

GEPHE SUMMARY

Gephebase Gene		GepheID
fog-2 (https://www.gephebase.org/search-criteria?/and+Gene+Gephebase=~fog-2^#gephebase-summary-title)	GP00000349	
Entry Status	Martin	Main curator
Published		

PHENOTYPIC CHANGE

Trait Category			
Physiology (https://www.gephebase.org/search-criteria?/and+Trait+Category=~Physiology^#gephebase-summary-title)		Trait	
Fertility (self-fertility; hermaphrodite spermatogenesis) (https://www.gephebase.org/search-criteria?/and+Trait=~Fertility+hermaphrodite+spermatogenesis^#gephebase-summary-title)		Trait State in Taxon A	
Caenorhabditis briggsae		Trait State in Taxon B	
Caenorhabditis elegans		Ancestral State	
Data not curated		Taxonomic Status	
Interspecific (https://www.gephebase.org/search-criteria?/and+Taxonomic+Status=~Interspecific^#gephebase-summary-title)			
Taxon A		Taxon B	
Caenorhabditis briggsae		Caenorhabditis elegans	
(https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=~Caenorhabditis+briggsae^#gephebase-summary-title)		(https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=~Caenorhabditis+elegans^#gephebase-summary-title)	
Common Name		Common Name	
-		-	
Synonyms		Synonyms	
Caenorhabditis briggsae Dougherty & Nigon, 1949		roundworm; Rhabditis elegans; Caenorhabditis elegans (Maupas, 1900); Rhabditis elegans Maupas, 1900	
Rank		Rank	
species		species	
Lineage		Lineage	
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Nematoda; Chromadorea; Rhabditida; Rhabditina; Rhabditomorpha; Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis		cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Nematoda; Chromadorea; Rhabditida; Rhabditina; Rhabditomorpha; Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis	
Parent		Parent	
Caenorhabditis () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=6237)		Caenorhabditis () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=6237)	
NCBI Taxonomy ID		NCBI Taxonomy ID	
6238		6239	
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=6238)		(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=6239)	
is Taxon A an Intraspecies?		is Taxon B an Intraspecies?	
No		No	

GENOTYPIC CHANGE

Generic Gene Name		UniProtKB Caenorhabditis elegans
fog-2		Q2YS43 (http://www.uniprot.org/uniprot/Q2YS43)
Synonyms		GenebankID or UniProtKB
CELE_Y113G7B.5; Y113G7B.5		BX284605 (https://www.ncbi.nlm.nih.gov/nucleotide/BX284605)
String		
6239.Y113G7B.5b		
(http://string-db.org/newstring.cgi/show_network_section.pl?identifier=6239.Y113G7B.5b)		
Sequence Similarities		
-		
GO - Molecular Function		
GO:0030371 : translation repressor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0030371)		
GO - Biological Process		
GO:0007283 : spermatogenesis (https://www.ebi.ac.uk/QuickGO/term/GO:0007283)		
GO:0017148 : negative regulation of translation (https://www.ebi.ac.uk/QuickGO/term/GO:0017148)		
GO:0040021 : hermaphrodite germ-line sex determination		

(<https://www.ebi.ac.uk/QuickGO/term/GO:0040021>)
GO:1900195 : positive regulation of oocyte maturation
(<https://www.ebi.ac.uk/QuickGO/term/GO:1900195>)
GO:2000738 : positive regulation of stem cell differentiation
(<https://www.ebi.ac.uk/QuickGO/term/GO:2000738>)
GO:2000648 : positive regulation of stem cell proliferation
(<https://www.ebi.ac.uk/QuickGO/term/GO:2000648>)

GO - Cellular Component

GO:0005737 : cytoplasm (<https://www.ebi.ac.uk/QuickGO/term/GO:0005737>)

Presumptive Null

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

Molecular Type

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

Aberration Type

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

Molecular Details of the Mutation

Gene birth by duplication of an ancestral gene ; FOG-2 binds the translational regulator GLD-1 and promotes spermatogenesis ; see also tra-2 entry and associated references

Experimental Evidence

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

Main Reference

fog-2 and the evolution of self-fertile hermaphroditism in *Caenorhabditis*. (2005) (<https://pubmed.ncbi.nlm.nih.gov/15630478>)

Authors

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Abstract

Somatic and germline sex determination pathways have diverged significantly in animals, making comparisons between taxa difficult. To overcome this difficulty, we compared the genes in the germline sex determination pathways of *Caenorhabditis elegans* and *C. briggsae*, two *Caenorhabditis* species with similar reproductive systems and sequenced genomes. We demonstrate that *C. briggsae* has orthologs of all known *C. elegans* sex determination genes with one exception: fog-2. Hermaphroditic nematodes are essentially females that produce sperm early in life, which they use for self fertilization. In *C. elegans*, this brief period of spermatogenesis requires FOG-2 and the RNA-binding protein GLD-1, which together repress translation of the tra-2 mRNA. FOG-2 is part of a large *C. elegans* FOG-2-related protein family defined by the presence of an F-box and Duf38/FOG-2 homogy domain. A fog-2-related gene family is also present in *C. briggsae*, however, the branch containing fog-2 appears to have arisen relatively recently in *C. elegans*, post-speciation. The C-terminus of FOG-2 is rapidly evolving, is required for GLD-1 interaction, and is likely critical for the role of FOG-2 in sex determination. In addition, *C. briggsae* gld-1 appears to play the opposite role in sex determination (promoting the female fate) while maintaining conserved roles in meiotic progression during oogenesis. Our data indicate that the regulation of the hermaphrodite germline sex determination pathway at the level of FOG-2/GLD-1/tra-2 mRNA is fundamentally different between *C. elegans* and *C. briggsae*, providing functional evidence in support of the independent evolution of self-fertile hermaphroditism. We speculate on the convergent evolution of hermaphroditism in *Caenorhabditis* based on the plasticity of the *C. elegans* germline sex determination cascade, in which multiple mutant paths yield self fertility.

Additional References

RELATED GEPHE

Related Genes

1 (tra-2) (<https://www.gephebase.org/search-criteria?/or+Taxon ID=^6238^/and+Trait=Fertility/or+Taxon ID=^6239^/and+Trait=Fertility/and+groupHaplotypes=true#gephebase-summary-title>)

Related Haplotypes

No matches found.

EXTERNAL LINKS

COMMENTS