SUMMARY

Each odorant receptor gene defines a unique type of olfactory receptor neuron (ORN) and a corresponding type of second-order neuron. Because each odor can activate multiple ORN types, information must ultimately be integrated across these processing channels to form a unified percept. Here, we show that, in Drosophila, integration begins at the level of second-order projection neurons (PNs). We genetically silence all the ORNs that normally express a particular odorant receptor and find that PNs postsynaptic to the silent glomerulus receive substantial lateral excitatory input from other glomeruli. Genetically confining odor-evoked ORN input to just one glomerulus reveals that most PNs postsynaptic to other glomeruli receive indirect excitatory input from the single ORN type that is active. Lateral connections between identified glomeruli vary in strength, and this pattern of connections is stereotyped across flies. Thus, a dense network of lateral connections distributes odor-evoked excitation between channels in the first brain region of the olfactory processing stream.

INTRODUCTION

All the olfactory receptor neurons (ORNs) that express the same odorant receptor gene respond similarly to odors, and all project to the same sphere of neuropil (glomerulus) in the brain (Buck, 1996). Furthermore, in most species, each second-order olfactory neuron receives direct synaptic input from only one glomerulus. Thus, each odorant receptor gene defines a unique parallel processing channel in the olfactory system.

There is good evidence for crosstalk between these glomerular channels very early in the olfactory system, as early as the level of second-order neurons. Anatomically, glomeruli are interconnected by a network of local interneurons. Functionally, synaptic connections have been demonstrated between interneurons and second-order principal neurons, both in the insect antennal lobe and in the mammalian olfactory bulb (Jahr and Nicoll, 1980; Ernst and Boeckh, 1983; Hoskins et al., 1986; Malun, 1991; Leitch and Laurent, 1996; MacLeod and Laurent, 1996; Stöcker et al., 1997; Christensen et al., 1998; Isaacson and Strowbridge, 1998; Python and Stöcker, 2002; Urban and Sakmann, 2002; Aungst et al., 2003; Hayar et al., 2004; Wilson et al., 2004; Wilson and Laurent, 2005). It seems likely that these lateral connections have a role in shaping the output of the olfactory bulb and the antennal lobe. However, none of these studies has directly assessed the net effect of glomerular crosstalk on second-order neurons during in vivo odor stimulation.

Understanding how the antennal lobe and olfactory bulb transform olfactory representations requires an in vivo characterization of interglomerular synaptic connections. Is the net effect of these lateral connections inhibitory or excitatory? How powerful are these connections compared to direct ORN inputs? How does the amount of lateral input to a second-order neuron vary across odor stimuli? What rules govern the pattern of synaptic connectivity between glomeruli? Is a single glomerulus only connected with glomeruli in its local vicinity? Is each glomerulus only connected to a few others, or is the pattern of connectivity more dense? Is there a correlation between glomerular connectivity and odor tuning? To what extent is the pattern of lateral connectivity stereotyped from one animal to the next?

The most direct way to characterize lateral input to a second-order olfactory neuron would be to silence its direct ORN inputs while preserving ORN input to another glomerulus or glomeruli. Any remaining odor responses in cells postsynaptic to a “silent” glomerulus should reflect purely lateral inputs. This type of experiment is currently not possible in vertebrates, so we have turned to the Drosophila antennal lobe. This structure shares the same basic architecture as the vertebrate olfactory bulb but presents several experimental advantages. First, the odorant receptor gene expressed by most Drosophila ORN types has already been identified and mapped onto spatially stereotyped glomeruli in the antennal lobe.
(Couto et al., 2005; Fishilevich and Vosshall, 2005; Hallem and Carlson, 2006). Second, the circuitry of the Drosophila antennal lobe develops properly in the absence of normal olfactory activity, and ORNs target the correct glomerulus even when they do not express a functional receptor (Dobritsa et al., 2003; Berdnik et al., 2006). Finally there are only ∼50 glomeruli in the Drosophila antennal lobe, as compared to ∼1000 in the mouse olfactory bulb (Laiissue et al., 1999). This makes it possible to record from second-order neurons (called projection neurons, or PNs) postsynaptic to an identified ORN input. Importantly, like olfactory bulb mitral cells in most vertebrates, PNs in the Drosophila antennal lobe receive direct ORN input from a single glomerulus.

Circumstantial evidence suggests that interglomerular shape pattern odor responses in Drosophila. A comparison of odor-evoked responses in ORNs and PNs corresponding to the same glomerulus revealed that the rank order of a PN’s odor preferences can be different from the preferences of its presynaptic ORNs (Wilson et al., 2004). This argues that a PN’s responses are not completely determined by its direct ORN inputs. Instead, it implies that PNs integrate input from multiple ORN types. However, this conclusion has been challenged by functional imaging studies (Ng et al., 2002; Wang et al., 2003).

Here, we use a variety of genetic manipulations and microdissections to remove direct ORN inputs to one or more glomeruli in the Drosophila antennal lobe. In vivo recordings from PNs postsynaptic to “silent” glomeruli reveal that these PNs receive lateral inputs from other glomeruli. The net effect of lateral synaptic inputs to PNs is predominately excitatory and can be strong enough to trigger a train of action potentials. In order to define the functional connectivity between identified glomeruli, we have generated flies with only a single active ORN type. Stimulation of just one ORN type is sufficient to recruit lateral excitatory inputs to other glomeruli, but this sensitivity also leads to saturation as more ORN types are activated. A single glomerulus provides indirect excitatory input to most, if not all, glomeruli, thus defining a dense network of lateral connections spanning the entire antennal lobe. However, some lateral excitatory connections are substantially stronger than others. Finally, we find that this pattern of connection strengths is relatively stereotyped across flies, suggesting that it may be genetically hardwired. These findings directly demonstrate that synaptic crosstalk between glomeruli can shape PN odor responses in vivo. Furthermore, our results reveal some of the fundamental rules governing these interglomerular interactions.

RESULTS

Olfactory Stimuli Trigger Lateral Excitatory Interactions among Glomeruli

The Drosophila antennal lobes receive olfactory input from two peripheral organs, the antennae and the maxillary palps. The palps contain ∼120 ORNs which fall into six distinct types not found in the antennae (de Bruyne et al., 1999; Goldman et al., 2005). The antennae contain an additional 43 ORN types not found in the palps (de Bruyne et al., 2001; Hallem et al., 2004). Thus, each glomerulus receives direct ORN input exclusively from either the palps or the antennae. Within the antennal lobe, the six palp glomeruli are intermingled with the 43 antennal glomeruli (Couto et al., 2005; Fishilevich and Vosshall, 2005). This anatomy provides a convenient way to independently manipulate inputs to two groups of glomeruli.

In order to determine whether interactions between glomeruli shape PN odor responses, we began by acutely severing the antennal nerves. This manipulation leaves the palp ORNs intact and allows us to test whether antennal PNs receive lateral input from palp glomeruli (Figure 1A). Consistent with previous reports (Berdnik et al., 2006), we found that ablating ORN input to some glomeruli does not induce morphological rearrangement of the remaining ORN axons, even five days postsurgery (Figure 1B). However, we cannot exclude the possibility that removing ORN input to some glomeruli induces more subtle synaptic plasticity of interglomerular connections. Therefore, we performed PN recordings immediately after severing the antennal nerves in order to rule out a role for any such plasticity.

Whole-cell patch-clamp recordings from PNs typically show abundant spontaneous synaptic input (Figure 1C). In contrast, PNs postsynaptic to antennal glomeruli in antennae-less flies show no spontaneous activity (Figures 1D and 1E). Nevertheless, odor stimulation of the maxillary palps evokes a depolarization in these PNs (n = 6 PNs in 6 flies; Figures 1D and 1E). The magnitude of this depolarization varied across cells but was sufficient to produce a train of spikes in a PN postsynaptic to glomerulus VCS, for example (Figure 1E). We confirmed that each of the PNs we recorded from innervated an antennal glomerulus by filling the cell with biocytin (see Experimental Procedures).

We averaged the membrane potential across six presentations of the same odor in the same cell (Figure 1F) before averaging across experiments (Figure 1G). All these odors strongly activate one or more palp ORN types (de Bruyne et al., 1999), and all elicited a substantial depolarization in each of the antennal PNs we recorded from. The solvent we use to dilute our odors (paraffin oil) evoked no response (Figure 1G). Removing both antennae and maxillary palps abolished odor-evoked depolarizations (n = 3), demonstrating that the maxillary palps mediate this response (Figure 1G).

These results demonstrate that antennal PNs receive lateral synaptic input from palp ORNs and that the net effect of this indirect input is excitatory. It is possible that some interglomerular synapses hyperpolarize PNs, but if so, this inhibition is evidently obscured by a larger excitatory component. It is important to note that ORN input was removed acutely (∼10–20 min before recording). Therefore, the lateral excitation we observed cannot reflect remodeling of the antennal lobe circuitry.
What is the total impact of all lateral synaptic input to a PN? In order to answer this question, we selectively silenced a single ORN type. By recording from a PN postsynaptic to a deafferented glomerulus reveals indirect input to that cell from palp ORNs.

Axon targeting of palp ORNs is not altered by removing antennal ORNs. Projections of confocal stacks through the antennal lobes (neuropil in magenta) show ORN axons labeled with CD8:GFP (green). Glomeruli targeted by palp ORNs are outlined. Arrow indicates maxillary nerve. Arrowheads indicate axons projecting to or from the midline; each ORN projects bilaterally. Scale bars = 20 μm.

With antennal nerve intact, a recording from an antennal PN (in glomerulus DM3) shows spontaneous and odor-evoked activity. Period of odor delivery is indicated by gray bar (500 ms).

Recording from an antennal PN (in glomerulus DM3) after severing antennal nerves. Spontaneous activity is abolished, but a small odor-evoked depolarization remains.

Recording from an antennal PN (in glomerulus VC3) after severing antennal nerves shows a large odor-evoked depolarization.

Odor-evoked depolarizations averaged across all experiments with severed antennal nerves (magenta). No depolarization is observed when both antennae and maxillary palps are removed (blue). Note that this analysis pools data from six PNs corresponding to different antennalglomeruli.

**Total Lateral Synaptic Excitation to a PN Is Substantial**

What is the total impact of all lateral synaptic input to a PN? In order to answer this question, we selectively silenced a single ORN type. By recording from PNs postsynaptic to the “silent” glomerulus while stimulating the antennae and palps with odors, we should be able to observe the total lateral input to that PN.

In order to silence a single ORN type, we used flies with mutations in one of two odorant receptor genes, Or43b and Or10a. Or43b is expressed in ORNs that project to glomerulus VM2, and Or10a is expressed in ORNs that project to glomerulus DL1 (Couto et al., 2005; Fishilevich and Vosshall, 2005). The Or43b<sup>1</sup> null allele was produced by gene targeting and has been described previously (Elmore et al., 2003). The Or10a<sup>f03694</sup> allele results from a pBac insertion in the gene (Thibault et al., 2004) but has not been characterized previously. We first verified that this mutation does not disrupt ORN axon targeting (see Figure S1A in the Supplemental Data available with this article online). We then confirmed that both the Or43b<sup>1</sup> and Or10a<sup>f03694</sup> mutations virtually abolish ORN odor responses (Figures S1B–S1D). We occasionally observed very weak responses to a few specific odors in...
both mutants (Figure S2). These weak responses may be mediated by Or83b, a protein with homology to odorant receptors that is expressed in most ORNs and which is required for trafficking receptors to the cell membrane (Larsson et al., 2004). We did not use these odors in our stimulus set when we analyzed PN responses in these mutants (Figures 2–4). Note that these mutations also reduce the rate of spontaneous ORN spikes (Figure S1C).

We then recorded from PNs postsynaptic to these “silent” ORNs (Figure 2A). In flies with the mutation that silences VM2 ORNs, we targeted VM2 PNs using an enhancer trap line (Tanaka et al., 2004) to specifically label these cells with GFP (NP5103-Gal4,UAS-CD8:GFP;Or43b1). We recorded from one PN per fly and confirmed the glomerular identity of each recorded PN by imaging the biocytin fill post hoc. As expected from
the decrease in spontaneous activity in mutant ORNs, spontaneous spiking in VM2 PNs was reduced compared to wild-type (Figures 2B and 2C). Similarly, in flies with mutant DL1 ORNs, we recorded from DL1 PNs using an enhancer trap line (Tanaka et al., 2004) to specifically label these cells with GFP (Or10af03694;+/+;NP3529-Gal4, UAS-nlsGFP). Again, spontaneous spiking was reduced in these PNs (Figures 2B and 2D). We confirmed that these PNs show normal dendrite morphology in the absence of functional ORNs (Figures 2C and 2D), consistent with previous reports (Wong et al., 2002; Berdnik et al., 2006).

In flies with silent VM2 ORNs, all VM2 PNs were depolarized by every odor we tested (n = 10 cells in 10 flies). Again, most responses were large enough to elicit a train of spikes. We also saw similar spiking responses to odors in cell-attached mode (prior to going whole-cell), demonstrating that these depolarizations are not an artifact of intracellular dialysis.

As expected, most odors evoked a larger response in wild-type flies than in mutant flies. For example, ethyl butyrate elicits vigorous activity in normal VM2 ORNs (Figure S1) and in wild-type VM2 PNs. When the VM2 ORNs are silenced, this odor elicits a much smaller response in VM2 PNs (Figure 3A1). Similarly, methyl salicylate evokes a very strong response in DL1 ORNs (Figure S1) and in wild-type DL1 PNs but only a small response in DL1 PNs postsynaptic to silent ORNs (Figure 3B1). We also noticed that in PNs postsynaptic to silent ORNs, odor-evoked responses were often more transient than in wild-type PNs (Figures 3A1 and 3B1).

Figure 3. Comparing Odor Responses in PNs Postsynaptic to Normal versus “Silent” ORNs
Representative responses of PNs postsynaptic to wild-type ORNs (left column) and PNs postsynaptic to nonfunctional ORNs (right column). Most odors elicit a substantially larger response when presynaptic ORNs are functional (A1, B1). However, some odors elicit a similar response with or without functional presynaptic ORNs (A2, B2). Gray bars indicate the 500 ms period of odor stimulation. Transient stimulus artifacts from the olfactometer (at the end of the odor stimulus period) were blanked in some traces. Traces in the same row have the same y axis.
Excitatory Interactions in Drosophila Antennal Lobe

Figure 4. Odor Tuning of PN Responses Postsynaptic to Normal versus “Silent” ORNs

(A and B) Tuning curves comparing odor responses of PNs postsynaptic to wild-type (magenta) and mutant ORNs (blue). PNs are postsynaptic to glomerulus VM2 (A) or DL1 (B). Each point represents firing rate over the 500 ms odor stimulus period, averaged across experiments (mean ± SEM). Note different y scales for magenta and blue symbols. Odors are arranged so the smallest wild-type responses are on the left and the largest are on the right. See Experimental Procedures for odor abbreviations.

(C) Tuning curves comparing odor responses of VM2 and DL1 PNs postsynaptic to mutant ORNs. Note correlated but not identical odor tuning.

Some odor responses, however, were relatively unaffected by odorant receptor mutations. For example, 2,3-butanedione evokes very little response in wild-type VM2 ORNs (Figure S1). When these ORNs are silenced, the response of VM2 PNs to this odor is virtually unaltered (Figure 3A2). Similarly, ethyl acetate does not excite wild-type VM2 ORNs (Figure S1). When these ORNs are silenced, butanedione evokes very little response in wild-type PNs but ethyl acetate is undiminished by silencing their presynaptic ORNs (Figure 3B2).

How does the size of total lateral input to a PN depend on the odor stimulus? We quantified odor responses by computing mean firing rates over the 500 ms duration of the odor stimulus, and we plotted these response magnitudes for each odor stimulus to produce tuning curves (Figures 4A and 4B). These plots show that different odors evoke different amounts of lateral excitatory input to each PN. However, the odor tuning of PNs post-synaptic to silent ORNs is completely different from the normal odor tuning of these PNs. For both glomeruli, the odor tuning of wild-type and mutant PNs showed no significant correlation (both comparisons Pearson’s $r^2 < 0.05$, $p > 0.4$).

We also noted that, in the absence of direct ORN inputs, both VM2 PNs and DL1 PNs are broadly tuned to odors. This suggests that each of these glomeruli receives indirect excitatory input from multiple ORN types, not just one or two. We wondered whether these two glomeruli receive indirect input from similar or different populations of ORNs. To assess this, we compared the responses of PNs in these two “silent” glomeruli to the same odors. Overall, the responses of these two PN types are significantly correlated (Figure 4C; Pearson’s $r^2 = 0.31$, $p < 0.05$). However, some odors elicit different amounts of lateral input to these glomeruli. For example, butyric acid elicits a larger response in VM2 than in DL1 (p = 0.05, t test, n = 6 for each glomerulus). These results are consistent with the idea that these two glomeruli receive indirect input from overlapping populations of ORNs but that the indirect inputs to these PNs are not identical.

Characterizing Lateral Input to Many Glomeruli Originating from a Single Glomerulus

Because total lateral excitatory input to VM2 and DL1 PNs is broadly tuned and significantly correlated, it is likely that two glomeruli receive indirect input from many of the same ORNs. This, in turn, implies that each ORN type provides indirect input to many glomeruli. To test this prediction directly, we designed experiments to measure the spread of excitation across the antennal lobe evoked by activation of a single ORN type. We used two approaches to selectively stimulate one ORN type (Figure 5). These two approaches have complementary strengths and weaknesses but yielded similar results. In the first method (experiment 1), we took advantage of a mutation in Or83b. This gene is expressed in most ORNs and encodes a chaperone protein required for trafficking odorant receptors to ORN dendrites (Larsson et al., 2004; Benton et al., 2006). Mutating Or83b abolishes odor responses in all maxillary palp ORNs (and many antennal ORNs; Figure S3). Thus, all ORN input to the antennal lobes is abolished by removing the antennae of mutant flies. In these flies, we then rescued normal function in the maxillary palp ORNs that project to glomerulus VA71. This was done by expressing Or83b under the control of the odorant receptor gene promoter corresponding to the VA71 ORNs (Or46-Gal4/UAS-Or83b; Or83b-GFP; Fishilevich and Vosshall, 2005). We verified the specificity of this rescue by making extracellular ORN recordings from the maxillary palps. Each sensillum in the palp contains exactly two ORNs, and a VA71 ORN is always paired with an ORN that projects to another glomerulus (de Bruyne et al., 1999; Goldman et al., 2005). In the palps of “rescued” flies, we encountered only silent sensilla, or sensilla containing exactly one spontaneously active ORN (n = 51 sensilla). This ORN always displayed odor tuning that matched the odor tuning of wild-type VA71 ORNs (Figures 5B and S4). We verified that the rescued ORNs correctly target glomerulus VA71 by coexpressing CD8:GFP with Or83b (Figure 5C). Furthermore, biocytin fills show that...
PNs postsynaptic to neighboring glomeruli do not inappropriately invade glomerulus VA7l (Figure 5C). This argues that the antennal lobe circuitry is grossly normal in the Or83b2 mutant.

This approach permits us to selectively stimulate exactly one ORN type. However, it has the drawback that most ORN types are inactive during the development of the fly, which could conceivably produce subtle changes in the antennal lobe circuitry. To address this issue, we used a second method to stimulate one ORN type under conditions in which almost all ORNs are normal and active throughout the life of the fly; this ensures that normal antennal lobe circuitry is preserved. In this approach (experiment 2, Figure 5D), we sought to identify a panel of odors that only stimulates one ORN type. Because most ORNs respond to many different odors, it is difficult to find such an odor set. To simplify the problem, we cut the antennal nerves just prior to recording, leaving only the six maxillary palp ORN types. Additionally, we used flies bearing the D85 mutation, which abolishes most odor responses in two of the six maxillary palp ORN types (Figures S5 and S6). Thus, in antennae-less flies in a D85 background, there are only four functional maxillary palp ORN types. We screened a large panel of odors and identified a set of 14 that exclusively stimulates the VM7 ORNs while evoking no response from the other three functional ORN types in this genotype. This is demonstrated by local field potential recordings from the maxillary palp. When VM7 ORNs are functional, all odors in this set evoke a local field potential response. These responses are abolished by the Or42aD4305 mutation (Thibault et al., 2004), which renders VM7 ORNs nonfunctional (Figure 5E; see also Figures S5 and S6). In summary, both experiments 1 and 2 permit selective stimulation of one ORN type.
In experiment 1 (genetically rescuing VA7l ORNs), we recorded from a total of 72 PNs postsynaptic to nonrescued glomeruli. PNs were selected at random from the dorsal cluster of PN cell bodies in the antennal lobe. Only one PN was recorded in each fly, which allowed us to unambiguously determine the glomerular identity of each PN after filling it with biocytin. Together, these 72 PNs targeted 24 out of the 49 glomeruli in the antennal lobe. In every one of these PNs, we observed a depolarization while stimulating the VA7l ORNs with odors. This implies that the VA7l ORNs broadcast indirect excitatory input to most glomeruli. As expected, the tuning of the lateral excitatory input to PNs always reflected the tuning of the VA7l ORNs (Figure 6A). Odors that excited VA7l ORNs produced depolarizations in PNs, whereas inhibition of VA7l ORNs led to a small hyperpolarization. This hyperpolarization probably represents an interruption in tonic lateral excitation driven by spontaneous action potentials in VA7l ORNs. In both experiment 1 (VA7l-only) and experiment 2 (VM7-only), the magnitude of lateral depolarization did not scale linearly with ORN firing rate. Odors that evoked only a small ORN response produced a near-maximal lateral depolarization in PNs. Odors that evoked a larger ORN response saturated the lateral circuitry. This is shown by plotting the area under the membrane potential deflection (computed after low-pass filtering the membrane potential) versus ORN firing rate (Figures 6C and 6D). The nonlinearity of these curves illustrates both the sensitivity and the saturation of interglomerular excitatory circuits. To confirm that the lateral depolarizations in experiment 2 are driven by the VM7 ORNs, we combined the \( D_85 \) mutation with an odorant receptor gene mutation (\( Or42af04305; D_85 \)). In this genotype, we saw essentially no odor responses in any PNs (Figure 6D, open circles, \( n = 3 \)).

These results demonstrate that ORN inputs from one glomeral processing channel can easily saturate the lateral excitatory circuitry of the antennal lobe. How are inputs from multiple ORN types integrated by this circuitry? We used an odor blend to investigate how odor-evoked signals from two ORN inputs are combined. In antennae-less \( D_85 \) flies, the odor 2-butanone (10\(^{-5}\) dilution) activates only the VM7 ORNs (Figure 5E), while the odorant methyl salicylate (10\(^{-2}\) dilution) activates only the VA7l ORNs (data not shown). In isolation, these stimuli

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**Figure 6. Odor Stimulation of One ORN Type Evokes Lateral Input to Many PNs**

(A) Experiment 1. Top, peristimulus-time histograms show odor responses of VA7l ORNs (n = 5). Bottom, average depolarizations recorded in PNs postsynaptic to glomeruli lacking direct ORN input (n = 72).

(B) Experiment 2. Top, peristimulus-time histograms showing odor responses of VM7 ORNs (n = 6). Bottom, average depolarizations recorded in PNs postsynaptic to glomeruli lacking direct ORN input (n = 15).

(C) Experiment 1. Average depolarization area plotted versus VA7l ORN firing rate. Each point represents a different odor. Curve is an exponential fit (only excitatory ORN responses are included in fit).

(D) Experiment 2. Average depolarization area plotted versus VM7 ORN firing rate for each odor (solid symbols). Curve is an exponential fit. In flies lacking functional VM7 ORNs (open symbols), odor-evoked depolarizations are virtually absent (\( Or42af04305; D_85 \); \( n = 3 \)).

All panels: values are mean ± SEM, averaged across experiments.
Inputs from Multiple ORN Types

Figure 7. Lateral Excitatory Circuits Sublinearly Summately Inputs from Multiple ORN Types

(A) Average depolarization evoked in antennal PNs by selective stimulation of VM7 ORNs with 2-butanol (10^{-5} dilution) in Δ85 flies. (B) Average depolarization evoked in antennal PNs by selective stimulation of VA7l ORNs with methyl salicylate (10^{-2} dilution) in Δ85 flies, recorded from the same PNs as in (A). (C) Comparing the average depolarization evoked by simultaneous stimulation of both ORN types (green), versus the predicted linear sum of stimulating each ORN type alone (black). (D) Depolarization quantified as the area under the membrane potential traces. What happens when these two inputs are integrated? When we blended the two odors to stimulate both VA7l and VM7 ORNs simultaneously in antennaeless Δ85 flies, we observed lateral depolarizations that were significantly larger than the response to either odor alone (Figure 7; p ≤ 0.01 for both comparisons, n = 8 PNs, paired t tests). This shows that multiple ORN inputs are effectively integrated by the lateral excitatory circuitry of the antennal lobe. However, the response to the blend was significantly smaller than that predicted by a linear sum of the responses evoked by stimulating each ORN type individually (Figure 7; p < 10^{-4}, paired t test). This demonstrates that saturation occurs across ORN input channels, not just within an ORN input channel.

For a given ORN input (either VA7l or VM7), different PNs showed substantially different amounts of lateral depolarization in response to the same odor. In experiment 1, for example, the response evoked by the strongest odor (4-methyl phenol) ranged from 1.7 mV to 11.8 mV. Responses in some PNs were large enough to elicit a few spikes, but responses in other PNs responses were much weaker. This suggests that different PNs receive different amounts of lateral excitatory input from any given glomerulus. Are these connections random, or are they a stereotyped function of glomerular identity? To address this question, we compared the responses of PNs corresponding to the same glomerulus recorded in different flies. In our data set from experiment 1, there were 11 glomeruli that we hit at least three times. In experiment 2, we recorded selectively from PNs in only three glomeruli by using an enhancer trap line to label these cells with GFP (NP3481-Gal4, UAS-CD8:GFP;+/+;Δ85). This allowed us to obtain multiple recordings from the same PN type.

Overall, we found that the strength of lateral excitatory connections was relatively stereotyped across flies. In quantitative terms, differences we observed between glomeruli were larger than the variability we observed for a given glomerulus across flies. For example, in experiment 1, selective stimulation of VA7l ORNs produced a significantly greater depolarization in VC4 PNs compared to VA1v PNs (Figures 8A–8C; p < 0.05; VC4, n = 7; VA1v, n = 8). In experiment 2, selective stimulation of VM7 ORNs produced a significantly greater depolarization in glomerulus DL5 than in glomerulus VM2 or DM6 (Figures 8D–8F; p < 0.05; DL5, n = 5; VM2, n = 3; DM6, n = 7). We also noted that the nonlinear relationship between the lateral depolarization and ORN firing rate had a similar sensitivity and exponential shape across different glomeruli; it is the saturation level that differs (Figures 8B and 8E). Together, these results demonstrate that the amount of lateral excitatory input to a PN depends on the identity of the glomerulus where its dendrite is located. The magnitude of lateral depolarization is not correlated with the input resistance of the cell (Pearson’s r^2 = 0.027), ruling out one trivial explanation for this systematic difference.

We next asked whether the magnitude of glomerular crosstalk varied systematically with distance. Figure 9 shows the relative connection strength for all 24 glomeruli we recorded from in experiment 1, mapped in relation to the location of the only glomerulus receiving direct ORN input (VA7l). Connection strength is defined as the relative response to 4-methyl phenol, which is the strongest stimulus in our odor set for the VA7l ORNs. This map illustrates that there is no obvious relationship between interglomerular distance and the strength of lateral excitatory connections. We also asked if there is a correlation between connection strength and the morphological class of the ORN type corresponding to each glomerulus. Previous studies have shown that ORNs housed in the three major morphological classes of sensilla (basi-conic, coeloconic, and trichoid) tend to project to nearby glomeruli and therefore define several zones in the antennal lobe (Couto et al., 2005; Fishilevich and Vosshall, 2005). Figure 9B shows that the zones defined by these sensillum classes receive similar overall levels of lateral excitation. Finally, we tested whether there is a relationship between the odor tuning of the normal ORN input to a glomerulus and the strength of excitatory lateral input it receives from VA7l. Comprehensive odor-tuning data for 16 of the colored glomeruli in Figure 9A has been...
compiled by other investigators (Hallem and Carlson, 2006). We used this data set to compute the correlation coefficient (Pearson’s r) between ORN odor responses for every pairwise combination of these 16 glomeruli and then plotted the difference in lateral depolarization for each pair as a function of this correlation. We found there is no relationship between glomerular tuning and depolarization strength (Figure 9C).

DISCUSSION

In this study, our goal was to observe the synaptic inputs to PNs arising from local antennal lobe circuits. We have used a variety of complementary strategies to remove direct ORN input to the PN we were recording from, meanwhile leaving other ORNs intact. These manipulations allowed us to directly observe lateral excitatory input to a PN originating from other glomeruli.

It is important to emphasize that this lateral excitation cannot be ascribed purely to compensatory rearrangement of the antennal lobe circuitry. This point is most forcefully demonstrated by experiments in which most or all ORNs are normal and active until we sever the antennal nerves immediately before recording (Figures 1 and 6B). In these experiments we recorded from PNs 10–20 min after removing the antennae and always observed odor-evoked lateral depolarizations. Hence, the circuitry mediating these responses must exist in normal flies prior to removing antennal input.

An Individual PN Integrates Lateral Inputs from Many ORN Types

Excitatory connections between glomeruli appear to be very dense, perhaps all-to-all. This conclusion is supported by four pieces of evidence. First, the magnitude of the depolarization we observed when almost all ORNs are intact (Figures 2–4) is larger than that observed when only the maxillary palp ORNs are intact (Figure 1), which in turn is larger than that observed when only a single ORN type is intact (Figures 5–9). This argues that most PNs receive indirect input from many ORN types. Second, when we restricted ORN input to a single glomerulus, every PN we recorded from (87 of 87 cells) received at least weak lateral input from that glomerulus. This implies that each ORN type broadcasts indirect input to most or all glomeruli. Third, the odor tuning of the total lateral input to a glomerulus is much broader than the odor tuning of a typical ORN. Fourth, the lateral input to VM2 PNs and DL1 PNs has a relatively similar (though not identical) odor-tuning profile. This suggests that large and
overlapping populations of ORNs provide indirect input to these two types of PNs. All-to-all connectivity is a parsimonious explanation for all these observations.

A Specific Matrix of Excitatory Connections

It should be noted that although lateral excitatory connectivity is dense and perhaps all-to-all, it is nevertheless selective. When we stimulated a single ORN type and recorded sequentially from PNs in different glomeruli, we found that each PN type receives a characteristically strong or weak lateral input from that ORN type. Furthermore, these characteristic connection strengths are relatively stereotyped across flies. This suggests that the synaptic connectivity of local interneurons in the antennal lobe may be genetically hardwired.

Notably, the strength of these lateral excitatory connections is not correlated with the distance between the target glomerulus and the location of the ORN inputs. This means that the spatial relationship between glomeruli does not limit the strength of their lateral interactions. This finding also argues that lateral excitation does not reflect spillover of excitatory neurotransmitter from the glomerulus receiving active ORN input, since in this case PNs closer to the active glomerulus would be expected to see a larger depolarization.

There is some tension between the idea that excitatory connection strengths between glomeruli are varied and the finding that VM2 and DL1 PNs see similarly tuned total lateral excitatory input. One possibility is that the lateral inputs to VM2 and DL1 PNs just happen to be unusually well correlated. Another possibility is that a given target glomerulus receives characteristically strong (or characteristically weak) indirect inputs from all ORN types. In this latter scenario, the strength of the lateral depolarization would vary across glomeruli, but its odor tuning would not.

Sensitivity and Saturation of Lateral Excitatory Circuits

The lateral excitatory circuits of the antennal lobe are remarkably sensitive to small levels of afferent input. We have shown that activating ORNs presynaptic to a single glomerulus produces a substantial lateral depolarization in many or all PNs. Moreover, the magnitude of the lateral depolarization arising from a single ORN type is extremely sensitive to small increases in ORN firing rate. Even an odor that evokes a very weak response in these ORNs (e.g., 1-butanol or geranyl acetate in Figure 6) still evokes substantial lateral excitation.

Another striking feature of lateral excitatory circuits is their saturability. In experiments where we stimulated only one ORN type, increasing the rate of incoming ORN spikes from 50 to 150 spikes/second had little effect on the amount of lateral excitatory input that was broadcast to other glomeruli. Furthermore, in experiments where we stimulated two ORN types, the combined effect of these two input channels was substantially less than the sum of each channel when stimulated individually. This type of saturation should tend to limit the magnitude of lateral excitatory synaptic input to a PN.

Figure 9. Lateral Excitation Is Broadly Distributed throughout the Antennal Lobe

(A) Schematic representation of all glomeruli in the antennal lobe, represented as three sections through the fly’s right lobe (modified from Laisanne et al. [1999]). Color indicates average relative depolarization measured in PNs postsynaptic to each glomerulus during selective stimulation of VA71 ORNs. Asterisk marks glomerulus VA71. We did not sample PNs postsynaptic to white glomeruli.

(B) Glomeruli postsynaptic to different morphological types of sensilla receive similar levels of lateral input. Graph compares average depolarization area (±SEM) evoked by 4-methyl phenol in glomeruli targeted by ORNs in basiconic sensilla (n = 14 glomeruli), coeloconic sensilla (n = 3 glomeruli), and trichoid sensilla (n = 6 glomeruli). Data on morphological types were taken from Couto et al. (2005).

(C) There is no relationship between odor tuning and strength of lateral input. Comprehensive odor-tuning data for 16 ORN types was taken from Hallem and Carlson (2006). For every possible pairwise combination of 16 glomeruli, the difference in the average lateral depolarization evoked by 4-methyl phenol in these two PN types is plotted versus the correlation between the odor tuning of the ORN inputs to those glomeruli.
Together, these results suggest that the impact of lateral excitatory connections might be strongly dependent on odor concentration. Testing this hypothesis will require comparing the sensitivity of direct and lateral inputs to a range of concentrations and understanding how these inputs are integrated by PNs.

A Cellular Substrate for Lateral Excitatory Connections

While this manuscript was under review, a report appeared that identified a novel population of cholinergic local neurons in the *Drosophila* antennal lobe (Shang et al., 2007). There is no direct evidence that these local neurons mediate the local excitatory connections we have observed, but this hypothesis seems plausible. Each cholinergic local neuron reportedly innervates most glomeruli, and this morphology could easily explain our observation that a single ORN type broadcasts excitatory input to most or all PNs. Interestingly, excitatory (glutamatergic) local neurons were also recently identified in the olfactory bulb (Aungst et al., 2003), although it is not known whether these cells make synapses onto mitral cells, the analog of antennal lobe PNs.

Shang et al. (2007) also independently provided evidence that PNs receive lateral excitatory input. As in our study (Figures 2–4), these investigators measured activity in PNs whose presynaptic ORNs have been silenced by an odorant receptor gene mutation. Complementary to our electrophysiological approach, Shang et al. (2007) used a genetically-encoded ecliptic pHluorin to monitor the balance of synaptic vesicle exocytosis and endocytosis at presynaptic sites in PN dendrites. They found that PNs whose presynaptic ORNs were silent still showed odor-evoked dendritic synaptopHluorin signals, implying that these PNs receive indirect excitatory input from other ORNs.

Lateral Inhibition in the *Drosophila* Antennal Lobe

Models of olfactory processing in the insect antennal lobe and the vertebrate olfactory bulb stress the importance of inhibitory connections between glomeruli (Mori et al., 1999; Laurent, 2002). What about lateral inhibition in the *Drosophila* antennal lobe? It is known that GABAergic interneurons ramify throughout the *Drosophila* antennal lobe and release GABA in response to odor stimulation (Stocker et al., 1997; Ng et al., 2002; Wilson and Laurent, 2005). *Drosophila* antennal lobe PNs have GABAA-like and GABAB-like receptors, and antagonists of these receptors disinhibit PN odor responses (Wilson and Laurent, 2005). Given this, it is perhaps surprising that we have not observed lateral synaptic inhibition in PNs.

Several considerations put this finding in perspective. First, although the lateral inputs we observe are dominated by excitation, it is possible that these responses reflect the integration of both excitatory and inhibitory inputs. As a result, inhibition could be masked by a larger postsynaptic excitation. Second, although our results are inconsistent with a dominant role for interglomerular post-synaptic inhibition of PNs, our findings do not preclude a role for interglomerular presynaptic inhibition of ORN axon terminals. Presynaptic inhibition of neurotransmitter release from ORN axons is a well-known phenomenon in the mammalian olfactory bulb (Ennis et al., 2001; McGann et al., 2005; Murphy et al., 2005; Wachowiak et al., 2005) and in the crustacean olfactory lobe (Wachowiak et al., 2002). In this study, we abolished or severely reduced direct ORN input to the PNs we were recording from, which necessarily prevents us from observing any substantial presynaptic inhibition.

It is worth noting that neither GABAA nor GABAB receptors can mediate the lateral depolarization we observe. Both GABAA and GABAB conductances are hyperpolarizing in PNs (Wilson and Laurent, 2005). And although GABAA and GABAB receptor antagonists together completely block GABA-evoked hyperpolarizations in PNs (Wilson and Laurent, 2005), they do not diminish the lateral depolarization we describe in this study (Figure S7). This result also demonstrates that the lateral depolarization does not represent disinhibition (inhibition of inhibitory input to PNs).

Implications of Lateral Excitatory Connections for Odor Processing

A significant transformation in odor responses occurs between the ORN and PN layer in the *Drosophila* olfactory system. First, the odor tuning of PNs can be broader than the odor tuning of their presynaptic ORNs (Wilson et al., 2004). This may reflect, in part, the effects of the lateral excitatory connections we have described in this study. Because we have observed that the odor tuning of lateral input to a PN is different from the odor tuning of its direct ORN input, it seems likely that these lateral inputs promote excitatory responses to odors that would not have otherwise excited that PN. A second feature of the ORN-to-PN transformation is that the rank order of PN odor preferences can differ from the odor preferences of their presynaptic ORNs (Wilson et al., 2004). Again, because the odor tuning of lateral input to a PN is different from the odor tuning of its direct ORN input, it seems likely that lateral excitatory connections between glomeruli contribute to this phenomenon.

However, it would be misleading to neatly assign different components of a PN’s odor response to direct versus lateral excitatory inputs. Direct and lateral excitation may coexist with pre- and postsynaptic inhibition, and all these inputs are likely to be integrated by PNs in a nonlinear fashion. Broad tuning in PNs could also reflect some non-linearity in ORN-to-PN connections.

In general, bridging the gap between cellular and systems neuroscience will require a deeper understanding of how neurons integrate complex synaptic inputs in vivo. Using a combination of genetic techniques and in vivo electrophysiology, we have begun to dissect the various synaptic interactions involved in odor processing in the *Drosophila* antennal lobe. Our strategy has been to eliminate one input to an identified neuron in order to
unmask other relevant interactions. Here, this approach has revealed broadly distributed but specific excitatory connections between glomeruli. Although the behavior of a neural circuit is ultimately a complex product of its components, some insight can nevertheless be gained by manipulating one element at a time, provided that appropriate genetic tools are available. In this respect, the *Drosophila* olfactory circuit represents a powerful system for understanding the synaptic and cellular computations performed on sensory stimuli that ultimately produce perception and behavior.

**EXPERIMENTAL PROCEDURES**

**Fly Stocks**

Flies were reared at room temperature on conventional cornmeal agar. All experiments were performed on adult female flies 2–7 days post-eclosion. Flies were kindly provided as follows: *Or10a-Gal4* (Barry Dickson); *Or43b* (Dean Smith); *NP3103-Gal4, NP3529-Gal4, and NP3481-Gal4* (Kei Ito and Liqun Luo); *UAS-Or83b-Gal4, Or83b-Gal4, Or83b*, *Or46a-Gal4, Or92a-Gal4, and Or42b-Gal4* (Leslie Vossshall); *UAS-DTl/CyO and UAS-DTl10* (Leslie Stevens); *UAS-CD8:GFP*, *UAS-CD8:DpFp*, and *UAS-CD8:GFP* were obtained from the Bloomington Stock Center. *Or10a*D03204 and *Or42a*D03205 are indexed on Flybase as *pBac* insertions and were obtained from Bloomington. We found that the *Or42a*D03205 stock had a second mutation that specifically affects the pb2A pb3B neurons (Figures S5 and S6). This mutation is on the third chromosome and most likely affects the odorant receptor genes Or86e and Or86d, which are expressed in the pb2A and pb3B neurons, respectively. We term this mutation *Δ58*. The *Or42a*D03205 and *Δ58* mutations were separated using standard genetic crosses. The genotypes used in all experiments are listed in Table S1.

**ORN Recordings**

Flies were immobilized in the trimmed end of a plastic pipette tip. A reference electrode filled with *Drosophila* saline was inserted into the eye, and a sharp saline-filled glass capillary (tip diameter < 1 μm) was inserted into a sensillum. Sensilla were visualized using an Olympus BX51WI microscope with a 50× water-immersion objective, IR-DIC optics, and a fluorescence attachment was used to obtain recordings under visual control. One neuron was recorded per brain, and the morphology of each cell was visualized post hoc with biocytin histochemistry. Histology with biocytin-streptavidin and nc82 antibody was performed as described previously (Wilson and Laurent, 2005), except that in the secondary incubation we used 1:250 goat anti-mouse/AlexaFluor 633 and 1:1000 streptavidin:AlexaFluor 568 (Molecular Probes). The nc82 antibody (used to outline glomerular boundaries) was obtained from the Developmental Studies Hybridoma Bank (University of Iowa). Histology with *Δ*CD8 antibody (Figures 1B, 5C, and S1A) was performed as described previously (Wilson and Laurent, 2005). Glomeruli were identified using published maps (Laisisse et al., 1999; Couto et al., 2005).

**Olfactory Stimulation**

Odors were diluted in paraffin oil at a ratio of 1:100 (except for the VM7-only experiments, see below). Odors in Figure 4 are benzaldehyde (BNZ), butyric acid (BUA), 2,3-butanediol (BUD), 1-butanol (BUT), cyclohexanone (CYH), cis-3-hexen-1-ol (CIS), ethyl butyrate (EB), ethyl acetate (ETA), geranyl acetate (GER), methyl salicylate (MSL), 3-methylthio-1-propanol (MTP), 4-methyl phenol (MPH), γ-valerolactone (VAL). Odors used for stimulation of rescued VA71 ORNs (Figures 6A and 6C, 8A–8C, and 9) were 4-methyl phenol, benzaldehyde, methyl salicylate, ethyl acetate, 1-butanol, cyclohexanone, 3-octanol, and paraffin oil. For selective stimulation of VM7 ORNs (Figures 5E, 6B, 6D, 7A, and 8D–8F), dilution ratios in paraffin oil were adjusted for most odors to achieve specific stimulation; odors were acetone (10−4 dilution), 2-butanol (10−3), ethyl acetate (10−3), geranyl acetate (10−3), hexanal (10−4), hexyl acetate (10−4), isomyrcene (10−4), methyl acetate (10−4), octanal (10−4), propanol (10−4), 2-pentanone (10−4), trans-2-hexenal (10−4), butyric acid (10−4), and paraffin oil. Odor-source details are at http://wilson.med.harvard.edu/odors.html. Odors were delivered with a custom-built olfactometer. A continuous stream of charcoal-filtered air (2.2 l/min) was directed over the fly. Switching of a three-way solenoid redirected 200 ml/min of this air through an odor vial, which rejoined the air stream 12 cm from the end of the odor tube. Thus, all odors were diluted 10-fold in air just before reaching the fly. All odor stimuli were applied for 500 ms. The odor tube was ~8 mm in diameter and terminated ~8 mm from the fly.

**Data Analysis**

Data were analyzed using custom software written in Igor Pro (WaveMetrics). In Figures 1, 6, 7, and 8, PN voltage traces were averaged over six repeated presentations of each odor and low-pass filtered offline at 13 Hz to remove spikes before averaging across experiments. Depolarization area was computed as the area under the baseline-zeroed membrane potential over a 500 ms window starting 100 ms after opening of the odor valve. Traces are presented as the mean ± SEM across experiments. Tuning curves in Figure 4 were generated by computing mean firing rate over the 500 ms odor presentation, minus the baseline firing rate.

**Supplemental Data**

The Supplemental Data for this article can be found online at http://www.neuron.org/cgi/content/full/54/1/89/DC1/.
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Excitatory Interactions between Olfactory Processing Channels in the *Drosophila* Antennal Lobe

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Table S1. Genotypes Used in Each Experiment

<table>
<thead>
<tr>
<th>Fig.</th>
<th>Description</th>
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<tbody>
<tr>
<td>1B</td>
<td><strong>UAS-CD8:GFP/+; Or83b-Gal4/+</strong></td>
</tr>
<tr>
<td>1C-G</td>
<td>w[1118]</td>
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</tbody>
</table>
| 2-4    | VM2: wild-type is NP5103-Gal4, UAS-CD8:GFP, mutant is NP5103-Gal4, UAS-CD8:GFP; Or43b[1]  
         | DL1: wild-type is NP3529-Gal4, UAS-nlsGFP, mutant is Or10a[f03694]; +/+; NP3529-Gal4, UAS-nlsGFP |
| 5C     | top: UAS-CD8:GFP/+; Or46a-Gal4/ UAS-Or83b; Or83b[2]; Or83b[2]  
         | bottom: Or46-Gal4/UAS-Or83b; Or83b[2]                                          |
| 5E     | VM7 ORNs functional: NP3481-Gal4, UAS-CD8:GFP; +/+; ∆85  
         | VM7 ORNs non-functional: Or42a[f04305]; ∆85                                    |
| 6A,C   | Or46a-Gal4/UAS-Or83b; Or83b[2]                                               |
| 6B,D   | NP3481-Gal4, UAS-CD8:GFP; +/+; ∆85                                            |
| 7      | NP3481-Gal4, UAS-CD8:GFP; +/+; ∆85                                            |
| 8A-C   | Or46a-Gal4/UAS-Or83b; Or83b[2]                                               |
| 8D-F   | NP3481-Gal4, UAS-CD8:GFP; +/+; ∆85                                            |
| 9      | Or46a-Gal4/UAS-Or83b; Or83b[2]                                               |
| S1A    | Or10a[f03694]; Or10a-Gal4; UAS-CD8:GFP                                       |
| S1C-D  | VM2: wild-type is GH146-Gal4, UAS-CD8:GFP or NP5103-Gal4, UAS-CD8:GFP, mutant is Or43b[1]  
         | DL1: wild-type is w[1118] or Or42b[EY14886]a or Or42b-Gal4/UAS-DiphTx; Or92a-Gal4/UAS-DiphTxb, mutant is Or10a[f03694] |
| S2     | same as Fig. S1                                                              |
| S3     | control: w[1118]  
         | mutant: Or83b[2]                                                              |
| S4     | control: Or33c-Gal4/+; UAS-DTl/+c or Or33c-Gal4/UAS-DTlc  
         | rescued: Or46-Gal4/UAS-Or83b; Or83b[2]                                         |
| S5     | wild-type: w[1118]  
         | mutant: Or42a[f04305]; ∆85                                                   |
| S6     | see italicized labels in figure                                              |

*a* pBac insertion in the Or42b gene (Bloomington Stock Center). We have observed that this mutation abolishes odor responses in ab1A ORNs, allowing us to count more accurately the spikes from ab1D (DL1) and ab1B (VA2) ORNs in response to odors that normally drive the ab1A ORNs strongly.
b We killed ab1A and ab1B ORNs by expressing diphtheria toxin selectively in these cells. This allowed us to more accurately detect the small spikes of ab1D (DL1) ORNs in response to odors that normally drive both either the ab1A or ab1B ORNs strongly.

c We killed pb2A ORNs by expressing diphtheria toxin selectively in these cells, allowing us to count more accurately the spikes from pb2B (VA7l) ORNs.
**Fig. S1. Mutations in odorant receptor genes virtually abolish odor responses in ORNs.**

(A) ORNs that normally express Or10a target glomerulus DL1 correctly in the Or10a<sup>559564</sup> mutant. Projection of a confocal stack through the antennal lobes (neuropil in magenta) shows ORN axons labeled with CD8:GFP (green). This result is consistent with a previous report that a functional odorant receptor is not required for correct ORN axon targeting in Drosophila (Dobritsa et al., 2003).

(B) Schematic of experiments in Fig. S1, panels C-D: extracellular recordings are performed from ORNs whose corresponding odorant receptor gene has been mutated.

(C) Rasters of odor-evoked responses from wild-type and mutant ORNs. Each row is a separate trial with the same odor. Gray bar = 500-ms period of odor delivery. Note that spontaneous activity is also reduced by these mutations.

(D) Average odor responses of wild-type and mutant ORNs. Each response is the mean ± SEM of 4-18 independent experiments in different flies. The firing rates for three odors (marked with ††) could not be determined in the mutant due to the fact that other ORNs in the same sensillum as DL1 ORNs respond very strongly to these stimuli.
Fig. S2: Weak responses to a few odors persist in mutant ORNs.
(A) Single-sensillum recordings. Top trace is a recording from an ab8 sensillum in an Or43b^1 fly. Symbols (●) indicate spikes originating from the neuron that normally expresses Or43b (the ab8A cell). Bottom trace is a recording from an ab1 sensillum in an Or10a^93864 fly. Symbols indicate spikes originating from the neuron that normally expresses Or10a (the ab1D cell). Although the odorant receptor gene corresponding to these ORNs has been mutated, they still show weak responses to a few odors.
(B) Rasters compare responses of control and mutant ORNs to the same odor. Mutant responses are very weak, but are consistent from trial to trial.
(C) Summary of odors eliciting a nonzero response in mutant ORNs. Mean response during the 500-msec odor stimulation period (±SEM), averaged over 4-12 experiments in different flies. Some of the same odors elicit nonzero responses in both types of mutant ORNs (especially 2-octanone and pentyl acetate). We have also observed very weak responses to some of these odors in Or22a mutant flies (Δhalo) while recording from ab3a ORNs. We have excluded all five odors shown in (C) from our analysis in Figs. 3-4 and Fig. S2.
Fig. S3: The Or83b\textsuperscript{2} mutation abolishes odor responses in the palps but not in the antennae.  
(A) Odor-evoked local field potentials recorded from the antenna in wild-type (w\textsuperscript{1118}) and mutant (Or83b\textsuperscript{2}) flies. Gray bar indicates 500-ms odor stimulation period.  
(B) Odor-evoked local field potentials recorded from the maxillary palp in w\textsuperscript{1118} and Or83b\textsuperscript{2} flies.  
(C) Extracellular recordings of spikes from single sensilla on the maxillary palp. All palp sensilla contain two ORNs, distinguishable by their spike shape and size. Top trace shows a recording from a palp basiconic type 3 sensillum (pb3). Large spikes come from the A neuron (pb3A, magenta) and small spikes from the B neuron (pb3B, blue). Lower trace shows a sensillum recording from an Or83b\textsuperscript{2} mutant; both spontaneous and odor-evoked spikes are absent. Odor stimulus is ethyl acetate in both traces.  
(D) Average firing rates evoked by four odors. These odors elicit ORN spikes in control sensilla, but no responses in Or83b\textsuperscript{2} mutant sensilla (n = 3 sensilla of each type for control, n = 23 sensilla total for mutant). Responses are computed over the 500-msec odor stimulus period, minus baseline firing rate. Mean ± SEM, averaged across experiments.
Fig. S4: Rescued VA71 ORNs recapitulate normal ORN odor responses.
Peristimulus-time histograms comparing odor responses in control versus "rescued" VA71 ORNs. Mean ± SEM, averaged across experiments, n ≥ 5 for each trace. Gray bar = 500-ms period of odor stimulation. Genotype of control flies is Or33c-Gal4++;UAS-DTric or Or33c-Gal4/UAS-DTric. (In order to count more accurately the spikes from VA71 ORNs in control sensilla, we expressed diphtheria toxin under the control of the Or33c-Gal4 driver in order to kill the other ORN in the pb2 sensillum.) Genotype of "rescued" flies is Or46-Gal4/UAS-Or83b;Or83b^2.
Fig. S5: Or42a^{D42D5}; Δ85 flies have a deficit in some maxillary palp ORNs but not antennal ORNs.

(A) Antennal local field potential recordings from w^{1118} (green; n = 5) and Or42a^{D42D5}; Δ85 (magenta; n=3). Mean ± SEM, averaged across experiments. Black bar = 500-ms period of odor stimulation.

(B) Quantification of field potential area shows that Or42a^{D42D5}; Δ85 flies have normal odor responses in the antenna. This rules out a general olfactory deficit in these flies.

(C) Maxillary palp local field potential recordings from w^{1118} (green) and Or42a^{D42D5}; Δ85 (magenta). Mean ± SEM, averaged across experiments. Black bar = odor.

(D) Quantification of maxillary palp field potential area shows that Or42a^{D42D5}; Δ85 flies have reduced maxillary palp ORN responses to most odors, as compared to w^{1118}.
Fig. 36: Electrophysiological characterization of the Δ85 and Or42a<sup>Δ4305</sup> mutations.

(A) Extracellular recordings from pb1 sensilla showing that the Or42a<sup>Δ4305</sup> mutation abolishes odor-evoked activity in the pb1A neurons (these ORNs target the VM7 glomerulus). Each tick represents a spike from either the “A” (blue) or “B” (green) ORN in the pb1 sensillum. Odor stimulation is indicated by the gray bar (500 msec).

(B) Recordings from pb2 sensilla showing that the Δ85 mutation severely reduces odor responses in the pb2A neuron. In the top trace, the pb2B neuron has been killed with diphtheria toxin to show only spikes from the pb2A neuron (Or46a-Gal4/UAS-DTI).

(C) Recordings from pb3 sensilla show that the Δ85 mutation eliminates odor-evoked activity in pb3B neurons.

(D) Summary of single sensillum recordings in Or42a<sup>Δ4305;Δ85</sup> and +/+;Δ85. Numbers in parentheses indicate the number of recordings.

*Odor responses in pb2A are dramatically reduced, but not eliminated. This ORN expresses two receptors, Or33c and Or85e. We hypothesize that the Δ85 mutation removes Or85e and that Or33c mediates the residual responses. This interpretation is consistent with the known odor tuning of Or33c and Or85e (Goldman et al., 2005).
Fig. S7: GABA receptor antagonists do not abolish lateral excitatory inputs to PNs.

(A) Recording from a VM2 PN postsynaptic to non-functional ORNs (genotype NP5103-Gal4, UAS-
CD8:GFP;Or43b'). The lateral depolarization is unaffected by 50 μM CGP54626, a concentration which it is effective at blocking GABAA receptors in Drosophila PNs (Wilson & Laurent 2005). Similar results were observed in other experiments (n=3). Also, CGP54246 did not affect the magnitude of the lateral depolarization in antennal PNs in experiments where only VA71 ORNs were stimulated (n=2, experimental design same as in Figs. 6A and 8A).

(B) Recording from a VM2 PN postsynaptic to non-functional ORNs (genotype NP5103-Gal4, UAS-
CD8:GFP;Or43b'). The lateral depolarization is unaffected by 10 μM picrotoxin, a concentration which it is effective at blocking GABAA receptors in Drosophila PNs (Wilson & Laurent 2005, see Supplemental Fig. 3 of that paper). Similar results were observed in other experiments (n=3). Also, picrotoxin did not diminish the lateral depolarization in antennal PNs in experiments where only VA71 ORNs were stimulated (n=4, experimental design same as in Figs. 6A and 8A). In some experiments, picrotoxin speeded the kinetics of the lateral depolarization but did not diminish it.