

## IDENTIFICATION

**Species:** *Setaria viridis*

**Locus:** Sevir.8G199200

**Gene Model:** Sevir.8G199200.2.p

**Description:** SvEXPB-13

**Family:** Beta Expansin

**3D structure:**



## GENOME DATABASES

Phytozome: [https://phytozome-next.jgi.doe.gov/info/Sviridis\\_v2\\_1](https://phytozome-next.jgi.doe.gov/info/Sviridis_v2_1)

KEGG:-

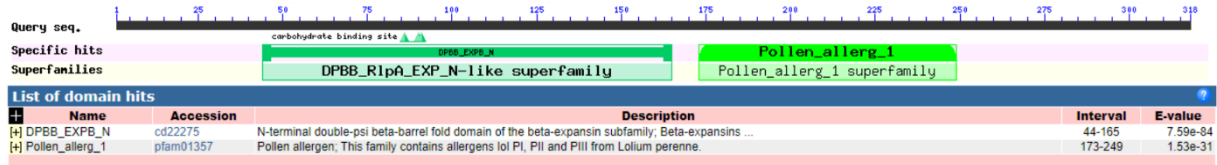
## EXTERNAL RESOURCES

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## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>SvEXPB-13

MASMIASSKMIALRALIFLLLVLVYGSCTRIVNFNASHITADPYWVAARATWYGAPTG  
AGPYDNGGACGFKNVNLPFSAMTSCGNQPLFKDGGKGCSCYQIRCLNHPACSGNP  
ETVAITDMNYYPVAKYHFDLSGTAFGALAKPGRNDELRHAGIIDIQFKRVPCIYPGQ  
MVTFHIEHGSNPNYLAVLVEFEDGDGDVVQVDLMEANSGWWTMRESWGSIWRLD  
TRRPLTAPFSLRITNESGQKLVA YQVIPANWAPNTYYRSNIQYQAFSSDDGLDIGSAA  
GLVISSAAGLDTKILGVIGLICLVLSRLHGTDVP\*

### CDS (coding sequence)

>SvEXPB-13

ATGGCATCCATGATCGCCTCGTCCAAGATGATTGCACTTCGTGCACTGATCTTCCCT  
CCTCCTTGTCTTGTATGGCTCATGCACTAGGATCGTCAACTTCAATGCCTCCCACA  
TCACCGCAGACCCCTACTGGGTGGCTGCCAGGGCAACTTGGTATGGTGCGCCAAC  
CGGCGCTGGCCCCTATGACAATGGTGGAGCTTGCGGGTTCAAGAACGTGAACCTG  
CCTCCTTTCTCGGCCATGACGTCATGCGGCAACCAGCCCCTGTTC AAGGATGGCA  
AGGGCTGCGGCTCGTGCTACCAGATAAGATGCCTCAACCACCCTGCATGCTCCGG  
CAATCCGGAGACGGTGGCAATCACTGACATGAACTACTACCCAGTCGCCAAGTAC  
CACTTTGACCTCAGCGGCACGGCGTTTGGTGCCTTGGCAAAGCCCGGCCGCAATG  
ATGAGCTCCGCCACGCCGGCATCATCGACATCCAGTTCAAGAGGGTGCCCTGCAT  
CTACCCCGGGCAGATGGTGACCTTCCACATCGAGCACGGCTCAAACCCTAACTAC  
TTGGCGGTGCTCGTTGAATTCGAAGATGGCGACGGCGATGTCGTCCAGGTGGACC  
TCATGGAGGCCAACTCCGGGTGGTGGACGCCGATGCGGGAGTCGTGGGGATCCA  
TCTGGAGGTTGGACACCAGACGACCGCTCACGGCCCCCTTCTCGCTGCGCATCAC  
CAACGAGTCTGGCCAGAAGCTGGTGGCTTACCAGGTCATCCCGGCCAACTGGGGC  
CCAACACCTACTACCGCTCCAACATCCAGTACCAGGCTTTTAGCTCTGATGATG  
GTCTGGATAATTGGCTCTGCTGCTGGACTGGTTATTAGCTCTGCTGCTGGACTGGAC  
ACGAAAATTCTCGGCGTTATTGGCTTGATTTGTTTGGTGTGTCTCGTTTGCATGG  
GACTGATGTGCCGTAG

## Nucleotide

>SvEXPB-13

GGATACGGAAGCGATCCAGGCCGTCCACTCCGACTCCGAATCGCAGACCGAACG  
GGCGAACGAGCCGCGAGACGGAGGCGCGGGGGCGGAGCGGATGGCGACCTAGG  
GTTCCGCCGATTCAAATCGGGCGGCGCGGGTGGCCATCCCCCGCCGGATCCGGGCA  
TCCGGCGACGGCGACTCGGGCGACGCGTCGACGGAGGGAGCCAGGGAGGCAGG  
AGAGGAGGAGCGAGCCGGCGTCCGGCAAGGGAGGGCGCGGCCCGCGCGGAGGAGA  
TCCGGCGGGCGGCAGCGGCGGCGGGTGGGCTGGCTCTCCGGCGGCGCCCTCAGC  
CGACAGCCTGGCGTAGCAGGAGCAGAGGAGCGACAGGGAAGCAGCCGACACCC  
AGCACCAAGCCGGTAGGCAAGCAGCAAGGTAGCTCACCAATTCAGTTATTCGTA  
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GAATGTAGGATTGTAGCGCATTGGTAGCGCATTGCGCATTGTTGTGAGAAAACAT  
GGAATGCACATTGGTAGCCGGCGGTAGGAATGTACATCTGTAAATTCGCAAGCA  
ACTAGTAATTTGTGAAGATGTTTTATTCAGTGTGAATGAAGATGGTATAATGGTA  
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GTATTGTTGTCTTATTTGAACTATTGTAAGTGTGTAAGCTCTCTTTTGAGGCAAT  
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TTAAGTTGATAATTTACATGAGATATTCAAAGTTTTGACTTATTGGACCACCCTA  
GCTTCAAATCCTATATAAGCGCGACTGACCTGATGCCTCCATTCCTTTCATGTTTT  
GCTCCAGAGCCATGTAACCCTCCACCAAAAAGACATCAGGTGCTCTGCGTTTTGCT  
CCAGAGCCATGTGACCCCACTACCAAAGGATATCCTGGCAAGGTGCTTCTCGGGC  
TGACCATCGAAACAGAGAGCTTCATGCCTGCCTAGTACTAGCAGCTAACGATCTA  
GCTAGTAAGATGGCATCCATGATCGCCTCGTCCAAGATGATTGCACTTCGTGCAC  
TGATCTTCCCTCCTTGTCTTGTATGGCTCATGCACTAGGATCGTCAACTTCAAT  
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GCTACCAGGTACGTTACTTGGTCAGAGATACTATTTCTAGGATGTGACAAGCTT  
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AACTAGATGAGGCACAGTGGACTAAATCCTTTCATTTGTTGCTGACCTTTCTCAGA  
TAAGATGCCTCAACCACCCTGCATGCTCCGGCAATCCGGAGACGGTGGCAATCAC  
TGACATGAACTACTACCCAGTCGCCAAGTACCACTTTGACCTCAGCGGCACGGCG  
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TCGACATCCAGTTCAAGAGGTCTGTGACATGCACTGCTTATTTTCATAGAACCAGA  
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CCAAATTACATATAACATGGCCTGTTCCCTGTAGGGTGCCCTGCATCTACCCCGGG  
CAGATGGTACCTTCCACATCGAGCACGGCTCAAACCCTAACTACTTGGCGGTGC  
TCGTTGAATTCGAAGATGGCGACGGCGATGTCGTCCAGGTGGACCTCATGGAGGC  
CAACTCCGGGTGGTGGACGCCGATGCGGGAGTCGTGGGGATCCATCTGGAGGTT  
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CCGTAGTTAATTTGCAGGTGACAAGTGATATGTCGGGTGTCCTATGCGGCTATGC  
TGTAGGAGGTGTTGTGTGTCTTCAACAATTTGATGATGAAAACCATAATATTCCT  
ATGAATTCGTTGATTCCTCTGCATTTGTTTTCCCCACCTCTGAAAACATAAAA  
AAAAACTCGTCAG