

## IDENTIFICATION

**Species:** *Oryza sativa* Kitaake

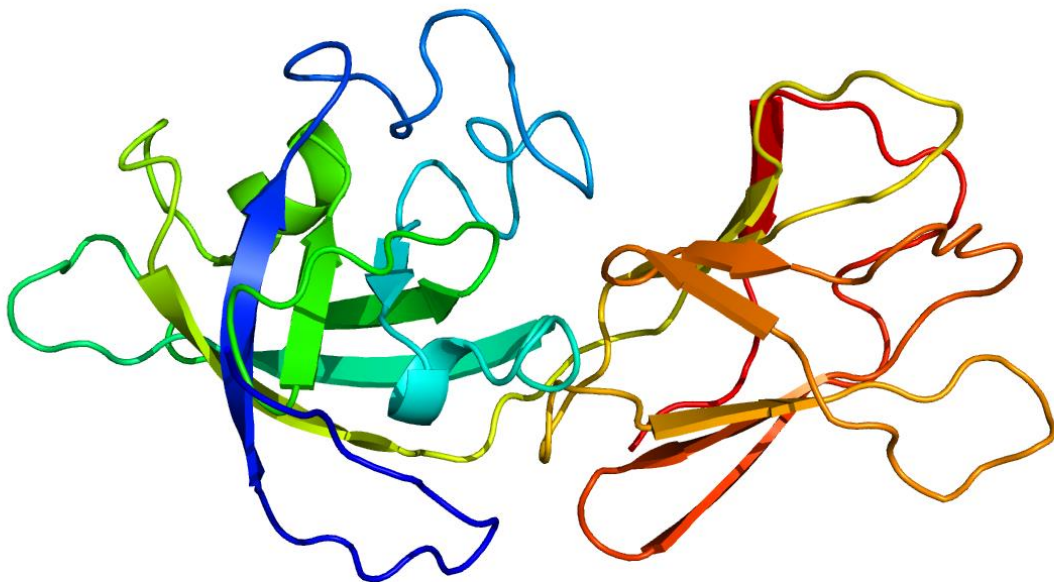
**Locus:** OsKitaake02g118800

**Gene Model:** OsKitaake02g118800.1.p

**Description:** OskEXPA-08

**Family:** Alpha Expansin

**3D structure:**



## GENOME DATABASES

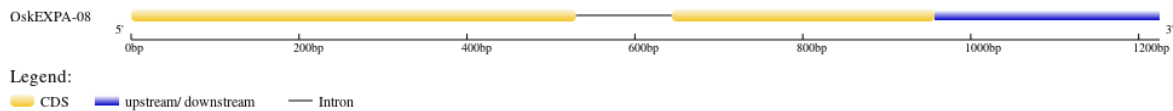
Phytozome: [https://phytozome-next.jgi.doe.gov/info/OsativaKitaake\\_v3\\_1](https://phytozome-next.jgi.doe.gov/info/OsativaKitaake_v3_1)

KEGG:-

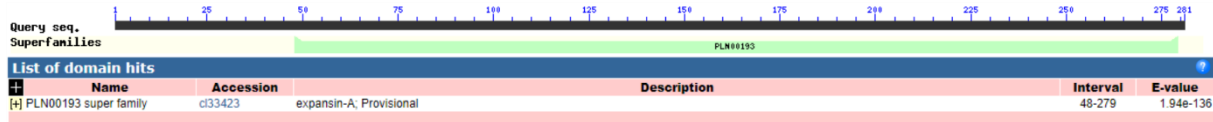
## EXTERNAL RESOURCES

[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-08

MAPARPFALLFLAVTVGFVLLTAADDSANATATTTTAMAPSSSTDDAAPPVWLKAH  
ATFYGGADASGTMGGACGYGDLYSQGYGTRNAALSTALFNDGASCGQCYKIACDR  
KRAPQWCRPGVTVTITATNFCPPNWDLPDNGGWCNPPRPHFDMAQPAWEKIGIYR  
AGIIPVIYQRVPCVKKGGVRFTHDHFNLVLVTNVATTGLIKSMDVMGSNSTDWL  
PMVRNWGANWHSLSYLTGQMLSFRVTNMDGQTLVFRNIVPSGWKFGQTFASKLQF  
K\*

### CDS (coding sequence)

>OskEXPA-08

ATGGCTCCAGCTCGACCTTTCGCGTTGCTGTTTCTCGCAGTCACGGTCCGGCTTCGT  
TCTGCTGACGGCGGCCGACGACTCGGCCAATGCGACGGCGACGACGACGACAGC  
GATGGCTCCGTCGTCATCGACCGACGACGCGGCTCCGCCCCTGTGGCTGAAGGCG  
CACGCGACGTTCTACGGCGGCGCGGACGCGTCCGGCACCATGGGAGGCGCGTGC  
GGCTACGGCGACCTTTACTCGCAGGGGTATGGGACGCGGAACGCGGCGCTGAGC  
ACGGCGCTGTTCAACGACGGCGCGTCTGCGGGCAGTGCTACAAGATCGCCTGCG  
ACCGCAAGAGGGCACCGCAGTGGTGCAGGCCC GGCGTCACGGTCACCATCACCG  
CCACCAACTTCTGCCCGCCCAACTGGGACCTCCCCAGCGACAACGGCGGCTGGTG  
CAACCCTCCACGGCCGCACTTCGATATGGCGCAGCCGGCCTGGGAGAAGATCGGT  
ATTTACCGTGCCGGCATCATCCCCGTCATCTACCAAAGGGTTCATGTGTAAAGA  
AGGGTGGTGTGCGGTTACCATCAACGGGCATGACTACTTCAATCTCGTTCTTGT  
GACCAACGTTGCAACCACCGGCTTGATCAAGTCGATGGATGTGATGGGCTCCAAC  
TCAACCGACTGGCTGCCAATGGTGAGAACTGGGGTGCAAATTGGCACTCGTTAT  
CGTATCTCACTGGACAGATGCTTTCCTTTAGGGTGACCAACATGGACGGCCAGAC  
GCTTGTCTTCAGAAACATTGTGCCGTCAGGATGGAAGTTCGGGCAAACATTTGCA  
AGCAAACCTGCAGTTCAAGTGA

### Nucleotide

>OskEXPA-08

ATGGCTCCAGCTCGACCTTTCGCGTTGCTGTTTCTCGCAGTCACGGTCCGGCTTCGT  
TCTGCTGACGGCGGCCGACGACTCGGCCAATGCGACGGCGACGACGACGACAGC  
GATGGCTCCGTCGTCATCGACCGACGACGCGGCTCCGCCCCTGTGGCTGAAGGCG

CACGCGACGTTCTACGGCGGGCGCGGACGCGTCGGGCACCATGGGAGGGCGCGTGC  
GGCTACGGCGACCTTTACTCGCAGGGGTATGGGACGCGGAACGCGGGCGCTGAGC  
ACGGCGCTGTTCAACGACGGCGCGTCGTGCGGGCAGTGCTACAAGATCGCCTGCG  
ACCGCAAGAGGGCACCGCAGTGGTGCAGGCCCGGCGTCACGGTCACCATCACCG  
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ACGGGACAAACACAAATCGTATTTGTTTTTGATATTGCACACCTATATGACAATA  
TATAAGACATATTTGTAACATGTATTTTGTGATGTTTTTCAGGGTTCATGTGTAA  
AGAAGGGTGGTGTGCGGTTACCATCAACGGGCATGACTACTTCAATCTCGTTCT  
TGTGACCAACGTTGCAACCACCGGCTTGATCAAGTCGATGGATGTGATGGGCTCC  
AACTCAACCGACTGGCTGCCAATGGTGAGAACTGGGGTGCAAATTGGCACTCGT  
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GACGCTTGTCTTCAGAAACATTGTGCCGTCAGGATGGAAGTTCGGGCAAACATTT  
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CAGATTAATTT