

GEPHE SUMMARY

StCDF1 (https://www.gephebase.org/search-criteria?/and+GeneGephebase=~StCDF1~#gephebase-summary-title)	Gephebase Gene	GP00001080	GepheID
	Entry Status	Martin	Main curator
	Published		

PHENOTYPIC CHANGE

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

GENOTYPIC CHANGE

Generic Gene Name		UniProtKB Arabidopsis thaliana	
CDF1		Q8W1E3 (http://www.uniprot.org/uniprot/Q8W1E3)	
Synonyms		GenebankID or UniProtKB	
cycling DOF factor 1; K19B1.4; K19B1_4; DOF5.5; At5g62430; MMI9.24		AGF33077 (https://www.ncbi.nlm.nih.gov/nuccore/AGF33077)	
String			
3702.AT5G62430.1 (http://string-db.org/newstring.cgi/show_network_section.pl?identifier=3702.AT5G62430.1)			
Sequence Similarities			
-			
GO - Molecular Function			
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			

(<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

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

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

Insertion Size

100-999 bp

Molecular Details of the Mutation

865bp insertion leading to a fusion 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; Smaczniak 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