

IDENTIFICATION

Species: *Citrus clementina*

Locus: Ciclev10032524

Gene Model: Ciclev10032524m

Description: CclEXPA-08

Family: Alpha Expansin

3D structure:



GENOME DATABASES

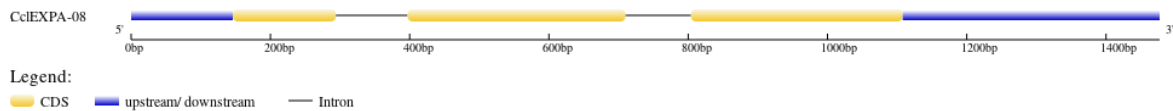
Phytozome: https://phytozome-next.jgi.doe.gov/info/Cclementina_v1_0

KEGG: <https://www.genome.jp/entry/T02982>

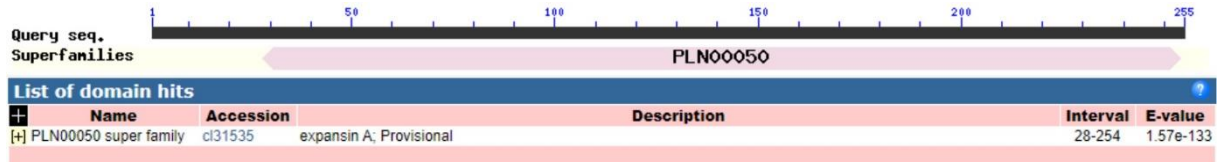
EXTERNAL RESOURCES

<https://www.citrusgenomedb.org/organism/Citrus/clementina>

GENE STRUCTURE



DOMAIN ARCHITECTURE



SEQUENCES

Peptide

>CclEXPA-08

MANIAASSSVALLLFVLNFCFRGTFGDYGGGWQSGHATFYGGGDASGTMGGACGY
GNLYSQGYGTNTAALSTALFNGLSCGSCYEMKCENDPKWCLPGSIIVTATNFCPPN
LALSNDNGGWCNPPLQHFDMAEPAFLQIAQYRAGIVPISFRRIPCAKKGIRFTVNGH
SYFNLVLTNVGGAGDVHSVSIKGSKTGWQAMSRNWGQNWQSN SYLNGQSLSFQV
TASDGRTVTSNNVVPANWQFGQTFEGGQF*

CDS (coding sequence)

>CclEXPA-08

ATGGCCAATATTGCAGCATCATCTAGCGTTGCTCTCCTTTTGTGGTCAATTTT
TGCTTTCGTGGCACATTTGGAGACTATGGTGGTGGGTGGCAAAGTGGCCATGCAA
CTTTCTATGGCGGGGGTGTATGCTTCCGGCACAATGGGTGGTGTCTGTGGGTATGG
CAATTTGTACAGCCAAGGCTATGGGACTAACACTGCAGCACTTAGTACCGCTCTA
TTCAACAATGGCCTAAGCTGTGGTTCATGCTATGAAATGAAATGTGAAAATGACC
CCAAGTGGTGCCTCCCCGGCTCCATCATTGTCACCGCCACCAACTTCTGCCACCT
AACCTTGCCCTGTCTAACGACAACGGCGGTTGGTGCATCCTCCCCTCCAGCACT
TTGACATGGCTGAGCCCGCTTTCTTGCAAATTGCCAATACCGCGCCGGCATCGT
CCCAATTCCTTCAGAAGGATCCCGTGTGCGAAGAAAGGAGGAATAAGGTTTACC
GTCAATGGACTCATACTTCAACCTGGTTTTGGTCACAAATGTCGGAGGAGCAG
GAGATGTACATTCAGTATCAATCAAGGGTTCAAAGACTGGATGGCAAGCAATGTC
AAGGAACTGGGGCCAAAATTGGCAGAGCAATTCTTATCTTAACGGCCAAAGTCTT
TCTTTCCAAGTGACAGCCAGTGACGGCAGGACTGTGACTAGCAACAATGTTGTGC
CTGCAAATTGGCAATTTGGGCAAACCTTTGAGGGTGGTCAGTTTTAG

Nucleotide

>CclEXPA-08

GGTGGCTGTAGCTTTACAAAGTCCGCATCTCACTGTCTATATAAAGGAGGCTCCA
TTCCCCCTGCACTAAGGCACTAAGCACTCAAAGCATCTCTTTTTCTTAACAAAA
AAACACATAAACATAAGCAGCCAGCAGTAGCAGCCATGGCCAATATTGCAGCAT
CATCTAGCGTTGCTCTCCTTTTGTGGTCAATTTTTGCTTTCGTGGCACATTTG

GAGACTATGGTGGTGGGTGGCAAAGTGGCCATGCAACTTTCTATGGCGGGGGTG
ATGCTTCCGGCACAATGGGTAAAGTGTATAAGTAACAAGTTAAAACATTTGTCTCT
TTTGTTGCATAACTTTTTGGTCAATTTGCCATGATGGCAATTGATGTTTTCTACTTG
GTACTGTAGGTGGTGGCTTGTGGGTATGGCAATTTGTACAGCCAAGGCTATGGGAC
TAACACTGCAGCACTTAGTACCGCTCTATTCAACAATGGCCTAAGCTGTGGTTCA
TGCTATGAAATGAAATGTGAAAATGACCCCAAGTGGTGCCTCCCCGGCTCCATCA
TTGTCACCGCCACCAACTTCTGCCACCTAACCTTGCCCTGTCTAACGACAACGGC
GGTTGGTGCAATCCTCCCCTCCAGCACTTTGACATGGCTGAGCCCGCTTTCTTGCA
AATTGCCCAATACCGCGCCGGCATCGTCCCAATTTCCCTCAGAAGGTAACAACAT
ATCACATTATAAAGTTGATGTCACTACACTTGTAAGCCTGTATGAATCTGATTTTG
ACATGTTTCTATCCGGATTTGTAATCAGGATCCCGTGTGCGAAGAAAGGAGGAAT
AAGGTTTACCGTCAATGGACACTCATACTTCAACCTGGTTTTGGTCACAAATGTC
GGAGGAGCAGGAGATGTACATTCAGTATCAATCAAGGGTTCAAAGACTGGATGG
CAAGCAATGTCAAGGAACTGGGGCCAAAATTGGCAGAGCAATTCTTATCTTAACG
GCCAAAGTCTTTCTTTCCAAGTGACAGCCAGTGACGGCAGGACTGTGACTAGCAA
CAATGTTGTGCCTGCAAATTGGCAATTTGGGCAAACCTTTGAGGGTGGTCAGTTT
TAGATTTTAAGATTTTAAAAAAGTGAATCAGAAATTGTATATTTAATTGAATTTG
GCATTGTGTTTGTGAGGGGAGAAAGGGTGAGAATAGGGGTAGGTGAGGCTGAGGC
CTGGCCGATGTGGCCTGTCTTGTGCTGTGGTGGTCTGTTGGCACCCGCTAGGCCT
CTATAAATAGAAATGATTGTGTTCTATATATAGTTATTACTCCAATTGAATGAACG
GGTATAATTCATTCAACGTTGTGATTTCTGGTTATCTTTTGAGACATCAATAAA
AGTTTATTGATAAAAAAATTTCAATCTTAATGTATAATTCAGAATTTTGTTTATGA
AAAATAAAAAATTTCCCTCTCCAAATTATTAACACTACGTA