

IDENTIFICATION

Species: *Physcomitrium patens*

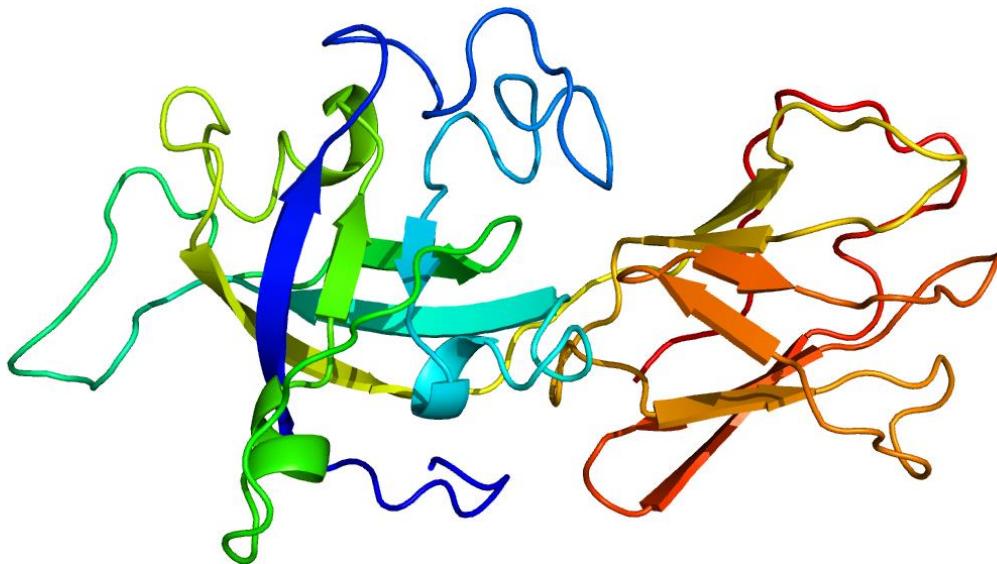
Locus: Pp3c7_12810V3

Gene Model: Pp3c7_12810V3.1.p

Description: PpEXPA-03

Family: Alpha Expansin

3D structure:



GENOME DATABASES

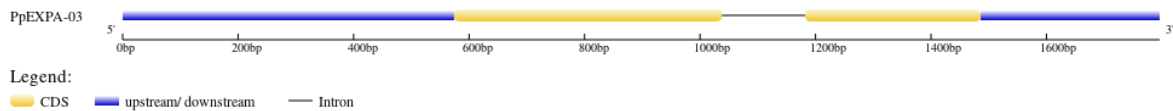
Phytozome: https://phytozome-next.jgi.doe.gov/info/Ppatens_v3_3

KEGG: <https://www.genome.jp/entry/gn:T01041>

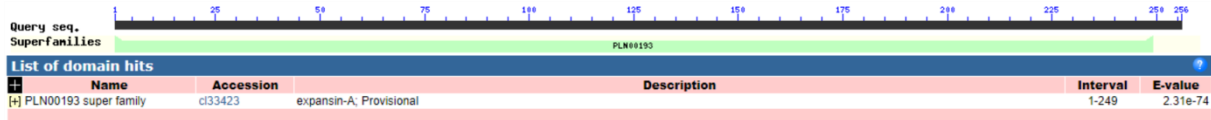
EXTERNAL RESOURCES

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GENE STRUCTURE



DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>PpEXPA-03

MSGLTRVEVTFVMFVATLVIPSVLGMMPVGRDAHITYYGGSPNGGGTQGGACAYQNT
FSLGYGAMTAALSSPLFEGGAACGACYQLQCKRVQETRTRVKNWCWSYSRTITITAT
NLCPPGSAGAWCDPPRHFDLTPAFLTLARREGGVAPVLYRRVKCVKRGGIRFTIG
GNPWFLMILHNVAGAGDVRAVRIKTPSTDWIPMYRNWGALWTVQRKLSGPLSFQIT
AGDRRQITINSAVGNAWKFGQTWEGHNFR*

CDS (coding sequence)

>PpEXPA-03

ATGAGCGGCCTAACTCGTGTGGAGGTCACCTTTTGTTCATGTTTGTGCTACGCTGGT
TATACCGTCGGTGCTTGGAAATGCCGGTTGGATGGCGCGATGCGCACATCACCTAT
TACGGGTCACCTAACGGTGGAGGAACACAGGGTGGCGCTTGC GCGTATCAGAAC
ACATTCTCGCTGGGCTACGGTGCCATGACTGCGGGCGTTGAGCTCGCCTTTGTTTGA
GGGTGGAGCGGCTTGTGGGGCTTGTACCAGCTTCAGTGCAAACGAGTGCAAGA
AACCCGTA CTGTGAAGAACTGGTGCTGGAGTTATTCTCGAACCATCACCATCACT
GCCACTAACCTCTGTCCACCAGGCTCTGCCGGGGCGTGGTGC GACCCTCCAAGGC
ACCATTTGACTTGACAATGCCTGCCTTCTAACCTGGCGAGACGTGAAGGTGG
TGTGGCTCCAGTGCTATACAGAAGGGTTAAATGTGTCAAACGCGGTGGAATTCGG
TTCACCATAGGAGGCAATCCTTGGTTTCTGATGATCCTCATCCACAACGTCGCTGG
CGCAGGCGATGTAAGGGCCGTTAGAATCAAGACCCCGTCCACTGACTGGATTCCC
ATGTACCGCAACTGGGGTGC GTTGTGGACAGTACAAAGGAACTCAGCGGGCCC
CTCTCATTTAGATCACTGCCGGCGACCGCCGTCAAATCACCATCAACAGTGCAG
TAGGCAATGCATGGAAGTTTGGACAAACCTGGGAGGGCCACA ACTTCCGCTGA

Nucleotide

>PpEXPA-03

GGCACAGATTTGATACTGTGGCTCCTTCGTCTGCGCAGTCAAGCGTAGCGTTCA
TAACTTCCCTTAGGAGAGAGGTTATCTATCATATATTGAGTTCGGCCAAAACCAC
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TCTTGTA ACTGTTGCTTGTCTCGTACTGTTACAGCAGTGTGAGGATTCAATCCG
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TTGAGCTCGCCTTTGTTTGAAGGTGGAGCGGCTTGTGGGGCTTGCTACCAGCTTCA
GTGCAAACGAGTGCAAGAAACCCGTAAGTGAAGAAGTGGTGGTGGAGTTATTCT
CGAACCATCACCATCACTGCCACTAACCTCTGTCCACCAGGCTCTGCCGGGGCGT
GGTGCGACCCTCCAAGGCACCATTTGCACTTGACAATGCCTGCCTTCCTAACCT
GGCGAGACGTGAAGGTGGTGTGGCTCCAGTGCTATACAGAAGGTACATTACCGC
GCTTTTGCCAATCTATGGTGTGGTCTGGTGCAGAATCTTAGAATTTTGTGCACAT
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AAATACAGTCTGTGGGTGCAGGGTTAAATGTGTCAAACGCGGTGGAATTCGGTTC
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CAGGCGATGTAAGGGCCGTTAGAATCAAGACCCCGTCCACTGACTGGATTCCCAT
GTACCGCAACTGGGGTGCCTTGTGGACAGTACAAAGGAAACTCAGCGGGCCCCT
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