

## IDENTIFICATION

**Species:** *Ananas comosus*

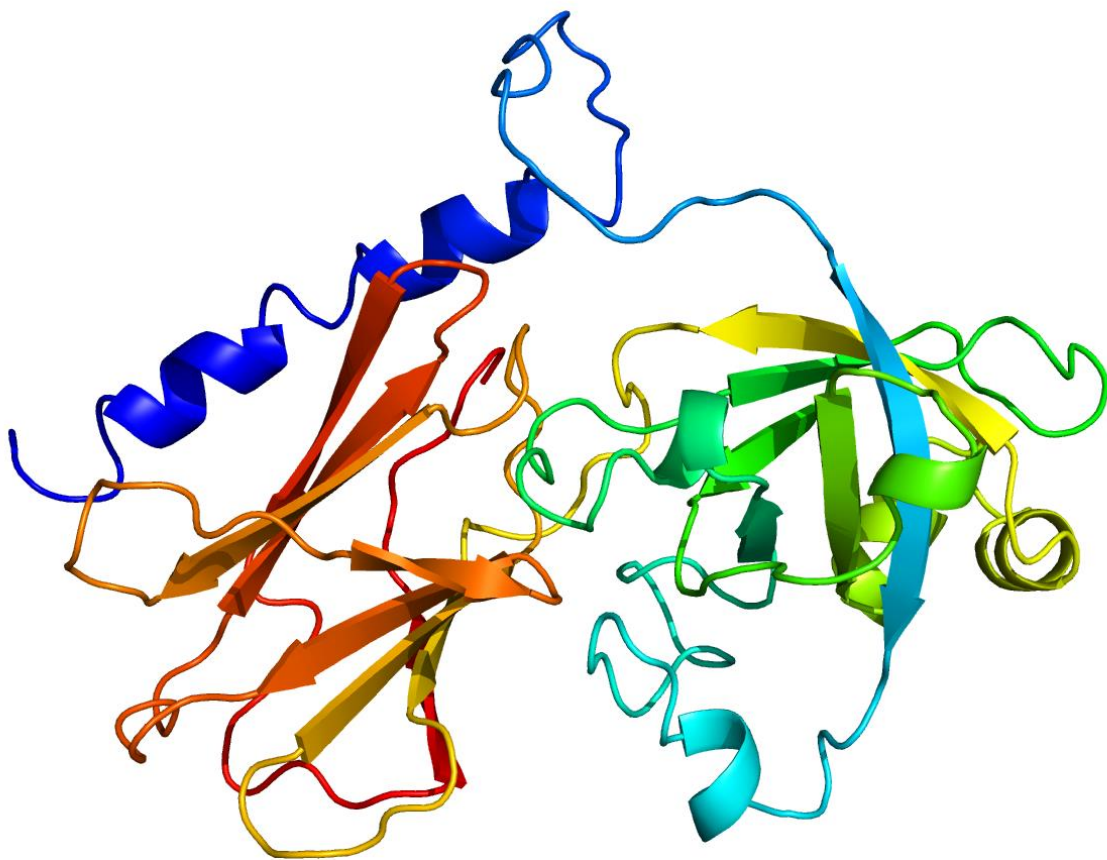
**Locus:** Aco011717

**Gene Model:** Aco011717.1

**Description:** AncEXPB-06

**Family:** Beta Expansin

**3D structure:**



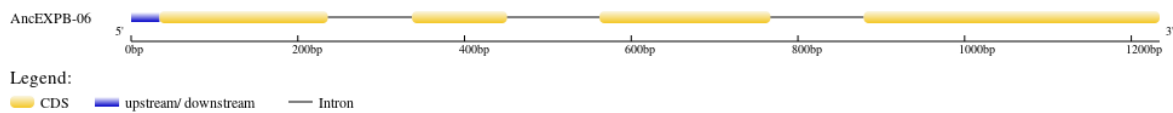
## GENOME DATABASES

Phytozome: [https://phytozome-next.jgi.doe.gov/info/Acomosus\\_v3](https://phytozome-next.jgi.doe.gov/info/Acomosus_v3)

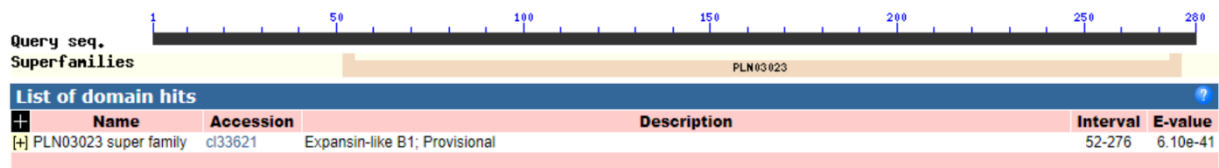
## EXTERNAL RESOURCES

[http://pineapple.angiosperms.org/pineapple/html/index.html#:~:text=The%20PGD%20\(Pineapple%20Genomics%20database,genomics%20and%20CAM%20pathway%20genes..](http://pineapple.angiosperms.org/pineapple/html/index.html#:~:text=The%20PGD%20(Pineapple%20Genomics%20database,genomics%20and%20CAM%20pathway%20genes..)

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>AncEXPB-06

MAPLPHLFSSSIFLLILSLFSLINPCFCFHPKKLSNYTYTNPSAWSSGGATWYGSPLG  
AGSDGGGYQDAVDKPPFSSMIAAGSPSIFKSGKGCYQVKCTGNASESPVTVVVTDE  
CPGGPCLDEPAHFDMSGTAFGAMARPGQADQLRNAGRIQIQYSRVECNHYHGIDVAF  
HVDGGSNPNYFAVLIEYEGGDGDLAAVDIMQQGSTSWVPMQQSWGAVWRLNSGPA  
MRPPFSLRLTSGKSDNTLVAKNVIPAGWCAGATYRSVVNYST\*

### CDS (coding sequence)

>AncEXPB-06

ATGGCTCCTCTTCTCCTCTTCTCCTCTTCTATTTTTCTCCTTATTCTCTCCTTATTT  
TCGCTCTCTCATTAATCCCTGCTTTTGTTCCTCCCAAAGAAGCTCTCGATATATATC  
ACCCTCAGCTTGGTCTAGTGGTGGAGCCCTGGTGGAAAGCCATTAGGAGCGGGGA  
GTGGGTGGTGCTTGTGGGTCAAGATGCGGTCGATAAGCCGCCGTTCTCGTCAATG  
ATAGCTGCAGGAAGCCCTTCGATATTCAAGTCCGGGAAGGGCTGCGGGGCTTGT  
ATCAGGTGAAATGCGGGCAATGCGGCATGTTTCGGAGAGCCCGGTGCGTCGTGGT  
CCGGAATGCCCGGCGGGCCGTGCCTCGGACCGCGCTTCGATGAGCGGGCGCGTT  
CGGAGCCATGGCGAGGCCAGGCCAAGCCGATCAGCTTCGAAGCCGGAAGGATCC  
AAATAGTAGCAGGGTGGAGTGCATCGGGATCGGTCGCGTTCATGTGGGGAGGT  
TCGACCGATTCGCCGTTTTGATCGAGTGAGGGAGGCGGGCGCTCGCGGGCGGTTCG  
ATATCATGCAGCAGGGATCGGTCGTGGGTTCGATGCAGCAGTCGTGGGGCGCCG  
TCTGGAGGCTGAATTCCGGTCCGGCGATGCGCGCCGTTCTCGCTGCGTGGTTCGGG  
GAAATCCGACCTCGTCGCGAAGAGTGATTCCGGCCGGGTGGTGCGCCGGCGCGCT  
CGCTCCGTGGTGATAGCCTAA

## Nucleotide

>AncEXPB-06

CCTTCTCCTTGCTGAGTATAAGCCAATGGCTCCTCTTCCTCCTCTTCTCCTCTTCTA  
TTTTTCTCCTTATTCTCTCCTTATTTTCGCTCTCACTCATTAAATCCCTGCTTTTGTTT  
CCCCAAAGAAGCTCTCGATATATATCACCCCTCAGCTTGGTCTAGTGGTGGAGCCC  
TGGTGGAAGCCATTAGGAGCGGGGAGTGGGTATAAATATGTATCATCTGTATAT  
CTATATATATGTATGTAGCTTGATCTCTCTCTCAATGTGTGCATATATATATATAT  
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AATGATAGCTGCAGGAAGCCCTTCGATATTCAAGTCCGGGAAGGGCTGCGGGGC  
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CGGCATGTTTCGGAGAGCCCGGTGCGTCGTGGTCCGGAATGCCCCGGCGGGCCGTG  
CCTCGGACCGCGCTTCGATGAGCGGGCGCGTTCGGAGCCATGGCGAGGCCAGGC  
CAAGCCGATCAGCTTCGAAACGCCGGAAGGATCCAAATAGTAGCAGGTGCGAAT  
TGAATCTGCGTGCTGATAAAAATATCGAGCTCAAATAAGAATTTGGAAAATTTAAA  
GAATAAATTTTAATTGTTTCCATGGTTTGTTCAGGGTGGAGTGCATCGGGATC  
GGTCGCGTTCATGTGGGGAGGTTCGACCGATTCGCCGTTTTGATCGAGTGAGG  
GAGGCGGGCGCTCGCGGCGGTCGATATCATGCAGCAGGGATCGGTTCGTGGGTTC  
CGATGCAGCAGTCGTGGGGCGCCGTCTGGAGGCTGAATTCGGTCCGGCGATGCG  
CGCCGTTCTCGCTGCGTGGTTCGGGGAAATCCGACCTCGTCGCGAAGAGTGATTCC  
GGCCGGGTGGTGCGCCGGCGCGCTCGCTCCGTGGTGATAGCCTAA