

GEPHE SUMMARY

	Gephebase Gene	GepheID
follistatin (https://www.gephebase.org/search-criteria/?and+Gene Gephebase=%22follistatin%22#gephebase-summary-title)	GP00002138	
	Entry Status	Main curator
Published	Martin	

PHENOTYPIC CHANGE

Trait #1	Trait Category	Trait
Morphology (https://www.gephebase.org/search-criteria/?and+Trait+Category=%22Morphology%22#gephebase-summary-title)		
Limb morphology (wing dimorphism) (https://www.gephebase.org/search-criteria/?and+Trait=%22Limb+morphology+(wing+dimorphism)%22#gephebase-summary-title)	Trait	
winged males (aphicarus allele)	Trait State in Taxon A	
wingless males	Trait State in Taxon B	

Trait #2	Trait Category	Trait
Behavior (https://www.gephebase.org/search-criteria/?and+Trait+Category=%22Behavior%22#gephebase-summary-title)		
Flight behavior (wing dimorphism) (https://www.gephebase.org/search-criteria/?and+Trait=%22Flight+behavior+(wing+dimorphism)%22#gephebase-summary-title)	Trait	
winged males (aphicarus allele); more active	Trait State in Taxon A	
wingless males; less active	Trait State in Taxon B	

Taxon A	Ancestral State	Taxonomic Status	
Intraspecific (https://www.gephebase.org/search-criteria/?and+Taxonomic+Status=%22Intraspecific%22#gephebase-summary-title)			
Taxon A	Latin Name	Taxon B	Latin Name
Acyrthosiphon pisum (https://www.gephebase.org/search-criteria/?and+Taxon+and+Synonyms=%22Acyrthosiphon+pisum%22#gephebase-summary-title)		Acyrthosiphon pisum (https://www.gephebase.org/search-criteria/?and+Taxon+and+Synonyms=%22Acyrthosiphon+pisum%22#gephebase-summary-title)	
pea aphid	Common Name	pea aphid	Common Name
Acyrthosiphum pisum; pea aphid; Acyrthosiphon pisum (Harris, 1776); Acyrthosiphum pisum species	Rank	Acyrthosiphum pisum; pea aphid; Acyrthosiphon pisum (Harris, 1776); Acyrthosiphum pisum species	Rank
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Panarthropoda; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Dicondylia; Pterygota; Neoptera; Paraneoptera; Hemiptera; Sternorrhyncha; Aphidomorpha; Aphidoidea; Aphididae; Aphidinae; Macrosiphini; Acyrthosiphon	Lineage	cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Panarthropoda; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Dicondylia; Pterygota; Neoptera; Paraneoptera; Hemiptera; Sternorrhyncha; Aphidomorpha; Aphidoidea; Aphididae; Aphidinae; Macrosiphini; Acyrthosiphon	Lineage
Acyrthosiphon () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 7028)	Parent	Acyrthosiphon () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 7028)	Parent
7029 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 7029)	NCBI Taxonomy ID	7029 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 7029)	NCBI Taxonomy ID
No	is Taxon A an Infraspecies?	No	is Taxon B an Infraspecies?

GENOTYPIC CHANGE

Fs	Generic Gene Name CG12955; CG12956; CG33466; dFol1; dfs; dFS; Dmel\CG33466; Fol1; fs; Dmel_CG33466	UniProtKB Drosophila melanogaster Q86NV3 (http://www.uniprot.org/uniprot/Q86NV3)
	Synonyms 7227.FBpp0089409 (http://string-db.org/newstring_cgi/show_network_section.pl?identifier=7227.FBpp0089409)	GenebankID or UniProtKB 0
	String	
	Sequence Similarities	
-		
	GO - Molecular Function GO:0048185 : activin binding (https://www.ebi.ac.uk/QuickGO/term/GO:0048185)	
	GO - Biological Process GO:0007275 : multicellular organism development (https://www.ebi.ac.uk/QuickGO/term/GO:0007275)	
	GO:0030154 : cell differentiation (https://www.ebi.ac.uk/QuickGO/term/GO:0030154)	
	GO:0030510 : regulation of BMP signaling pathway (https://www.ebi.ac.uk/QuickGO/term/GO:0030510)	
	GO:0030514 : negative regulation of BMP signaling pathway (https://www.ebi.ac.uk/QuickGO/term/GO:0030514)	
	GO:0032926 : negative regulation of activin receptor signaling pathway (https://www.ebi.ac.uk/QuickGO/term/GO:0032926)	
	GO:0032927 : positive regulation of activin receptor signaling pathway (https://www.ebi.ac.uk/QuickGO/term/GO:0032927)	
	GO - Cellular Component GO:0005576 : extracellular region (https://www.ebi.ac.uk/QuickGO/term/GO:0005576)	
	GO:0005615 : extracellular space (https://www.ebi.ac.uk/QuickGO/term/GO:0005615)	
	No (https://www.gephebase.org/search-criteria/?and+Presumptive+Null^No^#gephebase-summary-title)	Presumptive Null
	Gene Amplification (https://www.gephebase.org/search-criteria/?and+Molecular+Type^Gene+Amplification^#gephebase-summary-title)	Molecular Type
	Insertion (https://www.gephebase.org/search-criteria/?and+Aberration+Type^Insertion^#gephebase-summary-title)	Aberration Type
-		Insertion Size
	the api allele from winged males is a 120kb insertion that includes a copy of the follistatin gene	Molecular Details of the Mutation
	Linkage Mapping (https://www.gephebase.org/search-criteria/?and+Experimental+Evidence^Linkage+Mapping^#gephebase-summary-title)	Experimental Evidence
	A large genomic insertion containing a duplicated follistatin gene is linked to the pea aphid male wing dimorphism. (2020) (https://pubmed.ncbi.nlm.nih.gov/32141813)	Main Reference
	Li B; Bickel RD; Parker BJ; Saleh Ziabari O; Liu F; Vellichirammal NN; Simon JC; Stern DL; Brisson JA	Authors
	Wing dimorphisms have long served as models for examining the ecological and evolutionary tradeoffs associated with alternative phenotypes. Here, we investigated the genetic cause of the pea aphid (<i>Acyrtosiphon pisum</i>) male wing dimorphism, wherein males exhibit one of two morphologies that differ in correlated traits that include the presence or absence of wings. We mapped this trait difference to a single genomic region and, using third generation, long-read sequencing, we identified a 120 kb insertion in the wingless allele. This insertion includes a duplicated follistatin gene, which is a strong candidate gene in the minimal mapped interval to cause the dimorphism. We found that both alleles were present prior to pea aphid biotype lineage diversification, we estimated that the insertion occurred millions of years ago, and we propose that both alleles have been maintained in the species, likely due to balancing selection.	Abstract
	Â© 2020, Li et al.	Additional References

RELATED GEPHE

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

EXTERNAL LINKS

COMMENTS

@GeneDuplication @BalancingSelection ; ancient polymorphism in the pea aphid lineage

