

## GEPHE SUMMARY

	Gephebase Gene	GephelD
EPSPS ( <a href="https://www.gephebase.org/search-criteria?/and+Gene">https://www.gephebase.org/search-criteria?/and+Gene</a> Gephebase=^EPSPS^#gephebase-summary-title)	GP00001888	Main curator
Published	Entry Status	Courtier

## PHENOTYPIC CHANGE

	Trait Category		
Physiology ( <a href="https://www.gephebase.org/search-criteria?/and+Trait">https://www.gephebase.org/search-criteria?/and+Trait</a> Category=^Physiology^#gephebase-summary-title)	Trait		
Xenobiotic resistance (herbicides; glyphosate) ( <a href="https://www.gephebase.org/search-criteria?/and+Trait=^Xenobiotic+resistance+(herbicides;+glyphosate)^#gephebase-summary-title">https://www.gephebase.org/search-criteria?/and+Trait=^Xenobiotic+resistance+(herbicides;+glyphosate)^#gephebase-summary-title</a> )			
	Trait State in Taxon A		
Eleusine indica - sensitive	Trait State in Taxon B		
Eleusine indica - resistant	Ancestral State		
Taxon A	Taxonomic Status		
Intraspecific ( <a href="https://www.gephebase.org/search-criteria?/and+Taxonomic">https://www.gephebase.org/search-criteria?/and+Taxonomic</a> Status=^Intraspecific^#gephebase-summary-title)			
	Taxon A	Taxon B	
	Latin Name	Latin Name	
Eleusine indica ( <a href="https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Eleusine+indica^#gephebase-summary-title">https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Eleusine+indica^#gephebase-summary-title</a> )	Eleusine indica ( <a href="https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Eleusine+indica^#gephebase-summary-title">https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Eleusine+indica^#gephebase-summary-title</a> )		
goosegrass	Common Name	Common Name	
goosegrass; crabgrass; fowl foot grass; wire grass; yardgrass; Eleusine indica (L.) Gaertn.	Synonyms	goosegrass; crabgrass; fowl foot grass; wire grass; yardgrass; Eleusine indica (L.) Gaertn.	
species	Rank	Rank	
	Lineage	Lineage	
cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliopsida; Mesangiospermae; Liliopsida; Petrosaviidae; commelinids; Poales; Poaceae; PACMAD clade; Chloridoideae; Cynodonteae; Eleusininae; Eleusine		cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliopsida; Mesangiospermae; Liliopsida; Petrosaviidae; commelinids; Poales; Poaceae; PACMAD clade; Chloridoideae; Cynodonteae; Eleusininae; Eleusine	
	Parent	Parent	
Eleusine () - (Rank: genus) ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 4510">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 4510</a> )	Eleusine () - (Rank: genus) ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 4510">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 4510</a> )		
29674 ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 29674">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 29674</a> )	NCBI Taxonomy ID	NCBI Taxonomy ID	
	is Taxon A an Infraspecies?	is Taxon B an Infraspecies?	
No		No	

## GENOTYPIC CHANGE

	Generic Gene Name	UniProtKB Arabidopsis thaliana
At2g45300	P05466 ( <a href="http://www.uniprot.org/uniprot/P05466">http://www.uniprot.org/uniprot/P05466</a> )	GenebankID or UniProtKB
F4L23.19; At2g45300	Synonyms	
3702.AT2G45300.1 ( <a href="http://string-db.org/newstring_cgi/show_network_section.pl?identifier= 3702.AT2G45300.1">http://string-db.org/newstring_cgi/show_network_section.pl?identifier= 3702.AT2G45300.1</a> )	String	
	Sequence Similarities	
Belongs to the EPSP synthase family.		
	GO - Molecular Function	
GO:0003866 : 3-phosphoshikimate 1-carboxyvinyltransferase activity ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0003866">https://www.ebi.ac.uk/QuickGO/term/GO:0003866</a> )		
	GO - Biological Process	
GO:0009073 : aromatic amino acid family biosynthetic process ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0009073">https://www.ebi.ac.uk/QuickGO/term/GO:0009073</a> )		
GO:0009423 : chorismate biosynthetic process		

GO - Cellular Component

GO:0009507 : chloroplast (<https://www.ebi.ac.uk/QuickGO/term/GO:0009507>)

GO:0009570 : chloroplast stroma (<https://www.ebi.ac.uk/QuickGO/term/GO:0009570>)

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>)

Aberration Type

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

SNP Coding Change

Nonsynonymous

Molecular Details of the Mutation

Pro106Ser leading to 2-4-fold resistance

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	Pro	Ser	106

Main Reference

Glyphosate-resistant goosegrass. Identification of a mutation in the target enzyme 5-enolpyruylshikimate-3-phosphate synthase. (2002) (<https://pubmed.ncbi.nlm.nih.gov/12114580>)

Authors

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Abstract

The spontaneous occurrence of resistance to the herbicide glyphosate in weed species has been an extremely infrequent event, despite over 20 years of extensive use. Recently, a glyphosate-resistant biotype of goosegrass (*Eleusine indica*) was identified in Malaysia exhibiting an LD<sub>50</sub> value approximately 2- to 4-fold greater than the sensitive biotype collected from the same region. A comparison of the inhibition of 5-enolpyruylshikimate-3-phosphate synthase (EPSPS) activity by glyphosate in extracts prepared from the resistant (R) and sensitive (S) biotypes revealed an approximately 5-fold higher IC<sub>50</sub>(glyphosate) for the (R) biotype. Sequence comparisons of the predicted EPSPS mature protein coding regions from both biotypes revealed four single-nucleotide differences, two of which result in amino acid changes. One of these changes, a proline to serine substitution at position 106 in the (R) biotype, corresponds to a substitution previously identified in a glyphosate-insensitive EPSPS enzyme from *Salmonella typhimurium*. Kinetic data generated for the recombinant enzymes suggests that the second substitution identified in the (R) EPSPS does not contribute significantly to its reduced glyphosate sensitivity. *Escherichia coli* aroA- (EPSPS deficient) strains expressing the mature EPSPS enzyme from the (R) biotype exhibited an approximately 3-fold increase in glyphosate tolerance relative to strains expressing the mature EPSPS from the (S) biotype. These results provide the first evidence for an altered EPSPS enzyme as an underlying component of evolved glyphosate resistance in any plant species.

Additional References

## RELATED GEPHE

Related Genes

No matches found.

Related Haplotypes

1 (<https://www.gephebase.org/search-criteria?/or+Gene+Gephebase=%EPSPS%/and+Taxon+ID=%29674%/or+Gene+Gephebase=%EPSPS%/and+Taxon+ID=%29674%#gephebase-summary-title>)

## EXTERNAL LINKS

## COMMENTS