

## IDENTIFICATION

**Species:** *Oryza sativa* Kitaake

**Locus:** OsKitaake03g170000

**Gene Model:** OsKitaake03g170000.1.p

**Description:** OskEXPA-16

**Family:** Alpha Expansin

**3D structure:**



## GENOME DATABASES

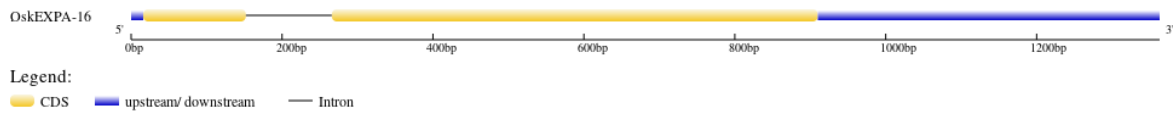
Phytozome: [https://phytozome-next.jgi.doe.gov/info/OsativaKitaake\\_v3\\_1](https://phytozome-next.jgi.doe.gov/info/OsativaKitaake_v3_1)

KEGG:-

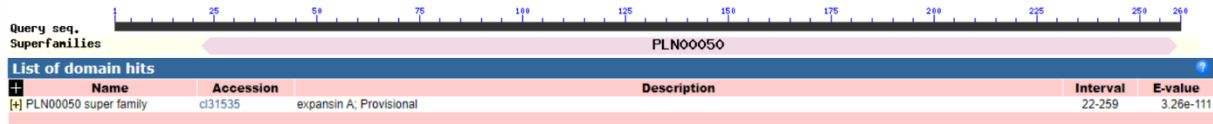
## EXTERNAL RESOURCES

[https://rice-genome-hub.southgreen.fr/bio\\_data/185326](https://rice-genome-hub.southgreen.fr/bio_data/185326)

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>OskEXPA-16

MAPPLLLLLASLLLVAARRALGLGLGQWQPGHATFYGGGDASGTMGGACGYGNLY  
SQGYGTSTAALSTALFNRLSCGSCYELRCAGDHRRSCLPGGATVTVTATNFCPPNY  
ALPSDGGGWCNPPRRHFDLAEP AFLRIARHAAGIVPVSFRRVACARKGGVRFTVNGH  
AYFNLVLVTNVGGAGDVRSLAVKGS GSGSRVGGRWQPMSRNWQNWQSNAYLDG  
KALSFRVTAGDGRSLTCADVAPAGWQFGQTFEGRQF\*

### CDS (coding sequence)

>OskEXPA-16

ATGGCGCCACCCCTGCTCCTCCTCGCTTCTCTCCTCCTCGTCGCAGCGCGGCG  
AGCCCTCGGCCTCGGCCTCGGCCAGTGGCAGCCCGGCCACGCCACGTTCTACGGC  
GGCGGCGACGCTCCGGCACGATGGGTGGCGCGTGCGGGTACGGGAACCTGTAC  
AGCCAGGGGTACGGCACGAGCACGGCGGCGCTGAGCACGGCGCTGTTCAACAGG  
GGCCTGAGCTGCGGCTCCTGCTACGAGCTCCGGTGC GCGGGCGACCACCGCCGGT  
CGTGCCTCCCCGGCGGCGCCACCGTGACGGTGACGGCGACCAACTTCTGCCCCCC  
GAACTACGCGCTCCCGAGCGACGGCGGCGGCTGGTGCAACCCACCGCGGCGGCA  
CTTCGACCTCGCCGAGCCGGCGTTCCTCCGCATCGCGCGGCACGCCGCCGGGATC  
GTCCCGGTCTCCTTCCGCCGCGTGGCGTGCGCCAGGAAGGGCGGCGTCAGGTTC A  
CCGTCAACGGCCACGCCTACTTCAACCTGGTGCTCGTCACCAACGTCGGCGGCGC  
CGGCGACGTGCGCTCCCTCGCCGTGAAGGGGTCCGGGTCCGGGTCCGCGCGTGGG  
GGGGCGGTGGCAGCCGATGTCGCGCAACTGGGGGCAGAACTGGCAGAGCAACGC  
GTACCTCGACGGCAAGGCGCTCTCCTTCCGCGTACCCGCCGGCGACGGCCGCTCC  
CTCACCTGCGCCGACGTGGCGCCC GCCGGCTGGCAGTTCGGCCAGACCTTCGAGG  
GCAGGCAGTTCTAG

### Nucleotide

>OskEXPA-16

GCACCAAAC TACCCCAATGGCGCCACCCCTGCTCCTCCTCGCTTCTCTCCTCC  
TCGTTCGAGCGCGGCGAGCCCTCGGCCTCGGCCTCGGCCAGTGGCAGCCCGGCCA  
CGCCACGTTCTACGGCGGCGGCGACGCTCCGGCACGATGGGTACGCACTGGCGC  
AACTGACACTGACAGTCTGACACAGATGGCGGCTTAGCTTACCTATGATGTCGC  
CGCTCATGGCGCCGCCGCCGCGCGCGGCGGCTTGTTCGGTGCAGGTGGCGCGTG

CGGGTACGGGAACCTGTACAGCCAGGGGTACGGCACGAGCACGGCGGGCGCTGAG  
CACGGCGCTGTTCAACAGGGGCCTGAGCTGCGGCTCCTGCTACGAGCTCCGGTGC  
GCGGGCGACCACCGCCGGTCGTGCCTCCCCGGCGGGCGCCACCGTGACGGTGACG  
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GCAACCCACCGCGGGCGGCACTTCGACCTCGCCGAGCCGGCGTTCCCTCCGCATCGC  
GCGGCACGCCGCCGGGATCGTCCCGGTCTCCTTCCGCCGCGTGGCGTGCGCCAGG  
AAGGGCGGGCGTCAGGTTACCGTCAACGGCCACGCCTACTTCAACCTGGTGCTCG  
TCACCAACGTCGGCGGGCGCCGGCGACGTGCGCTCCCTCGCCGTGAAGGGGTCCG  
GGTCCGGGTCGCGCGTGGGGGGGGCGGTGGCAGCCGATGTCGCGCAACTGGGGGC  
AGAACTGGCAGAGCAACGCGTACCTCGACGGCAAGGCGCTCTCCTCCGCGTCAC  
CGCCGGCGACGGCCGCTCCCTCACCTGCGCCGACGTGGCGCCCCGCCGGCTGGCAG  
TTCGGCCAGACCTTCGAGGGCAGGCAGTTCTAGATTAGTTTAGCCAAGAGGATTA  
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TAGTGAGGTCTAGTCGTCTAGTAGATGCTGAGGTTGCTGTGGCTCCTGCAAGAGA  
AGGCAAGGCATTTTCGCGCCCGCGTGTGTCAGAGGCCGGATGGTTTGCAGTTTGCTGC  
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TAATCATCTCTAATTTATTTATTTTCTCTATATACTTAAATTT