

## IDENTIFICATION

**Species:** *Brachypodium distachyon*

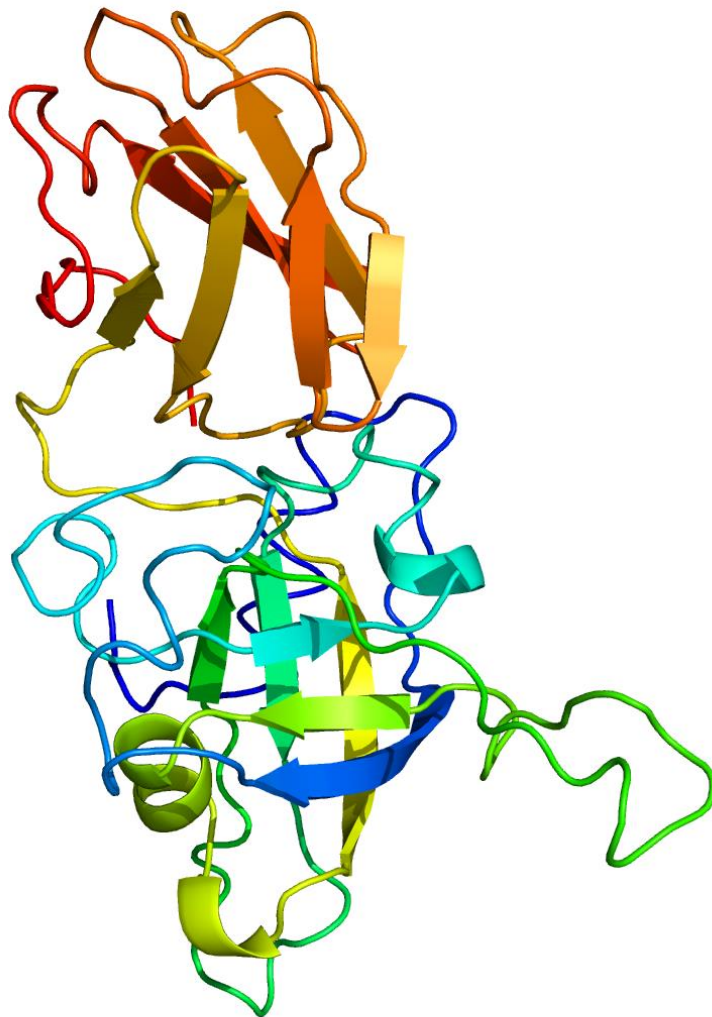
**Locus:** Bradi1g74710

**Gene Model:** Bradi1g74710.1.p

**Description:** BdEXPA-05

**Family:** Alpha Expansin

**3D structure:**



## GENOME DATABASES

Phytozome: [https://phytozome-next.jgi.doe.gov/info/Bdistachyon\\_v3\\_1](https://phytozome-next.jgi.doe.gov/info/Bdistachyon_v3_1)

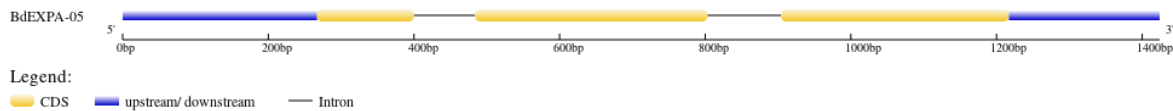
KEGG: <https://www.genome.jp/entry/T01717>

## EXTERNAL RESOURCES

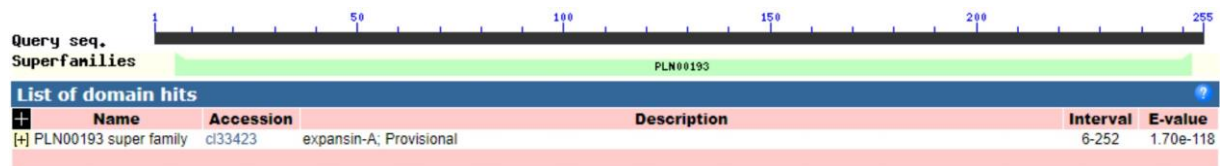
<https://brachypodium.org/>

[https://archive.gramene.org/species/brachypodium/brachypodium\\_intro.html](https://archive.gramene.org/species/brachypodium/brachypodium_intro.html)

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>BdEXPA-05

MTGKRSVRLFAAVLAVALCFAPAKSDWLPATATFYGEPDASGTMGGACGYGNLYD  
QGYGVSNAAALSTALFNDGASCGQCYLIICDTSKTGWCKPGTSATVSATNFCPPNWTL  
PNDNGGWCNPPRFHFDMSQPSWETIAIYRAGIVPVLYQQVKCWRYGGVRFTIAGFN  
YFELVLLTNVGGSGSVKSMAMVKGPN TGWIQMSRNWGANWQCLAGLEKQPLSFALT  
STGGQYLVFQDAVPAGWQFGQTFSTYRQFDY\*

### CDS (coding sequence)

>BdEXPA-05

ATGACGGGGAAACGTTCCGTGCGTCTGTTTCGCCGCTGTTCTGGCAGTAGCACTGT  
GCTTTGCGCCGGCCAAGTCCGACTGGCTTCCGGCCACCGCCACCTTCTACGGCGA  
ACCGGACGCCTCCGGCACCATGGGTGGTGC GTGGGTACGGGAACCTGTACGA  
CCAGGGTTACGGGGTGAGCAACGCGGCGCTGAGCACGGCGCTGTTCAACGACGG  
GGCGTCGTGCGGGCAGTGCTACCTGATCATCTGCGACACGAGCAAGACCGGGTG  
GTGCAAGCCC GGAACGTCCGCCACGGTGTCCGCCACCAACTTCTGCCCGCCCAAC  
TGGACGCTCCCCAACGACAACGGCGGCTGGTGCAATCCTCCCCGCTTCCACTTCG  
ACATGTCCCAACCTTCCTGGGAGACCATCGCCATCTACCGCGCCGGCATCGTCCC  
CGTCTCTACCAACAGGTGAAGTGTGGAGGTACGGAGGGGTGAGGTTACGATT  
GCAGGGTTCAACTACTTCGAGCTGGTGTGCTGACCAATGTCCGGAGGGAGCGGGT  
CGGTGAAGAGCATGGCCGTGAAGGGGCCTAACACGGGGTGGATCCAGATGTCCA  
GGA ACTGGGGGGCAAATTGGCAGTGCCTGGCTGGGCTGGAGAAGCAGCCGCTCA  
GCTTCGCGCTCACCTCCACCGGCGGGCAGTACCTCGTCTTCCAGGACGCCGTTCC  
CGCCGGGTGGCAGTTCGGACAGACCTTCAGCACCTATCGCCAGTTCGACTACTAG

### Nucleotide

>BdEXPA-05

TCCGCGGACGTTGCAGAGCCATGTGCAAACATACGATCGTTGGAACATCGGTTGC  
ATTTGCAGCAATTATTCCTCTCGATCCATCGAACGGAATCCAAGCCGGGCTCCTG  
AGTTCCCCTATTTAAGCAGCAGCAATGTCGCAATCATTCTCACCTCCAAAATCC  
AGTCGCCACAACATCCAAAGCTAGCTGCAAAGCCGTGCTACTTATCTTGAGCACA  
ACACACAGTCTTCTACTGCTAGCTGAGTCTGCGGAGTCCAGGAGAAGATGACGGG

GAAACGTTCCGTGCGTCTGTTTCGCCGCTGTTCTGGCAGTAGCACTGTGCTTTGCGC  
CGGCCAAGTCCGACTGGCTTCCGGCCACCGCCACCTTCTACGGCGAACCGGACGC  
CTCCGGCACCATGGGTACGCTAAAAATCAAGCCCCAACACCTCCGGAACGACC  
AAAATGGACGTAATAATGTGGTCTGATTTCTGAATATTTATACAGGTGGTGGCGTGT  
GGGTACGGGAACCTGTACGACCAGGGTTACGGGGTGAGCAACGCGGGCGCTGAGC  
ACGGCGCTGTTCAACGACGGGGCGTCGTGCGGGCAGTGCTACCTGATCATCTGCG  
ACACGAGCAAGACCGGGTGGTGCAAGCCCGGAACGTCCGCCACGGTGTCCGCCA  
CCAACTTCTGCCCGCCAACTGGACGCTCCCCAACGACAACGGCGGGCTGGTGCAA  
TCCTCCCCGCTTCCACTTCGACATGTCCCAACCTTCCTGGGAGACCATCGCCATCT  
ACCGCGCCGGC ATCGTCCCCGTCTCTACCAACAGTACGCAAATCTAGCTCGCGT  
CCATCAATCAA AATTGCTCCCAATAACACGATGGATGAACTCATGAACTGACGAT  
GTCTTGTGGGATGTGCGTTGTGCAGGGTGAAGTGCTGGAGGTACGGAGGGGTGA  
GGTTCACGATTGCAGGGTTCAACTACTTCGAGCTGGTGCTGCTGACCAATGTCCG  
AGGGAGCGGGTCGGTGAAGAGCATGGCCGTGAAGGGGCCTAACACGGGGTGGAT  
CCAGATGTCCAGGAACTGGGGGGCAAATTGGCAGTGCCTGGCTGGGCTGGAGAA  
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GACGCCGTCCCCGCCGGGTGGCAGTTCGGACAGACCTTCAGCACCTATCGCCAGT  
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CCTGTTAGAACGAACTGTGTAAAAGTCACTTATTCAA AATTCATCATTGATTATCT  
GAACGCCAAATATTTACATACTGGAGATTTTTCCAGCCATCCTAATACAG