

GEPHE SUMMARY

Gephebase Gene
phytochrome D (PHYD)

Entry Status
Published

GepheID
GP00000874

Main curator
Martin

PHENOTYPIC CHANGE

Trait #1

Trait Category
Morphology

Trait

Leaf morphology (increased petiole length)

Trait State in Taxon A

Arabidopsis thaliana

Trait State in Taxon B

Arabidopsis thaliana Ws

Trait #2

Trait Category
Morphology

Trait

Cotyledon morphology (reduced cotyledon area)

Trait State in Taxon A

-

Trait State in Taxon B

-

Trait #3

Trait Category
Morphology

Trait

Coloration (anthocyanin accumulation in seedling stems)

Trait State in Taxon A

-

Trait State in Taxon B

-

Trait #4

Trait Category
Physiology

Trait

Plant size (diminished effect of end-of-day pulse of far red light on hypocotyl elongation)

Trait State in Taxon A

-

Trait State in Taxon B

-

Trait #5

Trait Category
Morphology

Trait

Plant architecture (decrease in number of rosette leaves at onset of flowering)

Trait State in Taxon A

-

Trait State in Taxon B

-

Ancestral State

Taxon A

Taxonomic Status

Intraspecific

Taxon A

Latin Name

Arabidopsis thaliana

Common Name

thale cress

Synonyms

thale cress; mouse-ear cress; thale-cress; *Arabidopsis thaliana* (L.) Heynh.; *Arabidopsis thaliana* (thale cress); *Arabidopsis_thaliana*; *Arbisopsis thaliana*; thale kress

Rank

species

Lineage

cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; rosids; malvids; Brassicales; Brassicaceae; Camelineae; *Arabidopsis*

Parent

Arabidopsis () - (Rank: genus)

NCBI Taxonomy ID

3702

is Taxon A an Intraspecies?

No

Taxon B

Latin Name

Arabidopsis thaliana

Common Name

thale cress

Synonyms

thale cress; mouse-ear cress; thale-cress; *Arabidopsis thaliana* (L.) Heynh.; *Arabidopsis thaliana* (thale cress); *Arabidopsis_thaliana*; *Arbisopsis thaliana*; thale kress

Rank

species

Lineage

cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; rosids; malvids; Brassicales; Brassicaceae; Camelineae; *Arabidopsis*

Parent

Arabidopsis () - (Rank: genus)

NCBI Taxonomy ID

3702

is Taxon B an Intraspecies?

Yes

Taxon B Description

Arabidopsis thaliana Ws

GENOTYPIC CHANGE

Generic Gene Name

PHYD

Synonyms

DL4165C; FCAALL.323; phytochrome D; PHYTOCHROME D; At4g16250; dl4165c

String

3702.AT4G16250.1

Sequence Similarities

Belongs to the phytochrome family.

GO - Molecular Function

GO:0042802 : identical protein binding
GO:0042803 : protein homodimerization activity
GO:0000155 : phosphorelay sensor kinase activity
GO:0009881 : photoreceptor activity

GO - Biological Process

GO:0006355 : regulation of transcription, DNA-templated
GO:0018298 : protein-chromophore linkage
GO:0009585 : red, far-red light phototransduction
GO:0009584 : detection of visible light
GO:0017006 : protein-tetrapyrrole linkage

GO - Cellular Component

GO:0005634 : nucleus

Presumptive Null

Yes

Molecular Type

Coding

Aberration Type

Deletion

Deletion Size

10-99 bp

Molecular Details of the Mutation

14bp deletion causing premature stop codon

Experimental Evidence

Candidate Gene

Main Reference

UniProtKB *Arabidopsis thaliana*

P42497

GenebankID or UniProtKB

X76609

A deletion in the PHYD gene of the Arabidopsis Wassilewskija ecotype defines a role for phytochrome D in red/far-red light sensing. (1997)

Authors

Aukerman MJ; Hirschfeld M; Wester L; Weaver M; Clack T; Amasino RM; Sharrock RA

Abstract

The PHYD gene of the Wassilewskija (*Ws*) ecotype of *Arabidopsis* contains a 14-bp deletion (the phyD-1 mutation) beginning at amino acid 29 of the reading frame, resulting in translation termination at a nonsense codon 138 nucleotides downstream of the deletion end point. Immunoblot analyses showed that *Ws* lacks phyD but contains normal levels of phyA, phyB, and phyC. By backcrossing into the *Ws* and Landsberg erecta genetic backgrounds, we constructed sibling pairs of PHYD+ and phyD-1 lines and of phyB- PHYD+ and phyB- phyD- lines. Hypocotyl lengths after growth under white or red light increased sequentially in strains that were B+D+, B+D-, B-D+, and B-D-. In the *Ws* genetic background, an increase in petiole length, a reduction in cotyledon area and in anthocyanin accumulation in seedling stems, a diminished effect of an end-of-day pulse of far-red light on hypocotyl elongation, and a decrease in the number of rosette leaves at the onset of flowering were also seen sequentially in these lines. Thus, phyD, which is approximately 80% identical in amino acid sequence to phyB, acts in conjunction with phyB in regulating many shade avoidance responses. The existence of the apparently naturally occurring phyD-1 mutation indicates that phyD is not essential in some natural environments.

Additional References

RELATED GEPHE

Related Genes

8 (ACD6 = ACCELERATED CELL DEATH 6, ACS11, AGAMOUS-Like6, AtGA20ox1 (=GA5=Sd1), ERECTA, HUA2, PAPI, PAP2)

Related Haplotypes

No matches found.

COMMENTS

@Pleiotropy