

GEPHE SUMMARY

	Gephebase Gene		GepheID
Cryptochrome 2 (CRY2) EDI allele (https://www.gephebase.org/search-criteria?/and+Gene+Gephebase=^Cryptochrome 2 (CRY2) EDI allele^#gephebase-summary-title)		GP00000192	
	Entry Status	Martin	Main curator
Published			

PHENOTYPIC CHANGE

Trait #1	Trait Category
Physiology (https://www.gephebase.org/search-criteria?/and+Trait+Category=^Physiology^#gephebase-summary-title)	
Flowering time (https://www.gephebase.org/search-criteria?/and+Trait=^Flowering time^#gephebase-summary-title)	Trait
Arabidopsis thaliana	Trait State in Taxon A
Arabidopsis thaliana	Trait State in Taxon B

Trait #2	Trait Category
Morphology (https://www.gephebase.org/search-criteria?/and+Trait+Category=^Morphology^#gephebase-summary-title)	
Fruit shape (fruit length) (https://www.gephebase.org/search-criteria?/and+Trait=^Fruit shape (fruit length)^#gephebase-summary-title)	Trait
Arabidopsis thaliana	Trait State in Taxon A
Arabidopsis thaliana	Trait State in Taxon B

Trait #3	Trait Category
Morphology (https://www.gephebase.org/search-criteria?/and+Trait+Category=^Morphology^#gephebase-summary-title)	
Flower morphology (ovule number per fruit) (https://www.gephebase.org/search-criteria?/and+Trait=^Flower morphology (ovule number per fruit)^#gephebase-summary-title)	Trait
Arabidopsis thaliana	Trait State in Taxon A
Arabidopsis thaliana	Trait State in Taxon B

Trait #4	Trait Category
Physiology (https://www.gephebase.org/search-criteria?/and+Trait+Category=^Physiology^#gephebase-summary-title)	
Fertility (percentage of unfertilized ovules) (https://www.gephebase.org/search-criteria?/and+Trait=^Fertility (percentage of unfertilized ovules)^#gephebase-summary-title)	Trait
Arabidopsis thaliana	Trait State in Taxon A
Arabidopsis thaliana	Trait State in Taxon B

	Ancestral State
Data not curated	
	Taxonomic Status
Intraspecific (https://www.gephebase.org/search-criteria?/and+Taxonomic+Status=^Intraspecific^#gephebase-summary-title)	

Taxon A	Taxon B
Arabidopsis thaliana (#gephebase-summary-title) thale cress thale cress; mouse-ear cress; thale-cress; Arabidopsis thaliana (L.) Heynh.; Arabidopsis thaliana (thale cress); Arabidopsis_thaliana; Arbisopsis thaliana; thale kress species cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetales; rosids; malvids; Brassicales; Brassicaceae; Camelineae; Arabidopsis Arabidopsis () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=3701) 3702 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=3702) No is Taxon A an Intraspecies?	Arabidopsis thaliana (#gephebase-summary-title) thale cress thale cress; mouse-ear cress; thale-cress; Arabidopsis thaliana (L.) Heynh.; Arabidopsis thaliana (thale cress); Arabidopsis_thaliana; Arbisopsis thaliana; thale kress species cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetales; rosids; malvids; Brassicales; Brassicaceae; Camelineae; Arabidopsis Arabidopsis () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=3701) 3702 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=3702) No is Taxon B an Intraspecies?

GENOTYPIC CHANGE

CRY2	Generic Gene Name Q96524 (http://www.uniprot.org/uniprot/Q96524) AT-PHH1; ACRY2; cryptochrome 2; CRYPTOCHROME 2 APOPROTEIN; F19P19.14; F19P19_14; FHA; PHH1; SEL20; At1g04400 EU351972 (https://www.ncbi.nlm.nih.gov/nucore/EU351972)
3702.AT1G04400.1 (http://string-db.org/newstring.cgi/show_network_section.pl?identifier=3702.AT1G04400.1) String Belongs to the DNA photolyase class-1 family. GO:0005524 : ATP binding (https://www.ebi.ac.uk/QuickGO/term/GO:0005524) GO:0042802 : identical protein binding (https://www.ebi.ac.uk/QuickGO/term/GO:0042802) GO:0046872 : metal ion binding (https://www.ebi.ac.uk/QuickGO/term/GO:0046872) GO:0009882 : blue light photoreceptor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0009882) GO:0071949 : FAD binding (https://www.ebi.ac.uk/QuickGO/term/GO:0071949)	UniProtKB Arabidopsis thaliana GenebankID or UniProtKB Sequence Similarities GO - Molecular Function GO - Biological Process

(<https://www.ebi.ac.uk/QuickGO/term/GO:0042752>)
GO:0009909 : regulation of flower development
(<https://www.ebi.ac.uk/QuickGO/term/GO:0009909>)
GO:1901371 : regulation of leaf morphogenesis
(<https://www.ebi.ac.uk/QuickGO/term/GO:1901371>)
GO:0010075 : regulation of meristem growth
(<https://www.ebi.ac.uk/QuickGO/term/GO:0010075>)
GO:2000028 : regulation of photoperiodism, flowering
(<https://www.ebi.ac.uk/QuickGO/term/GO:2000028>)
GO:0009646 : response to absence of light
(<https://www.ebi.ac.uk/QuickGO/term/GO:0009646>)
GO:0010244 : response to low fluence blue light stimulus by blue low-fluence system
(<https://www.ebi.ac.uk/QuickGO/term/GO:0010244>)
GO:1902347 : response to strigolactone
(<https://www.ebi.ac.uk/QuickGO/term/GO:1902347>)
GO:0009414 : response to water deprivation
(<https://www.ebi.ac.uk/QuickGO/term/GO:0009414>)
GO:0010118 : stomatal movement (<https://www.ebi.ac.uk/QuickGO/term/GO:0010118>)
GO - Cellular Component
GO:0005737 : cytoplasm (<https://www.ebi.ac.uk/QuickGO/term/GO:0005737>)
GO:0005634 : nucleus (<https://www.ebi.ac.uk/QuickGO/term/GO:0005634>)
GO:0016604 : nuclear body (<https://www.ebi.ac.uk/QuickGO/term/GO:0016604>)
GO:0016605 : PML body (<https://www.ebi.ac.uk/QuickGO/term/GO:0016605>)
GO:0005773 : vacuole (<https://www.ebi.ac.uk/QuickGO/term/GO:0005773>)

No (<https://www.gephebase.org/search-criteria?/and+Presumptive Null=~No~#gephebase-summary-title>) Presumptive Null

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

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

Nonsynonymous SNP Coding Change

V367M Molecular Details of the Mutation

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

	Taxon A	Taxon B	Position
Codon	-	-	-
Amino-acid	-	-	-

A QTL for flowering time in Arabidopsis reveals a novel allele of CRY2. (2001) (<https://pubmed.ncbi.nlm.nih.gov/11726930>) Main Reference

El-Din El-Assal S; Alonso-Blanco C; Peeters AJ; Raz V; Koornneef M Authors

Variation of flowering time is found in the natural populations of many plant species. The underlying genetic variation, mostly of a quantitative nature, is presumed to reflect adaptations to different environments contributing to reproductive success. Analysis of natural variation for flowering time in Arabidopsis thaliana has identified several quantitative trait loci (QTL), which have yet to be characterized at the molecular level. A major environmental factor that determines flowering time is photoperiod or day length, the length of the light period, which changes across the year differently with geographical latitude. We identified the EDI locus as a QTL partly accounting for the difference in flowering response to the photoperiod between two Arabidopsis accessions: the laboratory strain Landsberg erecta (Ler), originating in Northern Europe, and Cvi, collected in the tropical Cape Verde Islands. Positional cloning of the EDI QTL showed it to be a novel allele of CRY2, encoding the blue-light photoreceptor cryptochrome-2 that has previously been shown to promote flowering in long-day (LD) photoperiods. We show that the unique EDI flowering phenotype results from a single amino-acid substitution that reduces the light-induced downregulation of CRY2 in plants grown under short photoperiods, leading to early flowering. Abstract

Linkage disequilibrium mapping of Arabidopsis CRY2 flowering time alleles. (2004) (<https://pubmed.ncbi.nlm.nih.gov/15280248>) Additional References

Pleiotropic effects of the Arabidopsis cryptochrome 2 allelic variation underlie fruit trait-related QTL. (2004) (<https://pubmed.ncbi.nlm.nih.gov/15248119>)

The Cape Verde Islands allele of cryptochrome 2 enhances cotyledon unfolding in the absence of blue light in Arabidopsis. (2003) (<https://pubmed.ncbi.nlm.nih.gov/14605225>)

RELATED GEPHE

13 (AGAMOUS-LIKE 50, EARLY FLOWERING 3(ELF3), FLC (Flowering Locus C), FLM (MAF1), Flowering locus T (FT), Frigida (FRI), Frigida like 1 (FRL1), Frigida like 2 (FRL2), MADS AFFECTING FLOWERING 2 (MAF2), SVP (SHORT VEGETATIVE PHASE), VIN3, GIBBERELLIC ACID REQUIRING 1 (GA1), HUA2) Related Genes

(<https://www.gephebase.org/search-criteria?/or+Taxon ID=~3702~/and+Trait=Flowering time/or+Taxon ID=~3702~/and+Trait=Fruit shape/or+Taxon ID=~3702~/and+Trait=Flower morphology/or+Taxon ID=~3702~/and+Trait=Fertility/and+groupHaplotypes=true#gephebase-summary-title>)

No matches found. Related Haplotypes

EXTERNAL LINKS

COMMENTS

@Pleiotropy @GxE