Title

A GAL80 collection to nullify transgenes

in *Drosophila* olfactory sensory neurons

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Summary

   Fruit flies recognize hundreds of ecologically relevant odors and respond

appropriately to them. The noise, redundancy and interconnectedness of the olfactory

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machinery complicate efforts to pinpoint the functional contributions of any component

neuron or receptor to behavior. Some contributions can only be elucidated in flies that

carry multiple mutations and transgenes, but the production of such flies is currently

labor-intensive and time-consuming. Here, we describe a set of transgenic flies that

express the *Saccharomyces cerevisiae* GAL80 in specific olfactory sensory neurons

(OrX-GAL80s). The GAL80s effectively and specifically subtract the activities of GAL4-

driven transgenes that impart anatomical, behavioral, and physiological phenotypes.

OrX-GAL80s can allow researchers to efficiently activate only one or a few types of

functional neurons in an otherwise nonfunctional olfactory background. Such

experiments will improve our understanding of the mechanistic connections between

odorant inputs and behavioral outputs at the resolution of only a few functional neurons.

Key Words

Olfaction; GAL80; Drosophila; Odorant Receptor; Sensory Neurons

Introduction

The olfactory system of *Drosophila melanogaster* is often the subject in studies of

memory, evolution, gene choice, development and odorant-induced behavior. It is a

good model system because of its relatively stereotyped neuronal circuitry, complex

behaviors and convenient genetic tools.

In *Drosophila*, most olfactory sensory neuron (OSN) typically expresses a single

odorant receptor (OR) from a genomic repertoire of 60 genes (Vosshall et al., 1999;

Robertson et al., 2003; Vosshall et al., 2000; Clyne et al., 1999; Goldman et al., 2005).

The promoter of an OR gene can thus be employed to label specific subsets of OSNs

with a particular transgene. ORs, which vary in sensitivity and specificity to a wide range

of different odorants, determine the firing kinetics and odor-response dynamics of each

OSN (de Bruyne et al., 1999; de Bruyne et al., 2001; Dobritsa et al., 2003; Elmore et al.,

2003; Hallem et al., 2004; Couto et al., 2005; Fishilevich & Vosshall, 2005; Hallem &

Carlson, 2006; Kreher et al., 2008).

Most OSNs express Odorant Receptor Co-Receptor (Orco), a highly conserved

member of the olfactory receptor family (Krieger et al., 2003; Vosshall and Hansson,

2011), in addition to a single selected OR. Though Orco usually does not contribute to

the structure of the odorant binding site (Nakagawa and Vosshall, 2009; Nichols and

Luetje, 2010; Jones et al., 2011; Jung et al., 2011), it is essential for odorant-invoked

signaling in flies. Without Orco, the co-expressed OR cannot localize to the dendritic

membrane or relay an odor-evoked signal (Larsson et al., 2004; Benton et al., 2006;).

Orco null flies are largely anosmic, though some chemosensation remains due to the

presence of ionotropic receptors and gustatory receptors, which do not require Orco to

function (Jones et al., 2007; Kwon et al., 2007; Benton et al., 2009; Silbering et al.,

2011). The Orco promoter is consequently a convenient device for the expression of

transgenes in most OSNs.

The olfactory organs, the antenna and maxillary palp, contain OSNs dendrites

within structures called sensilla. ORs and Orco are embedded in the dendritic

membrane. OSN axons project to the antennal lobes in the brain of the animal. Each

antennal lobe consists of ~50 globular synaptic sites called glomeruli. All OSNs on the

periphery that expresses the same OR converge onto their own unique glomerulus. For

example, all OSNs expressing Or22a will send axons to the DM2 glomerulus in the

antennal lobe while all OSNs expressing Or82a will send axons to the VA6 glomerulus

(**Figure 1**). The stereotyped organization of OSNs and their projections is known as the

olfactory sensory map (Stocker et al., 1990; Vosshall et al., 2000; Couto et al., 2005;

Fishilevich and Vosshall, 2005). The regularity of this map is a key feature that makes

Drosophila olfaction such a useful model, as any aberration to the typical pattern will be

apparent. The apparent simplicity of the map (**Figure 1**), however, obscures

mechanistic complexities that are yet to be discovered, in part because necessary tools

remain unavailable.

Drosophila geneticists have traditionally relied on genetic mutations or deletions

to understand how complex biological systems normally work. Most alleles are

recessive, so homozygotes must be bred over multiple generations. Achieving

homozygosity of a mutation while also adding transgenes to the system often requires

the creation of recombinant chromosomes produced after multiple generations of

crossing and PCR screening. Classical genetic strategies thus limit the number and

complexity of combinatorial genotypes that one can achieve. More challenging

experimental questions demand more facile and versatile genetic tools.

The GAL4/UAS gene regulation system has become a *defacto* standard in

studies of *Drosophila*. GAL4 is a yeast transcription activator that binds to the Upstream

Activating Sequence (UAS) and induces expression of downstream genes (Giniger et

al., 1985). By driving GAL4 expression from an OR promoter, specific expression of a

UAS-transgene can be obtained for any OSN subtype. An *OrX-GAL4* line exists for

almost every OR. This collection of GAL4 lines is a powerful toolbox since different

UAS-transgenes can be introduced into a line via conventional mating. For example,

human α-synuclein has been expressed in OSNs to model human Parkinson's disease

(Chen et al., 2014). Alternatively, protein expression levels can be knocked down using

any specified *UAS-RNAi* transgene.

A variety of existing compatible effectors can be used study different aspects of

neuronal communication. *UAS-Kir2.1* effector is used as an example in experiments

described below. This inward rectifier potassium channel electrically inactivates the

neurons that express it (Johns et al., 1999; Baines et al., 2001; Hodge, 2009). Similarly,

shibire[ts] or tetanus toxin can be used to silence synaptic communications (van der Bliek

and Meyerowitz, 1991; Chen et al., 1991; Sweeney et al., 1995; Baines et al., 1999;

Kitamoto, 2001; Kitamoto, 2002), *reaper/grim/hid* genes can be used to physically kill

neurons using their own apoptotic pathways (Abrams, 1999; Song and Steller, 1999), or

ricin toxin can be expressed ectopically to kill neurons. Conversely, neurons can be

selectively activated with *trp1a* or a variety of other channelrhodopsin transgenes

(Pulver et al., 2009; Boyden, 2011).

If GAL4 is a standard on-switch for nearly any desired transgene, GAL80 is the

logical off-switch. GAL80 binds GAL4 transcriptional activation domain, thereby

preventing recruitment of RNA polymerase (Ma and Ptashne, 1987). GAL80 crosses

are much more convenient than classical breeding approaches (**Figure 2**). In order to

have a single functional OSN in an otherwise silent olfactory system, the traditional

method uses an Orco null mutation (Larsson et al., 2004). In this genetic setup, Orco

mutant flies are anosmic, but function is restored to one OSN subset with *Or-GAL4,*

UAS-Orco transgenes (Fishilevich et al., 2005; Benton et al., 2006; Olsen et al., 2007;

DasGupta and Waddell, 2008; Hoare et al., 2008; 2011) (**Figure 2a**). An *Orco-GAL4,*

UAS-effector, Or-GAL80 method can be used instead (**Figure 2b**). Kir2.1 is used as an

example of an effector (Johns et al., 1999; Baines et al., 2001; Hodge, 2009;). Classical

breeding strategies (**Figure 2a**) may look less complicated on paper than GAL80

crosses (**Figure 2b**) but are actually more time-consuming and limited. The Orco

mutation must be homozygous. Since most *Drosophila* transgenes are embedded into

the same two chromosomes (2 or 3) recombination and PCR screening may be

required to achieve this homozygosity.

Neurons seldom operate autonomously, but rather groups of neurons coordinate

within a circuit to provide an organism with perception and behavior. An investigation of

the behavioral impact provided by a limited number of different functional neuronal

types would require additional genes. The elaboration of genotypes (**Figure 2**) to

restore pairs or groups of functional OSNs in a nonfunctional background normally

requires generations of crosses (followed by PCR screens for desired recombinants). A

GAL80 strategy can shorten this process by achieving similar results in only one or two

generations with no necessary recombinant creation. Furthermore, a GAL80 strategy

takes advantage of the interchangeable variety of existing *UAS-transgene* lines. Here

we describe a new collection of *OrX-GAL80* lines designed to complement existing *OrX-*

GAL4 lines, and demonstrate their potential utility for behavioral and neuroanatomical

studies of the *Drosophila* olfaction model.

Results

Design of GAL80 Constructs

The following criteria were used to choose OR promoters for the collection. i) The

ORs should be relevant to current research as shown by the number of studies that

used it. ii) The ORs should represent a variety of expression patterns (larval or adult,

antennae or maxillary palps, sensillary class etc.). iii) Finally, the ORs should reflect a

variety of different odorant response profiles. The promoter regions were defined based

largely on the work of Couto *et al*, 2005.

Equimolar expression of GAL4 and GAL80 is not always sufficient to effectively

eliminate GAL4 activity so the pBP-GAL80uW-6 vector was used. This vector contains a

modified GAL80 sequence, designed to increase the stability and expression of its gene

product (Pfeiffer et al. 2010). A few *OrX-GAL80*s were already made with this vector

and used effectively. Gao et al. (2015) pioneered the technique by creating a few OrX-

GAL80s. This work is a logical extension of their contributions and will make many

GAL80s freely available for the continued development of the field.

Testing GAL80 Efficacy and Specificity

GAL80 lines were created for the following odorant receptor promoters: Or7a,

Or9a, Or10a, Or13a, Or19a, Or22a, Or22b, Or33c, Or35a, Or42a, Or42b, Or43b,

Or47a, Or56a, Or59b, Or59c, Or67a, Or67d, Or71a, Or82a, Orco, Or85a, Or85b,

Or85c, and Gr21a. To examine GAL4 subtraction *in vivo*, *OrX-GAL80* flies were crossed

to flies with the genotype *OrX-GAL4, UAS-GFP*. OSNs expressing the same OR can be

identified from their specific glomerulus in the antennal lobe (**Figure 1**). *OrX-GAL4,*

UAS-GFP flies show robust expression of the GFP reporter gene in their respective

glomeruli. However, when *OrX-GAL80* is added to the genotype, GFP expression is

entirely absent, indicating a robust antagonism of GAL4 activity (**Figure 3**). Several of

these lines also have expression in larvae. GAL4 subtraction was examined in larval

brains using the *UAS-GFP* reporter gene. In larvae, GAL80 reduced but did not

eliminate GAL4 activity (**Figure S1a**).

The *OrX-GAL80* lines were checked to ensure they would not have aberrant

expression in untargeted OSN subtypes. The pBP-GAL80uW-6 vector contains a

Drosophila Synthetic Core Promoter (DSCP). DSCP is an effective means of using

enhancer elements to drive strong expression (Pfeiffer et al., 2008), but it could also

cause the GAL80s to have nonspecific or leaky expression. Therefore, a version of

pBP-GAL80uW-6 was cloned with the DSCP removed. However, when the DSCP was

absent, GAL80 expression was insufficient to subtract GAL4 activity (**Figure S1b**). A

few lines were tested to see if DSCP causes nonspecific GAL80 expression. For these

lines, an *OrY-GAL80* did not impede GAL4 activity of an *OrX-GAL4* neuron (**Figure**

S1c). Due to the uneven expression in an *Orco-GAL4, UAS-GFP* line, it could not be

determined if each OrX-GAL80 subtracts GAL4 from only one glomerulus in an

otherwise fully-labeled brain, but results shown in Figure S1c give reasonable

confidence that the GAL80s do not have widespread nonspecific expression. It can also

be noted that the GAL80 subtraction does not interfere with reporter gene expression in

a genetic system that does not use GAL4. When *Or22a-GAL80* is used in conjunction

with *Or22a-GFP*, containing no GAL4/UAS intermediary, the GFP is still expressed

(**Figure S1d**). These images, showing subtraction of reporter gene expression, confirm

that GAL4 activity is suppressed anatomically by the GAL80 lines.

To confirm GAL4 was suppressed physiologically by the GAL80s, Single

Sensillum Recordings (SSRs) were used to assay electrical activity of OSNs. *Gr21a-*

GFP was used to identify sensilla of interest without interfering with the

GAL4/UAS/GAL80 system. Gr21a neurons are housed in ab1 sensilla. Carbon Dioxide

exposure causes a robust response in Gr21a ab1C neurons (Jones et al., 2007; Kwon

et al., 2007). When *Gr21a-GFP* flies were exposed to CO_2, their ab1C sensillar neurons

showed robust responses (mean Δspikes/s=88, N=8 sensilla). Adding Kir2.1 to Gr21a

neurons (genotype *Gr21a-GFP, Gr21a-GAL4, UAS-Kir2.1)* greatly reduced spiking

responses to CO_2 (mean Δspikes/s=14, N=12 sensilla). When Gr21a-GAL80 was added

(genotype *Gr21a-GFP, Gr21a-GAL4, UAS-Kir2.1, Gr21a-GAL80),* responses to CO_2

were restored (mean Δspikes/s=94, N=6 sensilla) (**Figure 4**). These results confirm that

GAL80 is able to prevent GAL4-induced activity in a physiological experimental

paradigm.

Behavior

To demonstrate how experiments can be simplified using GAL80 reagents, the

GAL80s were used alongside a genetic strategy to create flies with a single functional

OSN (**Figure 2**). The transgenic genotypes are easier to construct than those created

by classical genetics, but the two systems are expected to be quite similar in behavior.

Insect olfactory neurons do not require odor-evoked activity to develop with complete

fidelity (Elmore and Smith, 2001; Wong et al., 2002; Dobritsa et al., 2003; Larsson et al.,

2004; Berdnik et al., 2006;), and therefore both Orco null and *Orco-GAL4, UAS-Kir2.1*

brains will have normal circuitry, despite their nonfunctional neurons. Orco null mutants

also have greatly diminished spontaneous activity, and Kir2.1-containing neurons are

expected to show little to no spontaneous firing (Hoare et al., 2008; Olsen et al., 2007).

A simple odor preference test was used to assay both types of adult flies. Flies

were allowed to walk freely in a circular arena with four air quadrants separated by air

flow but not physical barriers. After one minute, two quadrants along the diagonal

received an odorant (Aso and Rubin, 2016). The diagonal quadrants receiving odorant

were randomized. Flies were recorded and their motions tracked using Ctrax software

(Branson et al., 2009) (**Figure 5a**). Response index was measured in each frame.

Response Index= $Flies_{odor}$- $Flies_{air}$/Total Flies, so 0 indicates no response, +1 would be

total attraction, and -1 would be total aversion. Each N represented an experiment

involving 20-30 flies. Therefore, if an N had an RI of -0.5, about 23 of 30 flies were in

the air quadrant.

Or22a and Or82a were tested since they represent opposite ends of the odorant

tuning spectrum. Or22a is broadly tuned and its OSN responds to multiple odorants;

Isoamyl acetate (IAA) is one odorant that evokes a strong response. Or82a is narrowly

tuned, responding to only one known odorant, geranyl acetate (GAc) (de Bruyne et al.,

2001; Hallem et al., 2004; Fishilevich and Vosshall, 2005; Hallem and Carlson, 2006).

Or22a genotypes (**Figure 5b**) received 1:1000 IAA in mineral oil and Or82a genotypes

(**Figure 5c**) received 1:100 GAc in mineral oil. The mean response index for the middle

part of the odorant exposure (time = 75-105 seconds) was calculated. The positive

control was *Orco-GAL4* with no effector added; this line and the OrX-GAL80 lines were

made in a w^{1118} background. These flies find both odorants aversive, as shown by a

negative and significantly lower response index to all other genotypes ($p \leq .0001$ for IAA

and $p \leq .006$ for GAc). *Orco* null and *Orco-GAL4, UAS-Kir2.1* flies served as negative

controls and had response indices around 0, showing they are anosmic as expected.

Or22a-GAL4, UAS-Kir2.1 flies did not find IAA aversive, and *Or82a-GAL4, UAS-Kir2.1*

flies did not find GAc aversive (**Figure 5b and c**), suggesting that Or22a and Or82a are

each necessary for behavioral responses to their respective odorants.

As noted previously, two genotypes for each OR had only a single functional

OSN subset in an otherwise nonfunctioning olfactory system (**Figure 2**). One group

represents the classic strategy (*OrX-GAL4, UAS-Orco* in an Orco null background) and

the other represents the GAL80 strategy (*Orco-GAL4, UAS-Kir2.1, OrX-GAL80*). When

O22a or Or82a OSN function is restored, neither of the single-functional OSN

genotypes is sufficient alone to restore normal aversive behavior. However, for both

Or22a genotypes with IAA and Or82a genotypes with GAc, the behavioral variance for

the classical method vs the GAL80 method is comparable (F-test for the 75-105sec

window is $p=.44$ for the Or22a/IAA group and $p=.45$ for the Or82a/GAc group). Since

both groups have similar variance, this result supports the idea that the GAL80 strategy

can be adopted as a substitute for the more traditional and time-consuming breeding

strategy.

Discussion

The lack of aversive behavior in flies with a single functional OSN subtype is not

consistent with some previous studies that showed one functional OSN was sufficient to

restore aversive behavior. Fishilevish et al (2005) used larvae in their study to restore

aversion with a single functional OSN subtype, but the larval olfactory system may be

fundamentally different in this respect. Gao et al (2015) gave convincing evidence of

aversive restoration in adults, though they didn't use Or82a and they tested Or22a with

a different odorant (E2-hexenal) of a higher concentration. It could be that the principle

of single-OSN aversive behavior is highly dependent on the odorant and receptor used.

(Bhandawat et al., 2010) also showed that single glomerular activity is sufficient to

invoke a behavioral response, but that study was done using an intact and fully

functional olfactory background, so some neuronal cooperation may still have occurred.

DasGupta and Waddell (2008) provided evidence that a single functional OSN subtype

is sufficient to learn odor discrimination, but the experiments here do not extend to

assess learning or memory.

Despite some specific circumstances, the lack of aversive rescue is generally

consistent with the current models of odor coding by the olfactory system. I.e. a

coordinated effort of many OSNs is usually required to produce a behavioral output.

Paired neurons in sensilla can affect the firing dynamics of their neighbors in the

periphery (Dobritsa et al., 2003; Kazama and Wilson, 2009; Su et al., 2012), and

downstream neurons such as interneurons and projection neurons may rely on

synchronized input from multiple OSN types (Ng et al., 2002; Olsen et al., 2007; Chou

et al., 2010; Yaksi and Wilson, 2010; Acebes et al., 2011; Kazama et al., 2011; Wilson,

2011; Hong and Wilson, 2015). The behavioral conditions here give no effect and are

included as a demonstration of GAL80 possibilities. The hope is that additional

researchers will use the reagents and validate them in their own assays and use them

to study effects of interest.

The collection of GAL80 lines subtracts GAL4 activity efficiently and specifically

in OSNs. In anatomical studies, reporter gene expression from the GAL4/UAS system is

suppressed. Neurons silenced with Kir2.1 expression have normal firing capacity

restored when GAL4 is antagonized using the GAL80 lines. In behavioral assays, the

GAL80s have comparable variations to traditional methods that use mutant

backgrounds. However, using a GAL80 transgene will be more flexible than mutant

lines and less cumbersome than crafting the required recombinants as the complexity of

the genotype increases.

Though in some special circumstances, olfactory sensory neurons can produce

behaviors autonomously, this is not a widely applicable principle. Researchers

encounter a significant technical obstacle to the understanding of olfactory function if

they need to create genotypes with small groups of interacting neurons in isolation. The

tools presented here facilitate the activation or deactivation of combinations of particular

neurons, thereby overcoming this obstacle. The lines are available to order through

Bloomington Stock Center.

Author Contributions

Conceptualization, JE and IM; Methodology, JE; Formal Analysis, JE and AA;

Investigation, JE and AA; Writing—Original Draft, JE; Writing—Review and Editing, AA,

CP, IM; Visualization, JE and AA; Supervision, IM; Funding Acquisition, IM

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Figure Legends

Figure 1: Olfactory Sensory Map. Each neuron in the olfactory system expresses one

type of odorant receptor (OR). Or22a (teal) and Or82a (gold) are used here as

examples. Neurons usually exist in pairs or groups in sensilla within the olfactory

organs—antenna or maxillary palp. Neurons expressing the same OR are distributed

throughout the periphery, but project their axons onto the same glomerulus in the

antennal lobe of the brain. For example, all Or22a-expressing neurons synapse onto the

DM2 glomerulus while all Or82a-expressing neurons synapse onto the VA6 glomerulus.

Figure 2: Advantages of using a GAL80 approach over a null mutation. a) **Current**

method with available reagents. In order to examine a single type of Olfactory

Sensory Neuron (OSN) without interference from other OSNs, one can use an Orco null

mutant. Without Orco, ORs cannot reach the cell membrane or function properly. Orco

mutants are mostly anosmic (unable to smell.) A single OR can then be restored using

two transgenes, *OrX-GAL4* and *UAS-Orco*. *Or22a-GAL4* is shown here as an example.

This fly may require the making and validating of one or more recombinant

chromosomes, since the Orco mutation must be homozygous. In more complicated

systems, e.g. restoring more than one OSN, multiple recombinants would need to be

made and validated at a cost of several months of crossing. b) **Using a GAL80**. GAL80

is a potent GAL4 inhibitor. All olfactory neurons could be silenced using any number of

transgenes in an *Orco-GAL4, UAS-effector* (such as *UAS-Kir2.1*) genotype. A single

OSN subtype can then be restored using an *OrX-GAL80* (such as *Or22a-GAL80*). This

system requires no recombinant creation, and is amenable to the use of various

effectors or additional transgenes without requiring recombinant construction. (Receptor

appearance, orientation, and heterodimerization is based on previous designs by

Neuhaus et al. (2005), Benton et al. (2006), Smart et al. (2008), and Benton (2009).

Figure 3: OR-GAL80 reagents eliminate GAL4 activity. All antennal lobes are stained

with anti-nc82 (a general neuropil marker, grey) and anti-GFP (green). The orientation

of each image is dorsal-up, ventral-down, lateral-right, medial-left. Scale bars indicate

20µm. Each of the brains shown has the genotype *OrX-GAL4, UAS-GFP*. The specific

receptor promoter is given above each column. The top row in each set shows GFP

expression in these lines without GAL80. Notice how each neuron's target in the

antennal lobe glomeruli is expressing GFP. Each bottom row shows the brains

containing an additional *Or-GAL80* gene. Note how GAL80 effectively inhibits GAL4

activity, as seen by the elimination of GFP expression. The images are representative of

the 5-20 brains examined per genotype. GAL4 inactivation was 100% penetrant in one

day old female flies. Not shown: Or7a, Or9a, Or19a, Or22b, Or33c, Or56a, Or85b,

Or85c, and Orco.

Figure 4: Olfactory neuron responses towards CO_2 in Single Sensillum

Recordings (SSR). a) **Box plot of SSR responses.** Ab1C neurons in *Gr21a-GFP* flies

respond strongly to CO_2, adding *Kir2.1* reduces response to CO_2, and response is

restored when *Gr21a-GAL80* is added. Each circle shows response in an individual

sensillum, and filled squares indicate the means. b-d) **Examples of SSR traces** for: b)

Gr21a-GFP, c) *Gr21a-GFP, Gr21a-GAL4, UAS-Kir2.1*, and d) *Gr21a-GFP, Gr21a-*

GAL4, UAS-Kir2.1, Gr21a-GAL80.

Figure 5: Use of OR-GAL80 reagents in a behavioral experiment a) **Olfactory**

Arena Setup (Aso and Rubin, 2016)**.** Flies are freely walking in a circular enclosure.

The vertical and horizontal lines are added to indicate quadrant barriers, but no such

physical barriers exist in the arena. All four quadrants receive air for one minute, and

then two quadrants along a diagonal receive odorant for one minute. Diagonal quadrant

pairing is randomized. In the example shown, flies find the odorant aversive and

segregate into air quadrants. The colored outline around each fly was added during

tracking using Ctrax software (Branson et al., 2009). The response index in each frame

is measured. Response Index= $Flies_{odor}$- $Flies_{air}$/Total Flies. Each N represents a trial of

20-30 flies. E.g. A response index of -0.5 indicates ~23 of 30 flies are in the air

quadrants and the odorant is aversive. b) **Comparison of reagents in olfactory**

behavior using Isoamyl acetate (IAA). Flies are added to the arena as described in A.

Odorant is 1:1000 Isoamyl acetate (IAA) in mineral oil. Neurons expressing Odorant

Receptor 22a (Or22a) respond to IAA. The top panel shows the mean response index

of each genotype over the entire course of the trial. Dotted lines indicate the time

interval of the analysis shown in the bottom panel, time= 75-105 seconds. The bottom

panel shows the mean response index of flies over the middle part of the odorant

exposure. c) **Comparison of reagents in olfactory behavior using Geranyl acetate**

(GAc). Data are represented as described in B. Odorant is 1:100 Geranyl Acetate

(GAc). Neurons expressing Odorant Receptor 82a (Or82a) respond to the odorant GAc.

For both B and C: Positive control flies (red) respond to odorant with significantly more

aversion than every other genotype ($p \leq .0001$ for IAA and $p \leq .006$ for GAc). Negative

control flies (black and grey) cannot detect the odorant and have RIs around 0. The

Or22a and Or82a receptors are necessary for normal aversive behaviors to their

respective odorants (gold). Blue and Teal represent the genotypes as described in

Figure 2. In each, when only a single OSN subtype is functional, it is insufficient to

restore normal aversive behavior. But the two groups, i.e. using the GAL80 reagent

(blue) or using the traditional Orco mutant method (teal), have similar variance ranges

(F-test insignificant).

Materials and Methods

Fly Stocks

Flies were reared on standard cornmeal/molasses food and kept at 25C with a

16 hours on/8hours off light cycle. All lines were obtained from the Indiana University

Bloomington Stock Center and the Janelia Farm Research Campus. Any recombinants

made were validated with PCR.

Stock List:

592	Or7a-GAL4 #23907
593	Or7a-GAL4 #23908
594	Or9a-GAL4 #23918
595	Or10a-GAL4 #9944
596	Or13a-GAL4 #9946
597	Or13a-GAL4 #23886
598	Or19a-Gal4 #24617
599	Or22a-GAL4 #9951
600	Or22a-GAL4 #9952
601	Or22b-GAL4 #23891
602	Or33c-GAL4 #23893
603	Or35a-GAL4 #9967
604	Or42a-GAL4 #9970
605	Or42b-GAL4 #9971
606	Or43b-Gal4 #23894
607	Or46a-GAL4 #23291
608	Or47a-GAL4 #9981
609	Or56a-GAL4 #9988
610	Or59b-GAL4 #23897
611	Or59c-GAL4 #23899
612	Or67a-GAL4 #23904
613	Or67d-GAL4 #9998
614	Or71a-GAL4 #23121

Or82a-GAL4 #23125

Orco-GAL4 #23292

Orco-GAL4 #26818

Or85a-GAL4 #23133

Or85b-GAL4 #23911

Or85c-GAL4 #23913

Gr21a-GAL4 #24147

Or22a-mcd8::GFP #52620

Gr21a-mcd8::GFP #52619

Orco^{2} #23130

UAS-Orco #23145

UAS-mcd8::GFP #5130

UAS-mcd8::GFP #5137

UAS-Kir2.1 Janelia stock #3015545

UAS-Kir2.1 Janelia stock #3015298

UAS-Kir2.1::eGFP Janelia stock #BS00312

GAL80 Creation

Primers were designed to capture the entire promoters described by Couto et al.

(2005) (see **Table S1**). Promoters were amplified from genomic DNA using Q5 High

Fidelity PCR (NEB #M0491S) and added to entry vectors using the pENTR/D-TOPO

system (Invitrogen, 2012a). Recombination with the pBP-GAL80Uw-6 (Addgene

#26236) destination vector was done using the LR Clonase II system (Invitrogen,

2012b). To ensure no mutations, no gaps, and correct orientation, the complete

promoters were sequenced in the destination vector using the sequencing primers

shown in **Table S2**. PhiC31 site-directed transgenesis was performed by Genetivision

Inc. All GAL80 transgenes were inserted at the attP2 site.

Immunohistochemistry

Female adult brains were dissected one day after eclosion in cold S2 Schneider's

Insect Medium (Sigma Aldrich #S0146) and fixed while nutating for 55 minutes at room

temperature in 2mL 2%PFA (Electron Microscopy Sciences #15713) in protein loBind

Tubes (Eppendorf #022431102). Brains were washed 4x, 15min per wash while

nutating with 2mL PBT buffer (1xPBS, Cellgro #21-040, with 0.5% TritonX-100, Sigma

Aldrich #X100). Brains were then blocked with 200µL 5% Goat serum (ThermoFischer.

#16210064) in PBT for 90 minutes while nutating, upright. Block was removed and 200

µL primary antibodies in PBT were added for 4 hours at room temperature and then

transferred to 4C for 36-48 hours while nutating, upright. Primary antibodies: mouse α-

bruchpilot (Developmental Studies Hybridoma Bank. #nc82-s) at 1:30, rabbit α-GFP at

1:1000 (Thermo Fischer #A11122), or rabbit α-Tom at 1:500 (clontech #632393).

Monoclonal antibody nc82 identifies Bruchpilot. Bruchpilot can serve as a general

neuropil marker because it is required in synaptic zones (Wagh et al., 2006). Larval

brains were collected from third instar larvae and fixed in 4% PFA. Primary antibodies:

mouse α-neuroglian (Developmental Studies Hybridoma Bank. #BP104) at 1:50 and

rabbit α-GFP at 1:500. Brains were washed 4x, 15min per wash while nutating with 2mL

PBT. 200µL secondary antibodies in PBT were then added for 4 hours at room

temperature and then 3 overnights at 4C while nutating upright. Secondary antibodies:

AF568 goat α-mouse (Life Technologies #A11031) at 1:400 and AF488 goat α-rabbit

(ThermoFischer #A11034) at 1:800. Tubes were protected from light at all times after

secondary antibodies had been added. Brains were washed again 4x, 15min per wash

while nutating with 2mL PBT. Then washed with 1xPBS and mounted using Vectashield

mounting media (Vector Labs #H-1000). Confocal images were taken with Leica800

microscope.

Olfactory Arena

Female flies aged 3-6 days were cold-plate sorted 1-2 days before the assay.

The arena was setup and the assay performed as described in (Aso and Rubin, 2016)

sans the optogenetic components. Odorants were diluted in mineral oil (Sigma Aldrich,

Geranyl Acetate #173495, Isoamyl Acetate #306967). After trying starved vs unstarved

flies and multiple odorant concentrations (data not shown), it was determined unstarved

flies at concentrations of 10^{-3} Isoamyl acetate and 10^{-2} Geranyl acetate gave the most

consistent and robust behaviors. Flies in video recordings were tracked using Ctrax

software (Branson et al., 2009).

Single Sensillum Recordings

SSRs were performed as described in Lin et al 2015 (Lin and Potter, 2015). GFP

labeled ab1 sensilla were identified using a Zeiss AxioExaminer D1 compound

microscope with eGFP filter cube (FL Filter Set 38 HE GFP shift free). A glass recording

electrode filled with ringers solution (7.5g of NaCl+0.35g of KCl+0.279g of $CaCl_2$-$2H_2O$

in 1L of H_2O) was inserted into the base of an ab1 sensillum. CO_2 was delivered

through a tube ending with a Pasteur pipette that was inserted for 1 second into a hole

in a plastic pipette directed at the antenna. This plastic pipette (Denville Scientific Inc,

10ml pipette) carried a purified continuous air stream (8.3 ml/s) that used a stimulus

controller (Syntech) at the time of CO_2 delivery to correct for the increased air flow.

Signals were acquired and analyzed using AUTOSPIKE software (USB-IDAC System;

Syntech). Spikes were counted in a 500 ms window from 500 ms after CO_2 delivery and

multiplied by 2 to calculate spikes/second. Then, the spikes in 1000ms before CO_2

delivery were subtracted to calculate the increase in spike rate in response to CO_2

(Δspikes/second). For each genotype, 6 flies (4-8 days old) were tested, with 1-3

sensilla tested in each fly.

708   Figure 1

721   Figure 2

729   Figure 3

744   Figure 4

Response Index= Flies_odor - Flies_air/Total Flies

761   Figure 5