IDENTIFICATION

Species: *Manihot esculenta*

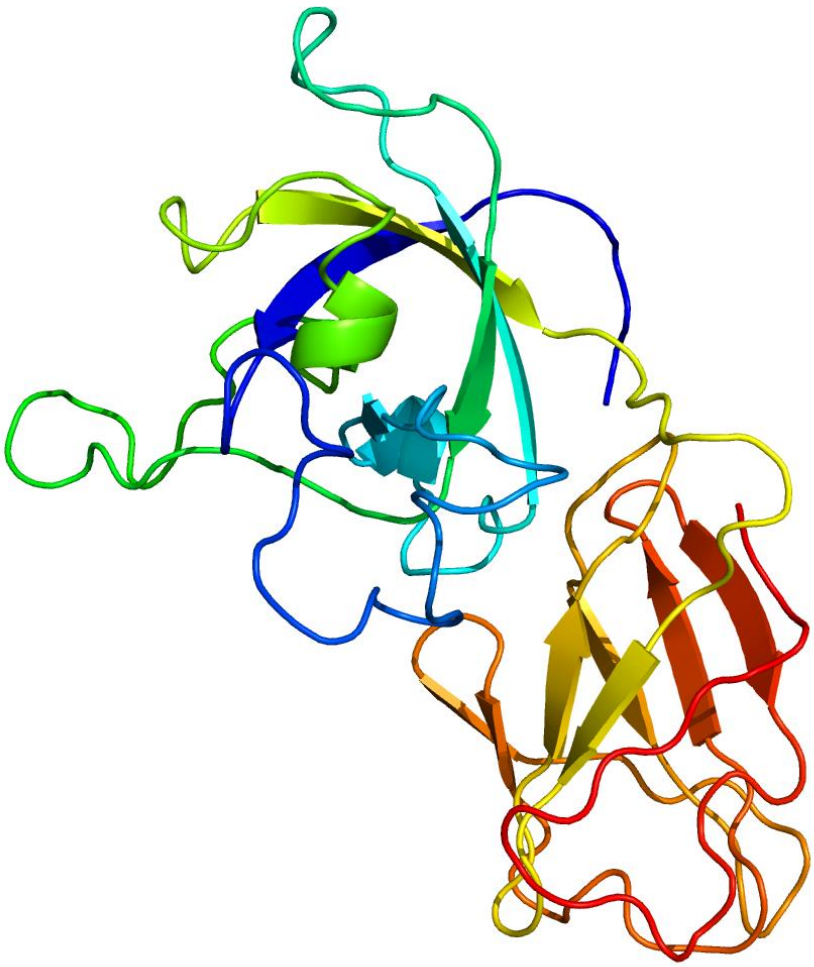
Locus: Manes.06G091300

Gene Model: Manes.06G091300.1

Description: MsEXPA-16

Family: Alpha Expansin

3D structure:



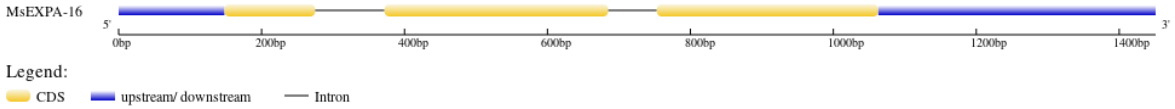
GENOME DATABASES

Phytozome: https://phytozome-next.jgi.doe.gov/info/Mesculenta_v7_1
KEGG: https://www.genome.jp/entry/T05761

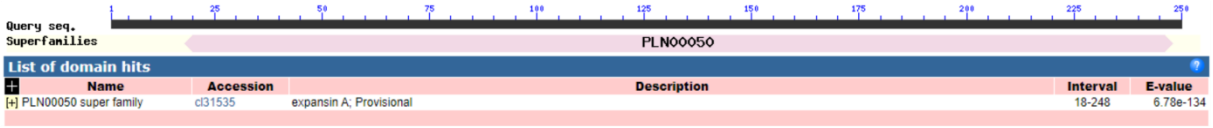
EXTERNAL RESOURCES

https://cassavagenome.org/
https://cassavabase.org/

GENE STRUCTURE



DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>MsEXPA-16

MAFAGLLFLGFLSLASFVSGDDGGWIDAHATFYGGGDASGTMGGACGYGNLYSQG
YGKDNAALSTALFNNGLSCGACFEIKCRDDPQWCLPGSIVVTATNFCPPNNALPNNA
GGWCNPPLHHFDLAQPVFERIAKFRAGIVPVSYRRIPCEKKGGIRFTINGHSYFNLILIT
NVGGAGDIHAVSVKGSKTGWQPMSRNWGQNWQSNAYLNGQSLSFMVTTSDGRSA
VCNDVVPARWSFGQTFITDQQFD*

CDS (coding sequence)

>MsEXPA-16

ATGGCATTTGCTGGGCTTCTTTTTCTGGGATTTCTTTCTTTAGCTTCATTCGTTTCT
GGGGATGATGGGGGCTGGATTGATGCTCATGCTACCTTCTATGGAGGTGGTGATG
CCTCTGGTACAATGGGTGGGGCTTGTGGGTATGGAAACCTATACAGCCAAGGTTA
CGGGAAAGACAATGCAGCTTTGAGCACAGCATTGTTCAACAATGGCTTGAGCTGT
GGAGCCTGTTTTGAAATAAAATGTAGGGATGACCCGCAGTGGTGCCTACCAGGCT
CAATTGTGGTTACAGCTACCAATTTCTGCCCACCGAACAACGCCCTCCCTAACAA
TGCCGGAGGCTGGTGCAATCCTCCTCTGCATCATTTTGATCTCGCTCAGCCTGTCT
TCGAGCGAATTGCTAAGTTTAGAGCAGGGATTGTGCCCGTGTCTTATAGAAGGAT
ACCATGCGAGAAGAAAGGAGGAATAAGATTCACAATCAATGGGCACTCATACTT
CAACTTAATACTGATAACCAACGTTGGAGGTGCTGGCGATATTCATGCAGTTTCA
GTAAAAGGGTCAAAAACTGGATGGCAACCCATGTCAAGAAACTGGGGTCAAAAC
TGGCAAAGCAACGCATACCTTAACGGACAAAGCCTCTCGTTTATGGTGACAACCA
GTGATGGACGCTCTGCGGTCTGTAATGATGTAGTACCAGCTAGGTGGTCGTTTGG
CCAGACCTTCATCACCGATCAACAGTTTGATTAA

Nucleotide

>MsEXPA-16

TTTCCTCACAAACAAATCCTTTCTCACTTTTCACTCGCAAATTAAGATTTAGCTGT
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TACACACGAGTTGTTGACCTTTTCTCTTTTGCAGGTGGGGCTTGTGGGTATGGAAA
CCTATACAGCCAAGGTTACGGGAAAGACAATGCAGCTTTGAGCACAGCATTGTTC

AACAATGGCTTGAGCTGTGGAGCCTGTTTTGAAATAAAATGTAGGGATGACCCGC
AGTGGTGCCTACCAGGCTCAATTGTGGTTACAGCTACCAATTTCTGCCCACCGAA
CAACGCCCTCCCTAACAATGCCGGAGGCTGGTGCAATCCTCCTCTGCATCATTTTG
ATCTCGCTCAGCCTGTCTTCGAGCGAATTGCTAAGTTTAGAGCAGGGATTGTGCC
CGTGTCTTATAGAAGGTATATCATACGTGCGATCCAGCCTTGTTTATAAAAAATTT
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CAACCCATGTCAAGAAACTGGGGTCAAAACTGGCAAAGCAACGCATACCTTAAC
GGACAAAGCCTCTCGTTTATGGTGACAACCAGTGATGGACGCTCTGCGGTCTGTA
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AGCAAATTTGGTTGGCCATGCAGCCGCTATTACCAGCCTTTTGCTGGTTGATGGC
GTAGCAACTTGTGACTTCCTCAACTCTTACCATTGCTGTTTAATAAAAAGTGTGCC
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