Natural variation at the *Drosophila melanogaster* Or22 odorant receptor locus is associated with changes in olfactory behaviour

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In insects, many critical olfactory behaviours are mediated by the large odorant receptor (Or) gene family, which determines the response properties of different classes of olfactory receptor neurons (ORNs). While ORN responses are generally conserved within and between *Drosophila* species, variant alleles of the *D. melanogaster* Or22 locus have previously been shown to alter the response profile of an ORN class called ab3A. These alleles show potential clinal variation, suggesting that selection is acting at this locus. Here, we investigated if the changes seen in ab3A responses lead to changes in olfactory-related behaviours. We show that variation at the Or22 locus and in the ab3A neurons are not fully compensated for by other ORNs and lead to overall changes in antennal odorant detection. We further show that this correlates with differences in odourant preference behaviour and with differences in oviposition site preference, with flies that have the chimaeric short allele strongly preferring to oviposit on banana. These findings indicate that variation at the Or22 locus leads to changes in olfactory-driven behaviours, and add support to the idea that the ab3A neurons are of especial importance to the ecology of *Drosophila* flies.

1. Introduction

Animals rely on their sense of smell to discriminate between odours in order to identify and locate mates, dangers and food sources. Flying insects need to be able to do all of this rapidly, and additionally use their chemosensory systems to identify the optimal places to lay their eggs. In insects, odour identity is combinatorially encoded through the inputs from a large number of classes of olfactory receptor neurons (ORNs) [1–3], whose responses are determined by the olfactory receptors they express [4–6]. One of the major families of olfactory receptors in insects is the rapidly evolving Or gene family, which encode highly divergent ligand-gated ion channels and primarily detect volatile odorants [7]. The odour responses of this family and their mapping to individual ORN classes have been best characterized in *Drosophila melanogaster*, where there are 62 Ors. In *D. melanogaster*, individual Ors/ORNs can be either broadly tuned (responding strongly to a wide range of odorants) or narrowly tuned (responding strongly to one or a few odorants), albeit with there being a continuum in the breadth of tuning rather than a strict dichotomy [8,9].

We previously reported one of the first described cases of within-species variation in responses of a *D. melanogaster* ORN, the ab3A neuron [10]. Among many ORN classes, this ORN is also one of only a few known to
show interspecies variation between closely related *Drosophila* species [11,12]. The olfactory receptor locus that determines ab3A response is the Or22 locus, and we showed that naturally occurring variation at this locus within *D. melanogaster* causes alterations to ab3A neuronal responses. Specifically, the Or22 locus variant found in the common laboratory strain Canton S has two genes encoding Ors at this locus, Or22a and Or22b (the long allele [10,13]), however the Or22b protein is not functional. In these flies, Or22a produces an ab3A response phenotype we call ab3A-1.

A major variant at this locus that has been found in wild populations [14–16] is a short allele in which a chimeraic receptor called Or22ab produces a different odour response profile that we call ab3A-2. We also identified a third phenotype with yet another different odour response profile (ab3A-3), in which there is still a long allele (with two gene copies) but Or22a is not expressed and instead a functional version of Or22b determines ab3A response [10]. In general, ab3A-1 neurons respond most strongly to the larger esters tested (e.g. ethyl hexanoate, methyl hexanoate and ethyl octanoate), ab3A-2 neurons to small acetate esters (e.g. propyl acetate, butyl acetate and pentyl acetate) and ab3A-3 neurons to smaller ethyl esters (e.g. ethyl propionate, ethyl butanoate and ethyl 2-methylbutanoate).

Frequencies of the long and short alleles at this locus have been proposed to be clinally varying in Australia, with the long allele being fully penetrant in the south and the short allele appearing at high frequency in the north [14]. Clinal variation can indicate that selection is acting at a locus [17,18]. For selection to be acting at an odorant receptor locus we might expect that the variation in neuron responses caused by the genetic variants has behavioural consequences. In support of this possibility, the ab3A neuron has already been implicated as an important modifier for host-preference behaviour in *D. melanogaster* and other *Drosophila* species [16,19–21]. Moreover, the Or22ab allele has been associated with altered host-seeking behaviour in a wild *D. melanogaster* population [16]. However, it is also possible that the observed clinal allele pattern in Australia is due instead to the Or22 locus being in linkage disequilibrium with another locus of ecological importance.

Here, we therefore investigated if the different Or22 variants cause changes in behaviours that may be important for individual fitness. We first show that broader antennal neuronal responses to odours are impacted by ab3A response variation. We further provide evidence of an association between ab3A phenotype and behavioural responses in two different olfactory behaviour paradigms. Taken together, our data strongly suggest that variation at the Or22 locus leads to changes in overall odorant detection by the antenna that are associated with changes in biologically significant behaviours.

### 2. Results

#### 2.1. Variation in ab3A phenotype alters overall antennal detection of ab3A ligands

The ab3A neuron is broadly tuned, with significant responses (greater than 50 spikes s$^{-1}$) described to at least 50 odorants [9]. The sensitivity of this neuron to different odorants varies among different strains [10,16], and the neuron is, like most ORNs, more specific in the odorants it responds to at lower odorant concentrations [22]. Most, if not all, of the odorants detected by ab3A neurons are also detected by other ORNs, for example, pentyl acetate is also detected by ab1A, ab5B, ab6A and ab7A neurons, and ethyl butanoate is also detected by ab1A, ab8A, ab8B, ab10A, ab3B and ab2B neurons [1,9]. We therefore wondered if we would be able to detect differences in response between strain with the different ab3A phenotypes at the level of large neuronal populations, or if perhaps some compensatory changes might occur. We measured responses from populations of neurons across the antenna using electroantennogram recordings (EAGs). Rather than measuring the response of a single neuron, an EAG signal represents the summed responses of all ORNs in the vicinity of the recording electrode [23] (see [1,24] for diagram of ORN distribution on the antenna).

We tested the EAG responses of fly lines isogenic (fully homozygous) for the second chromosome that individually exhibit the ab3A-1, ab3A-2, and ab3A-3 phenotypes to serial dilutions of odorants that we previously showed were able to distinguish between the three phenotypes at the single neuron level [10]. Odorant concentrations were selected based on our previous data on the ab3A responses to serial dilutions of these odorants, and we selected the lowest dose that gave the maximum response from the relevant phenotype and one dose either side of this. For ethyl hexanoate and ethyl butanoate, this gave a range from 10$^{-4}$ to 10$^{-2}$, while for isopentyl acetate this gave a range from 10$^{-5}$ to 10$^{-3}$. Paraffin oil was also tested as the solvent control.

We found significant differences in EAG responses between flies with the different ab3A phenotypes (figure 1; electronic supplementary material, table S1). Flies with the ab3A-1 phenotype show significantly higher EAG responses to the ab3A-1 ligand ethyl hexanoate at 10$^{-4}$ and 10$^{-3}$ than do ab3A-2 flies, but not ab3A-3 flies. ab3A-2 flies show significantly higher responses to the ab3A-2 ligand isopentyl acetate than ab3A-1 flies at all tested doses, and than ab3A-3 flies at the 10$^{-3}$ dose. ab3A-3 flies show higher responses to its ligand, ethyl butanoate, than both ab3A-1 and ab3A-2 flies at 10$^{-4}$, and ab3A-2 flies at 10$^{-3}$. We note that all three phenotypes have similar EAG responses at higher concentrations of each odorant, which is probably because individual ORN firing rates are known to reach a maximum and plateau [1,25]. These data thus show that the changes in ab3A response between the three different phenotypes result in changes in broader antennal neuronal responses.

#### 2.2. Flies with the ab3A-1 and ab3A-2 phenotypes have differing behavioural preferences for ethyl hexanoate

We then proceeded to investigate whether flies with the different ab3A phenotypes exhibit differences in olfactory behaviour responses. We were particularly interested in determining if we could identify behavioural differences between flies with the long and short Or22 allele. As mentioned, these may show latitudinal clinal distribution in Australia, with high frequencies of the short allele in the north and the long allele fully penetrant in the south. While both the ab3A-1 and ab3A-3 phenotypes are associated with the long allele, in our earlier study of this locus all isogenic lines that we derived from southern populations had
the ab3A-1 phenotype. We therefore tested for behavioural differences between the ab3A-1 and ab3A-2 phenotypes. We used ethyl hexanoate as our test odorant because this odorant is not detected by other neurons at lower concentrations [9,11,24], and ab3A-1 and ab3A-2 flies detect this odorant differently at the level of both the individual neuron [10] and the antenna (figure 1).

To measure olfactory preference behaviour, we used a two-choice cage assay adapted from Faucher et al. [26]. Flies were given a choice between two bottles with an attractive odorant added, one of which is dosed with the odorant of interest. A preference index (PI) was calculated from the number of flies in each bottle at the end of the assay. A negative PI indicates flies preferred the control bottle (C), while a positive PI indicates flies preferred the test bottle (T) with the odorant of interest added. We first tested the ethyl hexanoate preferences of flies from the ab3A-1 and ab3A-2 isogenic lines. We assessed the behavioural choices of males and females separately in order to determine if there was an effect of sex on behaviour. For both females (figure 2a) and males (figure 2b) we found that ab3A-2 flies were significantly more attracted to ethyl hexanoate than ab3A-1 flies (Mann–Whitney U of ab3A-1 versus ab3A-2: females \(p = 0.038\); males \(p = 0.004\)). For both isogenic lines we observed no significant difference in preference between males and females (ab3A-1 males versus females: \(p = 0.206\); ab3A-2 males versus females: \(p = 0.278\)). Given this, we used females only for all further experiments.

We note that the isogenic lines contain other differences in their genetic background aside from the genotype at the Or22 locus, and thus these could potentially contribute to the behavioural differences observed. Therefore, to further determine if variation at the Or22 locus leads to behavioural differences, we tested the same behavioural preference for ethyl hexanoate in females expressing either Or22a (ab3A-1) or Or22ab (ab3A-2) transgenes in the ‘empty neuron’ system [8]. We found a significant difference between the preferences of flies expressing Or22ab and those expressing Or22a (\(p = 0.014\)), with flies expressing Or22ab (ab3A-2) more attracted to ethyl hexanoate than those expressing Or22a (ab3A-1; figure 2c), as would be expected if the difference observed in the isogenic lines is due to the Or22 locus. We note that in this experiment the genetic background of the flies is not completely identical, as the background of the chromosome carrying the UAS transgene differs. Thus, while the combination of these two experiments is persuasive, we cannot definitively conclude from either alone that the behavioural differences we observe are solely due to the Or22 genotype.

We therefore performed a third experiment to further support that Or22 genotype influences attraction to ethyl hexanoate. In this case we tested attraction to ethyl hexanoate in females from two natural fly populations collected from the north and south of Australia. The population from the north (northern Queensland) was determined to have a 74.9% (±1.6%) frequency of the short allele, while the population from the south (southern Tasmania) only has the long allele present. We found that these two populations have significantly different preferences (Mann–Whitney \(U\), \(p = 0.003\)), with flies from the northern population, with the high frequency of Or22b, showing significantly higher attraction to ethyl hexanoate than flies from the southern population, which is fixed for the long allele (figure 2d). The combined data from these three different experiments thus strongly suggest that flies with the Or22ab variant are more strongly attracted to ethyl hexanoate than are flies in which Or22a determines ab3A response, and thus that genotype at the Or22 locus is associated with variation in olfactory preference behaviour.

2.3. Flies with the ab3A-1 and ab3A-2 phenotypes have different fruit oviposition preferences

Ethyl hexanoate and other odorants detected by the ab3A neuron are known to be released from a variety of fruits, including fruits on which D. melanogaster flies are known to
oviposit. Along the Australian east coast there are different geographical distributions of fruits grown due to different climates. For example, tropical fruits such as banana are grown in the north, whereas fruits that grow in temperate climates, such as apple, are grown predominantly in the south. We therefore wondered if there are differences between the ab3A phenotypes in preference of fruits for oviposition, which might lead to different phenotypes being favoured in different geographical regions, and might explain the observed cline in allele frequencies along the east coast.

To test for oviposition fruit preference, we designed an assay (adapted from that of Stensmyr et al. [27]) that gives females a choice between eight different fruit substrates on which to oviposit, as well as a control plate with no fruit added. Each fruit was homogenized and mixed with 1% agarose dyed blue to reduce any visual or textural influences on oviposition site choice. Based on the proportion of eggs laid on each fruit an oviposition PI was calculated that indicates the preference of the flies to oviposit on that particular fruit. For our eight fruits, we selected half that are ‘northern’ (tropical-growing) fruits (apple, apricot, pear and strawberry). This provided the flies with a wide variety of choice, and also allowed us to determine if flies preferred to oviposit on the northern or southern fruits. We used flies from the ab3A-1 and ab3A-2 isogenic lines in this experiment, as flies expressing Or22a and Or22ab in the empty neuron have very poor fertility (probably due to other genes within the chromosomal deletion in the empty neuron background strain [28]).

We found that the two isogenic lines showed significantly different preferences for northern and southern fruits, with ab3A-2 flies laying significantly more eggs on northern than southern fruits (figure 3c; Mann–Whitney U, \( p = 0.014 \)), whereas ab3A-1 flies showed no preference for one group over the other. When these data were broken down to analyse oviposition preference for the individual fruits we found that, while flies from the two isogenic lines showed similar preferences for ovipositing on six of the fruits, ab3A-1 flies had a stronger preference for apricot than did ab3A-2 flies (two-way ANOVA with Bonferroni post-tests, \( p<0.05 \)), and ab3A-2 flies had a stronger preference for banana than did ab3A-1 flies (\( p<0.001 \)). The difference in preference for banana was highly significant, and thus to add further evidence to this finding we performed a binary oviposition preference assay where flies had to select between banana and just one other fruit. For this we used apple as a fruit on which both lines laid well (compared to control plates, one-way ANOVA with Dunnett’s multiple comparison, ab3A-1 \( p<0.01 \), ab3A-2 \( p<0.05 \), but for which we had observed no preference difference between ab3A-1 and ab3A-2 flies (two-way ANOVA with Bonferroni post-tests, apple ab3A-1 v ab3A-2, \( p>0.05 \)). Supporting the initial finding, we found that ab3A-2 flies showed a strong preference for ovipositing on banana (on average laying 77% of their eggs on banana), whereas ab3A-1 flies did not discriminate between apple and banana (figure 3b; ab3A-1 \( p>0.05 \), ab3A-2 \( p<0.001 \)). These data therefore suggest that changes in ab3A phenotype correlate with altered oviposition site preference.

### 3. Discussion

Taken together, our data suggest that ab3A response variation due to changes at the \( D. \ melanogaster \) Or22 locus is associated with differences in two different olfactory-driven behaviours. Our finding that flies carrying the different Or22 variants show oviposition differences is especially intriguing. Oviposition decisions are crucial to fitness because \( Drosophila \) flies lay eggs on fermenting fruits where larvae then feed on yeast. It is thus possible that the high frequency of the Or22ab allele found in northern Australia reflects a selective advantage in these regions where different fruits are grown. This could be caused by differing sensitivity and behavioural responses to either the fruit-derived odours themselves, or due to odours produced by the yeasts and other microbes found in fermenting fruits, which could also differ in these geographical regions.

We note that it is, of course, possible that the differential distribution of Or22 alleles along the Australian east coast is not indicative of a role of this locus in selection. Instead, as \( D. \ melanogaster \) is a human commensal it may have been introduced to Australia multiple times, with different
frequencies of the different variants in the founding populations at their northern and southern introduction sites. It is also possible that the Or22 locus is in linkage disequilibrium with another locus of ecological importance, for example, one involved in heat adaptation. However, given that we find behavioural changes associated with the differences in ab3A response in transgenic flies where only this locus is different, it seems plausible that differences at this locus could lead to clinal distribution of these alleles.

Our work contributes to a growing body of evidence that, among the many classes of ORNs, the ab3A neurons are of particular importance to the ecological specialization of Drosophila flies. A recent study showed that a population of forest-dwelling wild D. melanogaster in Zimbabwe are seasonal specialists of the highly geographically restricted marula fruit [16]. The authors showed that this fruit releases high levels of the ester ethyl isovalerate, that this is detected by ab3A neurons, and that flies homozygous for Or22ab are more sensitive to this odorant than Canton-S flies. They further showed that removal of ab3A neuron function reduced the ability of flies to localize marula fruit, suggesting a key role for this neuron class in this behaviour. There is also significant evidence that the ab3A neuron plays a role in host-specialization in several other Drosophila species. D. sechellia are specialists of the morinda fruit, and their detection of this fruit is mediated by it releasing methyl hexanoate, which is predominantly detected by ab3A neurons [19]. D. erecta are seasonal specialists of the Pandanus fruit, and detection of this fruit is predominantly due to the release of 3-methyl-2-butenyl acetate, which is also primarily detected by ab3A neurons [20]. D. suzukii is a pest species that, unlike most other Drosophila species, prefers fresh rather than rotting fruit. The ab3A neurons of this species have been found to detect β-cyclocitrinal, a compound to which they are attracted that is released from leaves [21]. This ligand is not detected by the ab3A neurons of D. melanogaster, and the authors thus proposed that this change in ligand-specificity of the ab3A neuron is one of the factors involved in the switch of D. suzukii from rotting to fresh fruit.

Compared to other ORN classes, the ab3A neuron response profile is unusually variable across different Drosophila species [11,12], albeit this analysis of ORN types was not extensive. Paralleling this, in addition to showing genetic variation within D. melanogaster, the Or22 locus is also unique among the Ors in exhibiting high levels of genetic variation and extensive levels of copy number variation across species [15]. This suggests that the high level of genetic variability might provide opportunities for functional changes that

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**Figure 3.** Flies with different ab3A phenotypes show differences in fruit oviposition preference. (a) When presented with four fruits typically grown in the south and four typically grown in the north of Australia, isogenic lines with the ab3A-1 (blue) and ab3A-2 (orange) phenotype showed significantly different preferences for ovipositing on northern fruits (Mann–Whitney U, p = 0.014). Positive index indicates a preference for northern fruits. (b) Breaking down the data from (a) for the individual fruits, choices of ab3A-1 and ab3A-2 flies differed significantly for apricot and banana (2-way ANOVA with Bonferroni post-tests; apricot p < 0.05, banana p < 0.001) (c) When given a choice between only apple (red) and banana (yellow) as fruit substrates, ab3A-1 flies showed no differences in oviposition site choice, whereas ab3A-2 flies strongly preferred banana (2-way ANOVA with Bonferroni post-tests; ab3A-1 p > 0.05, ab3A-2 p < 0.001). All data are presented as the median + interquartile range, n = 12.
contribute to the evolution of differences in olfactory-driven behaviours. While naturally occurring changes in the sequence of some other Or genes have been implicated in changes in olfactory behaviour [29,30], our data provides one of the first insights into the causative links between Or genetic variation, neuronal response variation and behavioural changes. It will be of great interest in future to determine which Or22 genetic variants underpin shifts in host-specialization in other Drosophila species, and have thus contributed to the ability of these species to evolve and use different ecological niches.

4. Material and methods

4.1. Drosophila stocks

Four mass-bred populations of D. melanogaster, originally collected from Bowen, Queensland (19°58’S), Innisfail, Queensland (17°31’S), Northern Tasmania (41° S) and Southern Tasmania (43° S), were a kind gift from Carla Sgro, Monash University [31]. The allelic frequencies of the Southern Tasmania (43° S), were a kind gift from Carla Sgro, Monash University [31]. The allelic frequencies of the line were derived as described in Shaw et al. [31].

4.2. EAG recordings

Electroantennograms (EAGs) were recorded as per Tom et al. [32]. Briefly, a single fly was immobilized and a reference electrode inserted into the eye and a recording electrode placed on the surface of the antenna. Changes in voltage (mV) in response to 1 s stimulations with odorants were amplified using an active probe and a serial-IDAC amplifier (Syntech). EAGs represent the summed activity of a population of ORNs. Odour stimulation was by injecting volatiles from 5 ml syringes into an airstream blown over the preparation. All odorants were at highest available purity (greater than 98%, Sigma-Aldrich) and dissolved in paraffin oil to reach the specified dilution.

4.3. Two-choice cage assay

The two-choice cage assay was adapted from Faucher et al. [26]. A 305 x 305 x 305 mm cage was fitted with two 66 mm funnels placed 130 mm apart leading into 250 ml glass bottle traps. Bottles were filled with 40 ml solutions of 50% (vol/vol) apple cider vinegar (Cornwall’s) to attract flies.

The test odorant was added to one of the traps (10^-4 vol/vol for ethyl hexanoate). When testing isogenic lines and natural populations one hundred flies were starved on 1% agarose for 24 h before being released into the cage. When testing the flies expressing either Or22a or Or22ab in empty ab3A neurons between 50 and 80 flies were tested, due to the poor health of these flies. The cage was surrounded by an open-top black box to minimize directional visual stimuli and placed in a fume hood to provide ventilation. The assay was run for 17 h under natural light conditions, such that the dawn and dusk activity periods were captured. The side that contained the test odorant was alternated between every repetition. A PI was calculated as (T – C)/(T + C) where T is the number of flies in the test odorant-containing bottle and C is the total number of flies in the control bottle. The preference indices were then compared for significance using a Mann-Whitney U-test.

4.4. Oviposition behaviour assay

The oviposition assay was adapted from Stensmyr et al. [27]. We tested eight fruits representing those commonly grown in tropical northern Australia (banana, mango, pineapple and pomelo) and in temperate southern Australia (apple, apricot, pear and strawberry). Fruits were purchased from a vegetable market, aged for 3 days at room temperature and then homogenized and frozen. One gram of thawed homogenized fruit was added to the lid of a 35 mm tissue culture plate (Sarstedt) with two drops of blue food dye and 4 ml of 1% agarose minimizing both visual and textural differences. Once cooled and set, the plates were placed into a 235 x 235 x 120 mm plastic box with a damp sponge for humidity. A lid with holes for ventilation and left for 24 h at room temperature (22°C) before flies were added. Nine plates were placed in a 3 x 3 grid pattern; one for each of the eight fruits in a pseudorandomized distribution and a control plate containing water in the middle position. When only one apple and banana were tested the two fruits were distributed in a regular pattern over the eight plates, with the fruit in the corner plate alternating and the control plate remaining the same. Virgin females and males were kept separately for three days and then crossed. After 2 days 40 fertilized females were placed in a small cage with a food plate coated with yeast paste to induce laying. After 2 days laying on food plates, 30 females were gently sedated by cooling and released in the box. The assay was run for 24 h and the number of eggs laid on each plate was counted to calculate the proportion of eggs laid on each fruit out of the total.

Data accessibility. All data are available within the article or supporting electronic material, including the statistical analysis supporting this article as electronic supplementary material, table S1. The data are provided in electronic supplementary material [33].

Authors’ contributions. C.G.W. and M.d.B. conceived the experiments, interpreted the data and co-led the work. K.H.S. conceived the experiments, interpreted the data and performed the experiments. C.I.D. performed experiments. A.A. and T.K.J. interpreted the data. K.H.S. and C.G.W. wrote the paper.

Competing interests. Authors declare no conflicts of interest.

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