

GEPHE SUMMARY

<p>TAS1R1 (https://www.gephebase.org/search-criteria?/and+GeneGephebase=TAS1R1#gephebase-summary-title)</p> <p>Published</p>	<p>Gephebase Gene</p> <p>Entry Status</p>	<p>GP00001415</p> <p>Courtier</p>	<p>GepheID</p> <p>Main curator</p>
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PHENOTYPIC CHANGE

<p>Physiology (https://www.gephebase.org/search-criteria?/and+TraitCategory=Physiology#gephebase-summary-title)</p> <p>Taste sensitivity (umami) (https://www.gephebase.org/search-criteria?/and+Trait=Taste+sensitivity+(umami)#gephebase-summary-title)</p> <p>-</p> <p>Taxon A</p> <p>Interspecific (https://www.gephebase.org/search-criteria?/and+TaxonomicStatus=Interspecific#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>-</p> <p>-</p> <p>Taxon A</p> <p>Ursus maritimus (https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=Ursus+maritimus#gephebase-summary-title)</p> <p>polar bear</p> <p>Thalarcctos maritimus; polar bear; white bear species</p> <p>cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; Theria; Eutheria; Boreoeutheria; Laurasiatheria; Carnivora; Caniformia; Ursidae; Ursus</p> <p>Ursus () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=9639)</p> <p>29073 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=29073)</p> <p>No is Taxon A an Intraspecies?</p>	<p>Taxon B</p> <p>Latin Name</p> <p>Common Name</p> <p>Synonyms</p> <p>Rank</p> <p>Lineage</p> <p>Parent</p> <p>NCBI Taxonomy ID</p> <p>is Taxon B an Intraspecies?</p>	<p>Ailurus fulgens (https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=Ailurus+fulgens#gephebase-summary-title)</p> <p>lesser panda</p> <p>lesser panda; red panda species</p> <p>cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; Theria; Eutheria; Boreoeutheria; Laurasiatheria; Carnivora; Caniformia; Ailuridae; Ailurus</p> <p>Ailurus () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=9648)</p> <p>9649 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=9649)</p> <p>No is Taxon B an Intraspecies?</p>
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GENOTYPIC CHANGE

<p>Tas1r1</p> <p>TR1; T1r1; Gpr70; Tr1</p> <p>10090.ENSMUSP00000030792 (http://string-db.org/newstring.cgi/show_network_section.pl?identifier=10090.ENSMUSP00000030792)</p> <p>Belongs to the G-protein coupled receptor 3 family. TAS1R subfamily.</p> <p>GO:0004930 : G protein-coupled receptor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0004930)</p> <p>GO:0008527 : taste receptor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0008527)</p> <p>GO:0050917 : sensory perception of umami taste (https://www.ebi.ac.uk/QuickGO/term/GO:0050917)</p>	<p>Generic Gene Name</p> <p>Synonyms</p> <p>String</p> <p>Sequence Similarities</p> <p>GO - Molecular Function</p> <p>GO - Biological Process</p> <p>GO - Cellular Component</p>	<p>Q99PG6 (http://www.uniprot.org/uniprot/Q99PG6)</p> <p>()</p> <p>UniProtKB Mus musculus</p> <p>GenebankID or UniProtKB</p>
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GO:0005887 : integral component of plasma membrane
(<https://www.ebi.ac.uk/QuickGO/term/GO:0005887>)

Presumptive Null

Yes ([https://www.gephebase.org/search-criteria?/and+Presumptive Null=~Yes^#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Presumptive+Null=~Yes^#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

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

Deletion Size

1-9 bp

Molecular Details of the Mutation

1bp deletion (deletion of a C) in the sixth exon;

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))

Main Reference

Comparative genomics reveals convergent evolution between the bamboo-eating giant and red pandas. (2017) (<https://pubmed.ncbi.nlm.nih.gov/28096377>)

Authors

Hu Y; Wu Q; Ma S; Ma T; Shan L; Wang X; Nie Y; Ning Z; Yan L; Xiu Y; Wei F

Abstract

Phenotypic convergence between distantly related taxa often mirrors adaptation to similar selective pressures and may be driven by genetic convergence. The giant panda (*Ailuropoda melanoleuca*) and red panda (*Ailurus fulgens*) belong to different families in the order Carnivora, but both have evolved a specialized bamboo diet and adaptive pseudothumb, representing a classic model of convergent evolution. However, the genetic bases of these morphological and physiological convergences remain unknown. Through de novo sequencing the red panda genome and improving the giant panda genome assembly with added data, we identified genomic signatures of convergent evolution. Limb development genes *DYNC2H1* and *PCNT* have undergone adaptive convergence and may be important candidate genes for pseudothumb development. As evolutionary responses to a bamboo diet, adaptive convergence has occurred in genes involved in the digestion and utilization of bamboo nutrients such as essential amino acids, fatty acids, and vitamins. Similarly, the umami taste receptor gene *TAS1R1* has been pseudogenized in both pandas. These findings offer insights into genetic convergence mechanisms underlying phenotypic convergence and adaptation to a specialized bamboo diet.

Additional References

RELATED GEPHE

Related Genes

No matches found.

Related Haplotypes

1 ([https://www.gephebase.org/search-criteria?/or+Gene Gephebase=~TAS1R1^/and+Taxon ID=~29073^/or+Gene Gephebase=~TAS1R1^/and+Taxon ID=~9649^#gephebase-summary-title](https://www.gephebase.org/search-criteria?/or+Gene+Gephebase=~TAS1R1^/and+Taxon+ID=~29073^/or+Gene+Gephebase=~TAS1R1^/and+Taxon+ID=~9649^#gephebase-summary-title))

EXTERNAL LINKS

COMMENTS

TAS1R1 has become a pseudogene because of one nucleotide deletion in the sixth exon - as confirmed by Sanger sequencing of three additional individuals