

## IDENTIFICATION

**Species:** *Musa acuminata*

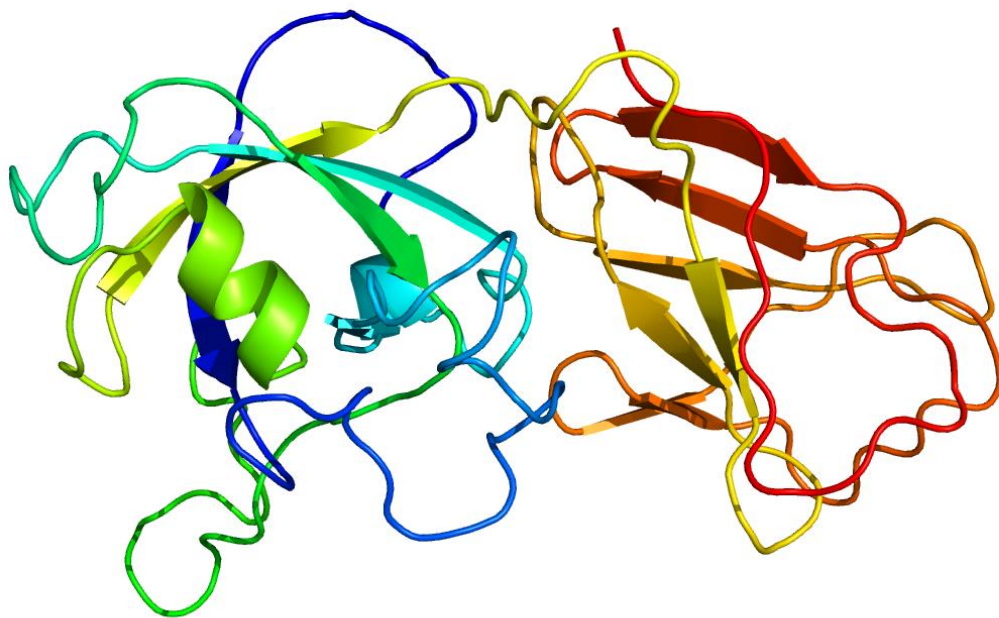
**Locus:** GSMUA\_Achr7P10570\_001

**Gene Model:** GSMUA\_Achr7P10570\_001

**Description:** MacEXPA-23

**Family:** Alpha Expansin

**3D structure:**



## GENOME DATABASES

Phytozome: [https://phytozome-next.jgi.doe.gov/info/Macuminata\\_v1](https://phytozome-next.jgi.doe.gov/info/Macuminata_v1)

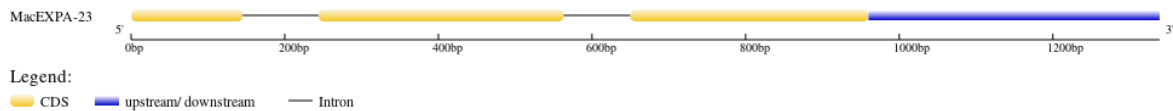
KEGG: <https://www.genome.jp/entry/T03447>

## EXTERNAL RESOURCES

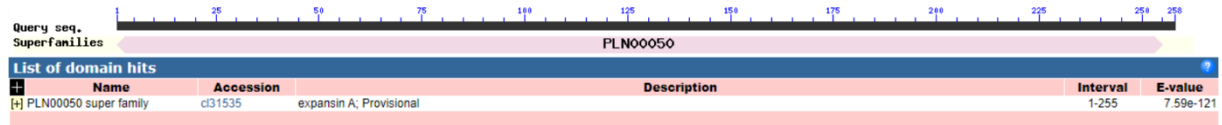
<https://banana-genome-hub.southgreen.fr/>

<https://musabase.org/>

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>MacEXPA-23

MALASFVSVALLALLAPVTARIPGLYAGGQWQNAHATFYGGSDASGTMGGACGYG  
NLYSQGYGVQTAALSTALFNEGQSCGACFEIKCAGDPQWCHGGSPSIFITATNFCPPN  
YALPSDDGGWCNPPRPHFDLSMPMFLKIAEYRAGIVPVSYRRVPCRRSGGIRFTINGF  
KYFNLVLITNVAGAGDIVRASVKGSRTGWMPMSRNWQNWQSNVAVLVGQSLSFRL  
TGSDRRTSTSWNIVPSNWQFGQTFEGKNFRT\*

### CDS (coding sequence)

>MacEXPA-23

ATGGCGTTGGCCTCCTTTGTCTCCGTCGCCCTCCTGGCTCTTCTGGCGCCGGTGAC  
TGCGCGCATTCCCGGTTTGTACGCCGGCGGTCAGTGGCAGAATGCCACGCCACC  
TTCTATGGCGGCAGCGACGCATCCGGGACCATGGGAGGGGGCGTGTGGGTATGGG  
AACCTGTACAGCCAGGGGTACGGGGTGCAGACGGCGGCTTTGAGCACGGCGTTG  
TTCAACGAGGGGCAGAGCTGCGGGGCGTGCTTCGAGATCAAGTGCGCGGGCGAT  
CCGCAGTGGTGCCACGGCGGGAGCCCCTCCATCTTCATCACAGCCACCAACTTCT  
GCCCGCCAACTACGCGCTCCCCTCGGACGATGGCGGGTGGTGCAACCCGCCCCG  
CCCTCACTTCGACCTCTCCATGCCATGTTCTCAAGATCGCCGAGTACCGCGCCG  
GCATCGTGCCCGTCTCGTACCGAAGGGTGCCGTGCAGGAGGTCGGGAGGGATCC  
GGTTCACCATAAATGGGTTCAAGTACTTCAACCTGGTGCTGATACCAACGTGGC  
CGGCGCCGGCGACATCGTGCGCGCCAGCGTGAAGGGGTCCC GCACTGGTTGGAT  
GCCATGTCCC GCAACTGGGGCCAGAACTGGCAGTCCAACGCCGTCTCGTCGGC  
CAGTCCCTCTCCTTCCGTCTCACC GGCAGCGACCGCCGCACCTCCACCTCCTGGAA  
CATCGTCCCCTCCA ACTGGCAGTTCGGCCAGACCTTCGAGGGCAAGAACTTCCGG  
ACCTGA

### Nucleotide

>MacEXPA-23

ATGGCGTTGGCCTCCTTTGTCTCCGTCGCCCTCCTGGCTCTTCTGGCGCCGGTGAC  
TGCGCGCATTCCCGGTTTGTACGCCGGCGGTCAGTGGCAGAATGCCACGCCACC  
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GCTATTGATCTGTGGCGTTGAGCTTCGAAAAA ACTTGAAGCCGATGAGATCTGCT

TGCTGTGGTTGGTGATTGTGCAGGAGGGGCGTGTGGGTATGGGAACCTGTACAGC  
CAGGGGTACGGGGTGCAGACGGCGGCTTTGAGCACGGCGTTGTTCAACGAGGGG  
CAGAGCTGCGGGGCGTGCTTCGAGATCAAGTGC GCGGGCGATCCGCAGTGGTGC  
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CAGGAGGTCGGGAGGGATCCGGTTCACCATAAATGGGTTCAAGTACTTCAACCTG  
GTGCTGATACCAACGTGGCCGGCGCCGGCGACATCGTGCGCGCCAGCGTGAAG  
GGGTCCCGCACTGGTTGGATGCCCATGTCCCGCAACTGGGGCCAGA ACTGGCAGT  
CCAACGCCGTCCTCGTCGGCCAGTCCCTCTCCTTCCGTCTCACCGGCAGCGACCGC  
CGCACCTCCACCTCCTGGAACATCGTCCCCTCCA ACTGGCAGTTCGGCCAGACCT  
TCGAGGGCAAGA ACTTCCGGACCTGACAACCCATTCCCGCGAGA ACTTCTCCACT  
AGTAATAGTAGTAGTATGTGTTAGTAGTGATGTCTGTGTGATTCGCGCCAAGCTTT  
TTGAATTCGGCGGTGCGACAATGGTAGTAGATATATGCTCCTTGGAATTCGTGAA  
AGCAAAGCAGGGGAGAGGGGAGAAGAGGTGTTTGTGATGTGCCTTTTGACCGTA  
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TGTCTCATACTGAGAGACTTCA