

GEPHE SUMMARY

MC4R (https://www.gephebase.org/search-criteria?/and+GeneGephebase=~MC4R~#gephebase-summary-title)		Gephebase Gene	GP00000638	GepheID
Published		Entry Status	Martin	Main curator

PHENOTYPIC CHANGE

Physiology (https://www.gephebase.org/search-criteria?/and+TraitCategory=~Physiology~#gephebase-summary-title)		Trait Category	
Sexual maturation (onset) (https://www.gephebase.org/search-criteria?/and+Trait=~Sexual maturation (onset)~#gephebase-summary-title)		Trait	
Xiphophorus maculatus		Trait State in Taxon A	
Xiphophorus maculatus		Trait State in Taxon B	
Data not curated		Ancestral State	
Intraspecific (https://www.gephebase.org/search-criteria?/and+TaxonomicStatus=~Intraspecific~#gephebase-summary-title)		Taxonomic Status	
Taxon A		Taxon B	
Xiphophorus maculatus (https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms=~Xiphophorus maculatus~#gephebase-summary-title)		Xiphophorus maculatus (https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms=~Xiphophorus maculatus~#gephebase-summary-title)	
southern platyfish		southern platyfish	
Platypoecilus maculatus; southern platyfish; Platypoecilus maculatus Guenther, 1866; Xiphophorus maculatus (Guenther, 1866)		Platypoecilus maculatus; southern platyfish; Platypoecilus maculatus Guenther, 1866; Xiphophorus maculatus (Guenther, 1866)	
species		species	
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Actinopterygii; Actinopteri; Neopterygii; Teleostei; Osteoglossocephalai; Clupeocephala; Euteleosteomorpha; Neoteleostei; Eurypterygia; Ctenosquamata; Acanthomorpha; Euacanthomorpha; Percomorphaceae; Ovalentaria; Atherinomorphae; Cyprinodontiformes; Cyprinodontoides; Poeciliidae; Poeciliinae; Xiphophorus		cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Actinopterygii; Actinopteri; Neopterygii; Teleostei; Osteoglossocephalai; Clupeocephala; Euteleosteomorpha; Neoteleostei; Eurypterygia; Ctenosquamata; Acanthomorpha; Euacanthomorpha; Percomorphaceae; Ovalentaria; Atherinomorphae; Cyprinodontiformes; Cyprinodontoides; Poeciliidae; Poeciliinae; Xiphophorus	
Xiphophorus () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 8082)		Xiphophorus () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 8082)	
8083 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 8083)		8083 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 8083)	
No		No	

GENOTYPIC CHANGE

mc4r		Generic Gene Name	UniProtKB Danio rerio
zMC4; MC4-R; si:dkey-151m6.62		Synonyms	GenebankID or UniProtKB
7955.ENSDDARP00000027547 (http://string-db.org/newstring.cgi/show_network_section.pl?identifier=7955.ENSDDARP00000027547)		String	AHC02893 (https://www.ncbi.nlm.nih.gov/nuccore/AHC02893)
Belongs to the G-protein coupled receptor 1 family.		Sequence Similarities	
GO:0004977 : melanocortin receptor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0004977)		GO - Molecular Function	
GO:0040008 : regulation of growth (https://www.ebi.ac.uk/QuickGO/term/GO:0040008)		GO - Biological Process	

GO:0007189 : adenylyate cyclase-activating G protein-coupled receptor signaling pathway
(<https://www.ebi.ac.uk/QuickGO/term/GO:0007189>)
GO:2000252 : negative regulation of feeding behavior
(<https://www.ebi.ac.uk/QuickGO/term/GO:2000252>)
GO:0097009 : energy homeostasis (<https://www.ebi.ac.uk/QuickGO/term/GO:0097009>)
GO:0006112 : energy reserve metabolic process
(<https://www.ebi.ac.uk/QuickGO/term/GO:0006112>)

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 (<a +no+"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Presumptive Null=">https://www.gephebase.org/search-criteria?/and+Presumptive Null="+No+"#gephebase-summary-title)	Molecular Type
Gene Amplification (<a +gene="" amplification+"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Molecular Type=">https://www.gephebase.org/search-criteria?/and+Molecular Type="+Gene Amplification+"#gephebase-summary-title)	Aberration Type
Complex Change (<a +complex="" change+"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Aberration Type=">https://www.gephebase.org/search-criteria?/and+Aberration Type="+Complex Change+"#gephebase-summary-title)	Molecular Details of the Mutation
unknown in the species used in crosses but a strong corelation of gene copy number is found with the phenotype in two closely related species	Experimental Evidence
Linkage Mapping (<a +linkage="" href="https://www.gephebase.org/search-criteria?/and+Experimental Evidence=" mapping+"#gephebase-summary-title"="">https://www.gephebase.org/search-criteria?/and+Experimental Evidence="+Linkage Mapping+"#gephebase-summary-title)	Main Reference
Determination of onset of sexual maturation and mating behavior by melanocortin receptor 4 polymorphisms. (2010) (https://pubmed.ncbi.nlm.nih.gov/20869245)	Authors
Lampert KP; Schmidt C; Fischer P; Volff JN; Hoffmann C; Muck J; Lohse MJ; Ryan MJ; Schartl M	Abstract

Polymorphisms in reproductive strategies are among the most extreme and complex in nature. A prominent example is male body size and the correlated reproductive strategies in some species of platyfish and swordtails of the genus *Xiphophorus*. This polymorphism is controlled by a single Mendelian locus (P) that determines the onset of sexual maturity of males. Because males cease growth after reaching puberty, this results in a marked size polymorphism. The different male size classes show pronounced behavioral differences (e.g., courtship versus sneak mating), and females prefer large over small males. We show that sequence polymorphisms of the melanocortin receptor 4 gene (*mc4r*) comprise both functional and non-signal-transducing versions and that variation in copy number of *mc4r* genes on the Y chromosome underlies the P locus polymorphism. Nonfunctional Y-linked *mc4r* copies in larger males act as dominant-negative mutations and delay the onset of puberty. Copy number variation, as a regulating mechanism, endows this system with extreme genetic flexibility that generates extreme variation in phenotype. Because *Mc4r* is critically involved in regulation of body weight and appetite, a novel link between the physiological system controlling energy balance and the regulation of reproduction becomes apparent.

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Additional References

RELATED GEPHE

	Related Genes
No matches found.	
	Related Haplotypes
No matches found.	

EXTERNAL LINKS

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