

## GEPHE SUMMARY

**Gephebase Gene**  
Acetylcholinesterase (Ace)

**Entry Status**  
Published

**GepheID**  
GP00002014

**Main curator**  
Courtier

## PHENOTYPIC CHANGE

**Trait Category**  
Physiology

**Trait**  
Xenobiotic resistance (insecticide)

**Trait State in Taxon A**  
Leptinotarsa decemlineata - sensitive

**Trait State in Taxon B**  
Leptinotarsa decemlineata - resistant

**Ancestral State**  
Taxon A

**Taxonomic Status**  
Intraspecific

### Taxon A

**Latin Name**  
*Leptinotarsa decemlineata*

**Common Name**  
Colorado potato beetle

**Synonyms**  
Leptinotarsa decimlineata; Stilodes decemlineata; Colorado potato beetle; Leptinotarsa decemlineata (Say, 1824)

**Rank**  
species

**Lineage**  
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Panarthropoda; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Dicondylia; Pterygota; Neoptera; Holometabola; Coleoptera; Polyphaga; Cucujiformia; Chrysomeloidea; Chrysomelidae; Chrysomelinae; Doryphorini; Leptinotarsa

**Parent**  
Leptinotarsa () - (Rank: genus)

**NCBI Taxonomy ID**  
7539

**is Taxon A an Intraspecies?**  
No

### Taxon B

**Latin Name**  
*Leptinotarsa decemlineata*

**Common Name**  
Colorado potato beetle

**Synonyms**  
Leptinotarsa decimlineata; Stilodes decemlineata; Colorado potato beetle; Leptinotarsa decemlineata (Say, 1824)

**Rank**  
species

**Lineage**  
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Protostomia; Ecdysozoa; Panarthropoda; Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Dicondylia; Pterygota; Neoptera; Holometabola; Coleoptera; Polyphaga; Cucujiformia; Chrysomeloidea; Chrysomelidae; Chrysomelinae; Doryphorini; Leptinotarsa

**Parent**  
Leptinotarsa () - (Rank: genus)

**NCBI Taxonomy ID**  
7539

**is Taxon B an Intraspecies?**  
No

## GENOTYPIC CHANGE

**Generic Gene Name**  
Ace

**Synonyms**  
AcChE; ace; ACE; ace-2; ache; AchE; AChE; CG17907; CHE; dAChE; dmAChE; DmAChE; Dmel\CG17907; Dm\_ace; FBgn0000024; l(3)26; l(3)87Ed

**String**  
7227.FBpp0289713

**Sequence Similarities**  
Belongs to the type-B carboxylesterase/lipase family.

**GO - Molecular Function**  
GO:0042803 : protein homodimerization activity  
GO:0003990 : acetylcholinesterase activity  
GO:0004104 : cholinesterase activity  
GO:0043199 : sulfate binding

**GO - Biological Process**  
GO:0006581 : acetylcholine catabolic process  
GO:0001507 : acetylcholine catabolic process in synaptic cleft  
GO:0007268 : chemical synaptic transmission

**UniProtKB Drosophila melanogaster**  
P07140

**GenebankID or UniProtKB**

GO:0042426 : choline catabolic process

GO:0042331 : phototaxis

**GO - Cellular Component**

GO:0005886 : plasma membrane

GO:0005737 : cytoplasm

GO:0031225 : anchored component of membrane

GO:0030054 : cell junction

GO:0043083 : synaptic cleft

**Presumptive Null**

No

**Molecular Type**

Coding

**Aberration Type**

SNP

**SNP Coding Change**

Nonsynonymous

**Molecular Details of the Mutation**

Ser>Gly (238 is the corresponding position in Torpedo)

**Experimental Evidence**

Candidate Gene

	Taxon A	Taxon B	Position
Codon	-	-	-
Amino-acid	Ser	Gly	238

**Main Reference**

[A Point Mutation of Acetylcholinesterase Associated with Azinphosmethyl Resistance and Reduced Fitness in Colorado Potato Beetle. \(1996\)](#)

**Authors**

Zhu KY; Lee SH; Clark JM

**Abstract**

A serine to glycine point mutation of acetylcholinesterase (AChE, EC 1.1.1.7) was identified in an azinphosmethyl-resistant strain of Colorado potato beetle [*Leptinotarsa decemlineata* (Say)]. The position of the mutation corresponds to Val 238 of the Torpedo AChE and represents the first amino acid residue to form the alpha-helix, alpha-E'1. The predicted secondary structure of the mutation-containing region of AChE suggested that the transition from the turn to the alpha-helix occurs sooner in the sequence when serine is replaced by glycine. Thus, conformational changes in the AChE due to the alpha-helix deformation were expected to impinge upon both the catalytic and the peripheral binding sites, resulting in the modification of the bindings of organophosphorus insecticides and other ligands to these sites. The mutation appeared to be associated with the fitness of the beetle. The intrinsic rate of increase of the azinphosmethyl-resistant (AZ-R) strain was relatively low when the beetles were reared on the Russet Burbank potato cultivar, but was relatively high when they were reared on the NDA 1725-1 potato cultivar. Because these two potato cultivars contain different amounts of steroidal glycoalkaloids (e.g., alpha-solanine and alpha-chaconine), the different fitness of the AZ-R strain on different potato cultivars may be partially attributed to the increased sensitivity of the azinphosmethyl-resistant form of AChE to the inhibition by alpha-solanine and reduced sensitivity to alpha-chaconine as previously reported.

**Additional References**

**RELATED GEPHE**

**Related Genes**

1 ([para \(kdr\)](#))

**Related Haplotypes**

1

**EXTERNAL LINKS**

**COMMENTS**

