The fly ur-host

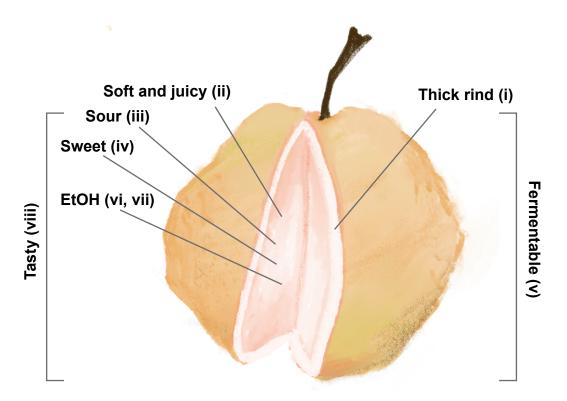


Figure S1. Putative characteristics of the ancestral host. Relates to Figure 1.

i) The citrus partiality indicates a fruit with thick rind [S1], ii) surrounding a soft and juicy pulp; allowing mobility of the larvae [S2]. iii) The fruit should be sour, since *D. melanogaster* preferentially lays eggs on acid-containing media [S3]. iv) The fruit should be sweet, given that *D. melanogaster* preferentially lays eggs on sugar rich substrates [S4]. v) The high sugar content would also ensure abundance of yeast – *D. melanogaster*'s favorite food [S5] – and enable rapid fermentation. vi) The fruit should have features that promote sustained high ethanol levels, under which *D. melanogaster* has a competitive advantage [S6]. vii) High ethanol levels also protect the larvae from parasitoid wasps [S7]. viii) The fruit should be palatable to humans, given that a shared human-fly preference would constitute the most direct route to commensalism. Drawing: Rakel Stensmyr.

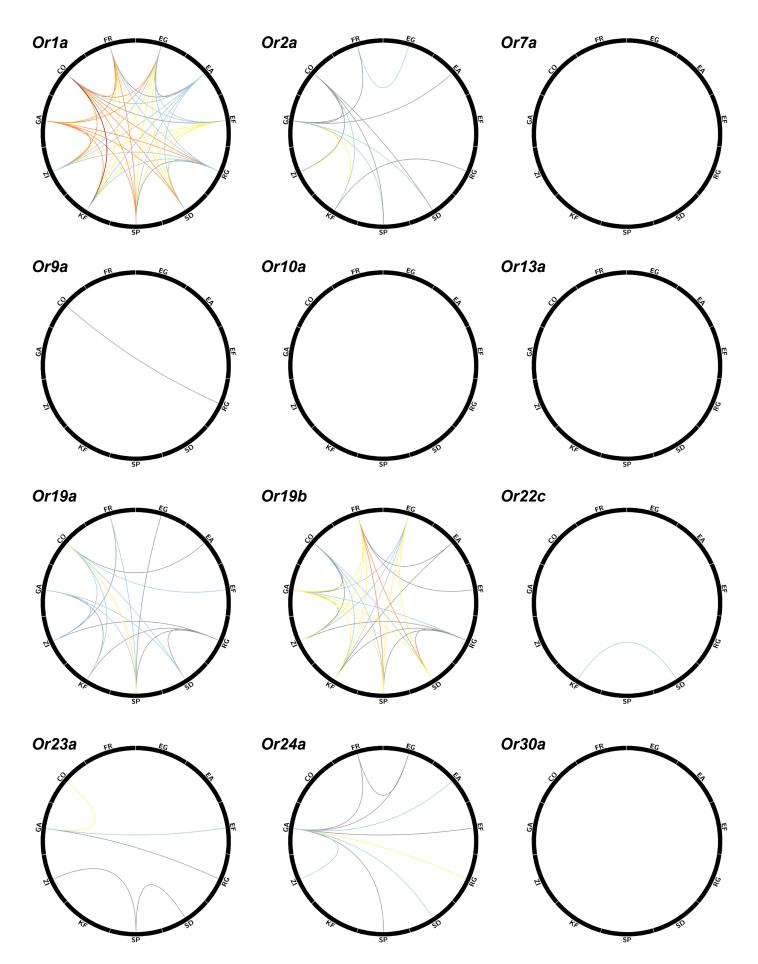


Figure S2. Local genetic differentiation within the OR family. Relates to Figure 3. Circos plots based on F_{ST} quantiles for all drosophila odorant receptors. Only connections between populations with unusually high F_{ST} values (elevated genetic differentiation) are shown. Color code as in Figure 3O.

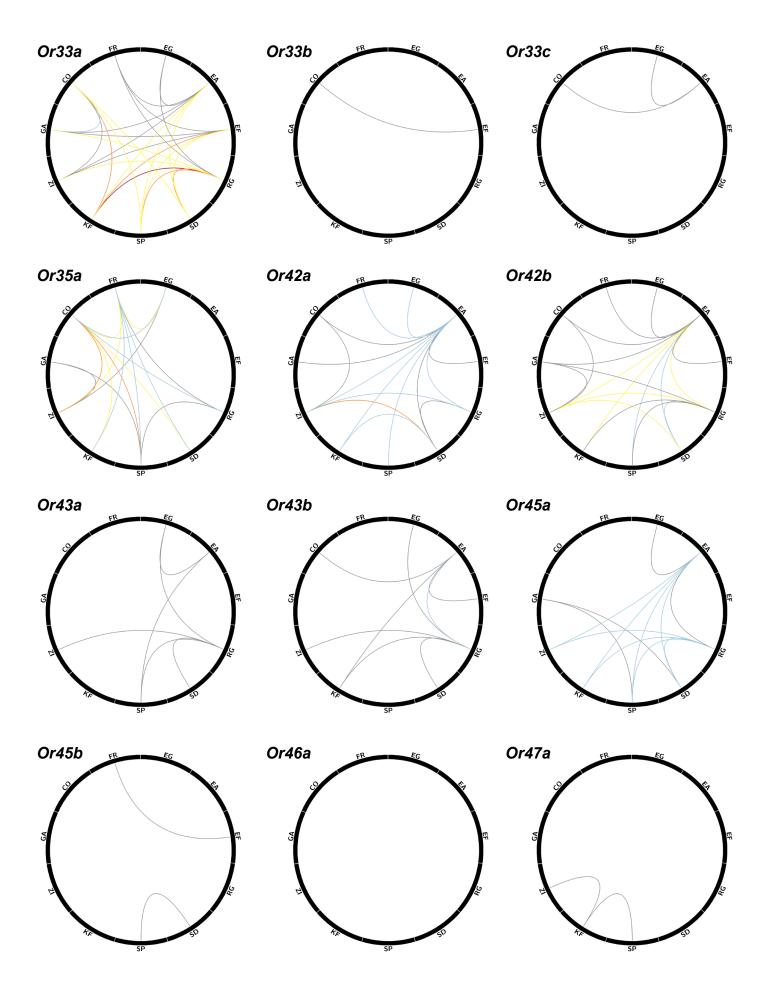


Figure S2. Local genetic differentiation within the OR family. Relates to Figure 3. ${\tt Continued}$

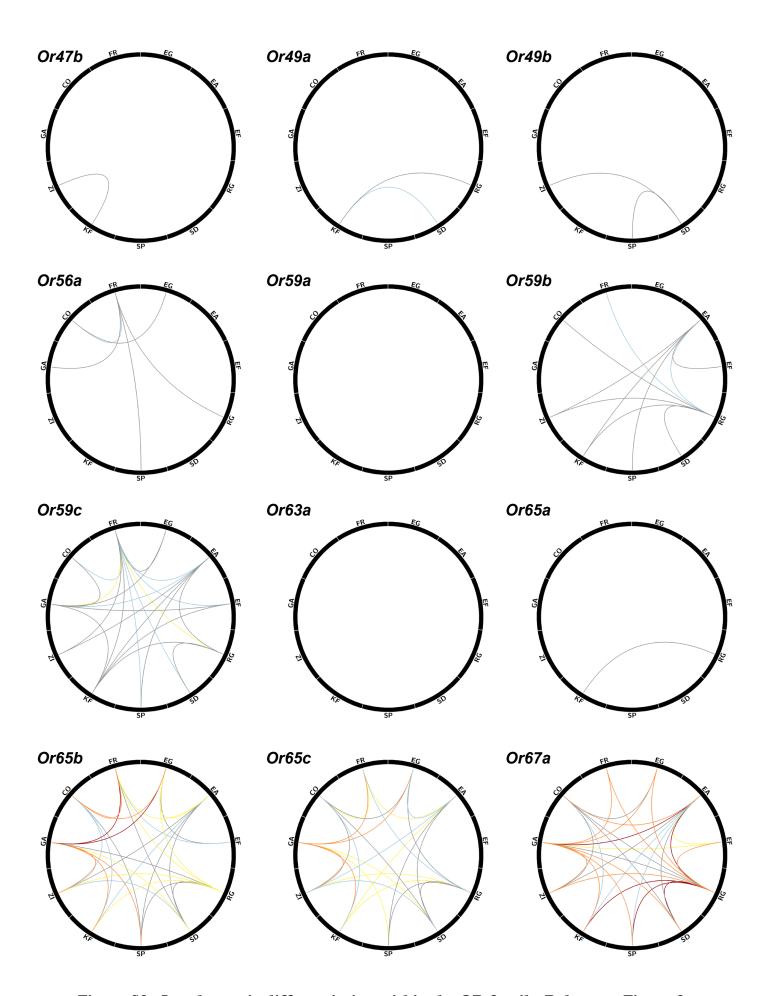


Figure S2. Local genetic differentiation within the OR family. Relates to Figure 3. $\,$ Continued

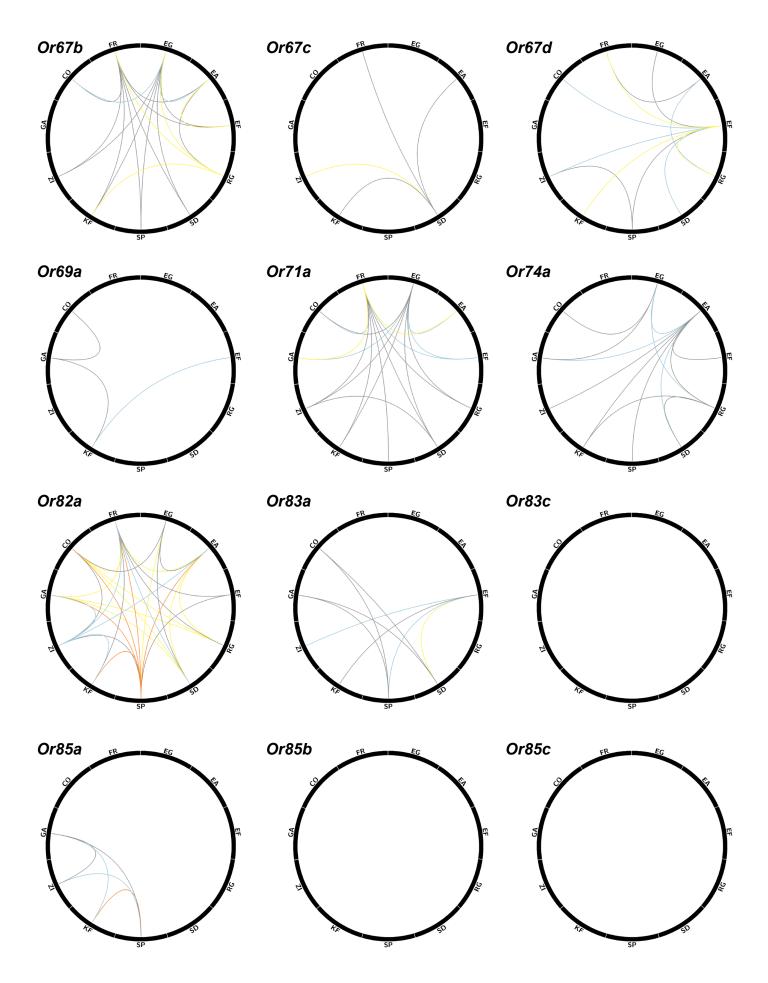


Figure S2. Local genetic differentiation within the OR family. Relates to Figure 3. ${\tt Continued}$

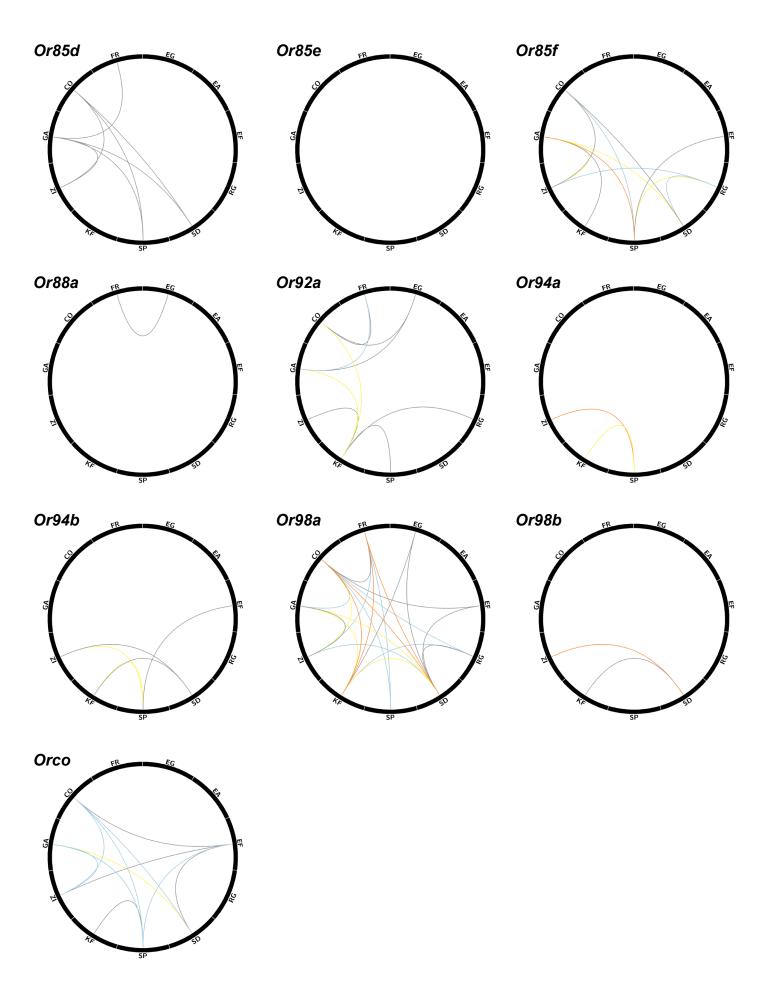


Figure S2. Local genetic differentiation within the OR family. Relates to Figure 3. $\,$ Continued

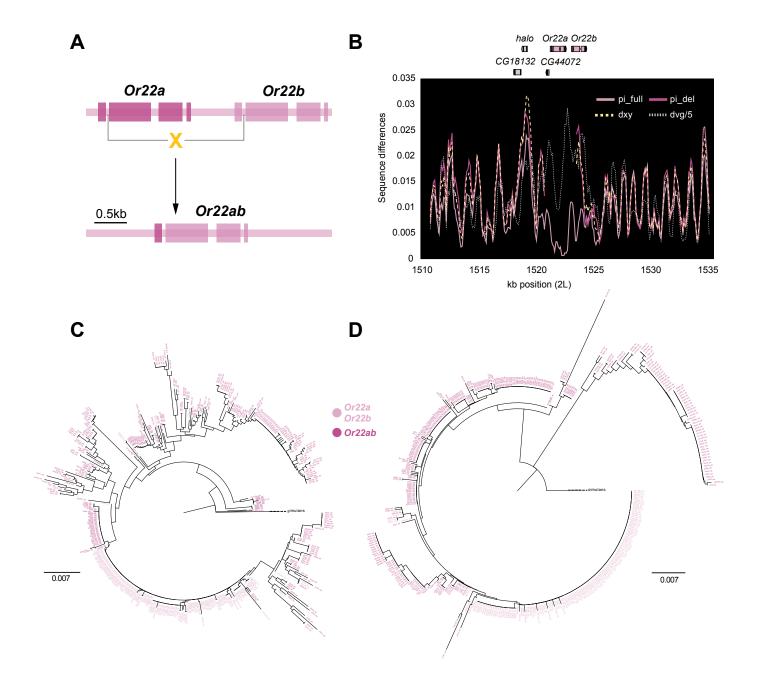


Figure S3. Genetic variation at the Or22 locus. Relates to Figure 3.

(A) The Or22a/Or22b locus, with the chimeric Or22ab deletion variant below, in D. melanogaster.
(B) Rates of pairwise sequence differences: among Zambia genomes carrying the full Or22a/Or22b haplotype (pi_full), among Zambia genomes carrying the deletion yielding the Or22ab fusion variant (pi_del), between Zambia full and Zambia deletion alleles (dxy) and average sequence divergence between Zambia D. melanogaster and the D. simulans reference (divided by 5 to show on the same scale).
(C) A neighbor joining tree for a 500 bp section of the Or22 region just upstream of the Or22ab deletion (1520.1 - 1520.6 kb), and (D) a comparable tree for a 500 bp region just downstream of this deletion (1522.9 - 1523.4 kb). Population labels are as in Figure 3O; "full" and "deletion" alleles are noted.

Supplemental references

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- **S6.** McKenzie, J. A., & Parsons, P. A. (1972). Alcohol tolerance: an ecological parameter in the relative success of *Drosophila melanogaster* and *Drosophila simulans*. Oecologia, 10, 373-388.
- **S7.** Lynch, Z.R., Schlenke, T.A., Morran, L.T., and De Roode, J.C. (2017). Ethanol confers differential protection against generalist and specialist parasitoids of *Drosophila melanogaster*. PloS one, 12, e0180182.