

IDENTIFICATION

Species: *Arabidopsis lyrata*

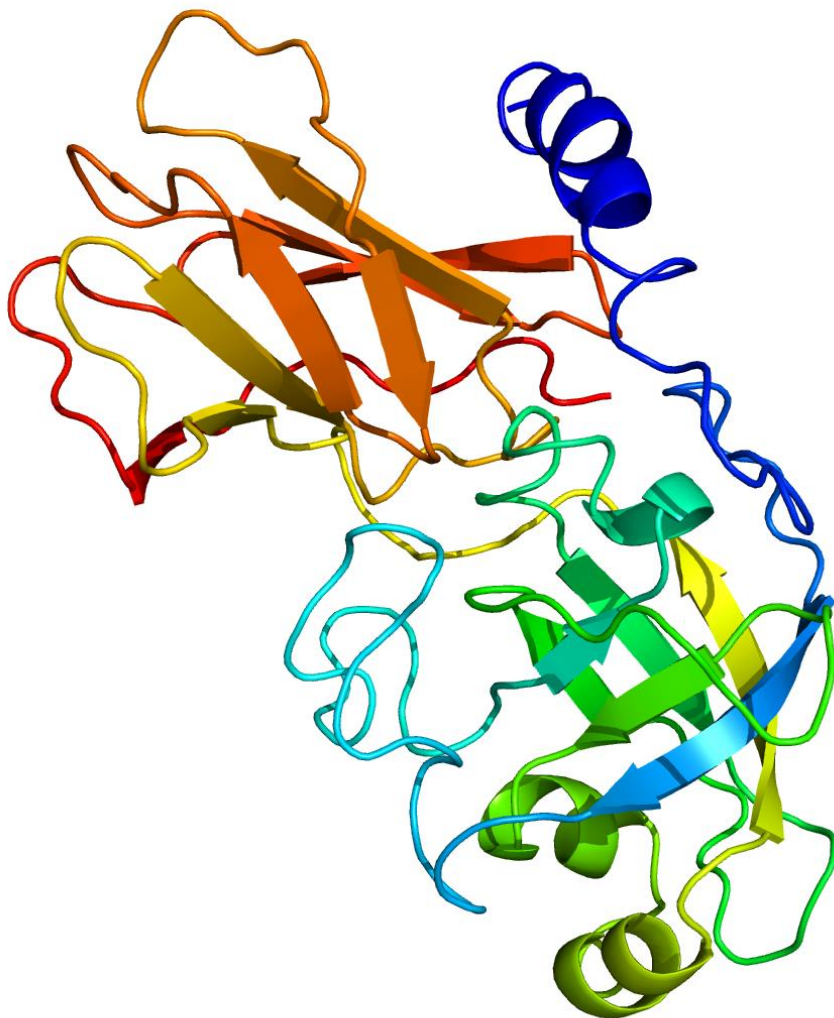
Locus: AL7G24350

Gene Model: AL7G24350.t1

Description: ALEXPB-03

Family: Beta Expansin

3D structure:



GENOME DATABASES

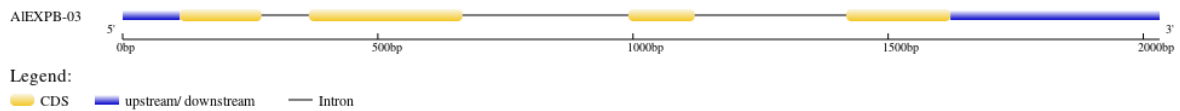
Phytozome: https://phytozome-next.jgi.doe.gov/info/Alyrata_v2_1

Kegg: <https://www.genome.jp/entry/T01578>

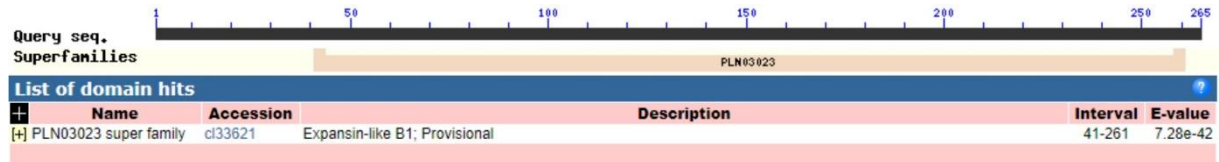
EXTERNAL RESOURCES

https://plants.ensembl.org/Arabidopsis_lyrata/Info/Index

GENE STRUCTURE



DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>AIEXPB-03

MQLSPVMLATLCIVLQLLIGSSALSTTNRHVSNSHWLPAVATWYGSPNGDGSDDGGA
CGYGTLDVVKPLHARVGA VNPILFKNGEGCGACYKVRCLDKSICSRRAVTVIITDECP
GCKTSTHFDLSGAVFGRLA IAGESGPLNRGLIPVIYRRTACKYRGKNIAFHVNEGS
TDFWLSLLVEFEDGE GDIGSMHIRQAGAREWLEMKHVWGANWCIVGGPLKGPFSIK
LTTLSAGKTLSATDVVPRNWAPKATYSSRLNFPVL*

CDS (coding sequence)

>AIEXPB-03

ATGCAGCTCTCTCCGGTCATGTTAGCCACACTCTGCATTGTCCTGCAGCTTCTAAT
TGGCTCGTCTGCTTTGTCTACGACCAATCGCCACGTGTCAAACCTCTCATTGGCTTC
CCGCAGTCGCTACATGGTACGGAAGCCCCAACGGCGACGGCAGCGACGGAGGAG
CGTGTGGTTACGGTACGTTGGTGGATGTGAAGCCGTTACATGCGAGAGTTGGAGC
GGTGAATCCTATTCTCTTCAAAAACGGTGAAGGTTGCGGCGCTTGTTACAAGGTT
CGGTGCTTGGACAAAAGCATTGTTCCCGGAGGGCCGTCACCGTTATAATCACCG
ATGAGTGTCCCGGCTGCTCTAAAACCAGCACTCACTTTGACCTCAGTGGTGCCGT
CTTCGGCCGGTTGGCTATTGCCGGAGAGTCTGGCCCTCTCCGTAACCGTGGGCTA
ATCCCCGTCATTTATCGCCGGACTGCATGCAAATATAGAGGGAAGAACATAGCAT
TCCATGTGAACGAAGGATCAACTGATTTTTGGCTATCTTTGCTGGTCGAGTTTGAA
GACGGAGAAGGAGACATTGGCTCCATGCACATTCGCCAAGCAGGAGCGAGGGAA
TGGTTAGAGATGAAGCATGTATGGGGAGCCA ACTGGTGCATCGTTGGAGGACCA
CTCAAGGGACCATTCTCCATAAAGCTCACCACTTTGTCTGCCGGTAAAACACTCT
CCGCCACCGACGTTGTCCCTAGAAACTGGGCTCCCAAAGCAACTTACAGTTCCCG
CCTCAATTTCTCCCCCGTTCTCTGA

Nucleotide

>AIEXPB-03

AATTTACACATAAAAACCTCACTCTCTCACTCTCTCACTCTCTCACTGTTCGGCGAAGT
AACATTTTCACTTTTCCGGCTCCGATTCTGCTCGGAAAAAACA ACTCCGGCAGAA
AATGCAGCTCTCTCCGGTCATGTTAGCCACACTCTGCATTGTCCTGCAGCTTCTAA

TTGGCTCGTCTGCTTTGTCTACGACCAATCGCCACGTGTCAAACCTCTCATTGGCTT
CCCGCAGTCGCTACATGGTACGGAAGCCCCAACGGCGACGGCAGCGACGGTAAT
TCTAAGTTAACACATAATTAACCACACTTCACAATCTAGTGATTA AAAACTAAAC
ACTAATATTAATCATATACCGGAATTGCGTAGGAGGAGCGTGTGGTTACGGTAC
GTTGGTGGATGTGAAGCCGTTACATGCGAGAGTTGGAGCGGTGAATCCTATTCTC
TTCAAAAACGGTGAAGGTTGCGGGCGCTTGTTACAAGGTTTCGGTGCTTGGACAAAA
GCATTTGTTCCCGGAGGGCCGTCACCGTTATAATCACCGATGAGTGTCCCGGCTG
CTCTAAAACCAGCACTCACTTTGACCTCAGTGGTGCCGTCTTCGGCCGGTTGGCTA
TTGCCGGAGAGTCTGGCCCTCTCCGTAACCGTGGGCTAATCCCCGTCATTTATCGC
CGGTACGTATATAAATAAATTCTCTTTTTTTCTTGATAAAGATATTAATATTCTC
ATTTATTAATGTTTATGCTCTTTCAATAACTAACTCGTGACAGTGCACGAGCAAAT
AGAAAAATACTAGTAGGAATTATTTAGGTAAGAGGCTAAACAGGGTTAAGATAA
TTATTAGATTTATTGCAATGGGTGTGAATGTATTAACAAAAGAGGTACCTTTAA
CGGCGATCTCGTTTTTTATGGTCCCAACCATTTTTTGACCATTCGATATCTTTTTCT
CACTGATATATTTTACTTTTTTCTTTTCTTTTGCATAATAATAAAAAAGGACTGCA
TGCAAATATAGAGGGAAGAACAATAGCATTCCATGTGAACGAAGGATCAACTGAT
TTTTGGCTATCTTTGCTGGTCGAGTTTGAAGACGGGAGAAGGAGACATTGGCTCCA
TGCACATTCGCCAAGTAACTCTCTTTTTCTTAATAACCAAACCTCGTGCACTCACT
TCTTTAATTTTCCTTTTTTCTTACCACAAAGTCACTTCCACTTCAAAGTCTTCTTT
TTTTTATTGTACTCCACTAAATAAGAAAAAGGAGGAAACGAATCTCAGTGGAGTG
AGCACATGTATGAGTCATTTATTGACCAACTAGCTAGAAAAGGCACACCACATCA
CTCAAAGTCTTCTGAAAAGTAACAATAAAACAGACTCAAAGTGTAAGACTGAT
GGGCTTTCCTTGTGTTTTGTGAATGTTTGGTGCAGGCAGGAGCGAGGGAATGGTTA
GAGATGAAGCATGTATGGGGAGCCAACTGGTGCATCGTTGGAGGACCACTCAAG
GGACCATTCTCCATAAAGCTCACCACTTTGTCTGCCGGTAAAACACTCTCCGCCA
CCGACGTTGTCCCTAGAAACTGGGCTCCCAAAGCAACTTACAGTTCCCGCCTCAA
TTTCTCCCCCGTTCTCTGAACCAACTCTTCTACATAGTCAAAGACTATTATTGTTC
CTATTATACTACTACTCGTGTGATCGTTTTTTGAAGAGAGAAACAGAGTGACTGC
TGGCCGTAGGATGGAAACAATAAGAAAAGTACATCCGAATCAACGGCTCTTGCG
TTCTCTCCTCTTCTTCTTGTGTGCTGTGGATTTGTGTGTGTGTTGTGATGTGTGCG
TGAGTGTGTTAGGAGTCACTGTAGCCATCTTCAAAGGGGAGATAGAGCTAGAAGT
AGTATTTGGTGGTGCATGTCTTTTGTTTTTTCGATATAATTCTTGTCACTATCTCT
TTCTATTATGGGACATGACATTATATATCATATGAATAATTTATCACTATGTCTTT
CTATTGTGGGACATTACATAATAATGTATCATATGACAA