

## IDENTIFICATION

**Species:** *Arabidopsis halleri*

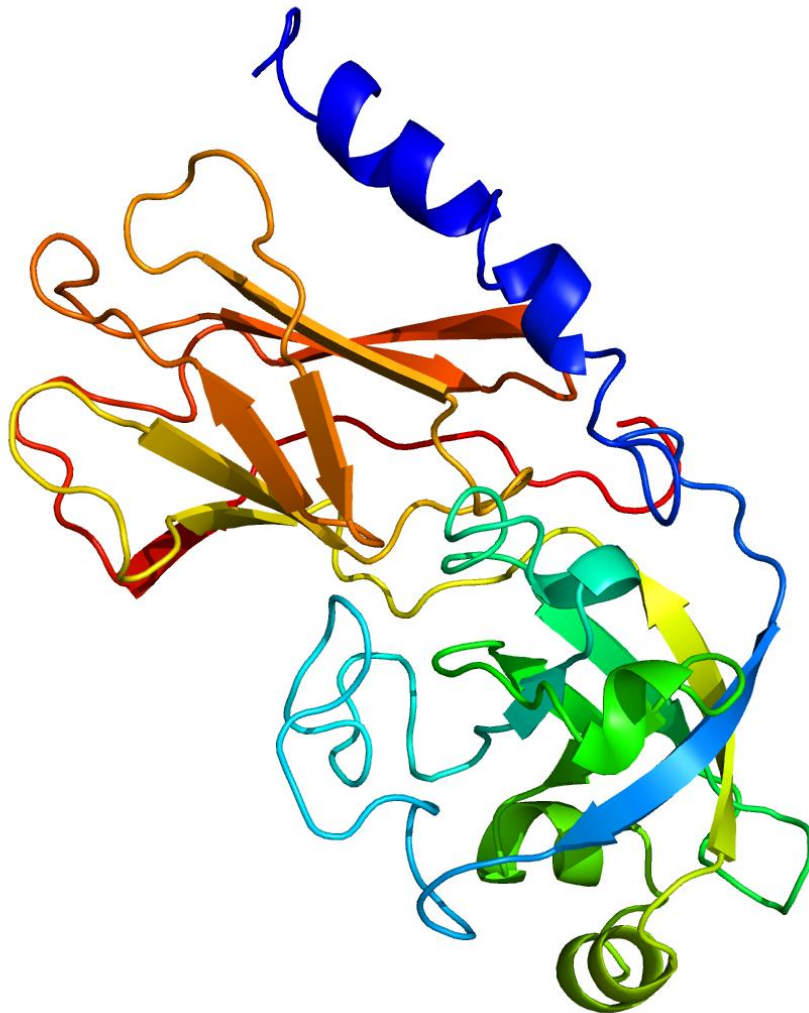
**Locus:** Araha.12096s0002

**Gene Model:** Araha.12096s0002.1

**Description:** AhEXPB-01

**Family:** Beta Expansin

**3D structure:**



## GENOME DATABASES

Phytozome: [https://phytozome-next.jgi.doe.gov/info/Ahalleri\\_v1\\_1](https://phytozome-next.jgi.doe.gov/info/Ahalleri_v1_1)

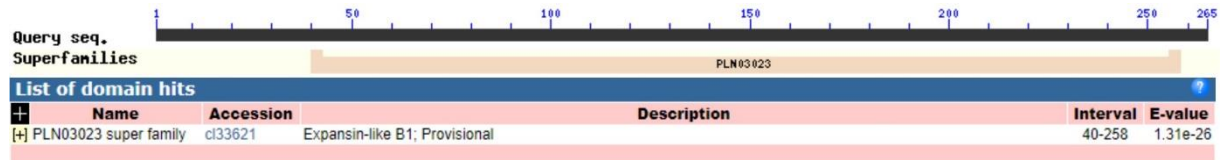
## EXTERNAL RESOURCES

[https://plants.ensembl.org/Arabidopsis\\_halleri/Info/Index](https://plants.ensembl.org/Arabidopsis_halleri/Info/Index)

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>AhEXPB-01

MASSSLKCLSSIVLTTFFAISLKPCSGHNKTHWNTAGITWYGDREGPGSTGGACGY  
GDAVAKHPYRCMVSAGGPSLFDGKGGACGYRLKCDHPLCTKKPIKVMISDECPGC  
TKEAVHFDLSGKAFGALAKRGKGDQLRNLGELKVS YKRACCKHPKAKIAIHVDSGA  
NPYYMSFAVKFANGDGNFACIEVQPAGGKYLKMEEMRSAVWRLSPGVPLKGFNIR  
LTSAVSGKKIVAKGVIPEKWSPGAIYHSKVNFPVQRKQK\*

### CDS (coding sequence)

>AhEXPB-01

ATGGCTTCCTCATCTCTCAAATGTTTGTCTTCTATCGTTGTTCTTACAACCTTCTTT  
GCAATCTCATTGAAGCCTTGTCTGGCCACAATAAAACCCACTGGAACACCGCCG  
GCATCACTTGGTATGGCGACCGCGAAGGTCCTGGCAGCACAGGAGGAGCTTGTG  
GATATGGTGATGCAGTGGCAAAGCACCCGTACAGATGTATGGTTTCAGCCGGAG  
GACCTTCATTGTTCAAAGATGGAAAGGGTTGTGGGGCATGTTACAGGCTTAAATG  
CGACCATCCGTTGTGCACGAAAAAGCCGATTAAGGTGATGATATCGGATGAGTGT  
CCCGGCTGTACGAAGGAGGCTGTTCAATTTGATCTTAGTGGCAAGGCCTTTGGTG  
CATTGGCCAAACGAGGCAAGGGCGATCAACTACGCAACCTTGGAGAACTAAAAG  
TTAGTTACAAACGTGCATGTTGCAAACACCCGAAGGCTAAGATCGCTATCCATGT  
CGACTCCGGAGCAAATCCTTACTACATGTCATTTCGAGTTAAGTTTGCAAACGGT  
GATGGGAACTTCGCTGCATCGAGGTCCAACCGGCAGGAGGAAAGTATTTGAAA  
ATGGAGGAAATGAGATCCGCTGTTTGGAGACTAAGCCCTGGTGTTCCTTTGAAGG  
GTCCGTTCAACATCAGGCTTACCTCCGCGGTCTCTGGTAAGAAGATCGTTGCGAA  
AGGCGTTATCCCTGAAAAGTGGAGCCCTGGTGCTATTTACCACTCTAAGGTTAAC  
TTCCCCGTTCAAAGGAAGCAGAAATGA

## Nucleotide

>AhEXPB-01

ATGGCTTCCTCATCTCTCAAATGTTTGTCTTCTATCGTTGTTCTTACAACCTTTCTTT  
GCAATCTCATTGAAGCCTTGTTCTGGCCACAATAAAACCCACTGGAACACCGCCG  
GCATCACTTGGTATGGCGACCGCGAAGGTCCTGGCAGCACAGGTAATTAATCAAA  
CCATTAATTCTTCCAATAAACTAATAAAACTTGAAAACATTTGTCCATATTTACCT  
ATGTACGGTTCGTAATAATATGTATACAGGAGGAGCTTGTGGATATGGTGATGCAG  
TGGCAAAGCACCCGTACAGATGTATGGTTTCAGCCGGAGGACCTTCATTGTTCAA  
AGATGGAAAGGGTTGTGGGGCATGTTACAGGGTATTTCTATAAAACATTTGCAAT  
TCATTGAAAATGATATTGTAATGTATAACAAATATACTAACCATGCATGTTTATAT  
GTTATAGCTTAAATGCGACCATCCGTTGTGCACGAAAAAGCCGATTAAGGTGATG  
ATATCGGATGAGTGTCCCGGCTGTACGAAGGAGGCTGTTCATTTTGATCTTAGTG  
GCAAGGCCTTTGGTGCATTGGCCAAACGAGGCAAGGGCGATCAACTACGCAACC  
TTGGAGAACTAAAAGTTAGTTACAAACGGTACGTATTGTAAAAAACCGAGTCGGT  
TCGGTTTGGTCAAGTTTTTGGATAGAAAATATATATATATATATATATATATATAT  
TTATATAAATATATATATATGCTTAATTCTAATCAATATCATACATATATATATAT  
GCAGTGCATGTTGCAAACACCCGAAGGCTAAGATCGCTATCCATGTGCGACTCCGG  
AGCAAATCCTTACTACATGTCATTCGCAGTTAAGTTTGCAAACGGTGATGGGAAC  
TTCGCCTGCATCGAGGTCCAACCGGCAGGAGGAAAGTATTTGAAAATGGAGGAA  
ATGAGATCCGCTGTTTGGGAGACTAAGCCCTGGTGTTCCTTTGAAGGGTCCGTTCA  
ACATCAGGCTTACCTCCGCGGTCTCTGGTAAGAAGATCGTTGCGAAAGGCGTTAT  
CCCTGAAAAGTGGAGCCCTGGTGCTATTTACCACTCTAAGGTTAACTTCCCCGTT  
AAAGGAAGCAGAAATGA