

## GEPHE SUMMARY

	Gephebase Gene	GephelD
ammonium transporter 2;1 ( <a href="https://www.gephebase.org/search-criteria/?and+Gene">https://www.gephebase.org/search-criteria/?and+Gene</a> Gephebase=^ammonium transporter 2;1^#gephebase-summary-title)	GP00001431	Main curator
Published	Entry Status	Prigent

## 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 (soil contamination; serpentine) ( <a href="https://www.gephebase.org/search-criteria/?and+Trait=^Xenobiotic+resistance+(soil+contamination;+serpentine)^#gephebase-summary-title">https://www.gephebase.org/search-criteria/?and+Trait=^Xenobiotic+resistance+(soil+contamination;+serpentine)^#gephebase-summary-title</a> )	
	Trait State in Taxon A
Arabidopsis arenosa Hochlantsch & Kasperstein populations not adapted to serpentine	Trait State in Taxon B
Arabidopsis arenosa Gulsen population adapted to serpentine	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	Latin Name	Taxon B	Latin Name
Arabidopsis arenosa ( <a href="https://www.gephebase.org/search-criteria/?and+Taxon+and+Synonyms=^Arabidopsis+arenosa^#gephebase-summary-title">https://www.gephebase.org/search-criteria/?and+Taxon+and+Synonyms=^Arabidopsis+arenosa^#gephebase-summary-title</a> )		Arabidopsis arenosa ( <a href="https://www.gephebase.org/search-criteria/?and+Taxon+and+Synonyms=^Arabidopsis+arenosa^#gephebase-summary-title">https://www.gephebase.org/search-criteria/?and+Taxon+and+Synonyms=^Arabidopsis+arenosa^#gephebase-summary-title</a> )	
-	Common Name	-	Common Name
	Synonyms		Synonyms
Arabis arenosa; Cardaminopsis arenosa; Arabidopsis arenosa (L.) Lawalree; Arabis arenosa (L.) Scop.; Cardaminopsis arenosa (L.) Hayek.	Arabis arenosa; Cardaminopsis arenosa; Arabidopsis arenosa (L.) Lawalree; Arabis arenosa (L.) Scop.; Cardaminopsis arenosa (L.) Hayek.		
species	Rank	species	Rank
	Lineage		Lineage
cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; rosids; malvids; Brassicales; Brassicaceae; Camelinae; Arabidopsis	cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; rosids; malvids; Brassicales; Brassicaceae; Camelinae; Arabidopsis		
Arabidopsis () - (Rank: genus) ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 3701">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 3701</a> )	Parent	Arabidopsis () - (Rank: genus) ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 3701">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 3701</a> )	Parent
38785 ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 38785">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 38785</a> )	NCBI Taxonomy ID	38785 ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 38785">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 38785</a> )	NCBI Taxonomy ID
Yes	is Taxon A an Infraspecies?	Yes	is Taxon B an Infraspecies?
	Taxon A Description		Taxon B Description
Arabidopsis arenosa Hochlantsch & Kasperstein populations not adapted to serpentine	Arabidopsis arenosa Gulsen population adapted to serpentine		

## GENOTYPIC CHANGE

AMT2	Generic Gene Name	UniProtKB Arabidopsis thaliana
	Synonyms	GenebankID or UniProtKB
ammonium transporter 2; AMMONIUM TRANSPORTER 2; AMMONIUM TRANSPORTER 2;1; AMT2;1; ATAMT2; F16M14.22; F16M14_22; At2g38290		0
3702.AT2G38290.1 ( <a href="http://string-db.org/newstring_cgi/show_network_section.pl?identifier= 3702.AT2G38290.1">http://string-db.org/newstring_cgi/show_network_section.pl?identifier= 3702.AT2G38290.1</a> )	String	
Belongs to the ammonia transporter channel (TC 1.A.11.2) family.	Sequence Similarities	
GO:0015398 : high-affinity secondary active ammonium transmembrane transporter activity ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0015398">https://www.ebi.ac.uk/QuickGO/term/GO:0015398</a> )	GO - Molecular Function	

## GO - Biological Process

GO:0009624 : response to nematode

(<https://www.ebi.ac.uk/QuickGO/term/GO:0009624>)GO:0015696 : ammonium transport (<https://www.ebi.ac.uk/QuickGO/term/GO:0015696>)

GO - Cellular Component

GO:0016021 : integral component of membrane

(<https://www.ebi.ac.uk/QuickGO/term/GO:0016021>)GO:0005886 : plasma membrane (<https://www.ebi.ac.uk/QuickGO/term/GO:0005886>)GO:0009506 : plasmodesma (<https://www.ebi.ac.uk/QuickGO/term/GO:0009506>)

Presumptive Null

Unknown ([https://www.gephebase.org/search-criteria?/and+Presumptive Null=%27Unknown%27#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Presumptive%20Null=%27Unknown%27#gephebase-summary-title))

Molecular Type

Unknown ([https://www.gephebase.org/search-criteria?/and+Molecular Type=%27Unknown%27#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Molecular%20Type=%27Unknown%27#gephebase-summary-title))

Aberration Type

Unknown ([https://www.gephebase.org/search-criteria?/and+Aberration Type=%27Unknown%27#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Aberration%20Type=%27Unknown%27#gephebase-summary-title))

Molecular Details of the Mutation

seven aa substitutions

Experimental Evidence

Association Mapping ([https://www.gephebase.org/search-criteria?/and+Experimental Evidence=%27Association Mapping%27#gephebase-summary-title](https://www.gephebase.org/search-criteria?/and+Experimental%20Evidence=%27Association%20Mapping%27#gephebase-summary-title))

Main Reference

Borrowed alleles and convergence in serpentine adaptation. (2016) (<https://pubmed.ncbi.nlm.nih.gov/27357660>)

Authors

Arnold BJ; Lahner B; DaCosta JM; Weisman CM; Hollister JD; Salt DE; Bomblies K; Yant L

Abstract

Serpentine barrens represent extreme hazards for plant colonists. These sites are characterized by high porosity leading to drought, lack of essential mineral nutrients, and phytotoxic levels of metals. Nevertheless, nature forged populations adapted to these challenges. Here, we use a population-based evolutionary genomic approach coupled with elemental profiling to assess how autotetraploid *Arabidopsis arenosa* adapted to a multichallenge serpentine habitat in the Austrian Alps. We first demonstrate that serpentine-adapted plants exhibit dramatically altered elemental accumulation levels in common conditions, and then resequence 24 autotetraploid individuals from three populations to perform a genome scan. We find evidence for highly localized selective sweeps that point to a polygenic, multtrait basis for serpentine adaptation. Comparing our results to a previous study of independent serpentine colonizations in the closely related diploid *Arabidopsis lyrata* in the United Kingdom and United States, we find the highest levels of differentiation in 11 of the same loci, providing candidate alleles for mediating convergent evolution. This overlap between independent colonizations in different species suggests that a limited number of evolutionary strategies are suited to overcome the multiple challenges of serpentine adaptation. Interestingly, we detect footprints of selection in *A. arenosa* in the context of substantial gene flow from nearby off-serpentine populations of *A. arenosa*, as well as from *A. lyrata*. In several cases, quantitative tests of introgression indicate that some alleles exhibiting strong selective sweep signatures appear to have been introgressed from *A. lyrata*. This finding suggests that migrant alleles may have facilitated adaptation of *A. arenosa* to this multihazard environment.

Additional References

## RELATED GEPHE

Related Genes

9 (calmodulin binding hydrolase, Caspary strip membrane domain protein 1, early responsive to dehydration stress protein 4, Ferroportin 2, high expression of osmotically responsive genes 2, K<sup>+</sup> uptake permease 9, LACCASE 8, sulfate transporter 1;1, two pore channel) ([https://www.gephebase.org/search-criteria?/or+Taxon ID=%2738785%27&Trait=Xenobiotic resistance&and+groupHaplotypes=true#gephebase-summary-title](https://www.gephebase.org/search-criteria?/or+Taxon%20ID=%2738785%27&Trait=Xenobiotic%20resistance&and+groupHaplotypes=true#gephebase-summary-title))

Related Haplotypes

No matches found.

## EXTERNAL LINKS

## COMMENTS