

GEPHE SUMMARY

<p>VGLL3 (vestigial-like family member 3 gene) (#gephebase-summary-title)</p> <p>Published</p>	<p>Gephebase Gene</p> <p>GP00001167</p> <p>Martin</p> <p>Entry Status</p>	<p>GepheID</p> <p>Main curator</p>
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PHENOTYPIC CHANGE

<p>Physiology (#gephebase-summary-title)</p> <p>Age of sexual maturity (#gephebase-summary-title)</p> <p>Salmo salar</p> <p>Salmo salar</p> <p>Data not curated</p> <p>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>Physiology</p> <p>Age of sexual maturity</p> <p>Salmo salar</p> <p>Salmo salar</p> <p>Data not curated</p> <p>Intraspecific</p>	
Taxon A		Taxon B	
<p>Salmo salar (#gephebase-summary-title)</p> <p>Atlantic salmon</p> <p>Atlantic salmon; Salmo salar Linnaeus, 1758</p> <p>species</p> <p>cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Actinopterygii; Actinopteri; Neopterygii; Teleostei; Osteoglossocephalai; Clupeocephala; Euteleostei; Protacanthopterygii; Salmoniformes; Salmonidae; Salmoninae; Salmo</p> <p>Salmo () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=8028)</p> <p>8030 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=8030)</p> <p>is Taxon A an Infrapopulation?</p> <p>No</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>Salmo salar (#gephebase-summary-title)</p> <p>Atlantic salmon</p> <p>Atlantic salmon; Salmo salar Linnaeus, 1758</p> <p>species</p> <p>cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Actinopterygii; Actinopteri; Neopterygii; Teleostei; Osteoglossocephalai; Clupeocephala; Euteleostei; Protacanthopterygii; Salmoniformes; Salmonidae; Salmoninae; Salmo</p> <p>Salmo () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=8028)</p> <p>8030 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=8030)</p> <p>is Taxon B an Infrapopulation?</p> <p>No</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>

GENOTYPIC CHANGE

<p>VGLL3</p> <p>VGL3; VGL-3</p> <p>9606.ENSPO0000381436 (http://string-db.org/newstring.cgi/show_network_section.pl?identifier=9606.ENSPO0000381436)</p> <p>Belongs to the vestigial family.</p> <p>-</p> <p>GO:0006357 : regulation of transcription by RNA polymerase II (https://www.ebi.ac.uk/QuickGO/term/GO:0006357)</p> <p>GO:0005634 : nucleus (https://www.ebi.ac.uk/QuickGO/term/GO:0005634)</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>A8MV65 (http://www.uniprot.org/uniprot/A8MV65)</p> <p>()</p> <p>UniProtKB Homo sapiens</p> <p>GenebankID or UniProtKB</p>
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Unknown (<https://www.gephebase.org/search-criteria?/and+Presumptive Null=^Unknown^#gephebase-summary-title>)

Presumptive Null

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

Molecular Type

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

Aberration Type

candidate amino-acid substitutions

Molecular Details of the Mutation

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

Experimental Evidence

Sex-dependent dominance at a single locus maintains variation in age at maturity in salmon. (2015) (<https://pubmed.ncbi.nlm.nih.gov/26536110>)

Main Reference

Barson NJ; Aykanat T; Hindar K; Baranski M; Bolstad GH; Fiske P; Jacq C; Jensen AJ; Johnston SE; Karlsson S; Kent M; Moen T; Niemelä E; Nome T; Näsje TF; Orell P; Romakkaniemi A; Sævið grov H; Urdal K; Erkinaro J; Lien S; Primmer CR

Authors

Abstract

Males and females share many traits that have a common genetic basis; however, selection on these traits often differs between the sexes, leading to sexual conflict. Under such sexual antagonism, theory predicts the evolution of genetic architectures that resolve this sexual conflict. Yet, despite intense theoretical and empirical interest, the specific loci underlying sexually antagonistic phenotypes have rarely been identified, limiting our understanding of how sexual conflict impacts genome evolution and the maintenance of genetic diversity. Here we identify a large effect locus controlling age at maturity in Atlantic salmon (*Salmo salar*), an important fitness trait in which selection favours earlier maturation in males than females, and show it is a clear example of sex-dependent dominance that reduces intralocus sexual conflict and maintains adaptive variation in wild populations. Using high-density single nucleotide polymorphism data across 57 wild populations and whole genome re-sequencing, we find that the vestigial-like family member 3 gene (*VGLL3*) exhibits sex-dependent dominance in salmon, promoting earlier and later maturation in males and females, respectively. *VGLL3*, an adiposity regulator associated with size and age at maturity in humans, explained 39% of phenotypic variation, an unexpectedly large proportion for what is usually considered a highly polygenic trait. Such large effects are predicted under balancing selection from either sexually antagonistic or spatially varying selection. Our results provide the first empirical example of dominance reversal allowing greater optimization of phenotypes within each sex, contributing to the resolution of sexual conflict in a major and widespread evolutionary trade-off between age and size at maturity. They also provide key empirical evidence for how variation in reproductive strategies can be maintained over large geographical scales. We anticipate these findings will have a substantial impact on population management in a range of harvested species where trends towards earlier maturation have been observed.

Additional References

The *vgll3* Locus Controls Age at Maturity in Wild and Domesticated Atlantic Salmon (*Salmo salar* L.) Males. (2015) (<https://pubmed.ncbi.nlm.nih.gov/26551894>)

RELATED GEPHE

No matches found.

Related Genes

No matches found.

Related Haplotypes

EXTERNAL LINKS

COMMENTS

@BalancingSelection @SexualTrait ; dominance ; complex trait