

## GEPHE SUMMARY

|  |  |                                    |
|--|--|------------------------------------|
| <p>early responsive to dehydration stress protein 4 (<a href="https://www.gephebase.org/search-criteria?/and+Gene+Gephebase=^early+responsive+to+dehydration+stress+protein+4^#gephebase-summary-title">https://www.gephebase.org/search-criteria?/and+Gene+Gephebase=^early+responsive+to+dehydration+stress+protein+4^#gephebase-summary-title</a>)</p> <p>Published</p> | <p>Gephebase Gene</p> <p>GP00001434</p> <p>Prigent</p> <p>Entry Status</p> | <p>GepheID</p> <p>Main curator</p> |
|--|--|------------------------------------|

## PHENOTYPIC CHANGE

|   |   |  |  |  |  |
|---|---|--|--|--|--|
| <p>Physiology (<a href="https://www.gephebase.org/search-criteria?/and+Trait+Category=^Physiology^#gephebase-summary-title">https://www.gephebase.org/search-criteria?/and+Trait+Category=^Physiology^#gephebase-summary-title</a>)</p> <p>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>)</p> <p>Arabidopsis arenosa Hochlantsch &amp; Kasperstein populations not adapted to serpentine</p> <p>Arabidopsis arenosa Gulsen population adapted to serpentine</p> <p>Taxon A</p> <p>Intraspecific (<a href="https://www.gephebase.org/search-criteria?/and+Taxonomic+Status=^Intraspecific^#gephebase-summary-title">https://www.gephebase.org/search-criteria?/and+Taxonomic+Status=^Intraspecific^#gephebase-summary-title</a>)</p> | <p>Trait Category</p> <p>Trait</p> <p>Trait State in Taxon A</p> <p>Trait State in Taxon B</p> <p>Ancestral State</p> <p>Taxonomic Status</p> | <p>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>)</p> <p>-</p> <p>Arabis arenosa; Cardaminopsis arenosa; Arabidopsis arenosa (L.) Lawalree; Arabis arenosa (L.) Scop.; Cardaminopsis arenosa (L.) Hayek.</p> <p>species</p> <p>cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; rosids; malvids; Brassicales; Brassicaceae; Camelineae; Arabidopsis</p> <p>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>)</p> <p>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>)</p> <p>is Taxon A an Intraspecies?</p> <p>Yes</p> <p>Arabidopsis arenosa Hochlantsch &amp; Kasperstein populations not adapted to serpentine</p> | <p>Taxon A</p> <p>Latin Name</p> <p>Common Name</p> <p>Synonyms</p> <p>Rank</p> <p>Lineage</p> <p>Parent</p> <p>NCBI Taxonomy ID</p> <p>is Taxon A an Intraspecies?</p> <p>Taxon A Description</p> | <p>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>)</p> <p>-</p> <p>Arabis arenosa; Cardaminopsis arenosa; Arabidopsis arenosa (L.) Lawalree; Arabis arenosa (L.) Scop.; Cardaminopsis arenosa (L.) Hayek.</p> <p>species</p> <p>cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; rosids; malvids; Brassicales; Brassicaceae; Camelineae; Arabidopsis</p> <p>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>)</p> <p>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>)</p> <p>is Taxon B an Intraspecies?</p> <p>Yes</p> <p>Arabidopsis arenosa Gulsen population adapted to serpentine</p> | <p>Taxon B</p> <p>Latin Name</p> <p>Common Name</p> <p>Synonyms</p> <p>Rank</p> <p>Lineage</p> <p>Parent</p> <p>NCBI Taxonomy ID</p> <p>is Taxon B an Intraspecies?</p> <p>Taxon B Description</p> |
|---|---|--|--|--|--|

## GENOTYPIC CHANGE

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|--|--|--|
| <p>ERD4</p> <p>early-responsive to dehydration 4; T4K22.4; T4K22_4; At1g30360</p> <p>3702.AT1G30360.1 (<a href="http://string-db.org/newstring.cgi/show_network_section.pl?identifier=3702.AT1G30360.1">http://string-db.org/newstring.cgi/show_network_section.pl?identifier=3702.AT1G30360.1</a>)</p> <p>Belongs to the CSC1 (TC 1.A.17) family.</p> <p>GO:0005227 : calcium activated cation channel activity (<a href="https://www.ebi.ac.uk/QuickGO/term/GO:0005227">https://www.ebi.ac.uk/QuickGO/term/GO:0005227</a>)</p> | <p>Generic Gene Name</p> <p>Synonyms</p> <p>String</p> <p>Sequence Similarities</p> <p>GO - Molecular Function</p> | <p>UniProtKB Arabidopsis thaliana</p> <p>Q9C8G5 (<a href="http://www.uniprot.org/uniprot/Q9C8G5">http://www.uniprot.org/uniprot/Q9C8G5</a>)</p> <p>()</p> <p>GenebankID or UniProtKB</p> |
|--|--|--|

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>)GO:0016020 : membrane (<https://www.ebi.ac.uk/QuickGO/term/GO:0016020>)GO:0005774 : vacuolar membrane (<https://www.ebi.ac.uk/QuickGO/term/GO:0005774>)GO:0009507 : chloroplast (<https://www.ebi.ac.uk/QuickGO/term/GO:0009507>)GO:0005773 : vacuole (<https://www.ebi.ac.uk/QuickGO/term/GO:0005773>)GO:0009941 : chloroplast envelope (<https://www.ebi.ac.uk/QuickGO/term/GO:0009941>)Unknown (<https://www.gephebase.org/search-criteria?/and+Presumptive Null=^Unknown^#gephebase-summary-title>)

Presumptive Null

Unknown (<https://www.gephebase.org/search-criteria?/and+Molecular Type=^Unknown^#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

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

Experimental Evidence

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

Main Reference

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

Authors

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, multitrait 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.

Abstract

Additional References

## RELATED GEPHE

Related Genes

9 (ammonium transporter 2;1, calmodulin binding hydrolase, Casparian strip membrane domain protein 1, 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=^38785^/and+Trait=Xenobiotic resistance/and+groupHaplotypes=true#gephebase-summary-title>)

Related Haplotypes

No matches found.

## EXTERNAL LINKS

## COMMENTS