

## IDENTIFICATION

**Species:** *Oryza brachyantha*

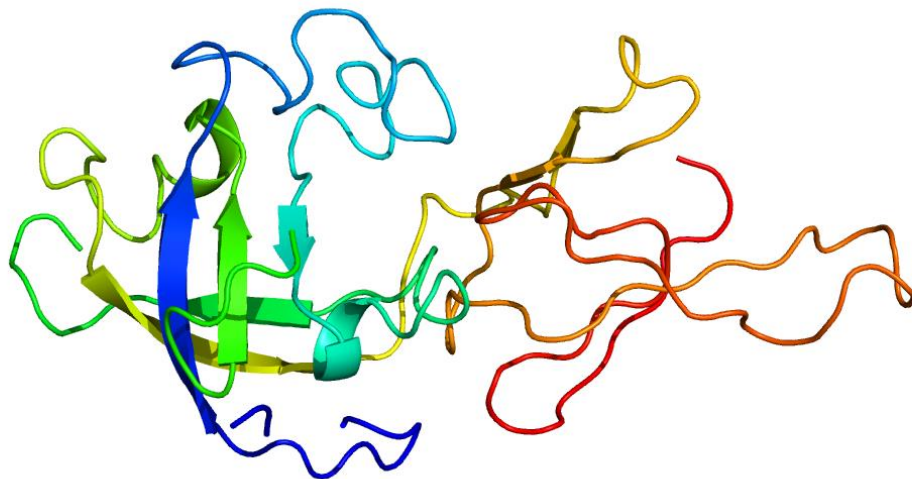
**Locus:** XP\_006651432

**Gene Model:** XP\_006651432.1

**Description:** ObEXPA-15

**Family:** Alpha Expansin

**3D structure:**



## GENOME DATABASES

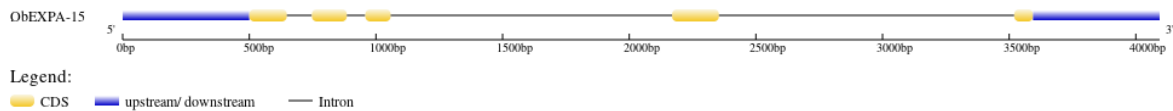
NCBI: [https://www.ncbi.nlm.nih.gov/genome/10862?genome\\_assembly\\_id=1593936](https://www.ncbi.nlm.nih.gov/genome/10862?genome_assembly_id=1593936)

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

## EXTERNAL RESOURCES

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

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>ObEXPA-15

MAALLAVLFLLSLSSPSSSSAAAAGYAPGRWTNAHATFYGGADASGTMGGACGY  
GNTYSQGYGADTAALSTAMFADGLSCGACFEVRGAGGGGGRWCNPPLHHLDL SQP  
AFLRIARYQAGIVPVS YRRVACRRKGGMRFTINGHSYFNLVLVSNVGGAGDVHAVA  
VKGGGGGRAARWQPTARDRGPNRQRIDAVPADQASEMFIGIGFTCVRS

### CDS (coding sequence)

>ObEXPA-15

ATGGCGGCTTGCTAGCGGTGCTCTTCTTGCTCTCGCTGTCGTCGCGGTCGTCGTC  
CTCCGCGGCGGCGGCGGCCGGGTACGCCCCGGCCGCTGGACCAACGCCACGC  
CACCTTCTACGGCGGCGCCGACGCTCCGGCACCATGGGTGGCGCGTGGCGGTAC  
GGCAACACGTACAGCCAGGGGTACGGCGCGGACACGGCGGCGCTGAGCACGGCG  
ATGTTTCGCGGACGGGCTGAGCTGCGGGCGCCTGCTTCGAGGTGCGGGGCGCCGGC  
GGCGGGGGGGGGCGGTGGTGCAACCCGCCGCTCCACCACCTGGACCTCTCCAG  
CCGGCCTTCCCTCCGCATCGCCCGCTACCAGGCCGGCATCGTCCCCGTCTCATAACCG  
CCGCGTGGCGTGCCGGAGGAAGGGCGGGATGAGGTTACCATCAACGGGCACTC  
CTACTTCAACCTGGTGCTGGTGAGCAACGTGGGCGGCGCCGGCGACGTGCACGCG  
GTGGCGGTGAAGGGCGGCGGCGGCGGAGGGCGGCGCGGTGGCAGCCACGGCC  
CGGGACCGGGGCCGAACCGGCAGAGGATAGACGCCGTCGCCGCTGATCAGGCT  
TCGGAGATGTTTATAGGGATAGGTTTACGTGCGTGCGTTCTTAG

### Nucleotide

>ObEXPA-15

TAGGTGGATGAGCGTAACATTGAGTATGTAATGGTGTGCACGCGTGTGTACGTCT  
TCCGTGCAATAAAAAAATCCCACATCTTTTCGCTTATGCTTATAAGAAAAAATAT  
TTGATTTTAATGTTTTTACTATAATTTATTTTCAGCTCTAACTTTTAAATCACTA  
AAAAATATGTTTACTTTTAGTTTACAAATATATTGTTTAAACATTTCCATAAAAA  
GACCAACCATTACCGTTGAGTTGGTAGGCAGAGAAAGACACGCACATTACACA  
TATCATAGAAAGACAAGCATGCATGCAAGCGTACGTGCAGTTAAAAAGTGTTAA  
ACCAGTGACATATTCTTCGAAAAATAAATCTACAACTTTTCTGTGCTAGCCAG  
CTATATGTGCGCGTGCCTATTTATACCCCCACACGAACGAACGAACGATCGATGC  
CAATGTAGTAGACACGAGCCTACTTTAAGCTCTCTACTTCTCGATCGCGCGTAC



AATTGCACAAGTTTATGTGACCTATGTATAGAGAAATACGAAGCAATTGATAAAT  
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AAACTTATATTAGATTCATCTAAAAAATTTTGGACATAAAAATTACACACTCCTC  
CTATAGAAATAAAAAAATTTATATTAGGAACACATATTTTTGCTCCTAATTTTAGG  
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GTTGGAGATATATTTTCCAGTCAAATTCTTAAAATTCAGTTTTATCATCAAATCTA  
GATATCTCTTTGAGATGCTCAATACTTCCCATTTTCTCGAAGGAAGCGGATCATCT  
GGGCAGTTAAGCTCTTAGATTTTCTAACGGCTTTGTGCTTTCAGGGATAGACGCC  
GTCCCCGCTGATCAGGCTTCGGAGATGTTTATAGGGATAGGGTTTACGTGCGTGC  
GTTCTTAGAGGAGAGTGCGTGCGTTGTGAGCGTCTATATTGTAATGTTCTCTTA  
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ATAAACTACGATAAAAAATATCAAAATTTTCGGTCTTTAGTTCATTTAGTTCGGTT  
TCTCAGTTTTGGTTATTTTTGTCCACTCCTATGCGTGTGTATATAAGTGCATAA  
ACAATTTTAAGGCTCTATGTACTTCTTATTGACTATGGGTTATCGAAAGTTTATA  
GCAGGGTAGGAGACAGTGTTTAGGCGATGGTACTTCATTAATTTTTTTTCAACGG  
TTAGGATTC