

GEPHE SUMMARY

| | | | |
|--|---|---------------------------------|------------------------------------|
| <p>avpr1a (https://www.gephebase.org/search-criteria?/and+Gene+Gephebase=^avpr1a^#gephebase-summary-title)</p> <p>Published</p> | <p>Gephebase Gene</p> <p>Entry Status</p> | <p>GP00000129</p> <p>Martin</p> | <p>GepheID</p> <p>Main curator</p> |
|--|---|---------------------------------|------------------------------------|

PHENOTYPIC CHANGE

| | | | |
|--|---|--|---|
| <p>Behavior (https://www.gephebase.org/search-criteria?/and+Trait+Category=^Behavior^#gephebase-summary-title)</p> <p>Male sexual fidelity (monogamous regime) (https://www.gephebase.org/search-criteria?/and+Trait=^Male+sexual+fidelity+(monogamous+regime)^#gephebase-summary-title)</p> <p>Microtus ochrogaster</p> <p>Microtus ochrogaster</p> <p>Data not curated</p> <p>Intraspecific (https://www.gephebase.org/search-criteria?/and+Taxonomic+Status=^Intraspecific^#gephebase-summary-title)</p> | <p>Trait Category</p> <p>Trait</p> <p>Trait State in Taxon A</p> <p>Trait State in Taxon B</p> <p>Ancestral State</p> <p>Taxonomic Status</p> | <p>Behavior</p> <p>Male sexual fidelity (monogamous regime)</p> <p>Microtus ochrogaster</p> <p>Microtus ochrogaster</p> <p>Data not curated</p> <p>Intraspecific</p> | <p></p> <p></p> <p></p> <p></p> <p></p> <p></p> <p></p> |
|--|---|--|---|

| Taxon A | Latin Name | Taxon B | Latin Name |
|--|--|--|--|
| Microtus ochrogaster (https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Microtus+ochrogaster^#gephebase-summary-title) | Microtus ochrogaster | Microtus ochrogaster (https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Microtus+ochrogaster^#gephebase-summary-title) | Microtus ochrogaster |
| prairie vole | prairie vole | prairie vole | prairie vole |
| prairie vole; prairie voles | prairie vole; prairie voles | prairie vole; prairie voles | prairie vole; prairie voles |
| species | species | species | species |
| cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; Theria; Eutheria; Boreoeutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Cricetidae; Arvicolinae; Microtus | cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; Theria; Eutheria; Boreoeutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Cricetidae; Arvicolinae; Microtus | cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; Theria; Eutheria; Boreoeutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Cricetidae; Arvicolinae; Microtus | cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; Theria; Eutheria; Boreoeutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Cricetidae; Arvicolinae; Microtus |
| Microtus (meadow voles) - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=10053) | Microtus (meadow voles) - (Rank: genus) | Microtus (meadow voles) - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=10053) | Microtus (meadow voles) - (Rank: genus) |
| 79684 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=79684) | 79684 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=79684) | 79684 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=79684) | 79684 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=79684) |
| is Taxon A an Intraspecies? | | is Taxon B an Intraspecies? | |
| No | | No | |

GENOTYPIC CHANGE

| | | | |
|--|--|---|--|
| <p>Avpr1a</p> <p>V1a; AVPR; V1aR; Avpr1</p> <p>10090.ENSMUSP00000020323 (http://string-db.org/newstring.cgi/show_network_section.pl?identifier=10090.ENSMUSP00000020323)</p> <p>Belongs to the G-protein coupled receptor 1 family. Vasopressin/oxytocin receptor subfamily.</p> <p>GO:0017046 : peptide hormone binding (https://www.ebi.ac.uk/QuickGO/term/GO:0017046)</p> <p>GO:0031894 : V1A vasopressin receptor binding (https://www.ebi.ac.uk/QuickGO/term/GO:0031894)</p> <p>GO:0005000 : vasopressin receptor activity</p> | <p>Generic Gene Name</p> <p>Synonyms</p> <p>String</p> <p>Sequence Similarities</p> <p>GO - Molecular Function</p> | <p>Q62463 (http://www.uniprot.org/uniprot/Q62463)</p> <p>()</p> <p></p> <p></p> <p></p> <p></p> <p></p> <p></p> | <p>UniProtKB Mus musculus</p> <p>GenebankID or UniProtKB</p> |
|--|--|---|--|

(<https://www.ebi.ac.uk/QuickGO/term/GO:0005000>)

GO - Biological Process

- GO:0010033 : response to organic substance
(<https://www.ebi.ac.uk/QuickGO/term/GO:0010033>)
- GO:0007186 : G protein-coupled receptor signaling pathway
(<https://www.ebi.ac.uk/QuickGO/term/GO:0007186>)
- GO:0008284 : positive regulation of cell proliferation
(<https://www.ebi.ac.uk/QuickGO/term/GO:0008284>)
- GO:0019722 : calcium-mediated signaling
(<https://www.ebi.ac.uk/QuickGO/term/GO:0019722>)
- GO:0032870 : cellular response to hormone stimulus
(<https://www.ebi.ac.uk/QuickGO/term/GO:0032870>)
- GO:0042631 : cellular response to water deprivation
(<https://www.ebi.ac.uk/QuickGO/term/GO:0042631>)
- GO:0007625 : grooming behavior (<https://www.ebi.ac.uk/QuickGO/term/GO:0007625>)
- GO:0002125 : maternal aggressive behavior
(<https://www.ebi.ac.uk/QuickGO/term/GO:0002125>)
- GO:0042711 : maternal behavior (<https://www.ebi.ac.uk/QuickGO/term/GO:0042711>)
- GO:0014902 : myotube differentiation
(<https://www.ebi.ac.uk/QuickGO/term/GO:0014902>)
- GO:0007621 : negative regulation of female receptivity
(<https://www.ebi.ac.uk/QuickGO/term/GO:0007621>)
- GO:0051970 : negative regulation of transmission of nerve impulse
(<https://www.ebi.ac.uk/QuickGO/term/GO:0051970>)
- GO:0043084 : penile erection (<https://www.ebi.ac.uk/QuickGO/term/GO:0043084>)
- GO:0045777 : positive regulation of blood pressure
(<https://www.ebi.ac.uk/QuickGO/term/GO:0045777>)
- GO:0030307 : positive regulation of cell growth
(<https://www.ebi.ac.uk/QuickGO/term/GO:0030307>)
- GO:0032849 : positive regulation of cellular pH reduction
(<https://www.ebi.ac.uk/QuickGO/term/GO:0032849>)
- GO:0007204 : positive regulation of cytosolic calcium ion concentration
(<https://www.ebi.ac.uk/QuickGO/term/GO:0007204>)
- GO:0014049 : positive regulation of glutamate secretion
(<https://www.ebi.ac.uk/QuickGO/term/GO:0014049>)
- GO:0010460 : positive regulation of heart rate
(<https://www.ebi.ac.uk/QuickGO/term/GO:0010460>)
- GO:0031394 : positive regulation of prostaglandin biosynthetic process
(<https://www.ebi.ac.uk/QuickGO/term/GO:0031394>)
- GO:0035815 : positive regulation of renal sodium excretion
(<https://www.ebi.ac.uk/QuickGO/term/GO:0035815>)
- GO:0003084 : positive regulation of systemic arterial blood pressure
(<https://www.ebi.ac.uk/QuickGO/term/GO:0003084>)
- GO:0045907 : positive regulation of vasoconstriction
(<https://www.ebi.ac.uk/QuickGO/term/GO:0045907>)
- GO:0001992 : regulation of systemic arterial blood pressure by vasopressin
(<https://www.ebi.ac.uk/QuickGO/term/GO:0001992>)
- GO:0051412 : response to corticosterone
(<https://www.ebi.ac.uk/QuickGO/term/GO:0051412>)
- GO:0010035 : response to inorganic substance
(<https://www.ebi.ac.uk/QuickGO/term/GO:0010035>)
- GO:0035176 : social behavior (<https://www.ebi.ac.uk/QuickGO/term/GO:0035176>)
- GO:0042713 : sperm ejaculation (<https://www.ebi.ac.uk/QuickGO/term/GO:0042713>)
- GO:0021537 : telencephalon development
(<https://www.ebi.ac.uk/QuickGO/term/GO:0021537>)

GO - Cellular Component

- GO:0005886 : plasma membrane (<https://www.ebi.ac.uk/QuickGO/term/GO:0005886>)
- GO:0005887 : integral component of plasma membrane
(<https://www.ebi.ac.uk/QuickGO/term/GO:0005887>)
- GO:0031410 : cytoplasmic vesicle (<https://www.ebi.ac.uk/QuickGO/term/GO:0031410>)

Unknown (<https://www.gephebase.org/search-criteria/?and+Presumptive Null=~Unknown^#gephebase-summary-title>) Presumptive Null

Cis-regulatory (<https://www.gephebase.org/search-criteria/?and+Molecular Type=~Cis-regulatory^#gephebase-summary-title>) Molecular Type

Unknown (<https://www.gephebase.org/search-criteria/?and+Aberration Type=~Unknown^#gephebase-summary-title>) Aberration Type

unknown Molecular Details of the Mutation

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

Sexual fidelity trade-offs promote regulatory variation in the prairie vole brain. (2015) (<https://pubmed.ncbi.nlm.nih.gov/26659055>) Main Reference

Okhovat M; Berrio A; Wallace G; Ophir AG; Phelps SM Authors

Abstract

Individual variation in social behavior seems ubiquitous, but we know little about how it relates to brain diversity. Among monogamous prairie voles, levels of vasopressin receptor (encoded by the gene *avpr1a*) in brain regions related to spatial memory predict male space use and sexual fidelity in the field. We find that trade-offs between the benefits of male fidelity and infidelity are reflected in patterns of territorial intrusion, offspring paternity, *avpr1a* expression, and the evolutionary fitness of alternative *avpr1a* alleles. DNA variation at the *avpr1a* locus includes

polymorphisms that reliably predict the epigenetic status and neural expression of avpr1a, and patterns of DNA diversity demonstrate that avpr1a regulatory variation has been favored by selection. In prairie voles, trade-offs in the fitness consequences of social behaviors seem to promote neuronal and molecular diversity.

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Additional References

Complex selection on a regulator of social cognition: Evidence of balancing selection, regulatory interactions and population differentiation in the prairie vole Avpr1a locus. (2018)
(<https://pubmed.ncbi.nlm.nih.gov/29218792>)

RELATED GEPHE

No matches found.

Related Genes

No matches found.

Related Haplotypes

EXTERNAL LINKS

COMMENTS

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