

IDENTIFICATION

Species: *Panicum hallii* HAL

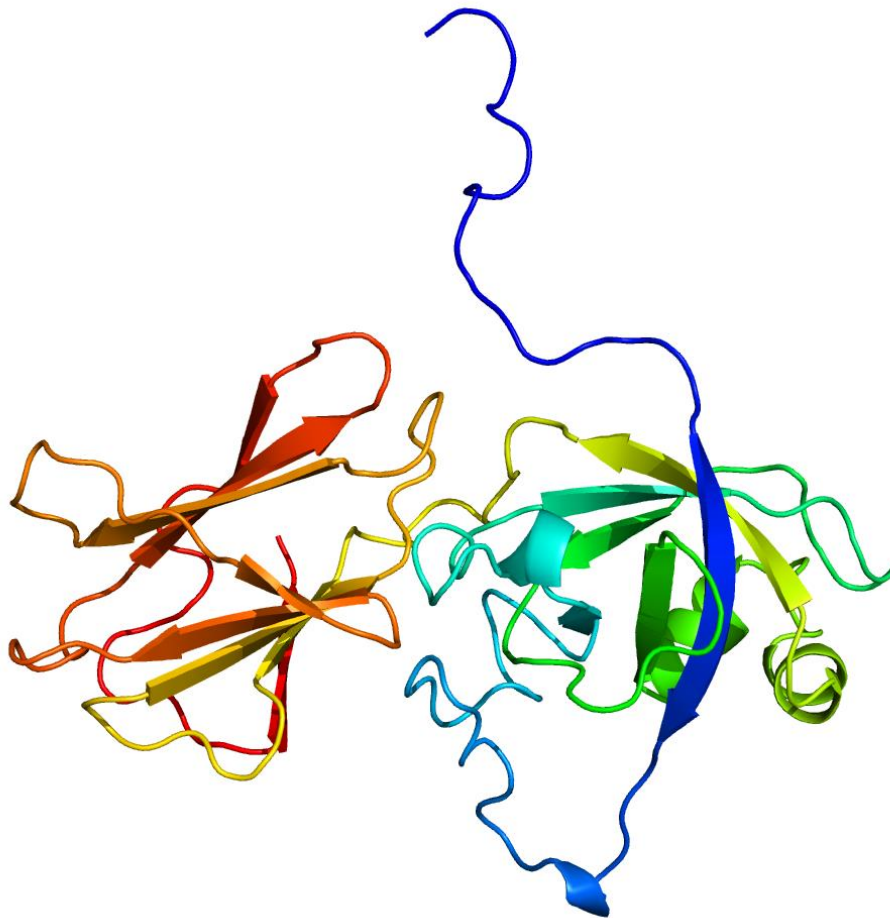
Locus: PhHAL.3G448100

Gene Model: PhHAL.3G448100.1.p

Description: PhhEXPB-08

Family: Beta Expansin

3D structure:



GENOME DATABASES

Phytozome: https://phytozome-next.jgi.doe.gov/info/PhaliiHAL_v2_1

KEGG:-

EXTERNAL RESOURCES

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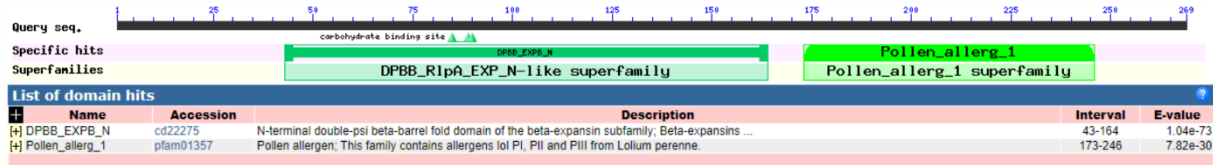
GENE STRUCTURE



Legend:

Exon

DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>PhhEXPB-08

MAAASTHLVAVAVVLAALVGGAWCGPPKVPPGKNISADCDGKWLEAKATWYGKP
TGAGPDDNGGACGYKEV NKAPFNMGACGNSPIFKDGLGCGSCYEIKCDKPAECSG
EPVIVYITDMNYEPIAA YHFDLAGTAFGAMA KKGEEKLRKAGIIDMQFRRVKCKYP
ADTKIAFHVEKGCNPNYLALLVKYAAGDGDIVGVDIKEKGAKEYQSLKHSWGAIWR
MDAPKPIKGPISIRITSEGGKTLEQEDVIPEGWKPD TLYPSKLQF*

CDS (coding sequence)

>PhhEXPB-08

ATGGCGGCGGCGTTCGACGCATCTTGTGCGGTGGCGGTGGTGCTCGCGGCGCTGG
TGGGCGGCGCATGGTGC GGTCGCCCAAGGTTCCCCGGGCAAGAACATCTCGGC
AGACTGCGACGGCAAGTGGCTGGAGGCCAAGGCGACATGGTACGGCAAGCCGAC
AGGCGCGGGGCCGACGACAACGGCGGCGCCTGCGGGTACAAGGAGGTGAACA
AGGCTCCCTTCAACGGCATGGGGGCGTGCGGCAACTCGCCCATCTTCAAGGACGG
CCTCGGCTGCGGCTCCTGCTACGAGATCAAGTGC GACAAGCCCGCCGAGTGCTCG
GGCGAGCCCGTCATCGTCTACATCACCGACATGAACTACGAGCCCATCGCCGCCT
ACCACTTCGACCTGGCCGGCACGGCCTTTGGAGCCATGGCCAAGAAGGGGGAGG
AGGAGAAGCTGCGCAAGGCGGGCATCATCGACATGCAGTTCCGCCGCGTCAAGT
GCAAGTACCCGGCCGACACCAAGATCGCCTTCCACGTCGAGAAGGGCTGCAACC
CCA ACTACCTGGCGCTGCTCGTCAAGTACGCCGCCGGCGACGGCGACATCGTCCG
CGTCGACATCAAGGAGAAGGGCGCCAAAGAGTACCAGTCCCTGAAGCACTCTTG
GGGCGCCATCTGGAGGATGGACGCCCCCAAGCCGATCAAGGGCCCCATCTCCATC
CGCATCACCGAGCGAGGGAGGCAAAACGCTCGAACAGGAGGATGTCATCCCCGAA
GGCTGGAAGCCCGACACCCTTACCCCTCCAAGCTCCAGTTCTGA

Nucleotide

>PhhEXPB-08

ATGGCGGCGGCGTTCGACGCATCTTGTGCGGTGGCGGTGGTGCTCGCGGCGCTGG
TGGGCGGCGCATGGTGC GGTCGCCCAAGGTTCCCCGGGCAAGAACATCTCGGC
AGACTGCGACGGCAAGTGGCTGGAGGCCAAGGCGACATGGTACGGCAAGCCGAC

AGGCGCGGGGCCCCGACGACAACGGCGGGCGCCTGCGGGTACAAGGAGGTGAACA
AGGCTCCCTTCAACGGCATGGGGGCGTGCGGCAACTCGCCCATCTTCAAGGACGG
CCTCGGCTGCGGCTCCTGCTACGAGATCAAGTGCGACAAGCCCGCCGAGTGCTCG
GGCGAGCCCGTCATCGTCTACATCACCGACATGAACTACGAGCCCATCGCCGCCT
ACCACTTCGACCTGGCCGGCACGGCCTTTGGAGCCATGGCCAAGAAGGGGGAGG
AGGAGAAGCTGCGCAAGGCGGGCATCATCGACATGCAGTTCCGCCGCGTCAAGT
GCAAGTACCCGGCCGACACCAAGATCGCCTTCCACGTCGAGAAGGGGCTGCAACC
CCAACCTACCTGGCGCTGCTCGTCAAGTACGCCGCCGGCGACGGCGACATCGTCGG
CGTCGACATCAAGGAGAAGGGCGCCAAAGAGTACCAGTCCCTGAAGCACTCTTG
GGGCGCCATCTGGAGGATGGACGCCCCCAAGCCGATCAAGGGCCCCATCTCCATC
CGCATCACAGCGAGGGAGGCAAAACGCTCGAACAGGAGGATGTCATCCCCGAA
GGCTGGAAGCCCGACACCCTCTACCCCTCCAAGCTCCAGTTCTGA