

GEPHE SUMMARY

	Gephebase Gene	GephelD
esterase isozyme E3 (https://www.gephebase.org/search-criteria/?and+Gene Gephebase=^esterase isozyme E3^#gephebase-summary-title)	GP00000294	Main curator
	Entry Status	Martin
Published		

PHENOTYPIC CHANGE

	Trait Category	
Physiology (https://www.gephebase.org/search-criteria/?and+Trait Category=^Physiology^#gephebase-summary-title)	Trait	
Xenobiotic resistance (insecticide) (https://www.gephebase.org/search-criteria/?and+Trait=^Xenobiotic+resistance+(insecticide)^#gephebase-summary-title)	Trait State in Taxon A	
Lucilia cuprina	Trait State in Taxon B	
Lucilia cuprina	Ancestral State	
Data not curated	Taxonomic Status	
Intraspecific (https://www.gephebase.org/search-criteria/?and+Taxonomic Status=^Intraspecific^#gephebase-summary-title)		
Taxon A		Taxon B
Lucilia cuprina (https://www.gephebase.org/search-criteria/?and+Taxon+and+Synonyms=^Lucilia+cuprina^#gephebase-summary-title)	Latin Name	Latin Name
Australian sheep blowfly	Common Name	Common Name
Australian sheep blowfly; greenbottle fly; Lucilia cuprina (Wiedemann, 1830)	Synonyms	Synonyms
species	Rank	Rank
	Lineage	Lineage
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Panarthropoda; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Dicondylia; Pterygota; Neoptera; Holometabola; Diptera; Brachycera; Muscomorpha; Eremoneura; Cyclorrhapha; Schizophora; Calyptratae; Oestroidea; Calliphoridae; Luciliinae; Lucilia		
Lucilia () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 7374)	Parent	Parent
7375 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 7375)	NCBI Taxonomy ID	NCBI Taxonomy ID
No	is Taxon A an Infraspecies?	is Taxon B an Infraspecies?
	No	

GENOTYPIC CHANGE

LcaE7	Generic Gene Name	UniProtKB Lucilia cuprina
-	Synonyms	GenebankID or UniProtKB
-	String	
	Sequence Similarities	
Belongs to the type-B carboxylesterase/lipase family.		
	GO - Molecular Function	
GO:0016787 : hydrolase activity (https://www.ebi.ac.uk/QuickGO/term/GO:0016787)		
	GO - Biological Process	
	GO - Cellular Component	
		Presumptive Null
No (https://www.gephebase.org/search-criteria/?and+Presumptive+Null=^No^#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%20Type=%Coding%23gephebase-summary-title))

Aberration Type

SNP ([https://www.gephebase.org/search-criteria/?and+Aberration Type=%SNP%#gephebase-summary-title](https://www.gephebase.org/search-criteria/?and+Aberration%20Type=%SNP%23gephebase-summary-title))

SNP Coding Change

Nonsynonymous

Molecular Details of the Mutation

Gly137Asp

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%20Evidence=%Candidate%20Gene%23gephebase-summary-title))

	Taxon A	Taxon B	Position
Codon	-	-	-
Amino-acid	-	-	-

Main Reference

A single amino acid substitution converts a carboxylesterase to an organophosphorus hydrolase and confers insecticide resistance on a blowfly. (1997)
(<https://pubmed.ncbi.nlm.nih.gov/9207114>)

Authors

Newcomb RD; Campbell PM; Ollis DL; Cheah E; Russell RJ; Oakeshott JG

Abstract

Resistance to organophosphorus (OP) insecticides is associated with decreased carboxylesterase activity in several insect species. It has been proposed that the resistance may be the result of a mutation in a carboxylesterase that simultaneously reduces its carboxylesterase activity and confers an OP hydrolase activity (the "mutant ali-esterase hypothesis"). In the sheep blowfly, *Lucilia cuprina*, the association is due to a change in a specific esterase isozyme, E3, which, in resistant flies, has a null phenotype on gels stained using standard carboxylesterase substrates. Here we show that an OP-resistant allele of the gene that encodes E3 differs at five amino acid replacement sites from a previously described OP-susceptible allele. Knowledge of the structure of a related enzyme (acetylcholinesterase) suggests that one of these substitutions (Gly137 --> Asp) lies within the active site of the enzyme. The occurrence of this substitution is completely correlated with resistance across 15 isogenic strains. In vitro expression of two natural and two synthetic chimeric alleles shows that the Asp137 substitution alone is responsible for both the loss of E3's carboxylesterase activity and the acquisition of a novel OP hydrolase activity. Modeling of Asp137 in the homologous position in acetylcholinesterase suggests that Asp137 may act as a base to orientate a water molecule in the appropriate position for hydrolysis of the phosphorylated enzyme intermediate.

Additional References

RELATED GEPHE

Related Genes

No matches found.

Related Haplotypes

1 ([https://www.gephebase.org/search-criteria/?or+Gene Gephebase=%esterase isozyme E3% and+Taxon ID=%7375% or+Gene Gephebase=%esterase isozyme E3% and+Taxon ID=%7375%#gephebase-summary-title](https://www.gephebase.org/search-criteria/?or+Gene%20Gephebase=%esterase%20isozyme%20E3%20%26%20Taxon%20ID=%7375%20%26%20Gene%20Gephebase=%esterase%20isozyme%20E3%20%26%20Taxon%20ID=%7375%23gephebase-summary-title))

EXTERNAL LINKS

COMMENTS