

## IDENTIFICATION

**Species:** *Oryza brachyantha*

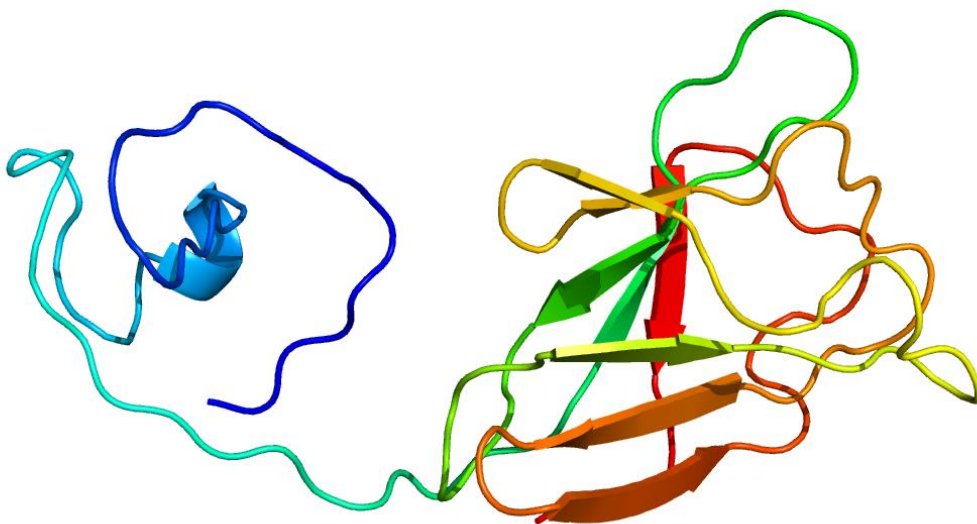
**Locus:** XP\_015693713

**Gene Model:** XP\_015693713.1

**Description:** ObEXPA-25

**Family:** Alpha Expansin

**3D structure:**



## GENOME DATABASES

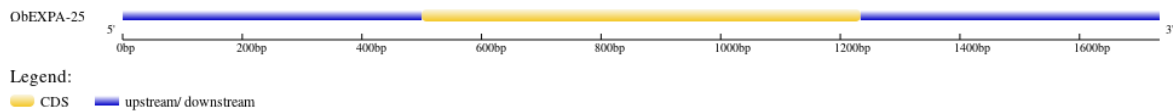
NCBI: [https://www.ncbi.nlm.nih.gov/genome/10862?genome\\_assembly\\_id=1593936](https://www.ncbi.nlm.nih.gov/genome/10862?genome_assembly_id=1593936)

KEGG: <https://www.genome.jp/entry/T02995>

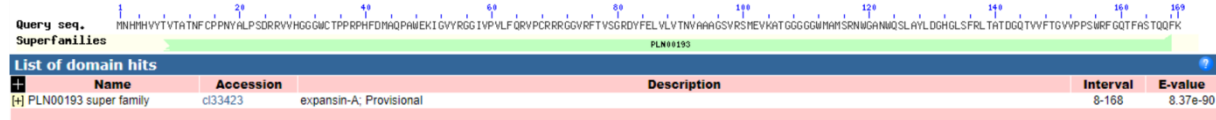
## EXTERNAL RESOURCES

<https://rice-genome-hub.southgreen.fr/organism/1941498>

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>ObEXPA-25

MNHMHVYTVTATNFCPPNYALPSDRRVVHGGGWCTPPRPHFDMAQPAWEKIGVYRGGIVPVLFRVPCRRRGGVRFVTVSGRDYFELVLVTNVAAAGSVRSMEVKATGGGGGWMAMSRNWGANWQSLAYLDGHGLSFRLTATDGQTVVFTGVVPPSWRFGQTFASQQFK

### CDS (coding sequence)

>ObEXPA-25

ATGAATCACATGCATGTATACACCGTGACGGCCACCAACTTCTGCCCGCCCAACTACGCGCTCCCAGCGACCGGCGGGTGGTGCACGGCGGCGGGTGGTGCACCCCGCGCGGCCGCACTTCGACATGGCGCAGCCGGCGTGGGAGAAGATCGGCGTGTACGGGGCGGCATCGTGCCGGTGCTGTTCCAGCGCGTCCCCTGCCGGAGGCGCGGCGGCGTGCGGTTACGGTGAGCGGGAGGGACTACTTCGAGCTGGTGTGTTGACGAA CGTGGCGGCGGCCGGGTCGGTGCGGTCGATGGAGGTGAAGGCCACGGGCGGCGGCGGGCGGCTGGATGGCCATGTCGCGCAACTGGGGCGCCAATTGGCAGTCGCTGGCGTACCTCGACGGCCACGGCCTGTCGTTCCGGCTCACCGCCACCGACGGCCAGACGGTCGTCTTACCGGCGTTCGTGCCGCGTTCGTGGAGGTTCCGGCCAGACGTTCCGCCAGCACCAGCAGTTCAAGTGATTGAATGAATTTAGATAATATCAGAAAGTCTTACACGGTGGCGGACTCAGAAAGTGGTAGTTTCAGTTTCAGGTTTGGTTGGCCGTGGCGTGCTTGACCGCTGCTCGATCTGTTACTAGCAGCCCAGCAACGATTTGTGCTCTGC AATCGTGTGATTTCGATCGATGATTTGTATTGACAAGACATGCCCATGTATCAATGCGATACCAACTACTTTGCAA

### Nucleotide

>ObEXPA-25

CCTTGATCTGAATGATTGATCTCTTCTTCCTTGATTGCCATCCTGAAGAAACAAACAGAGGCAGATTTCAGAACGTACCAAATCATCATGCTGCCTGTAGCAGAGCTAGTGACAGTGTGGCAAACCTGTTTAATTCCAAGGAAAATAGATCGGCATGACTGTCTCCCTTCTGCGTCATGTGCCAACAAGGTTGGCACCAACTTATCATCTGCAAGTGCACTAGCGAACATGGCAGATCATACTACGTACGTACGTAGGTACAGTGAGTTGCTCTTCACTTCAAGTGATCTGAAGCTGATGACGACAGAGGCATGCATGGTTCGGTGCAGCAGCAACAGCAGCAGAACTTGGCTCGAGTTAAAGCTGGATAGCAAGCGTACGTACG

AACGTCCTCTCTGTTCTGGCCGATTTATCGATGGATCATGGATGGATGGATCGAG  
CAGAACATGCATGGGACGGACTGATCATGACATTGACATGCCTTCTTCCACACAC  
GAGTAGATGAATCACATGCATGTATACACCGTGACGGCCACCAACTTCTGCCCCG  
CCA ACTACGCGCTCCCGAGCGACGGCGGGTGGTGCACGGCGGGTGGTG  
CACCCCGCCGCGGCCGCACTTCGACATGGCGCAGCCGGCGTGGGAGAAGATCGG  
CGTGTACCGGGGCGGCATCGTGCCGGTGCTGTTCCAGCGCGTCCCCTGCCGGAGG  
CGCGGCGGCGTGCGGTTACGGTGAGCGGGAGGGACTACTTCGAGCTGGTGCTG  
GTGACGAACGTGGCGGGCGGCCGGGTTCGGTGCGGTCGATGGAGGTGAAGGCCACG  
GGCGGCGGCGGCGGCTGGATGGCCATGTTCGCGCAACTGGGGCGCCAATTGGCAG  
TCGCTGGCGTACCTCGACGGCCACGGCCTGTCGTTCCGGCTCACCGCCACCGACG  
GCCAGACGGTCGTCTTACCGGCGTCGTGCCGCCGTCGTGGAGGTTCCGGCCAGAC  
GTTTCGCCAGCACCCAGCAGTTCAAGTGATTGAATGAATTTAGATAATATCAGAAA  
GTCTTACACGGTGGCGGACTCAGAAAGTGGTAGTTTCAGTTTCAGGTTTGGTTGG  
CCGTGGCGTGCTTGCACCGCTGCTCGATCTGTTACTAGCAGCCCGCCAACGATTT  
GTGCTCTGCAATCGTGTTGATTCGATCGATGATTTGTATTGACAAGACATGCCCAT  
GTATCAATGCGATACCAACTACTTTGCAAATTAATCACATCAGTTTAATTTGGATG  
TTTTCTTACCTTTTCACTCCCTTTCTTAATTTGTTTTTCATTCCTTTCCACTTTCTTG  
ATTTATTTTTCAACTTTTACCCCCCTTTCTTAATTTATTTTTTCGCTCCCTAGCTTTCT  
GCTTTCTTAATTTATTTATCACCTTTTCACTCCCTTTGTATTTTCAGTTTATACCGT  
TACTACCTTTCTTAATTTATTTTTCTTTTCCCCTTTCTTATTTGTTTGTTTTTTCAA  
ATTATTTTTACATGTACTCCCTTTTTTAATTTATTTTTTACCTTTTACTCCCTCGCTT  
CCTTTCTCAATTTTATATCCATTCAATTTGTTTTTACCTTTTACTCCCTTTCTTAATT  
TGTTTTTGTTTTCCCATGCTATGCTGTATTTTTCTTTTCTTCATTTCTCATTTTACCC  
ATGCTACAATTTCTCCATTTCTTACTTTTTTTTTTAATTTTTGCTAAGCTTTTATTCCA  
ATTTTACA