

GEPHE SUMMARY

	Gephebase Gene	GephelD
StCDF1 (https://www.gephebase.org/search-criteria?/and+Gene Gephebase=^StCDF1^#gephebase-summary-title)	GP00001079	Main curator
Published	Entry Status	Martin

PHENOTYPIC CHANGE

	Trait Category	
Physiology (https://www.gephebase.org/search-criteria?/and+Trait Category=^Physiology^#gephebase-summary-title)	Trait	
Latitudinal adaptation (https://www.gephebase.org/search-criteria?/and+Trait=^Latitudinal adaptation^#gephebase-summary-title)	Trait State in Taxon A	
Solanum tuberosum - wild	Trait State in Taxon B	
Solanum tuberosum - domesticated	Ancestral State	
Taxon A	Taxonomic Status	
Domesticated (https://www.gephebase.org/search-criteria?/and+Taxonomic Status=^Domesticated^#gephebase-summary-title)		
Taxon A		Taxon B
	Latin Name	Latin Name
Solanum tuberosum (https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Solanum tuberosum^#gephebase-summary-title)	Solanum tuberosum (https://www.gephebase.org/search-criteria?/and+Taxon+and+Synonyms=^Solanum tuberosum^#gephebase-summary-title)	
potato	Common Name	Common Name
Solanum tuberosum subsp. tuberosum; potato; potatoes; Solanum tuberosum L.	Synonyms	Synonyms
species	Rank	Rank
cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; asterids; lamiids; Solanales; Solanaceae; Solanoideae; Solaneae; Solanum	Lineage	Lineage
Solanum () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 4107)	Parent	Parent
4113 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 4113)	NCBI Taxonomy ID	NCBI Taxonomy ID
No	is Taxon A an Infraspecies?	is Taxon B an Infraspecies?

GENOTYPIC CHANGE

CDF1	Generic Gene Name	UniProtKB Arabidopsis thaliana
cycling DOF factor 1; K19B1.4; K19B1_4; DOF5.5; At5g62430; MM19.24	Synonyms	GenebankID or UniProtKB
3702.AT5G62430.1 (http://string-db.org/newstring_cgi/show_network_section.pl?identifier= 3702.AT5G62430.1)	String	AGF33077 (https://www.ncbi.nlm.nih.gov/nuccore/AGF33077)
-	Sequence Similarities	
GO:0003700 : DNA-binding transcription factor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0003700) GO:0043565 : sequence-specific DNA binding (https://www.ebi.ac.uk/QuickGO/term/GO:0043565) GO:0046872 : metal ion binding (https://www.ebi.ac.uk/QuickGO/term/GO:0046872) GO:0003677 : DNA binding (https://www.ebi.ac.uk/QuickGO/term/GO:0003677) GO:0000978 : RNA polymerase II proximal promoter sequence-specific DNA binding	GO - Molecular Function	

(<https://www.ebi.ac.uk/QuickGO/term/GO:0000978>)

GO:0001210 : transcriptional repressor activity, metal ion regulated sequence-specific DNA binding (<https://www.ebi.ac.uk/QuickGO/term/GO:0001210>)

GO - Biological Process

GO:0006355 : regulation of transcription, DNA-templated

(<https://www.ebi.ac.uk/QuickGO/term/GO:0006355>)

GO:0010228 : vegetative to reproductive phase transition of meristem

(<https://www.ebi.ac.uk/QuickGO/term/GO:0010228>)

GO:0045892 : negative regulation of transcription, DNA-templated

(<https://www.ebi.ac.uk/QuickGO/term/GO:0045892>)

GO:0009908 : flower development (<https://www.ebi.ac.uk/QuickGO/term/GO:0009908>)

GO:0048510 : regulation of timing of transition from vegetative to reproductive phase

(<https://www.ebi.ac.uk/QuickGO/term/GO:0048510>)

GO:0009658 : chloroplast organization

(<https://www.ebi.ac.uk/QuickGO/term/GO:0009658>)

GO - Cellular Component

GO:0005634 : nucleus (<https://www.ebi.ac.uk/QuickGO/term/GO:0005634>)

Presumptive Null

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

Molecular Type

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

Aberration Type

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

Insertion Size

1-9 bp

Molecular Details of the Mutation

+7bp insertion resulting in frameshift and truncated protein

Experimental Evidence

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

Main Reference

Naturally occurring allele diversity allows potato cultivation in northern latitudes. (2013) (<https://pubmed.ncbi.nlm.nih.gov/23467094>)

Authors

Kloosterman B; Abelenda JA; Gomez Mdel M; Oortwijn M; de Boer JM; Kowitwanich K; Horvath BM; van Eck HJ; Smacniak C; Prat S; Visser RG; Bachem CW

Abstract

Potato (*Solanum tuberosum* L.) originates from the Andes and evolved short-day-dependent tuber formation as a vegetative propagation strategy. Here we describe the identification of a central regulator underlying a major-effect quantitative trait locus for plant maturity and initiation of tuber development. We show that this gene belongs to the family of DOF (DNA-binding with one finger) transcription factors and regulates tuberization and plant life cycle length, by acting as a mediator between the circadian clock and the StSP6A mobile tuberization signal. We also show that natural allelic variants evade post-translational light regulation, allowing cultivation outside the geographical centre of origin of potato. Potato is a member of the Solanaceae family and is one of the world's most important food crops. This annual plant originates from the Andean regions of South America. Potato develops tubers from underground stems called stolons. Its equatorial origin makes potato essentially short-day dependent for tuberization and potato will not make tubers in the long-day conditions of spring and summer in the northern latitudes. When introduced in temperate zones, wild material will form tubers in the course of the autumnal shortening of day-length. Thus, one of the first selected traits in potato leading to a European potato type is likely to have been long-day acclimation for tuberization. Potato breeders can exploit the naturally occurring variation in tuberization onset and life cycle length, allowing varietal breeding for different latitudes, harvest times and markets.

Additional References

RELATED GEPHE

Related Genes

No matches found.

Related Haplotypes

1 (<https://www.gephebase.org/search-criteria?/or+Gene+Gephebase=^StCDF1^/and+Taxon+ID=^4113^/or+Gene+Gephebase=^StCDF1^/and+Taxon+ID=^4113^#gephebase-summary-title>)

EXTERNAL LINKS

COMMENTS