

GEPHE SUMMARY

anthocyanin2 (an2) (https://www.gephebase.org/search-criteria?/and+Gene Gephebase="anthocyanin2 (an2)"#gephebase-summary-title)	Gephebase Gene	GP00000087	GepheID
Published	Entry Status	Martin	Main curator

PHENOTYPIC CHANGE

Morphology (https://www.gephebase.org/search-criteria?/and+Trait Category="Morphology"#gephebase-summary-title)	Trait Category		
Coloration (fruit) (<a "="" href="https://www.gephebase.org/search-criteria?/and+Trait=">https://www.gephebase.org/search-criteria?/and+Trait=" Coloration (fruit)"#gephebase-summary-title)	Trait		
Capsicum annuum	Trait State in Taxon A		
Capsicum chinense	Trait State in Taxon B		
Data not curated	Ancestral State		
Domesticated (https://www.gephebase.org/search-criteria?/and+Taxonomic Status="Domesticated"#gephebase-summary-title)	Taxonomic Status		

Taxon A	Latin Name	Taxon B	Latin Name
Capsicum annuum (https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms="Capsicum annuum"#gephebase-summary-title)	Capsicum chinense (https://www.gephebase.org/search-criteria?/and+Taxon and Synonyms="Capsicum chinense"#gephebase-summary-title)		
-	Common Name	-	Common Name
Capsicum annuum L.; Capsicum annum; Capsicum capsicum species	Synonyms	Scotch bonnet; bonnet pepper; habanero; piri piri; rocotillo; Capsicum chinense Jacq. species	Synonyms
cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; asterids; lamiids; Solanales; Solanaceae; Solanoideae; Capsiceae; Capsicum	Rank		Rank
Capsicum (peppers) - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4071)	Lineage	Capsicum (peppers) - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4071)	Lineage
4072 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4072)	Parent	80379 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=80379)	Parent
No	NCBI Taxonomy ID	No	NCBI Taxonomy ID
	is Taxon A an Intraspecies?		is Taxon B an Intraspecies?

GENOTYPIC CHANGE

AN2	Generic Gene Name	A4GRU8 (http://www.uniprot.org/uniprot/A4GRU8)	UniProtKB Petunia integrifolia
-	Synonyms	0	GenebankID or UniProtKB
-	String		
-	Sequence Similarities		
-	GO - Molecular Function		
GO:0003677 : DNA binding (https://www.ebi.ac.uk/QuickGO/term/GO:0003677)	GO - Biological Process		
-	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
Cis-regulatory (https://www.gephebase.org/search-criteria?/and+Molecular Type="Cis-regulatory"#gephebase-summary-title)			

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

Aberration Type

unknown

Molecular Details of the Mutation

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

Experimental Evidence

The A locus that controls anthocyanin accumulation in pepper encodes a MYB transcription factor homologous to Anthocyanin2 of Petunia. (2004) (<https://pubmed.ncbi.nlm.nih.gov/14997303/>)

Main Reference

Borovsky Y; Oren-Shamir M; Ovadia R; De Jong W; Paran I

Authors

Pepper plants containing the dominant A gene accumulate anthocyanin pigments in the foliage, flower and immature fruit. We previously mapped A to pepper chromosome 10 in the F(2) progeny of a cross between 5226 (purple-fruited) and PI 159234 (green-fruited) to a region that corresponds, in tomato, to the location of Petunia anthocyanin 2 (An2), a regulator of anthocyanin biosynthesis. This suggested that A encodes a homologue of Petunia An2. Using the sequences of An2 and a corresponding tomato expressed sequence tag, we isolated a pepper cDNA orthologous to An2 that cosegregated with A. We subsequently determined the expression of A by Northern analysis, using RNA extracted from fruits, flowers and leaves of 5226 and PI 159234. In 5226, expression was detected in all stages of fruit development and in both flower and leaf. In contrast, A was not expressed in the sampled tissues in PI 159234. Genomic sequence comparison of A between green- and purple-fruited genotypes revealed no differences in the coding region, indicating that the lack of expression of A in the green genotypes can be attributed to variation in the promoter region. By analyzing the expression of the structural genes in the anthocyanin biosynthetic pathway in 5226 and PI 159234, it was determined that, similar to Petunia, the early genes in the pathway are regulated independently of A, while expression of the late genes is A-dependent.

Abstract

Additional References

RELATED GEPHE

No matches found.

Related Genes

1 ([https://www.gephebase.org/search-criteria?/or+Gene Gephebase=^anthocyanin2 \(an2\)^/and+Taxon ID=^4072^/or+Gene Gephebase=^anthocyanin2 \(an2\)^/and+Taxon ID=^80379^#gephebase-summary-title](https://www.gephebase.org/search-criteria?/or+Gene Gephebase=^anthocyanin2 (an2)^/and+Taxon ID=^4072^/or+Gene Gephebase=^anthocyanin2 (an2)^/and+Taxon ID=^80379^#gephebase-summary-title))

Related Haplotypes

EXTERNAL LINKS

COMMENTS