

IDENTIFICATION

Species: *Phalaenopsis equestris*

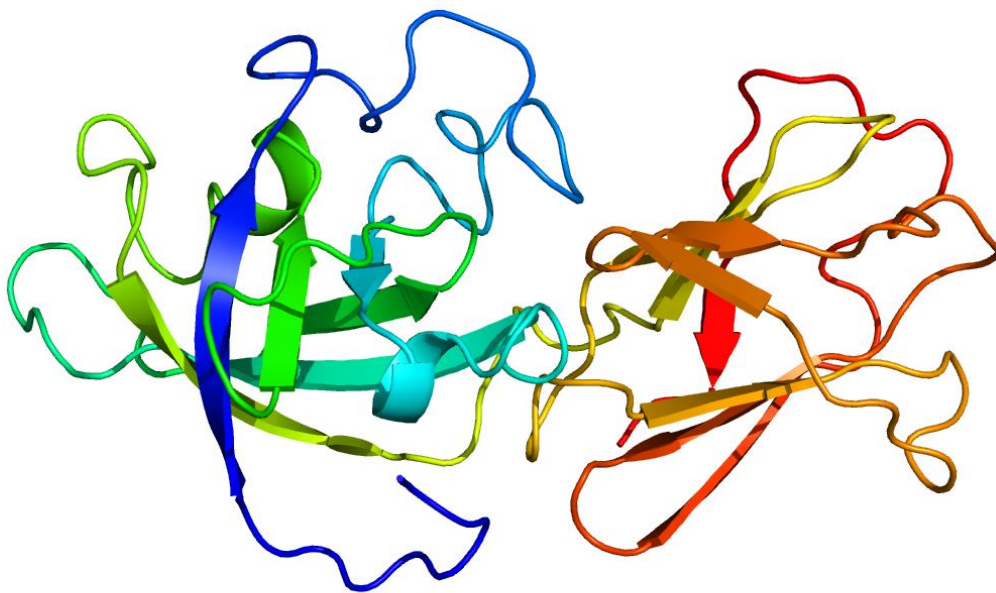
Locus: XP_020583030

Gene Model: XP_020583030.1

Description: PeqEXPA-06

Family: Alpha Expansin

3D structure:



GENOME DATABASES

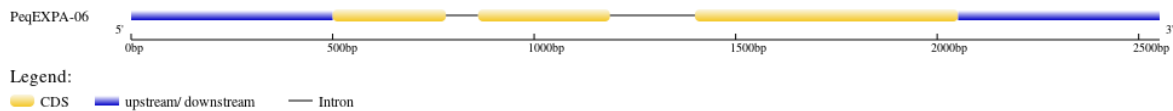
NCBI: <https://www.ncbi.nlm.nih.gov/genome/?term=Phalaenopsis+equestris>

KEGG: <https://www.genome.jp/entry/T05601>

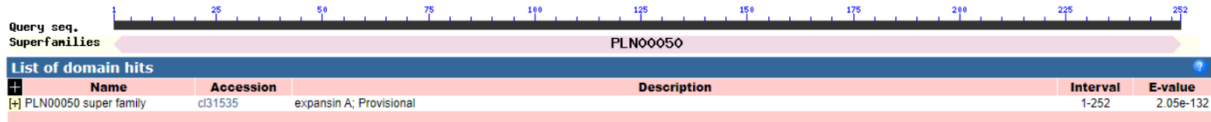
EXTERNAL RESOURCES

http://orchidbase.itps.ncku.edu.tw/est/Phalaenopsis_2019.aspx

GENE STRUCTURE



DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>PeqEXPA-06

MKHLALILIFVLSILRAVHGGGEEGWISAHATFYGGGDASGTMGGACGYGNLYSQG
YGTNTAALSTALFNGLSCGSCYEIMCANERQWCLPGTIVVTATNLCPPNYALPYTN
GGRGSPLMRLPLEHFDLSEPVFLRIAQYKAGIVPVQYRRVPCVKKGGVRFTHINGHSYF
NLVLITNVAGAGDVVAVSIKGSKTGWQPMSRNWQNWQSNLNLNGQSLSFKVTTS
DGRCVTSMDVIPSSWSFGQTFSGGQF

CDS (coding sequence)

>PeqEXPA-06

CCACCTTCACTCCCCATCGCCCTCCTTCTATAAATGCAGCTCTCCTCCCTCCCT
CCCTCCTCACCTTTCTCCTCTTTACTCGTAAAGACTCTTTTTAAGAACCAATCTTC
TTTCTTTAAGAATCTAGCCATTCTCCATTGCTGGGAAATGAAGCATCTTGCTTTAA
TCCTCATCTTTGTTCTGTCAATCCTCAGAGCTGTTCATGGAGGAGGAGAAGAAGG
ATGGATAAGTGCTCATGCTACCTTCTATGGAGGTGGCGATGCATCGGGTACCATG
GGAGGGGCTTGTGGATATGGAAATCTTTACAGCCAGGGCTATGGGACGAACACT
GCAGCTCTGAGCACTGCGTTGTTTAAACAATGGACTGAGCTGTGGATCATGCTATG
AGATTATGTGTGCTAATGAAAGGCAATGGTGTCTTCCCTGGTACTATTGTGGTTACT
GCAACCAACTTATGCCCGCCAAACTATGCTCTTCCGTACACTAATGGCGGGCGGG
GGTCGCCCTCATGCGGCTTCCATTGGAGCATTTTGATCTCTCTGAACCAGTTTTT
CTTCGTATTGCTCAGTATAAGGCCGGAATTGTTCCAGTTCAGTATAGAAGGGTGC
CCTGTGTTAAGAAGGGAGGCGTCCGGTTCACCATCAACGGCCACTCCTACTTCAA
CCTTGTTCTCATTACCAACGTCGCCGGCGCCGGCGACGTGGTGGCTGTCTCCATCA
AGGGCTCTAAAACCGGATGGCAGCCGATGAGCCGAAACTGGGGTCAAACTGGC
AGAGCAACAGCAACCTCAACGGCCAATCCCTGTCCTTCAAGGTCACCACAAGCG
ATGGCCGGTGTGTTACATCCATGGACGTCATCCCTTCCCTCCTGGAGCTTCGGCCAA
ACTTTCTCCGGCGGCCAGTTCTGAGAAAATAAATTTTCATCACTTTCAATCTAACTC
TACCATTACTGATATTAGTAAGCTTCTCCGTGTCATGTTGATTGCTGCTTTGAAAT
GGCAATCAATGGCGTTTAATTATCTTACTACTAAGTTGTTGCTGATCTTCTTCTCT
CCTTTTTGCCTTAAATATTGTGGCAGAGGAGGGCGCCGCATAAATAGTGGTGCAA
GGAGGCATTTTTATTGGATTTTGATTTTAATGGAGTGGTGGTGGAGGATTGGAATG
GCTGAGGTGAACATGGCTGAGCTTGTGCGCCCGCCATTAAATCCAACACTTTTAA
TTTATATGTAGAGGGTATTTGTGATGTAATCTGCTTTTAT

Nucleotide

>PeqEXPA-06

GCCGACCGAAAGGACAAATTACTAGCTGCGAAATTTGCCTCCATGCATTCCTTTT
TAAAATCTCCTCTGTTTCCTCTTTCTTTCAAATAAAAATAAAAAATATATATATATA
TAAATAATAATAATTTAAAATTAATGTCTTCTAATTCGTAATTAGTGATAATTTTG
AGAATTTGTTAACATTAATCGTAATTAAGACCAGCATATTCTTATAATAATTCTTA
ATCTAAAATATTCCTTCCTTTATCTGCGTACTGCAGCTTTATTCTTTCTACTACCAT
GCATTACAATTAACCTCAGCTGATGCAATAAAGGCATGAAACCACTTGAGCCTC
AGTTCGGTGGACAGTTGTACTTTTGGTTTTTTCTCCTCATAGTTGCAGTTATCTGTT
AAGAAATTTTTGTTCTGCTGGCCATTTGCAAAAATGTACATTACATTA AAAAATAT
AAAATTGATAATTTTCTTTATTA ACTTGTGATATTATGATCTTGTCTTCTCCACC
TTCCTCCCCCATCGCCCTCCTCTTCTATAAATGCAGCTCTCCTCCCTCCCTCCCTC
CTCACCTTTCTCCTCTTTACTCGTAAAGACTCTTTTTAAGAACCAAATCTTCTTTCT
TTAAGAATCTAGCCATTCTCCATTGCTGGGAAATGAAGCATCTTGCTTTAATCCTC
ATCTTTGTTCTGTCAATCCTCAGAGCTGTTTCATGGAGGAGGAGAAGAAGGATGGA
TAAGTGCTCATGCTACCTTCTATGGAGGTGGCGATGCATCGGGTACCATGGGTAA
GTAAAAAAAATCAAAGAGAATAAACTTTTCTGATCTCAAAATTAGATTTTGAAAT
GATAGTTGTTTGTAATTGAAGGAGGGGCTTGTGGATATGGAAATCTTTACAGCCA
GGGCTATGGGACGAACACTGCAGCTCTGAGCACTGCGTTGTTTAAACAATGGACTG
AGCTGTGGATCATGCTATGAGATTATGTGTGCTAATGAAAGGCAATGGTGTCTTC
CTGGTACTATTGTGGTTACTGCAACCAACTTATGCCCGCCAAACTATGCTCTTCCG
TACACTAATGGCGGGCGGGGGTCCGCCCTCCTCATGCGGCTTCCATTGGAGCATT
TGATCTCTCTGAACCAGTTTTTCTTCGTATTGCTCAGTATAAGGCCGGAATTGTT
CAGTTCAGTATAGAAGGTAATGATGAAATTTAGTTCTTCGAATGTTTAAAATTC
AACCTTGCATCGAAACAAAGATGAATGAAGAGGAATACTGGGAAGAGGAAGATG
AAAAAGATGAAATTTTACATTTACATTCCTTTAATTGATTCTCTCAACCTAACAC
ATAACTTAACTCCCCAAAAATTCTTTCTTTTAAATAAAAATCTGGATTTTAAAA
TCGCAGGGTGCCCTGTGTTAAGAAGGGAGGCGTCCGGTTCACCATCAACGGCCAC
TCCTACTTCAACCTTGTCTCATTACCAACGTCGCCGGCGCCGGCGACGTGGTGGC
TGTCTCCATCAAGGGCTCTAAAACCGGATGGCAGCCGATGAGCCGAAACTGGGG
TCAAAACTGGCAGAGCAACAGCAACCTCAACGGCCAATCCCTGTCTTCAAGGTC
ACCACAAGCGATGGCCGGTGTGTTACATCCATGGACGTCATCCCTTCCCTCCTGGA
GCTTCGGCCAAACTTTCTCCGGCGGCCAGTTCTGAGAAAATAAATTTTCATCACTTT
CAATCTAACTCTACCATTACTGATATTAGTAAGCTTCTCCGTGTCATGTTGATTGC
TGCTTTGAAATGGCAATCAATGGCGTTTAAATTATCTTACTACTAAGTTGTTGCTGA
TCTTCTTCTCTCCTTTTTTGCCTTAAATATTGTGGCAGAGGAGGGCGCCGCATAATA
GTGGTGCAAAGGAGGCATTTTTATTGGATTTTGATTTTAAATGGAGTGGTGGTGGAG
GATTGGAATGGCTGAGGTGAACATGGCTGAGCTTGTCCGCCGCCATTAATCCAA
ACTACTTTAATTTATATGTAGAGGGTATTTGTGATGTAATCTGCTTTTATTTTGATC
ATATATATCTATACATAGCTTTAAGATATTTTGTGTTTTTTTTGGGTGATATTACAC
ATGTGAGAACAGATTTCTCTTATATAAGATGAGTTGAGCAATTAGGGTAAACAAA
CACAACAATCAACAAAACAATCAAGAAAAAACACAAATATTTAATGAGGTTCGA
CAATATGCCTGAGTCATCTAGAGTAATGATGGGTGGAGGCCTTTTTTTTTAAAAA
AAAAAATCGTGCATTTTCTTAGCGTTACAAATAATATATATATATATATATATATG
TAATTA AAAAACCCTAAGCAATCGTGCCGCAGGGCATTGCCCCCACCCTGACT
CACTTTACGAACTCTCTCAATTAGACAGATGTTGGAGCGGAGCTTAGCGAGTCA

TGATCTGTCCCTTATCTTGTCTATGAGTCATACCTAACAAGGTGATCTATTTGAGT
GGCTTCAATTTCTTAAAAAATAAATTATAAAGCTATGGAGAAGAAGCCAAC