

GEPHE SUMMARY

		Gephebase Gene		GepheID
MATE1/AltSB/SbMATE (https://www.gephebase.org/search-criteria?/and+Gene)			GP00000565	
Gephebase="MATE1/AltSB/SbMATE"#gephebase-summary-title)				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		
Metal tolerance (https://www.gephebase.org/search-criteria?/and+Trait="Metal)				
tolerance"#gephebase-summary-title)		Trait State in Taxon A		
Sorghum bicolor- aluminum-tolerant SC283		Trait State in Taxon B		
Sorghum bicolor - aluminum-sensitive BR007		Ancestral State		
Data not curated		Taxonomic Status		
Domesticated (https://www.gephebase.org/search-criteria?/and+Taxonomic)				
Status="Domesticated"#gephebase-summary-title)				
Taxon A		Taxon B		
	Latin Name		Latin Name	
Sorghum bicolor		Sorghum bicolor		
(https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms="Sorghum bicolor"#gephebase-summary-title)		(https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms="Sorghum bicolor"#gephebase-summary-title)		
	Common Name		Common Name	
sorghum		sorghum		
	Synonyms		Synonyms	
Andropogon sorghum; Sorghum bicolor subsp. bicolor; Sorghum nervosum; Sorghum saccharatum; Sorghum vulgare; sorghum; broomcorn; milo; Andropogon sorghum (L.) Brot.; Sorghum bicolor (L.) Moench; Sorghum nervosum Besser ex Schult.; Sorghum saccharatum (L.) Moench; Sorghum vulgare Pers.; Sorghum bicolor milo; Sorghum_bicolor		Andropogon sorghum; Sorghum bicolor subsp. bicolor; Sorghum nervosum; Sorghum saccharatum; Sorghum vulgare; sorghum; broomcorn; milo; Andropogon sorghum (L.) Brot.; Sorghum bicolor (L.) Moench; Sorghum nervosum Besser ex Schult.; Sorghum saccharatum (L.) Moench; Sorghum vulgare Pers.; Sorghum bicolor milo; Sorghum_bicolor		
	Rank		Rank	
species		species		
	Lineage		Lineage	
cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliopsida; Mesangiospermae; Liliopsida; Petrosaviidae; commelinids; Poales; Poaceae; PACMAD clade; Panicoideae; Andropogonodae; Andropogoneae; Sorghinae; Sorghum		cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliopsida; Mesangiospermae; Liliopsida; Petrosaviidae; commelinids; Poales; Poaceae; PACMAD clade; Panicoideae; Andropogonodae; Andropogoneae; Sorghinae; Sorghum		
	Parent		Parent	
Sorghum () - (Rank: genus)		Sorghum () - (Rank: genus)		
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4557)		(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4557)		
	NCBI Taxonomy ID		NCBI Taxonomy ID	
4558		4558		
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4558)		(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4558)		
	is Taxon A an Intraspecies?		is Taxon B an Intraspecies?	
Yes		Yes		
	Taxon A Description		Taxon B Description	
Sorghum bicolor- aluminum-tolerant SC283		Sorghum bicolor - aluminum-sensitive BR007		

GENOTYPIC CHANGE

	Generic Gene Name		UniProtKB
-		0	
	Synonyms		GenebankID or UniProtKB
-		ABS89149 (https://www.ncbi.nlm.nih.gov/nucleotide/ABS89149)	
	String		
-			
	Sequence Similarities		
-			
	GO - Molecular Function		
-			
	GO - Biological Process		
-			
	GO - Cellular Component		

Unknown (<https://www.gephebase.org/search-criteria?/and+Presumptive Null=^Unknown^#gephebase-summary-title>)

Presumptive Null

Cis-regulatory (<https://www.gephebase.org/search-criteria?/and+Molecular Type=^Cis-regulatory^#gephebase-summary-title>)

Molecular Type

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

Aberration Type

unknown

Molecular Details of the Mutation

Linkage Mapping (<https://www.gephebase.org/search-criteria?/and+Experimental Evidence=^Linkage Mapping^#gephebase-summary-title>)

Experimental Evidence

A gene in the multidrug and toxic compound extrusion (MATE) family confers aluminum tolerance in sorghum. (2007) (<https://pubmed.ncbi.nlm.nih.gov/17721535>)

Main Reference

Magalhaes JV; Liu J; Guimarães CT; Lana UG; Alves VM; Wang YH; Schaffert RE; Hoekenga OA; Piñeros MA; Shaff JE; Klein PE; Carneiro NP; Coelho CM; Trick HN; Kochian LV

Authors

Abstract

Crop yields are significantly reduced by aluminum toxicity on highly acidic soils, which comprise up to 50% of the world's arable land. Candidate aluminum tolerance proteins include organic acid efflux transporters, with the organic acids forming non-toxic complexes with rhizosphere aluminum. In this study, we used positional cloning to identify the gene encoding a member of the multidrug and toxic compound extrusion (MATE) family, an aluminum-activated citrate transporter, as responsible for the major sorghum (*Sorghum bicolor*) aluminum tolerance locus, Alt(SB). Polymorphisms in regulatory regions of Alt(SB) are likely to contribute to large allelic effects, acting to increase Alt(SB) expression in the root apex of tolerant genotypes. Furthermore, aluminum-inducible Alt(SB) expression is associated with induction of aluminum tolerance via enhanced root citrate exudation. These findings will allow us to identify superior Alt(SB) haplotypes that can be incorporated via molecular breeding and biotechnology into acid soil breeding programs, thus helping to increase crop yields in developing countries where acidic soils predominate.

Additional References

RELATED GEPHE

No matches found.

Related Genes

No matches found.

Related Haplotypes

EXTERNAL LINKS

COMMENTS