mohamed et al., 2019)).

Data and statistical analysis for functional imaging

The LSM file obtained from the imaging software ZEN, was processed and analyzed in Fiji. 40 frames for each odor (40 frames X 14 odors) were opened and stitched together using 'Image>Stacks>Tools>Concatenate' command. The file combining all the frames for all odors were movement corrected using 'Plug in>Registration>Stackreg>Rigid body' command. The entire LH area was encircled and cropped and divided into 65 pixels, each having $30 \times 30 \mu\text{m}^2$ dimension. Using 'Analyze>Tools>ROI manager>more>multi measure' command fluorescence intensity for every pixel for every frame for every odor was obtained. This huge data set was further processed using Excel software. Average $\Delta F/F$ was calculated for 10-18 frames for each pixel for each odor.

The $\Delta F/F$ for 65 pixels for 14 odors for one replicate was aligned in table format and processed using 'past' statistical software. The correlation coefficient for each odor pair using Euclidean distance was analyzed.

2-Photon-mediated laser transection

Transection of either the iACT or mACT tract of PNs was performed using *GHI46-QF*, *QUAS-td tomato* as a guiding landmark. Transection of the PN tracts was done in one brain hemisphere of each fly only. The chosen area for transection was located using 925 nm of laser, close to the LH but far enough not to damage the neurites branching in the LH neuropil. ~60 mW of 760 nm laser was provided at the chosen area at a pulse for 2-3 s for 2-3 times with 1 s time interval (Strutz et al., 2014). Successful transection was verified after visualization of a small bubble. As a control, the LH from the other, untreated brain hemisphere was imaged to stimulation with benzaldehyde

before and after laser ablation to rule out that the laser transection of the specific targeted area caused any non-specific damage to other brain areas.

Behavioral assay

T-maze experiments with 4-7 days old flies were carried out for the olfactory preference assay. ~30 flies were starved for 24 hrs in each vial containing wet filter paper. Mated males and females were used for the behavioral assay. All odors were used at a concentration of 10^{-3} , which is the same as used for the functional imaging experiments. At each arm of the T-maze small filter papers containing either 6 μ l of mineral oil (control) or an odor (odorant) were placed. Briefly cold anaesthetized flies were released at the junction of horizontally placed T maze. After ~30 min flies in each side of the arm were counted. The preference index was calculated as (odorant-control)/total number of flies (i.e., 30). Each odor has ~20 replicates.

References:

- Aungst, J. L., Heyward, P. M., Puche, A. C., Karnup, S. V., Hayar, A., Szabo, G., & Shipley, M. T. (2003). Centre-surround inhibition among olfactory bulb glomeruli. *Nature*.
<https://doi.org/10.1038/nature02185>
- Badel, L., Ohta, K., Tsuchimoto, Y., & Kazama, H. (2016). Decoding of Context-Dependent Olfactory Behavior in *Drosophila*. *Neuron*, 91(1), 155–167.
<https://doi.org/10.1016/j.neuron.2016.05.022>
- Bates, A.S., Schlegel, P., Roberts, R.J.V., Drummond, N., Tamimi, I.F.M., Turnbull, R., Zhao, X., Marin, E.C., Popovici, P.D., Dhawan, S.,....., Jefferis, G.S. (2020). Complete Connectomic Reconstruction of Olfactory Projection Neurons in the Fly Brain. *Current Biology : CB*, 30(16), 3183–3199.e6. <https://doi.org/10.1016/J.CUB.2020.06.042>
- Beshel, J., & Zhong, Y. (2013). Graded encoding of food odor value in the *Drosophila* brain. *Journal of Neuroscience*, 33(40), 15693–15704. <https://doi.org/10.1523/JNEUROSCI.2605-13.2013>

- 541 Bhandawat, V., Olsen, S. R., Gouwens, N. W., Schlieff, M. L., & Wilson, R. I. (2007). Sensory
542 processing in the *Drosophila* antennal lobe increases reliability and separability of ensemble
543 odor representations. *Nature Neuroscience*, 10(11), 1474–1482.
544 <https://doi.org/10.1038/nn1976>
- 545 Chen, T.-W., Wardill, T. J., Sun, Y., Pulver, S. R., Renninger, S. L., Baohuan, A., Schreiter, E. R.,
546 Kerr, R. a, Orger, M. B., Jayaraman, V., Looger, L. L., Svoboda, K., & Kim, D. S. (2013).
547 Ultrasensitive fluorescent proteins for imaging neuronal activity. *Nature*, 499(7458), 295–
548 300. <https://doi.org/10.1038/nature12354>
- 549 Das, A., Chiang, A., Davla, S., Priya, R., Reichert, H., VijayRaghavan, K., & Rodrigues, V.
550 (2011a). Identification and analysis of a glutamatergic local interneuron lineage in the adult
551 *Drosophila* olfactory system. *Neural Systems & Circuits*, 1(1), 4.
552 <https://doi.org/10.1186/2042-1001-1-4>
- 553 Das Chakraborty, S., & Sachse, S. (2021). Olfactory processing in the lateral horn of *Drosophila*.
554 *Cell and Tissue Research* 2021 383:1, 383(1), 113–123. [https://doi.org/10.1007/S00441-](https://doi.org/10.1007/S00441-020-03392-6)
555 020-03392-6
- 556 Das, S., Sadanandappa, M. K., Dervan, A., Larkin, A., Lee, J. A., Sudhakaran, I. P., Priya, R.,
557 Heidari, R., Holohan, E. E., Pimentel, A., Gandhi, A., Ito, K., Sanyal, S., Wang, J. W.,
558 Rodrigues, V., & Ramaswami, M. (2011b). Plasticity of local GABAergic interneurons
559 drives olfactory habituation. *Proceedings of the National Academy of Sciences*, 108(36),
560 E646–E654. <https://doi.org/10.1073/pnas.1106411108>
- 561 Dolan, M.-J., Frechter, S., Bates, A. S., Dan, C., Huoviala, P., Roberts, R. J., Schlegel, P.,
562 Dhawan, S., Tabano, R., Dionne, H., Christoforou, C., Close, K., Sutcliffe, B., Giuliani, B.,
563 Li, F., Costa, M., Ihrke, G., Meissner, G. W., Bock, D. D., ... Jefferis, G. S. (2019).
564 Neurogenetic dissection of the *Drosophila* lateral horn reveals major outputs, diverse
565 behavioural functions, and interactions with the mushroom body. *eLife*, 8, 1–45.
566 <https://doi.org/10.7554/eLife.43079>
- 567 Fan, P., Manoli, D. S., Ahmed, O. M., Chen, Y., Agarwal, N., Kwong, S., Cai, A. G., Neitz, J.,
568 Renslo, A., Baker, B. S., & Shah, N. M. (2013). Genetic and neural mechanisms that inhibit

569 drosophila from mating with other species. *Cell*. <https://doi.org/10.1016/j.cell.2013.06.008>

570 Feinberg, E. H., VanHoven, M. K., Bendesky, A., Wang, G., Fetter, R. D., Shen, K., &
571 Bargmann, C. I. (2008). GFP Reconstitution Across Synaptic Partners (GRASP) defines
572 cell contacts and synapses in living nervous systems. *Neuron*, 57(3), 353–363.
573 <https://doi.org/10.1016/J.NEURON.2007.11.030>

574 Fişek, M., & Wilson, R. I. (2014). Stereotyped connectivity and computations in higher-order
575 olfactory neurons. *Nature Neuroscience*, 17(2), 280–288. <https://doi.org/10.1038/nn.3613>

576 Frechter, S., Bates, A. S., Tootoonian, S., Dolan, M. J., Manton, J., Jamasb, A. R., Kohl, J.,
577 Bock, D., & Jefferis, G. (2019). Functional and anatomical specificity in a higher olfactory
578 centre. *ELife*, 8, 1–39. <https://doi.org/10.7554/eLife.44590>

579 Galizia, C. G. (2014). Olfactory coding in the insect brain: Data and conjectures. *European*
580 *Journal of Neuroscience*, 39(11), 1784–1795. <https://doi.org/10.1111/ejn.12558>

581 Grabe, V., & Sachse, S. (2017). Fundamental principles of the olfactory code. *BioSystems*, 164,
582 94–101. <https://doi.org/10.1016/j.biosystems.2017.10.010>

583 Hildebrand, J. G., & Shepherd, G. M. (1997). Mechanisms of olfactory discrimination:
584 Converging evidence for common principles across phyla. *Annual Review of Neuroscience*,
585 20(1), 595–631. <https://doi.org/10.1146/annurev.neuro.20.1.595>

586 Huoviala, P., Dolan, M., Love, F., Frechter, S., Roberts, R., Mitrevica, Z., Schlegel, P., Bates, A.
587 S., Aso, Y., Rodrigues, T., Cornwall, H., Stensmyr, M., Bock, D., Rubin, G., Costa, M., &
588 Jefferis, G. (2018). Neural circuit basis of aversive odour processing in *Drosophila* from
589 sensory input to descending output. *BioRxiv*. <https://doi.org/10.1101/394403>

590 Jeanne, J. M., Fişek, M., & Wilson, R. I. (2018). The Organization of Projections from Olfactory
591 Glomeruli onto Higher-Order Neurons. *Neuron*, 98(6), 1198–1213.e6.
592 <https://doi.org/10.1016/j.neuron.2018.05.011>

593 Jefferis, G. S. X. E., Potter, C. J., Chan, A. M., Marin, E. C., Rohlfig, T., Maurer, C. R., & Luo,
594 L. (2007). Comprehensive Maps of *Drosophila* Higher Olfactory Centers: Spatially
595 Segregated Fruit and Pheromone Representation. *Cell*, 128(6), 1187–1203.

<https://doi.org/10.1016/j.cell.2007.01.040>

Knaden, M., Strutz, A., Ahsan, J., Sachse, S., & Hansson, B. S. (2012). Spatial Representation of Odorant Valence in an Insect Brain. *Cell Reports*, 1(4), 392–399.

<https://doi.org/10.1016/j.celrep.2012.03.002>

Koon, A. C., Ashley, J., Barria, R., DasGupta, S., Brain, R., Waddell, S., Alkema, M. J., & Budnik, V. (2010). Autoregulatory and paracrine control of synaptic and behavioral plasticity by octopaminergic signaling. *Nature Neuroscience* 2010 14:2, 14(2), 190–199.

<https://doi.org/10.1038/nn.2716>

Korsching, S. (2002). Olfactory maps and odor images. In *Current Opinion in Neurobiology*.

[https://doi.org/10.1016/S0959-4388\(02\)00348-3](https://doi.org/10.1016/S0959-4388(02)00348-3)

Liang, L., Li, Y., Potter, C. J., Yizhar, O., Deisseroth, K., Tsien, R. W., & Luo, L. (2013). GABAergic Projection Neurons Route Selective Olfactory Inputs to Specific Higher-Order Neurons. *Neuron*, 79(5), 917–931. <https://doi.org/10.1016/j.neuron.2013.06.014>

Liu, W. W., & Wilson, R. I. (2013). Glutamate is an inhibitory neurotransmitter in the *Drosophila* olfactory system. *Proceedings of the National Academy of Sciences*, 110(25), 10294–10299. <https://doi.org/10.1073/pnas.1220560110>

MacWilliam, D., Kowalewski, J., Kumar, A., Pontrello, C., Ray, A. (2018). Signaling Mode of the Broad-Spectrum Conserved CO₂ Receptor Is One of the Important Determinants of Odor Valence in *Drosophila*. *Neuron*, 97(5), 1153–1167.e4.

<https://doi.org/10.1016/J.NEURON.2018.01.028>

Mahr, A., & Aberle, H. (2006). The expression pattern of the *Drosophila* vesicular glutamate transporter: A marker protein for motoneurons and glutamatergic centers in the brain. *Gene Expression Patterns*, 6(3), 299–309. <https://doi.org/10.1016/J.MODGEP.2005.07.006>

Marin, E. C., Jefferis, G. S. X. E., Komiyama, T., Zhu, H., & Luo, L. (2002). Representation of the Glomerular Olfactory Map in the *Drosophila* Brain. *Cell*, 109(2), 243–255.

[https://doi.org/10.1016/S0092-8674\(02\)00700-6](https://doi.org/10.1016/S0092-8674(02)00700-6)

Min, S., Ai, M., Shin, S. A., & Suh, G. S. B. (2013). Dedicated olfactory neurons mediating

attraction behavior to ammonia and amines in *Drosophila*. *Proceedings of the National Academy of Sciences of the United States of America*, 110(14).
<https://doi.org/10.1073/pnas.1215680110>

Mohamed, A. A. M., Retzke, T., Das Chakraborty, S., Fabian, B., Hansson, B. S., Knaden, M., & Sachse, S. (2019). Odor mixtures of opposing valence unveil inter-glomerular crosstalk in the *Drosophila* antennal lobe. *Nature Communications*, 10(1).
<https://doi.org/10.1038/s41467-019-09069-1>

Münch, D., & Galizia, C. (2016). DoOR 2.0--Comprehensive Mapping of *Drosophila melanogaster* Odorant Responses. *Scientific Reports*, 6. <https://doi.org/10.1038/SREP21841>

Ng, M., Roorda, R., Lima, S., Zemelman, B., Morcillo, P., Miesenböck, G. (2002). Transmission of olfactory information between three populations of neurons in the antennal lobe of the fly. *Neuron*, 36(3), 463–474. [https://doi.org/10.1016/S0896-6273\(02\)00975-3](https://doi.org/10.1016/S0896-6273(02)00975-3)

Parnas, M., Lin, A. C., Huetteroth, W., & Miesenböck, G. (2013). Odor Discrimination in *Drosophila*: From Neural Population Codes to Behavior. *Neuron*, 79(5), 932–944.
<https://doi.org/10.1016/j.neuron.2013.08.006>

Root, C., Semmelhack, J., Wong, A., Flores, J., Wang, J. (2007). Propagation of olfactory information in *Drosophila*. *Proceedings of the National Academy of Sciences of the United States of America*, 104(28), 11826–11831. <https://doi.org/10.1073/PNAS.0704523104>

Ruta, V., Datta, S. R., Vasconcelos, M. L., Freeland, J., Looger, L. L., & Axel, R. (2010). A dimorphic pheromone circuit in *Drosophila* from sensory input to descending output. *Nature*, 468(7324), 686–690. <https://doi.org/10.1038/nature09554>

Schmucker, M., de Bruyne, M., Hähnel, M., & Schneider, G. (2007). Predicting olfactory receptor neuron responses from odorant structure. *Chemistry Central Journal* 2007 1:1, 1(1), 1–10.
<https://doi.org/10.1186/1752-153X-1-11>

Schmucker, M., & Schneider, G. (2007). Processing and classification of chemical data inspired by insect olfaction. *Proceedings of the National Academy of Sciences*, 104(51), 20285–20289. <https://doi.org/10.1073/PNAS.0705683104>

- 650 Seki, Y., Dweck, H. K. M., Rybak, J., Wicher, D., Sachse, S., & Hansson, B. S. (2017).
651 Olfactory coding from the periphery to higher brain centers in the Drosophila brain. *BMC*
652 *Biology*, 15(1), 18–22. <https://doi.org/10.1186/s12915-017-0389-z>
- 653 Shearin, H. K., Quinn, C. D., Mackin, R. D., Macdonald, I. S., & Stowers, R. S. (2018). t-
654 GRASP, a targeted GRASP for assessing neuronal connectivity. *Journal of Neuroscience*
655 *Methods*, 306, 94–102. <https://doi.org/10.1016/j.jneumeth.2018.05.014>
- 656 Stocker, R. F., Heimbeck, G., Gendre, N., & De Belle, J. S. (1997). Neuroblast ablation in
657 Drosophila P[GAL4] lines reveals origins of olfactory interneurons. *Journal of*
658 *Neurobiology*, 32(5), 443–456. [https://doi.org/10.1002/\(SICI\)1097-](https://doi.org/10.1002/(SICI)1097-4695(199705)32:5<443::AID-NEU1>3.0.CO;2-5)
659 [4695\(199705\)32:5<443::AID-NEU1>3.0.CO;2-5](https://doi.org/10.1002/(SICI)1097-4695(199705)32:5<443::AID-NEU1>3.0.CO;2-5)
- 660 Strutz, A., Soelter, J., Baschwitz, A., Farhan, A., Grabe, V., Rybak, J., Knaden, M., Schmuker,
661 M., Hansson, B. S., & Sachse, S. (2014). Decoding odor quality and intensity in the
662 Drosophila brain. *ELife*, 3, e04147. <https://doi.org/10.7554/eLife.04147>
- 663 Tanaka, N. K., Awasaki, T., Shimada, T., & Ito, K. (2004). Integration of chemosensory
664 pathways in the Drosophila second-order olfactory centers. *Current Biology*, 14(6), 449–
665 457. <https://doi.org/10.1016/j.cub.2004.03.006>
- 666 Taniguchi, H., Gollan, L., Scholl, F. G., Mahadomrongkul, V., Dobler, E., Limthong, N., Peck,
667 M., Aoki, C., & Scheiffele, P. (2007). Silencing of neuroligin function by postsynaptic
668 neurexins. *Journal of Neuroscience*. <https://doi.org/10.1523/JNEUROSCI.0032-07.2007>
- 669 Varela, N., Gaspar, M., Dias, S., & Vasconcelos, M. L. (2019). Avoidance response to CO₂ in
670 the lateral horn. *PLoS Biology*, 17(1), 1–26. <https://doi.org/10.1371/journal.pbio.2006749>
- 671 Wang, J. W., Wong, A. M., Flores, J., Vossahl, L. B., & Axel, R. (2003). Two-photon calcium
672 imaging reveals an odor-evoked map of activity in the fly brain. *Cell*, 112(2), 271–282.
673 [https://doi.org/10.1016/S0092-8674\(03\)00004-7](https://doi.org/10.1016/S0092-8674(03)00004-7)
- 674 Wilson, R. I., Turner, G. C., & Laurent, G. (2004). Transformation of Olfactory Representations
675 in the Drosophila Antennal Lobe. *Science (New York, N.Y.)*, 303(5656), 366–370.
676 <https://doi.org/10.1126/science.1090782>

- Wong, A. M., Wang, J. W., & Axel, R. (2002). Spatial representation of the glomerular map in the *Drosophila* protocerebrum. *Cell*, 109(2), 229–241. [https://doi.org/10.1016/S0092-8674\(02\)00707-9](https://doi.org/10.1016/S0092-8674(02)00707-9)
- Xing, G., Li, M., Sun, Y., Rui, M., Zhuang, Y., Lv, H., Han, J., Jia, Z., & Xie, W. (2018). Neurexin–neuroligin 1 regulates synaptic morphology and functions via the WAVE regulatory complex in *Drosophila* neuromuscular junction. *ELife*, 7, 1–23. <https://doi.org/10.7554/eLife.30457>
- Zhang, K., Guo, J., Peng, Y., Xi, W., Guo, A. (2007). Dopamine-mushroom body circuit regulates saliency-based decision-making in *Drosophila*. *Science (New York, N.Y.)*, 316(5833), 1901–1904. <https://doi.org/10.1126/SCIENCE.1137357>

Figure 1: Selective labeling of LHNs based on their neurotransmitter identities. (A) Photoactivation of *UAS-C3PA-GFP* expressed under the control of different Gal4 lines with various neurotransmitter and neuromodulator identities (*Cha-Gal4*, *GAD1-Gal4*, *dVGlut-Gal4*, *TH-Gal4* and *tdc2-Gal4*) reveals subsets of cholinergic, GABAergic, glutamatergic, octopaminergic and dopaminergic LHNs (left to right). Position of the cell bodies are demarcated with yellow arrowheads. (B) Immunohistochemistry of photoactivated brains with *dVGlut-Gal4* driving *UAS-C3PA-GFP* reveals different cell body clusters of glutamatergic LHNs found at various locations in different focal planes. *GH146-QF* driving *QUAS-td tomato* in the background are depicted in red and nc82 in blue. Position of the cell bodies are shown with yellow arrowheads. Scale bars = 10 μ m.

Figure 2: Odor response properties of glutamatergic LHNs and uPNs. (A) Representative images of calcium responses of glutamatergic LHNs in the LH brain area in three focal planes, evoked by various odors (repulsive odors: acetophenone, geosmin; attractive odors: 2,3

butanedione, propionic acid; pheromone: cVA). Right panel shows the innervation pattern of uPNs labeled by GH146 in three different focal planes, used as a landmark to maintain comparable focal planes for functional imaging across different animals. (B) Grid approach to analyze odor-evoked responses of glutamatergic LHNs. Representative images of the grid onto uPN labeling (upper panel), used as a background landmark, odor-evoked responses of glutamatergic LHNs (middle panel) and analyzed $\Delta F/F$ of each pixels from glutamatergic LHNs in the LH area (lower panel). (C) Scatter plot showing calcium responses of each pixel of glutamatergic LHNs at middle plane to repeated presentation of an odor stimulus. Different colored dots depict responses obtained from different animals. (D) Box plot represents comparison of odor-evoked responses of glutamatergic LHNs at middle plane, between repeated and across odor stimuli (student t-test, *** $p < 0.001$). (E) Comparison of correlation coefficients of uPN and LHN responses at three different focal planes. Different odors seem to be better segregated at the uPN level than at the level of LHNs (student t-test, *** $p < 0.001$). (F) Cluster analysis of LHN odor responses based on linkage distance ($n = 7$, ANOSIM, sequential Bonferroni significance, $p = 0.0001$). (G) Cluster analysis of uPN odor responses based on linkage distance ($n = 8$; ANOSIM, sequential Bonferroni significance, $p = 0.0001$). Both cluster analyses show that the majority of replicates for each odor are grouped together.

Figure 3: Segregation of hedonic valence in the LH. (A) Comparison of the correlation coefficients of odor responses of glutamatergic LHNs between the odors that share the same valence (i.e. within valence) and odors having an opposing valence (i.e. across valence). Responses to odors sharing the same valence are significantly more similar than those across valence (student

t-test, *** $p < 0.001$, * $p < 0.05$) (B) PCA analysis of odor responses of glutamatergic LHNs shows that attractive (green) and aversive (red) odors are significantly segregated in the LH (one way ANOSIM, $p = 0.0166$). (C) Response amplitudes of glutamatergic LHNs to individual attractive and aversive odors. (D) Comparison of the overall response strength of glutamatergic LHNs and uPNs to all attractive and aversive odors (student t-test, *** $p < 0.001$) (E) PCA analysis of odor responses of uPNs shows that attractive (green) and aversive (red) odors are significantly segregated in the LH (one way ANOSIM, $p = 0.0049$). (F) Response amplitudes of uPNs to individual attractive and aversive odors. (G) Scatter plot of olfactory preference indices of all tested odors to the response strengths of glutamatergic LHNs. The solid trend line demarcates significant correlation between the two parameters (Spearman Rank Correlation, $p = 0.04$). (H) Scatter plot of olfactory preference indices of all tested odors to the response strengths of uPNs, which does not show a significant correlation (dotted trend line). (I) Scatter plot of the response strength of glutamatergic LHNs for individual odors to the overall OSN activity. (J) Scatter plot of the response strength of glutamatergic LHNs for individual odors to the number of ORs activated. (K) Scatter plot of the response strength of glutamatergic LHNs and the response strengths of uPNs for individual odors.

Figure 4: Polarity and morphological connectivity of glutamatergic LHNs with uPNs and mPNs. (A) Expression of syp GC3 (left panel) and homer GC3 (right panel) in glutamatergic LHNs suggesting the presence of pre- as well as postsynapses in the LH. (B) Membrane-targeted CD4-GRASP between glutamatergic LHNs and uPNs (left panel) or mPNs (right panel). (C) The synaptic protein neurexin tagged to GRASP (Nrx-GRASP) was employed to confirm synaptic connectivity of glutamatergic LHNs with uPNs (left panel) or mPNs (right panel) in the LH. Scale bars = 10 μ m.

Figure 5: Glutamatergic LHNs receive major excitatory input from uPNs and an odorant selective inhibition from mPNs. Upper panel schematic represents the experimental approach: the iACT tract was laser transected while odor-evoked responses were monitored from glutamatergic LHNs. (A, A' and A'') Representative images and graphical comparison of response evoked by vinegar in glutamatergic LHNs before and after laser transection of the iACT, across three different planes. (B, B' and B'') Representative images and graphical comparison of response evoked by benzaldehyde in glutamatergic LHNs before and after laser transection of the iACT. Lower panel schematic represents the experimental approach: the mACT tract was laser transected while odor-evoked responses were measured from glutamatergic LHNs. (C, C' and C'') Representative images and graphical comparison of vinegar-evoked responses of glutamatergic LHNs before and after laser transection of the mACT. (D, D' and D''). Representative images and graphical comparison of benzaldehyde-evoked response of glutamatergic LHNs before and after laser transection of the mACT (paired t-test *** $p < 0.001$, * $p < 0.05$).

Figure 6: mPN-mediated inhibition facilitates odor specificity in glutamatergic LHNs. (A and A') Schematic represents the experimental approach: odor-evoked responses of glutamatergic LHNs were monitored to repeated odor presentations (1 and 2) before and after laser transection of the mACT (3) for the odors vinegar and benzaldehyde. (B and B') Comparison of the correlation coefficients between the repeated odor responses before laser ablation to those after the transection for vinegar and benzaldehyde (within odor comparison). (C) Comparison of the correlation coefficients across vinegar and benzaldehyde before and after microlesion (across odor comparison; paired t-test, *** $p < 0.001$, ** $p < 0.005$, * $p < 0.05$). (D) Schematic summarizing the observed connectivity between PNs and glutamatergic LHNs in the LH: Glutamatergic LHNs consists of both, LHLNs and LHONs. uPNs (green) provide the major excitatory input to glutamatergic LHNs (yellow) while they might also receive feedback input from LHNs. mPNs provide an odorant selective inhibition to glutamatergic LHNs. mPNs are known to encode odor valence in the LH (Strutz et al., 2014), uPNs and glutamatergic LHNs encode odor identity as well

782 as odor valence. The table illustrates that uPNs encode an improved odor identity than postsynaptic
 783 glutamatergic LHNs whereas hedonic valence is equally maintained at the level of uPNs and
 784 glutamatergic LHNs in the LH.

785 **Figure 5- figure supplement 1: Laser transaction does not affect activity of glutamatergic**
 786 **LHNs in the intact brain hemisphere.** Left panel shows the comparison of benzaldehyde-evoked
 787 activity in glutamatergic LHNs in the intact hemisphere, before and after laser ablation of the iACT
 788 in the other, treated brain hemisphere (paired t-test, $p = 0.19$). Right panel shows the comparison
 789 of benzaldehyde-evoked activities in glutamatergic LHNs in the intact hemisphere, before and
 790 after laser ablation of the mACT in the other hemisphere (paired t-test, $p = 0.25$).