## GEPHE SUMMARY

Gephebase Gene Acetylcholinesterase (Ace-1) (https://www.gephebase.org/search-criteria?/and+Gene Gephebase=^Acetylcholinesterase (Ace-1)^#gephebase-summary-title) Entry Status Published	GP0000036 Martin
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	
Aphis gossypii	
Trait State in Taxon B Aphis gossypii	
Ancestral State	
Taxonomic Status	
Intraspecific (https://www.gephebase.org/search-criteria?/and+Taxonomic Status=^Intraspecific^#gephebase-summary-title)	
Taxon A	
Latin Name	
Aphis gossypii	Aphis gossypii
(https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms=^Aphis	(https://www.gephebase.org
gossypii^#gephebase-summary-title)	gossypii^#gephebase-summ
Common Name	
cotton aphid Synonyms	cotton aphid
cotton aphid; melon aphid; Aphis gossypii Glover, 1877; Aphis gossyppi Rank	cotton aphid; melon aphid; /
species	species

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; Aphidini; Aphis; Aphis Parent

Aphis () - (Rank: subgenus)	
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwta	ıx.cgi?id= 464929 )
	NCBI Taxonomy ID
80765	
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwta	ıx.cgi?id= 80765 )
	is Taxon A an Infraspecies?
No	

# GENOTYPIC CHANGE

Generic Gene Na	me
Ace	
Synony AcChE: ace: ACE: ace-2: ache: AchE: AChE: CG17907: CHE: dAChE: dmAChE:	ms
DmAChE; Dmel\CG17907; Dm_ace; FBgn0000024; I(3)26; I(3)87Ed	
Stri	ng
7227.FBpp0289713	
(http://string-db.org/newstring_cgi/show_network_section.pl?identifier= 7227.FBpp02897	13
)	
Sequence Similarit	ies
Belongs to the type-B carboxylesterase/lipase family.	
GO - Molecular Functi	on
GO:0042803 : protein homodimerization activity	
(https://www.ebi.ac.uk/QuickGO/term/GO:0042803)	
GO:0003990 : acetylcholinesterase activity	
(https://www.ebi.ac.uk/QuickGO/term/GO:0003990)	
GO:0004104 : cholinesterase activity (https://www.ebi.ac.uk/QuickGO/term/GO:000410	)4)
GO:0043199 : sulfate binding (https://www.ebi.ac.uk/QuickGO/term/GO:0043199)	

00036			

Taxon B

Aprils gossypii
(https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms=^Aphis
gossypii^#gephebase-summary-title)
5 , 7 5 7 7 7
Common Name
cotton aphid
Synonyms
cotton aphid; melon aphid; Aphis gossypii Glover, 1877; Aphis gossyppi
Rank
species
Lineage
5
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; Aphidini; Aphis; Aphis
Parent
Aphis () - (Rank: subgenus)
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 464929 )
NCBI Taxonomy ID
80765
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 80765 )
is Taxon B an Infraspecies?
No

P07140 (http://www.uniprot.org/uniprot/P07140)

UniProtKB Drosophila melanogaster

GenebankID or UniProtKB

ALE67001 (https://www.ncbi.nlm.nih.gov/nuccore/ALE67001)

Main curator

Latin Name

GO - Biological Process
GO:0006581 : acetylcholine catabolic process
(https://www.ebi.ac.uk/QuickGO/term/GO:0006581)
GO:0001507 : acetylcholine catabolic process in synaptic cleft
(https://www.ebi.ac.uk/QuickGO/term/GO:0001507)
GO:0007268 : chemical synaptic transmission
(https://www.ebi.ac.uk/QuickGO/term/GO:0007268)
GO:0042426 : choline catabolic process
(https://www.ebi.ac.uk/QuickGO/term/GO:0042426)
GO:0042331 : phototaxis (https://www.ebi.ac.uk/QuickGO/term/GO:0042331)
GO - Cellular Component
GO:0005886 : plasma membrane (https://www.ebi.ac.uk/QuickGO/term/GO:0005886)
GO:0005737 : cytoplasm (https://www.ebi.ac.uk/QuickGO/term/GO:0005737)
GO:0031225 : anchored component of membrane
(https://www.ebi.ac.uk/QuickGO/term/GO:0031225)
GO:0030054 : cell junction (https://www.ebi.ac.uk/QuickGO/term/GO:0030054)
GO:0043083 : synaptic cleft (https://www.ebi.ac.uk/QuickGO/term/GO:0043083)

Mutation #1	
	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)	Molecular Type
	Aberration Type
Insertion (https://www.gephebase.org/search-criteria?/and+Aberration Type=^Insertion^#gephebase-summary-title)	
	Insertion Size
1-10 kb	Molecular Details of the Mutation
Duplications; A302S and S431F = S331F	Molecular Details of the Matation
	Experimental Evidence
Candidate Gene (https://www.gephebase.org/search-criteria?/and+Experimental Evidence=^Candidate Gene^#gephebase-summary-title)	
Extensive Ace2 duplication and multiple mutations on Ace1 and Ace2 are related with high level of organophosphates resistance in Aphis gossypii.	Main Reference (2014)
(https://pubmed.ncbi.nlm.nih.gov/22489048)	
Shang Q; Pan Y; Fang K; Xi J; Wong A; Brennan JA; Cao C	Authors
Shang Q; Pan T; Pang R; Al J; Wong A; Drennan JA; Cao C	Abstract
Aphis gossypii (Glover) has been found to possess multiple mutations in the acetylcholinesterase (AChE) gene (Ace) that might involve target site	
expression of AChEs reveals that the resistant Ace1 (Ace1R) and Ace2 (Ace2R) were significantly less inhibited by eserine, omethoate, and malaox	con than the susceptible Ace1 (Ace1S)
and Ace2 (Ace2S). Furthermore, in both the mutant and susceptible AChEs, Ace2 was significantly less sensitive to eserine, omethoate, and malao	
suggested that both the mutant Ace1 and Ace2 were responsible for omethoate resistance, while the mutant Ace2 played a major role in insecticide and transcription level of Ace2 were 1.52- and 1.88-fold higher in the ORR strain than in the OSS strain. Furthermore, the DNA copy number and t	
significantly higher than that of Ace1 in either OSS or ORR strains, demonstrating the involvement of Ace2 gene duplication in resistance. Thus, the	
resistance in cotton aphids appears to have evolved through a combination of multiple mutations and extensive Ace2R gene duplication.	
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Additional References Identification of mutations conferring insecticide-insensitive AChE in the cotton-melon aphid, Aphis gossypii Glover. (2004) (https://pubmed.ncbi.nlm.nih.gov/15373812) Biochemical evidence that an S431F mutation in acetylcholinesterase-1 of Aphis gossypii mediates resistance to pirimicarb and omethoate. (2004) (https://pubmed.ncbi.nlm.nih.gov/15532677)

Two amino acid substitutions in acetylcholinesterase associated with pirimicarb and organophosphorous insecticide resistance in the cotton aphid, Aphis gossypii Glover (Homoptera: Aphididae). (2004) (https://pubmed.ncbi.nlm.nih.gov/15373811)

Mutation #2					
			Presumptive Null		
No (https://www.gephebase.org/search-criteria?/and+Presumptive	• Null=^No^#gephebase-su	mmary-title)	Molecular Type		
Coding (https://www.gephebase.org/search-criteria?/and+Molecu	lar Type=^Coding^#gephel	base-summary-title)			
SNP (https://www.gephebase.org/search-criteria?/and+Aberration	Aberration Type				
Nonsynonymous			SNP Coding Change		
			Molecular Details of the Mutation		
Duplications; A302S and S431F			Experimental Evidence		
Candidate Gene (https://www.gephebase.org/search-criteria?/and+Experimental Evidence=^Candidate Gene^#gephebase-summary-title)					
	Taxon A	Taxon B	Position		
Codon	-	-	-		
Amino-acid	Ala	Ser	302		

### Shang Q; Pan Y; Fang K; Xi J; Wong A; Brennan JA; Cao C

Additional References

Abstract Aphis gossypii (Glover) has been found to possess multiple mutations in the acetylcholinesterase (AChE) gene (Ace) that might involve target site insensitivity. In vitro functional expression of AChEs reveals that the resistant Ace1 (Ace1R) and Ace2 (Ace2R) were significantly less inhibited by eserine, omethoate, and malaoxon than the susceptible Ace1 (Ace1S) and Ace2 (Ace2S). Furthermore, in both the mutant and susceptible AChEs, Ace2 was significantly less sensitive to eserine, omethoate, and malaoxon than Ace1. These results suggested that both the mutant Ace1 and Ace2 were responsible for omethoate resistance, while the mutant Ace2 played a major role in insecticide resistance. The DNA copy number and transcription level of Ace2 were 1.52- and 1.88-fold higher in the ORR strain than in the OSS strain. Furthermore, the DNA copy number and transcription level of Ace1 in either OSS or ORR strains, demonstrating the involvement of Ace2 gene duplication in resistance. Thus, the authors conclude that omethoate resistance in cotton aphids appears to have evolved through a combination of multiple mutations and extensive Ace2R gene duplication.

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ldentification of mutations conferring insecticide-insensitive AChE in the cotton-melon aphid, Aphis gossypii Glover. (2004) (https://pubmed.ncbi.nlm.nih.gov/15373812) Insecticide resistance traits differ among and within host races in Aphis gossypii. (2010) (https://pubmed.ncbi.nlm.nih.gov/19908228)

Mutation #3			
			Presumptive Null
No (https://www.gephebase.org/search-criteria?/and+Presum	otive Null=^No^#gephebase-summa	ry-title)	Molecular Type
Coding (https://www.gephebase.org/search-criteria?/and+Mo	lecular Type=^Coding^#gephebase-	summary-title)	··· · · · · · · · · · · · · · · · · ·
SNP (https://www.gephebase.org/search-criteria?/and+Aberra	ation Type=^SNP^#gephebase-sumr	nary-title)	Aberration Type
N			SNP Coding Change
Nonsynonymous			Molecular Details of the Mutation
Duplications; A302S and S431F			Experimental Evidence
Candidate Gene (https://www.gephebase.org/search-criteria?	/and+Experimental Evidence=^Cand	idate Gene^#gephebase-summary-title)	
	Taxon A	Taxon B	Position

	Taxon A	Taxon B	Position	
Codon	-	-	-	
Amino-acid	Ser	Phe	431	

Extensive Ace2 duplication and multiple mutations on Ace1 and Ace2 are related with high level of organophosphates resistance in Aphis gossypii. (2014) (https://pubmed.ncbi.nlm.nih.gov/22489048)

#### Shang Q; Pan Y; Fang K; Xi J; Wong A; Brennan JA; Cao C

Aphis gossypii (Glover) has been found to possess multiple mutations in the acetylcholinesterase (AChE) gene (Ace) that might involve target site insensitivity. In vitro functional expression of AChEs reveals that the resistant Ace1 (Ace1R) and Ace2 (Ace2R) were significantly less inhibited by eserine, omethoate, and malaoxon than the susceptible Ace1 (Ace1S) and Ace2 (Ace2S). Furthermore, in both the mutant and susceptible AChEs, Ace2 was significantly less sensitive to eserine, omethoate, and malaoxon than Ace1. These results suggested that both the mutant Ace1 and Ace2 were responsible for omethoate resistance, while the mutant Ace2 played a major role in insecticide resistance. The DNA copy number and transcription level of Ace2 were 1.52- and 1.88-fold higher in the ORR strain than in the OSS strain. Furthermore, the DNA copy number and transcription level of Ace1 in either OSS or ORR strains, demonstrating the involvement of Ace2 gene duplication in resistance. Thus, the authors conclude that omethoate resistance in cotton aphids appears to have evolved through a combination of multiple mutations and extensive Ace2R gene duplication.

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Additional References Identification of mutations conferring insecticide-insensitive AChE in the cotton-melon aphid, Aphis gossypii Glover. (2004) (https://pubmed.ncbi.nlm.nih.gov/15373812) Insecticide resistance traits differ among and within host races in Aphis gossypii. (2010) (https://pubmed.ncbi.nlm.nih.gov/19908228)

## RELATED GEPHE

**Related** Genes

Main Reference

Authors

Abstract

3 (Acetylcholinesterase (Ace-2), nAChR, para (kdr)) (https://www.gephebase.org/search-criteria?/or+Taxon ID=^80765^/and+Trait=Xenobiotic resistance/and+groupHaplotypes=true#gephebase-summary-title)

Related Haplotypes

1 (https://www.gephebase.org/search-criteria?/or+Gene Gephebase=^Acetylcholinesterase (Ace-1)^/and+Taxon ID=^80765^/or+Gene Gephebase=^Acetylcholinesterase (Ace-1)^/and+Taxon ID=^80765^#gephebase-summary-title)

# COMMENTS

@SeveralMutationsWithEffect