

IDENTIFICATION

Species: *Sphagnum fallax*

Locus: Sphfalx0229s0003

Gene Model: Sphfalx0229s0003.1.p

Description: SfEXPB-05

Family: Beta Expansin

3D structure:



GENOME DATABASES

Phytozome: https://phytozome-next.jgi.doe.gov/info/Sfallax_v0_5

KEGG:-

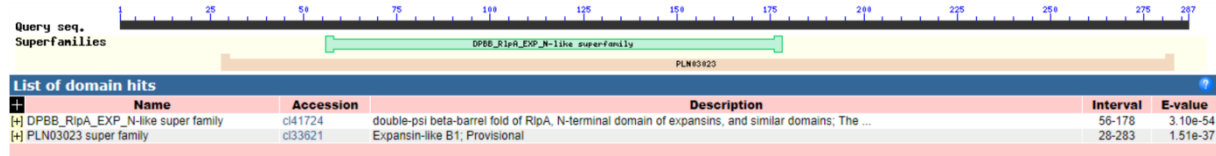
EXTERNAL RESOURCES

-

GENE STRUCTURE



DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>SfEXPB-05

MGVARFQQERLDDLLQLLQLRLLGVAILLVLLPCCINCQQQLSSSLLAQNGYDIDWQ
YGHATWYGNPYGDGSDGGACGYTSLNTPYGSNVAAGSAAVFSNGQGCGICYDVK
CTYSICNSQPTRIVITDFCPGGVYCSTDEVAFDLSGSAMD SLAVPGLESTLRDFGQYDI
QYMRVPCDYVGQNVAFVAVDAGSSPYWLSFAVR YEGGPGDIESVMIRQSGSYDWVA
MQHNWGASWMLIDYSGQAFRGPYDVQITAKLNGHSLIAWNAIPDYFQPGATYNSSV
QFLY*

CDS (coding sequence)

>SfEXPB-05

ATGGGCGTCGCGAGATTCCAGCAGGAGCGTCTTGATGATCTGCTGCAGCTGTTGC
AGCTTCGACTTCTGGGTGTGGCGATTCTTCTAGTCCTTCTCCTTGCTGCATTA
ACTGTCAGCAGCAGCTGAGTTCTAGTCTTCTTGACAGAATGGCTACGACATCGACTG
GCAGTATGGACATGCGACTTGGTATGGAAATCCTTACGGAGACGGCTCAGACGG
AGGAGCATGTGGGTATACCTCCCTGGAGAATACGCCATACGGCTCCAACGTCGCG
GCTGGAAGTGCTGCCGTTTTCTCCAATGGTCAAGGATGCGGAATCTGCTATGATG
TGAAATGCACATACTCAATTTGCAACTCGCAACCGACCAGAATCGTTATCACGGA
TTTCTGTCCC GCGGTGTGTATTGTAGCACCGACGAAGTTGCATTTGATCTCAGCG
GCAGTGCCATGGATAGCTTGGCCGTTCCCGGCCTGGAATCGACTCTACGCGACTT
CGGCCAATACGACATTCAGTACATGAGGGTGCCATGCGACTATGTTGGACAGAAC
GTTGCGTTCGCGGTTGATGCAGGCTCTAGTCCATACTGGCTCTCGTTTGCAGTCAG
ATACGAGGGAGGCCAGGCGACATCGAGTCTGTCATGATCCGACAGTCTGGAAG
CTACGATTGGGTGGCGATGCAACACA ACTGGGGTGCAAGTTGGATGTTGATCGAC
TACTCTGGACAAGCTTCCGAGGTCCTTACGACGTT CAGATCACAGCCAAGCTCA
ACGGCCATTCTCTCATCGCTGGAATGCCATTCCGGACTACTTCCAACCGGGAGC
TACCTACAATTCTAGCGTGCAGTTCTTATACTAG

Nucleotide

>SfEXPB-05

TTCTTCTTCTTGGTCTCTAATCCAAATCCAAATTTGACTTCTGATCTCGATTTGCA
GCAGCAGCAGCAGCAAGAAGCAGATTCTTCTGGAAGAGATCAAAACACTAAATA

TTATTCCTTCTCAGAAGAAATTATTTTATATTACTAGACTTGATCAAATGTGGTGG
TAATCTGCAAAAAATTGTCAACCCATCATGATCATCCCCAAGATTGGGTCACTGA
TAGAACAAGAAAATTACCCACAACACTCATGTACACTACAAGAGTAGCAGGAGTCC
AGAATTTTATATAAATCAGCTTTCTGTGCTGCAATCGTTGTCGCCCTCCATCTAGAG
AGTACAGTGTCTGATCAATAGCTAGCAGCTCGCGAGTCCAGTAGTCTGAGTCGTC
GCTGCTGCTGTTGCTGCAGTGATTGAAGCGGATGTTGCATTATTATAGCTCATCAT
CTAGTGTATCGCTTATAACCGTGATTACCATTGGGTTTCTGCCATTCTTCTTTTTTTT
TCTTTTTTTTTTCTTGGAGATTTTCTCTTGAATCCGTCTAGCTGTAGTTTGCGGTAG
CTGTAAGAGATGGGCGTCGCGAGATTCCAGCAGGAGCGTCTTGATGATCTGCTGC
AGCTGTTGCAGCTTCGACTTCTGGGTGTGGCGATTCTTCTAGTCTTCTTCTTGTCT
GCATTAAGTGTGAGCAGCAGCTGAGTCTAGTCTTCTTGCACAGAATGGCTACGA
CATCGACTGGCAGTATGGACATGCGACTTGGTATGGAAATCCTTACGGAGACGGC
TCAGACGGTGAGAGTTTGTTTACACTGTTTGATCACTAGTGCAAAATTTCTTCTCT
GATCAGCTTGCCACCACGACGAGTATTAAGTACTCGCAAGCAGTATTCGACTGAT
ATAAAGATGCAGTGAAGCGAAGCCTTCGCAAATGCTGCAGCACAACATCTGAAG
AAAGCTAGCTAGCTAGAGGCCGACTAATTACCTCTCATTTCATGTGCAGGAGGAGC
ATGTGGGTATACCTCCCTGGAGAATACGCCATACGGCTCCAACGTGCGGGCTGGA
AGTGCTGCCGTTTTCTCCAATGGTCAAGGATGCGGAATCTGCTATGATGTATGTA
ATGGATCGTCTCGTCTCACACTACCTGCAGGATCCAAAGTAGCTTGCTCTAGCTA
GACTGACCAGTTTGAAGAAGATTGATCAGGCATGTTGTAGCTAGCTCAAAGATTG
CTGCATGCATGCATACATACATCCTCTAGCTAGCTAGTACTGTGATGATCACTGAT
CAGTTCTACTGAAGCAGCTTTGGTTCGAGATCATATATACACGTAATGCGCTGCTG
AGATCATCCGTGCTCGATCTATGCGTGCAGGTGAAATGCACATACTCAATTTGCA
ACTCGCAACCGACCAGAATCGTTATCACGGATTTCTGTCCCGGCGGTGTGTATTG
TAGCACCGACGAAGTTGCATTTGATCTCAGCGGCAGTGCCATGGATAGCTTGGCC
GTTCCCGGCTGGAATCGACTCTACGCGACTTCGGCCAATACGACATTCAGTACA
TGAGGTACAGATTGTGATCTCGATCTCCTTGCCCTAGATCCTGCATGCTGTCCTCAA
ACTGGAACCCGGAAGAAGAGGAAGAATTTGGGTTCTGCAAGCTCACAGTCGA
TGTTGGAGCTCGCTGCTGCATCAGTGCGAACTCTTGTGACACCAGCTGCTGCAT
CGTTTGTCCGCTTACAATGCGTTTGTGTGTGTGTGTGTGTACTACTGCACTAGGGT
GCCATGCGACTATGTTGGACAGAACGTTGCGTTTCGCGGTTGATGCAGGCTCTAGT
CCATACTGGCTCTCGTTTGCAGTCAGATACGAGGGAGGCCAGGCGACATCGAGT
CTGTCATGATCCGACAGGTAACAAATTACATGCCAGCTCCTGCCGCACCCAATAT
TTCTTTACAATAATTATCTTCTGTAATTCTTGGTGAAGCTTTGCAATATCTGATTCT
GCTGCAAGAGCAAAAGATGATGAGACGTTTAGGAAGCAATTCATTTGCGTCTTGA
TGGTTCTGTGATTTTGTGTTGTGGTTTTGGATATCGATGAACTGCACAGTCTGGAAG
CTACGATTGGGTGGCGATGCAACACAACCTGGGGTGCAAGTTGGATGTTGATCGAC
TACTCTGGACAAGCTTTCCGAGGTCCTTACGACGTTTCAGATCACAGCCAAGCTCA
ACGGCCATTCTCTCATCGCCTGGAATGCCATTCCGGACTACTTCCAACCGGGAGC
TACCTACAATTCTAGCGTGCAGTTCTTATACTAGAAAGAAACAAGCCCATGACC
CATCTGCAAATTCATGTGCAAACTAGACGCACTAGCTAGCTCGCTGCAGATCCTC
ATGATCATCAGCTGCTGCTGCTGCTGCAACATTTCTGATAGAATTCATAATAGCT
ACCTCGCTAGCTTGTAGCGTCTCATGATGCTTTCTAGATGCATTTGGAAGAGAA
CCATCAACCATAAATCGATCTAGTACAGCTTGAAAAGATCAGGAGGATCCGCGA
AGATCAATCTGGAGCCGTTTTGTTGTGTAACAGGCCAGAGCTTTACAAGTAGATT
TGTATGCAGGTGCTATGCAGATTTTTGTTTCTAGAGACTGCTTTTTGTCTACATGA
TGAACAGTGCAGTCCAGATCAGAATCACTGCCATCCAGCTGCACAGCTTGGCCT

GCAATGGACTTACTTCTTCACAAAAATTGATGATCAGATGTTGATCCATTTGTAAT
TTTATCATCAAGAGAATCTC