

GEPHE SUMMARY

| | | | |
|--|----------------|------------|--------------|
| | Gephebase Gene | | GepheID |
| Odorant receptor 3 (OR3) (https://www.gephebase.org/search-criteria?/and+Gene) | | GP00000749 | |
| Gephebase="Odorant receptor 3 (OR3)"#gephebase-summary-title) | | | Main curator |
| Published | Entry Status | Martin | |

PHENOTYPIC CHANGE

| | | |
|--|------------------------|--|
| | Trait Category | |
| Behavior (https://www.gephebase.org/search-criteria?/and+Trait) | | |
| Category="Behavior"#gephebase-summary-title) | | |
| | Trait | |
| Olfactory behavior (pheromone) (<a (pheromone)"#gephebase-summary-title"="" behavior="" href="https://www.gephebase.org/search-criteria?/and+Trait=" olfactory="">https://www.gephebase.org/search-criteria?/and+Trait="Olfactory behavior (pheromone)"#gephebase-summary-title) | | |
| | Trait State in Taxon A | |
| Ostrinia nubilalis | | |
| | Trait State in Taxon B | |
| Ostrinia furnacalis | | |
| | Ancestral State | |
| Data not curated | | |
| | Taxonomic Status | |
| Interspecific (https://www.gephebase.org/search-criteria?/and+Taxonomic) | | |
| Status="Interspecific"#gephebase-summary-title) | | |

| Taxon A | Latin Name | Taxon B | Latin Name |
|--|-----------------------------|--|-----------------------------|
| Ostrinia nubilalis (https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms="Ostrinia nubilalis"#gephebase-summary-title) | | Ostrinia furnacalis (<a furnacalis"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms=" ostrinia="">https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms="Ostrinia furnacalis"#gephebase-summary-title) | |
| | Common Name | | Common Name |
| European corn borer | | Asian corn borer | |
| | Synonyms | | Synonyms |
| Pyralis nubilalis; Pyrausta nubilalis; European corn borer; Ostrinia nubilalis (Hubner, 1796) | | Asian corn borer; Ostrinia furnacalis (Guenee, 1854); Ostrinia furnacalis | |
| | Rank | | Rank |
| species | | species | |
| | Lineage | | Lineage |
| cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Panarthropoda; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Dicondylia; Pterygota; Neoptera; Holometabola; Amphimesnoptera; Lepidoptera; Glossata; Neolepidoptera; Heteroneura; Ditrysia; Obtectomera; Pyraloidea; Crambidae; Pyraustinae; Ostrinia | | cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Panarthropoda; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Dicondylia; Pterygota; Neoptera; Holometabola; Amphimesnoptera; Lepidoptera; Glossata; Neolepidoptera; Heteroneura; Ditrysia; Obtectomera; Pyraloidea; Crambidae; Pyraustinae; Ostrinia | |
| | Parent | | Parent |
| Ostrinia () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=29056) | | Ostrinia () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=29056) | |
| | NCBI Taxonomy ID | | NCBI Taxonomy ID |
| 29057 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=29057) | | 93504 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=93504) | |
| | is Taxon A an Intraspecies? | | is Taxon B an Intraspecies? |
| No | | No | |

GENOTYPIC CHANGE

| | | | |
|---|-------------------------|--|------------------------------|
| | Generic Gene Name | | UniProtKB Ostrinia nubilalis |
| OR3 | | D3J5H6 (http://www.uniprot.org/uniprot/D3J5H6) | |
| | Synonyms | | GenebankID or UniProtKB |
| - | | AFK30395 (https://www.ncbi.nlm.nih.gov/nucleotide/AFK30395) | |
| | String | | |
| - | | | |
| | Sequence Similarities | | |
| Belongs to the insect chemoreceptor superfamily. Heteromeric odorant receptor channel (TC 1.A.69) family. | | | |
| | GO - Molecular Function | | |
| GO:0005549 : odorant binding (https://www.ebi.ac.uk/QuickGO/term/GO:0005549) | | | |
| GO:0004984 : olfactory receptor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0004984) | | | |
| | GO - Biological Process | | |
| GO:0007165 : signal transduction (https://www.ebi.ac.uk/QuickGO/term/GO:0007165) | | | |
| | GO - Cellular Component | | |
| GO:0016021 : integral component of membrane | | | |

(<https://www.ebi.ac.uk/QuickGO/term/GO:0016021>)
GO:0005886 : plasma membrane (<https://www.ebi.ac.uk/QuickGO/term/GO:0005886>)

Presumptive Null

No ([https://www.gephebase.org/search-criteria?/and+Presumptive Null=~No^#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Presumptive+Null=~No^#gephebase-summary-title))

Molecular Type

Coding ([https://www.gephebase.org/search-criteria?/and+Molecular Type=~Coding^#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Molecular+Type=~Coding^#gephebase-summary-title))

Aberration Type

SNP ([https://www.gephebase.org/search-criteria?/and+Aberration Type=~SNP^#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Aberration+Type=~SNP^#gephebase-summary-title))

SNP Coding Change

Nonsynonymous

Molecular Details of the Mutation

A148T

Experimental Evidence

Candidate Gene ([https://www.gephebase.org/search-criteria?/and+Experimental Evidence=~Candidate Gene^#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Experimental+Evidence=~Candidate+Gene^#gephebase-summary-title))

| | Taxon A | Taxon B | Position |
|------------|---------|---------|----------|
| Codon | - | - | - |
| Amino-acid | - | - | - |

Main Reference

Single mutation to a sex pheromone receptor provides adaptive specificity between closely related moth species. (2012) (<https://pubmed.ncbi.nlm.nih.gov/22891317>)

Authors

Leary GP; Allen JE; Bunger PL; Luginbill JB; Linn CE; Macallister IE; Kavanaugh MP; Wanner KW

Abstract

Sex pheromone communication, acting as a prezygotic barrier to mating, is believed to have contributed to the speciation of moths and butterflies in the order Lepidoptera. Five decades after the discovery of the first moth sex pheromone, little is known about the molecular mechanisms that underlie the evolution of pheromone communication between closely related species. Although Asian and European corn borers (ACB and ECB) can be interbred in the laboratory, they are behaviorally isolated from mating naturally by their responses to subtly different sex pheromone isomers, (E)-12- and (Z)-12-tetradecenyl acetate and (E)-11- and (Z)-11-tetradecenyl acetate (ACB: E12, Z12; ECB: E11, Z11). Male moth olfactory systems respond specifically to the pheromone blend produced by their conspecific females. In vitro, ECB(Z) odorant receptor 3 (OR3), a sex pheromone receptor expressed in male antennae, responds strongly to E11 but also generally to the Z11, E12, and Z12 pheromones. In contrast, we show that ACB OR3, a gene that has been subjected to positive selection ($d_{i\%} = 2.9$), responds preferentially to the ACB E12 and Z12 pheromones. In *Ostrinia* species the amino acid residue corresponding to position 148 in transmembrane domain 3 of OR3 is alanine (A), except for ACB OR3 that has a threonine (T) in this position. Mutation of this residue from A to T alters the pheromone recognition pattern by selectively reducing the E11 response $\hat{\sim}^{1/4}$ 14-fold. These results suggest that discrete mutations that narrow the specificity of more broadly responsive sex pheromone receptors may provide a mechanism that contributes to speciation.

Additional References

RELATED GEPHE

No matches found.

Related Genes

No matches found.

Related Haplotypes

EXTERNAL LINKS

COMMENTS

@SexualTrait