

## IDENTIFICATION

**Species:** *Sorghum bicolor* Rio

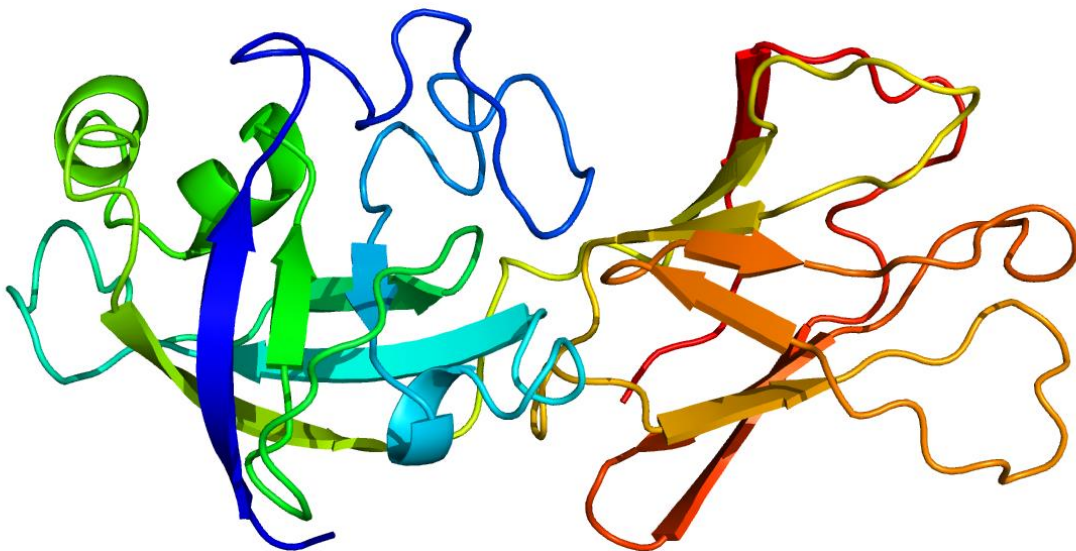
**Locus:** SbRio.01G579200

**Gene Model:** SbRio.01G579200.1.p

**Description:** SbrEXPB-23

**Family:** Beta Expansin

**3D structure:**



## GENOME DATABASES

Phytozome: [https://phytozome-next.jgi.doe.gov/info/SbicolorRio\\_v2\\_1](https://phytozome-next.jgi.doe.gov/info/SbicolorRio_v2_1)

KEGG:-

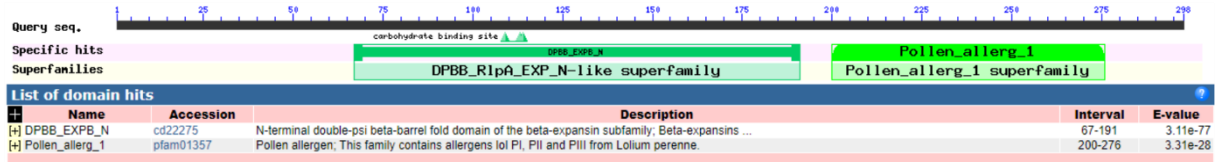
## EXTERNAL RESOURCES

<https://www.sorghumbase.org/post/sorghum-bicolor-rio>

## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>SbrEXPB-23

MDGWRRRVGSWIGAEQRKAAGHSLMGSLSSLSCLLVAARAAAVSLLLAYCAAAAV  
 NYNTSDAAALQWGNARATWYGQPNGAGPYDNGGACGFKKVNQYPFMAMTSCGN  
 QPLYRDGKGCSCYKIRCSSSKHAACSGRTETVITDMNYTPGVAPYHFDLSGTAFG  
 KLAKPGRNDELRRAGIIDIQFTRVPCEFPGLKVG FHVVEEGSSQVYFAVLVEYENDGDG  
 VVQVDLMEKGSRRWTPMRHSWGSIWRLDSNHRLQPPFSIRTRSDSGKTLVARDVIP  
 LNWRPNTFYRSIVQYSS\*

### CDS (coding sequence)

>SbrEXPB-23

ATGGATGGATGGAGGAGGAGTAGGTTGGATCGGAGCAGAGCAAAGAAA  
 GGCGGCCGGGCATAGTTTAATGGGGTCCCTGTCTCGCTCTCTGCTGCTAGTG  
 GCGGCAAGGGCCGCGCCGTCTCCCTCTGCTGGCCTATTGCGCCGCGCCGCCG  
 TGA ACTACAACACGAGCGATGCCGCCGCTTGCAGTGGGGCAACGCCAGGGCCA  
 CCTGGTACGGCCAGCCCAACGGTGCCGGGCCCTACGACAACGGCGGCGCTTGCG  
 GGTTCAAGAAGGTGAACCAAGTACCCGTTTCATGGCCATGACCTCGTGCGGCAACCA  
 GCCGCTGTACCGCGACGGCAAAGGCTGCGGTTCTGCTACAAGATCAGGTGCTCC  
 AGCTCCAAGCACGCCGCTGCTCCGGCCGACCGAGACGGTGGTGATCACGGAC  
 ATGAACTACACCCCGGGCGTGGCGCCCTACCACTTCGACCTCAGCGGCACCGCCT  
 TCGGCAAGCTGGCCAAGCCCGGCCGCAACGACGAGCTCCGCCGCGCGGGGATCA  
 TCGACATCCAGTTCACCAGGGTGCCCTGCGAGTTCCTCGGGCCTCAAGGTTCGGCTT  
 CCATGTCGAGGAGGGCTCCAGCCAGGTCTACTTCGCCGTGCTGGTTCGAGTACGAG  
 AACGGAGACGGCGACGTCGTGCAGGTGGACCTCATGGAGAAGGGCAGCAGCCGC  
 CGGTGGACGCCCATGCGCCACTCCTGGGGATCCATCTGGCGCCTCGACTCCAACC  
 ACCGCCTGCAGCCGCCCTTCTCCATCCGCACCCGAAGCGACTCCGGCAAGACGCT  
 CGTCGCACGCGACGTCATCCCACTCAACTGGAGGCCAAACACATTCTACAGATCA  
 ATCGTCCAGTACTCGTCGTGA

## Nucleotide

>SbrEXPB-23

GGGTGGGGTGGGGTGGGGTGGGCAGGGAGATGCGCCGCCAAACCCCAACATGGC  
AGTGGCAGTGGCAGGCATGTGTGTAGCAATGCATGCAATGCAAAAAACAAACA  
CACCACACTGCTGCATGCCATTGCCTGCCTTCCTTCCTTTAGCTAGCTTGCATTAT  
ATACTGGATGGATGGATGGAGGAGGAGTATGTATTGTCTAGCTAGTACCTCGATC  
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TGGATCGGAGCAGAGCAAAGAAAGGCGGCCGGGCATAGTTTAATGGGGTCCCTG  
TCCTCGCTCTCCTGCCTGCTAGTGGCGGCAAGGGGCCGCCGCGTCTCCCTCCTGCT  
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CAGTGGGGCAACGCCAGGGCCACCTGGTACGGCCAGCCCAACGGTGCCGGGCCC  
TACGACAACGGTATGTATGTATGTATGTATGTATGTATGTACTGTCGTATATAT  
TATGATCTGTATGCGTGCATTTGTTGGTTGATCGATGGCTCACCGTCACCGGACCG  
CCATGGCTGCTGCTGCTCCATATATATGCAGGCGGCGCTTGCGGGTTCAAGAAGG  
TGAACCAGTACCCGTTTCATGGCCATGACCTCGTGC GGCAACCAGCCGCTGTACCG  
CGACGGCAAAGGCTGCGGTTCCCTGCTACAAGATCAGGTGCTCCAGCTCCAAGCAC  
GCCGCCTGCTCCGGCCGCACCGAGACGGTGGTGATCACGGACATGAACTACACCC  
CGGGCGTGGCGCCCTACCACTTCGACCTCAGCGGCACCGCCTTCGGCAAGCTGGC  
CAAGCCCGGCCGCAACGACGAGCTCCGCCGCGCGGGGATCATCGACATCCAGTT  
CACCAGGGTGCCCTGCGAGTTCCCGGGCCTCAAGGTCGGCTTCCATGTCGAGGAG  
GGCTCCAGCCAGGTCTACTTCGCCGTGCTGGTCGAGTACGAGAACGGAGACGGC  
GACGTCGTGCAGGTGGACCTCATGGAGAAGGGCAGCAGCCGCCGGTGGACGCC  
ATGCGCCACTCCTGGGGATCCATCTGGCGCCTCGACTCCAACCACCGCCTGCAGC  
CGCCCTTCTCCATCCGCACCCGAAGCGACTCCGGCAAGACGCTCGTCGCACGCGA  
CGTCATCCCACTCAACTGGAGGCCAAACACATTCTACAGATCAATCGTCCAGTAC  
TCGTCGTGAACTCATCCATCCATCATTTGGCATTAAATTTGCTGCTCATTAGTTC  
ATATCTCTCTCTCGAGTCTCGATCGATCGATAACTATGAGTATTGATGAGCAATA  
AGCTGTATTATCTGTCTTACACATATGAATGTACATGTGTCTTCATTCCATCTTTGT  
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TGTATTGTTAGCATCTACTATATACTGTACTGCTGCTAGTAATACCTTATTAGG