

GEPHE SUMMARY

<p>Hd1 (<a +hd1+"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Gene+Gephebase=">https://www.gephebase.org/search-criteria?/and+Gene+Gephebase="+Hd1+"#gephebase-summary-title)</p> <p>Published</p>	<p>Gephebase Gene</p> <p>Entry Status</p>	<p>GP00001723</p> <p>Courtier</p>	<p>GepheID</p> <p>Main curator</p>
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PHENOTYPIC CHANGE

<p>Physiology (<a +physiology+"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Trait+Category=">https://www.gephebase.org/search-criteria?/and+Trait+Category="+Physiology+"#gephebase-summary-title)</p> <p>Flowering time (<a +flowering+time+"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Trait=">https://www.gephebase.org/search-criteria?/and+Trait="+Flowering+time+"#gephebase-summary-title)</p> <p>Oryza sativa - many cultivars</p> <p>Oryza sativa - many cultivars</p> <p>Taxon A</p> <p>Domesticated (<a +domesticated+"#gephebase-summary-title"="" href="https://www.gephebase.org/search-criteria?/and+Taxonomic+Status=">https://www.gephebase.org/search-criteria?/and+Taxonomic+Status="+Domesticated+"#gephebase-summary-title)</p>	<p>Trait Category</p> <p>Trait</p> <p>Trait State in Taxon A</p> <p>Trait State in Taxon B</p> <p>Ancestral State</p> <p>Taxonomic Status</p>	<p>Oryza sativa</p> <p>rice</p> <p>rice; red rice; <i>Oryza sativa</i> L.</p> <p>species</p> <p>cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; Liliopsida; Petrosaviidae; commelinids; Poales; Poaceae; BOP clade; Oryzoideae; Oryzaceae; Oryzinae; Oryza</p> <p>Oryza () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4527)</p> <p>4530 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4530)</p> <p>No is Taxon A an Intraspecies?</p>	<p>Taxon B</p> <p>Oryza sativa</p> <p>rice</p> <p>rice; red rice; <i>Oryza sativa</i> L.</p> <p>species</p> <p>cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; Liliopsida; Petrosaviidae; commelinids; Poales; Poaceae; BOP clade; Oryzoideae; Oryzaceae; Oryzinae; Oryza</p> <p>Oryza () - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4527)</p> <p>4530 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=4530)</p> <p>No is Taxon B an Intraspecies?</p>
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GENOTYPIC CHANGE

<p>HD1</p> <p>Hd1; SE1; OsHd1; Os06g0275000; LOC_Os06g16370; P0038C05.23; P0676F10.34</p> <p>39947.LOC_Os06g16370.1 (http://string-db.org/newstring.cgi/show_network_section.pl?identifier=39947.LOC_Os06g16370.1)</p> <p>Belongs to the CONSTANS family.</p> <p>GO:0003700 : DNA-binding transcription factor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0003700)</p> <p>GO:0008270 : zinc ion binding (https://www.ebi.ac.uk/QuickGO/term/GO:0008270)</p> <p>GO:0003677 : DNA binding (https://www.ebi.ac.uk/QuickGO/term/GO:0003677)</p> <p>GO:0030154 : cell differentiation (https://www.ebi.ac.uk/QuickGO/term/GO:0030154)</p> <p>GO:0045892 : negative regulation of transcription, DNA-templated</p>	<p>Generic Gene Name</p> <p>Synonyms</p> <p>String</p> <p>Sequence Similarities</p> <p>GO - Molecular Function</p> <p>GO - Biological Process</p>	<p>UniProtKB Oryza sativa subsp. japonica</p> <p>Q9FDX8 (http://www.uniprot.org/uniprot/Q9FDX8)</p> <p>BAB17628 (https://www.ncbi.nlm.nih.gov/nucleotide/BAB17628)</p> <p>GenebankID or UniProtKB</p>
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(<https://www.ebi.ac.uk/QuickGO/term/GO:0045892>)
 GO:0009908 : flower development (<https://www.ebi.ac.uk/QuickGO/term/GO:0009908>)
 GO:0009909 : regulation of flower development
 (<https://www.ebi.ac.uk/QuickGO/term/GO:0009909>)
 GO:0048579 : negative regulation of long-day photoperiodism, flowering
 (<https://www.ebi.ac.uk/QuickGO/term/GO:0048579>)
 GO:0048571 : long-day photoperiodism
 (<https://www.ebi.ac.uk/QuickGO/term/GO:0048571>)
 GO:0048576 : positive regulation of short-day photoperiodism, flowering
 (<https://www.ebi.ac.uk/QuickGO/term/GO:0048576>)
 GO:0048572 : short-day photoperiodism
 (<https://www.ebi.ac.uk/QuickGO/term/GO:0048572>)

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

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

Deletion Size

1-9 bp

Molecular Details of the Mutation

1bp deletion at position 321Å ; presumptive protein truncated

Experimental Evidence

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

Main Reference

Variations in Hd1 proteins, Hd3a promoters, and Ehd1 expression levels contribute to diversity of flowering time in cultivated rice. (2009) (<https://pubmed.ncbi.nlm.nih.gov/19246394>)

Authors

Takahashi Y; Teshima KM; Yokoi S; Innan H; Shimamoto K

Abstract

Rice is a facultative short-day plant, and molecular genetic studies have identified the major genes involved in short-day flowering. However, the molecular mechanisms promoting the diversity of flowering time in cultivated rice are not known. We used a core collection of 64 rice cultivars that represent the genetic diversity of 332 accessions from around the world and studied the expression levels and polymorphisms of 6 genes in the short-day flowering pathway. The RNA levels of Heading date 3a (Hd3a), encoding a floral activator, are highly correlated with flowering time, and there is a high degree of polymorphism in the Heading date 1 (Hd1) protein, which is a major regulator of Hd3a expression. Functional and nonfunctional alleles of Hd1 are associated with early and late flowering, respectively, suggesting that Hd1 is a major determinant of variation in flowering time of cultivated rice. We also found that the type of Hd3a promoter and the level of Ehd1 expression contribute to the diversity in flowering time and Hd3a expression level. We evaluated the contributions of these 3 factors by a statistical analysis using a simple linear model, and the results supported our experimental observations.

Additional References

RELATED GEPHE

Related Genes

9 (DTH2, EARLY FLOWERING 3/Hd17, Hd6a, PRR37 pseudoresponse regulator protein 37, se5, Early flowering1 (EL1), HEADING DATE 1, Ehd1 (Response regulator), Ghd7)
 (<https://www.gephebase.org/search-criteria?/or+Taxon ID=~4530^/and+Trait=Flowering time/and+groupHaplotypes=true#gephebase-summary-title>)

Related Haplotypes

5 (<https://www.gephebase.org/search-criteria?/or+Gene Gephebase=~Hd1^/and+Taxon ID=~4530^/or+Gene Gephebase=~Hd1^/and+Taxon ID=~4530^#gephebase-summary-title>)

EXTERNAL LINKS

COMMENTS