

## IDENTIFICATION

**Species:** *Panicum hallii*

**Locus:** Pahal.9G388400

**Gene Model:** Pahal.9G388400.1.p

**Description:** PhEXPB-22

**Family:** Beta Expansin

**3D structure:**



## GENOME DATABASES

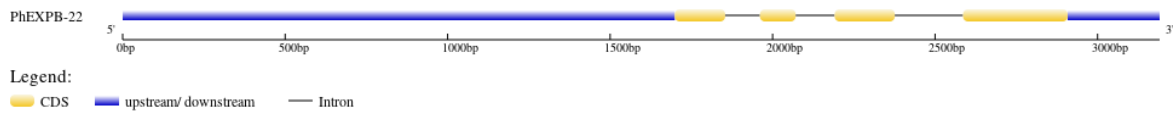
Phytozome: [https://phytozome-next.jgi.doe.gov/info/Phallii\\_v3\\_1](https://phytozome-next.jgi.doe.gov/info/Phallii_v3_1)

KEGG: <https://www.genome.jp/entry/T07366>

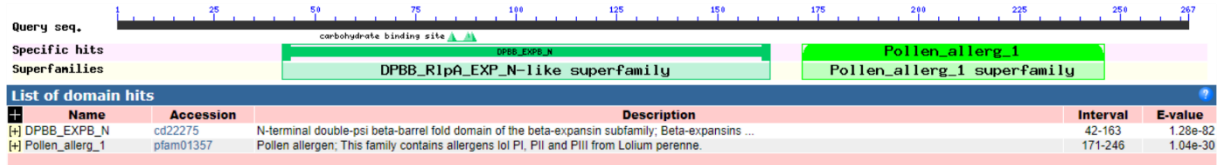
## EXTERNAL RESOURCES

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## GENE STRUCTURE



## DOMAIN ARCHITECTURE



## SEQUENCES

### Peptide

>PhEXPB-22

MASSSSKAAALAALLFSLLVTYGSCARPVSFNASAFTADPNWEAARATWYGAPTGA  
GPDDDGGACGFKNVNLPPFSAMTSCGNEPLFKDGGKCGSCYQIRCTNHAACSGNPET  
VIITDMNYYPVAKYHFDLSGTAFGAMAKPGRSDELLHAGIIDIQFKRVPCNYPGQKV  
TFHVEEGSNAVYLA VLVEFEDGDGDVVQVDLMEANSGSWAPMRESWGSIW RMSDN  
HRLQAPFSLRITNESGKQLVASNVIPADWVPNTYYRSIIQY\*

### CDS (coding sequence)

>PhEXPB-22

ATGGCCTCCTCCTCTTCCAAGGCTGCTGCACTTGCAGCACTACTCTTCTCCCTCCT  
TGTCACGTATGGCTCGTGCCTCGGCCGGTGAGCTTCAACGCCTCCGCCTTACC  
GCCGACCCCAACTGGGAGGCCGCCAGGGCCACCTGGTACGGCGCGCCACCGGC  
GCTGGTCTTGACGACGACGGTGGCGCCTGCGGGTTCAAGAATGTCAACCTGCCCG  
CGTTCTCGGCCATGACGTCGTGCGGCAACGAGCCACTGTTCAAGGACGGCAAGG  
GCTGCGGATCCTGCTACCAGATACGGTGCCTAACCACGCTGCGTGCTCCGGCAA  
CCCGGAGACGGTGATCATCACCGACATGA ACTACTACCCGGTCGCCAAGTACCAC  
TTCGACCTCAGCGGCACGGCCTTCGGCGCCATGGCCAAGCCCGGCCGACGACG  
AGCTCCTCCATGCCGGCATCATCGACATCCAGTTCAAGAGGGTGCCCTGCAACTA  
CCCCGGGCAGAAGGTGACGTTCCACGTCGAGGAGGGCTCGAACGCCGTCTACCT  
GGCGGTGCTCGTCGAGTTCGAGGACGGCGATGGCGACGTGGTGCAGGTGGACCT  
GATGGAGGCCAACTCCGGGTCGTGGGCGCCGATGCGCGAGTCTTGGGGATCCAT  
CTGGAGGATGGACTCCAACCACCGGCTGCAGGGCGCCCTTCTCGCTGCGCATCACC  
AACGAGTCCGGCAAGCAGCTGGTGGCCAGCAACGTCATCCCGGCCGACTGGGTG  
CCCAACACCTACTACCGCTCCATCATCCAGTACTAG

### Nucleotide

>PhEXPB-22

CGCGCTACACTCGTGTTCATCCTGTTTTCTTGTTCCTCGGGGGAAGGCCCAACAACA  
CCGTATGCCAGAGTCTGAGTACCCCTAATCTATGGTACTGTCTCTCGTGGATTAT  
GGATGGTGTGATTTTTAGGAGAGAAAAGGGCAAGCCCTCATTAACTCCCATC

ACGGAGAAGAAAGTGATCCAAGTTCTACAGCATCCCAGCACAGGCACAGAATTG  
AGCGTAGACCCAGGCACAAGCACAGCCAAAACCTTGAGGTGTCTCACCCCTCGCCG  
GAGGTGATGAACTGATGTACCTTGTGCTGCCATGTTTAGTACATCATGTAAGTAA  
GTGTTACATGCGTATGTGCTCAGCATGCTGCTACACTGTGTCATCGATCGACGCG  
GTCTCGTGGACGGTCACCTCGATTTTAATTACCCGTTCCCTGCCTGCTACACTACTA  
GCATCATTTTTATTTCATATTTTGGCTCGTTTCTATGTACCCAAATATGCGTGCCCGGT  
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GCATTTTATTTCCTAGCCATAGGCAGGGAGGAGGTGAGTAAGGCTTGGCCGCAAG  
AAGGGCAGAGCCACCTAGCCGTTCCCGGACATCATTTCGGCCAATTTCCCTCGAATGG  
TGCAGCATCTCAGAATAGAATATTTCTCTCAAATTGGATTACCTCAAGATGACTT  
GATTCAGTACACTGAACCAAACACAGGGACCACCCTTCATTTCAGGATCATCCCCT  
TCACGTACCATTTCGCATGATGCACCAAGCACTGCCTGCATGTTCAACTTGGTCCA  
GCCGACCAAAAAGGCATCAGGATAGGAGCGAGGACTCGCGGACTCGGTAGTAGT  
ACCCTGCAAGACAGTCGCATCTCTAGAAAATTGAGGCCCATGCGTTTCCCGTGGC  
ACAACTTGTAGGGCGGCTTCTACGCGCGGCATACTGGACTCTGCATTCACTTT  
GCACCACGCATTTCTTCCAATTCCCGACGGACAGCAACATGGTGTAAATGGTGT  
CGCCGTATCGTTGGATTAGTACCAACTTGGCAACTTCCAGCGCAACTGAAACGGA  
TTTGGTATCTCTTTACCCAGCTCTGTATTGCAGTTACAGTACCTACGCATCATTGA  
GGGGTTTGGTAGACAGAATCCATAACAATGATGACCCGGCAATGATGGTGCCCTCC  
CATGATTCAATTGCAGACCTGTCTCTCCTTCTGTATCCGGCTGTGCGCGAATCTCC  
CGCTCACGACACCCGGGACAGGAGTCTTGAAAATGTTCCAAGGATCACTGGATCA  
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ATCCCAGCGCACAACTCCTTCTCCTGTCTGCCCGGTGGTAGTAGTACTAGCTACCC  
TCGCAAGCTTGCAGCTTGTCCGACTTAGCTTGTAGCAGCGCTAGTTGCCTCGCTA  
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CGGCGCGCCACCAGGCGCCGGCCCATACGACGACGGTAAGTTTCGTGTGTAGATG  
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TGTAAACTTGCATCCATTTGCGTGTTCATGCAGGTGGCGCCTGTGGATTCAAGAA  
CGTTAACCTGCCGCCGTTCTCGTCCATGACGTGCTGCGGCAACCAGCCCCTGTTCA  
AGGACGGCAAGGGCTGCGGCTCCTGCTACCAGGTAGATTAATTGATTGGCCATCA  
AACAAGTTAAAGCTCTGCTGCCGCGGCCGTAATGGAAACGGAAAGCTCAAGCAC  
AGTTCTGAGTTTCGACCCACTCTGTTTTGTTTTCTTTTTTCAGATACGATGTACCAA  
CCACGCTGCGTGCTCCGGCAACCCGGAGACGGTGATCATCACCGACATGAACTAC  
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CTAATCTCTGGAGGCCTGATGACAAGACTGATGGAAACATGTCACATGCAGGGTG  
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CCCATGTACCTGGCGGTGCTCGTCGAGTTCGAGGACGGCGACGGCGACGTGGTGC  
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GGGGATCCATCTGGAGGATGGACTCCAACCACCGGCTGCAGGCGCCCTTCTCGCT  
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CAACTGGGCGCCCAACACCTACTACCGCTCCATCATCCAGTACTAGCCCTGCTAC  
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GGAGGCCGGTTTGGATGTGCTTTCTCGCCCGGTGCCAATGTATTTTTCAGTGTGAT  
CATCAAGACTATAATTGGAAAACCTGGTTCTGTGTGGCATTTCAGAACGGACAT