

IDENTIFICATION

Species: *Physcomitrium patens*

Locus: Pp3c11_12000V3

Gene Model: Pp3c11_12000V3.1.p

Description: PpEXPA-18

Family: Alpha Expansin

3D structure:



GENOME DATABASES

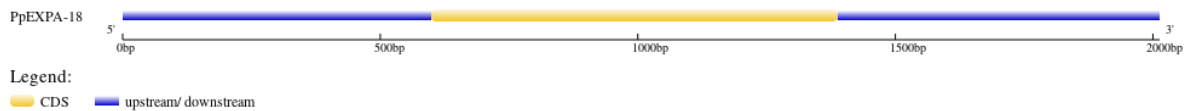
Phytozome: https://phytozome-next.jgi.doe.gov/info/Ppatens_v3_3

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

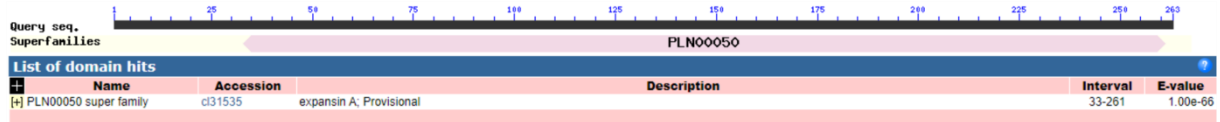
EXTERNAL RESOURCES

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GENE STRUCTURE



DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>PpEXPA-18

MKMDARSGMKLAVVLLVHLALCASPVFGAPPGGWINGRITYYGSPNGGGTQGGAC
GYQNTVSLGYGFMTAALSTTLFKGGAACGACYQLQCAPVSETPSGLLKRNWCWQV
GRSILVTATNLCPPGSSGGWCNPPQHFDLPMPAFLALARREGGVVPVYYRKNCA
RKGIRFTVGGNPWFLMILIHNVGGAGDVVAVKVKCPTSGWYDMYRNWGALWTV
QKKMNGPLSFAITSDGRTVTTYNAVGNGWKFGQTWEQAQYR*

CDS (coding sequence)

>PpEXPA-18

ATGAAGATGGACGCTCGATCCGGTATGAAGCTCGCGGTGGTCTGCTCGTTCACC
TGGCGCTCTGCGCGTCTCCTGTGTTCCGGTGCCCCGCCTGGAGGATGGATAAATGG
GCGCATCACGTA CTATGGTTCGCCAAATGGCGGAGGGACGCAGGGCGGTGCTTG
CGGGTATCAGA AACTGTATCGTTGGGATATGGCTTTATGACAGCAGCACTGAGT
ACCACTCTGTTTAAAGGTGGAGCGGCCTGTGGTGCGTGCTACCAGCTGCAGTGTG
CCCCAGTCAGCGAAACCCCGAGTGGGTTACTCAAGAGGAACTGGTGCTGGCAGG
TGGGTAGAAGCATTCTCGTCACCGCTACCAACCTCTGTCCTCCAGGCTCTTCAGG
AGGATGGTGTAACCTCCGCAGCACCATTTGACTTGCCAATGCCTGCCTTCTTG
CCCTCGCTAGGCGTGAAGGGGGTGTGTTCCAGTCTACTACCGGAAGGTGAATTG
TGCGAGGAAGGGTGAATTCGATTCAGTGTGGGAGGCAATCCTTGTTCTTGATG
ATCCTTATTCACAACGTTGGCGGGGCAGGTGATGTGGTAGCTGTCAAGGTTAAGT
GCCCCACCTCCGGATGGTATGACATGTACCGGAACTGGGGTGCATTATGGACTGT
GCAGAAGAAGATGAATGGGCCACTTTCCTTCGCGATCACAACGAGCGACGGGCG
CACGGTGACCACCTACAATGCGGTGGGCAACGGTTGGAAGTTCGGGCAGACCTG
GGAGGGCGCCAATACAGGTGA

Nucleotide

>PpEXPA-18

GGGCTGCAGCGTTGGGTTTGGATATAAGCCAGTTGGGGGGACGTCTTCAACGCAT
CAATGCAGCAGCGCCATCTTGTCAACTGGGGGAGAGAGTTTGCATCGGGGTCCGG
ATTCATGTCATGAGGGTGGTATCTTGAGGGGTTGTGTTGAGTATTGGCCCTGGCC
AGATCTTCAGTCGCTCACTTGGCTCTTGTAAGTTGTCTGGACTGATCCACAGGTGC
AGTTTTGTGTCTTGGGAATCAAACCTGGGTTGTGTTGGTTTCCACGATGAATTCATT

CGCTTCTAGAGTATAACAATATTCAAAGCATTGGGGTACTAAGTAGAAGTATCCAA
GTTTGTCCCTTGATAAAAAGAGATTCTAGTTGCGGTGACAAGATTACAGAGAATTTT
GAAACGGCAATTTGAACCACTGCTACTTTTGAAGAACTGTCGAGTGTTTGAAGAA
TATCAACCAGTATGAAAAGGATGTTAATTTTCGTGAAAAGCCATGGTGTGTGGTCC
ACCGAGCGAAACGTGTGTAGATCAAACAGATGGTTCGGCATGAATCTGGATTATT
TGTTTCTGATTTGTTAATCAACGGTACTGAATTGCAGCCTTGACATCATGAAGATG
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GCGCGTCTCCTGTGTTTCGGTGCCCCGCCTGGAGGATGGATAAATGGGCGCATCAC
GTACTATGGTTCGCCAAATGGCGGAGGGACGCAGGGCGGTGCTTGCGGGTATCA
GAACACTGTATCGTTGGGATATGGCTTTATGACAGCAGCACTGAGTACCACTCTG
TTTAAAGGTGGAGCGGCCTGTGGTGCCTGCTACCAGCTGCAGTGTGCCCCAGTCA
GCGAAACCCCGAGTGGGTTACTCAAGAGGAACTGGTGTGCGCAGGTGGGTAGAA
GCATTCTCGTCACCGCTACCAACCTCTGTCTCCAGGCTCTTCAGGAGGATGGTGT
AACCTCCGCAGCACCATTTTCGACTTGCCAATGCCTGCCTTCTTGGCCCTCGCTAG
GCGTGAAGGGGGTGTGTTCCAGTCTACTACCGGAAGGTGAATTGTGCGAGGAA
GGGTGGAATTCGATTCAGTGTGGGAGGCAATCCTTGGTTCCTGATGATCCTTATTC
ACAACGTTGGCGGGGCAGGTGATGTGGTAGCTGTCAAGGTTAAGTGCCCCACCTC
CGGATGGTATGACATGTACCGAACTGGGGTGCATTATGGACTGTGCAGAAGAA
GATGAATGGGCCACTTTCCTTCGCGATCACAACGAGCGACGGGGCGCACGGTGACC
ACCTACAATGCGGTGGGCAACGGTTGGAAGTTCGGGCAGACCTGGGAGGGCGCC
CAATACAGGTGATCGACGCGTACTTCCAGGTTCGAGAGAGAATTCGACGGACGG
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GCTGAGCTTGCATCAGTCCGGTAGGCCGCGCTAGAGATCGAAAGGCACGGAACGT
CGAATATTATCCTCTCGGCACAGAGTCGAGGGTCTCTTCTTGCCAGTCTTTCGGAG
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AGACTCTGCTTCTTCCACCTGTTTTGT