



**Figure S1.** The carotenoid biosynthesis and cleavage pathway in plants (modified from Moise et al. [9] and Colasuonno et al. [21]). PSY, phytoene synthase; PDS, phytoene desaturase; ZDS,  $\zeta$ -carotene desaturase; ZISO, 15-cis- $\zeta$ -carotene isomerase; CRTISO, carotene isomerase; LCYE, lycopene  $\epsilon$ -cyclase; LCYB, lycopene  $\beta$ -cyclase; CYCB, chromoplast-specific lycopene  $\beta$ -cyclase; BCH,  $\beta$ -carotene hydroxylase; ECH,  $\epsilon$ -carotene hydroxylase; ZEP, zeaxanthin epoxidase; VDE, violaxanthin de-epoxidase; NXY, neoxanthin synthase; CCS, capsanthin-capsorubin synthase; CCD7, (9',10') carotenoid cleavage dioxygenases; CCD8, (13,14) carotenoid cleavage dioxygenases; CYP711A1, cytochrome P450-type enzyme; NCED, nine-cis-epoxycarotenoid dioxygenase.

The sequence of the cloned *CYC-B* gene:

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CCACTTCTCTTGACCTCATTCCTCGCTCTAATGGCCACCCCTGCTCCGGCCATTCCACCGCCAC
CATCCGCCGCCAAACTTCCAATTCTCCACTCCTCCGGCTCTCCATTCTCCAAAACC
CATTACCCATCTCAAATAATCTCTCCAAAATCCACAGCAGCAAGTTGGCAACTTCTCG
ACTTAAAACCAGAGGCAAAACCCGAGTATTGCACTTCGATCTCCATCAATTGACCCGTCAAC
CCGGTCTCGCTTAGATGTGATCATCATCGGGACCGGCCGGCTTCGCCCTGCGGAGCAA
CTTTCTCGCTACGGCATTAAAGGTATGCTGCGTTGATCCTCTCCTTTCTATGTGCCAAGTAAC

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TATGGAGTTGGGTGAAGAATTGAAAGCTGAATCTGAAAGTTGCTGGACAAATATGGC  
CTATGGCTCTGTCATGTGAATGATAGTAAGACTAAGTITGGACCGCCCTATGCCAGAGTC  
AGTAGGAAGAAACTCAAGACTTGTGCTGGAGAGGTGTCTCTCGAATGGGTTCAATTTCATA  
GGGCCAAGGTTGGAAAATCGAACACGAAGAGTCGAGTCTCGATTTGTGATGATGGAA  
TGAGCTCAAGGCAAGCTGATTGATGCTAGTGGTTGCAAGCAGTTCTGAGACTATGAG  
AAGCCTAGGAACCATGGATATCAGATTGCTCATGGTATCTGGCTGAAGTCCAAGACACCCCT  
TTGATTGGATAAGATGCTCTGATGGATTGGAGAGATCCCATCTCGAAACGAGCCTTATTG  
CGCACTAGTAATTCTAGATTCCAACCTTTTGATGCAATGCCGTTGATTGAACTTGGT  
TTGGAAGAAACTCGCTTGTAGTAGGCCGGTGTCTTATATGGAGATTAAGAAACGAATGG  
TTGCAAGGCTAAGGCATTGGGATTAGAGTGAAGAGGGTAATTGAAGAGGGAGAAGTGTG  
TCCCAATGGGGGTCCGCTTCGATCCCCAACGTGTGATGCCAATTGGAGGGACTCTGG  
GGTGGTTCACCCCTCGACTGGTACATGGTGGCTCGGACCATGGCTCTAGCCCCAGTATTGG  
GAAGCCATTGCAGAGTGCCTGGCTCAACCAGAACATGATCCGAGGGCAGCCGTTATCATAGA  
GCGTGGAAATGGCTTGTGCCAATTGAGAGGAGATGCACGAGGGACTTTACTCATTTGGT  
AGACTTTGTGAAGCTTGATCTGAATGGGAGTAGAAGCTTCTTGACGCTTCTTGACTGG  
CCCTATTACTGCCAAGGCTTTATCGTCAAGGCTGTCTAAGAGAGCTGCTTGTGAGCTT  
ATCTCTGTTGGCCGAGCCTCCGCCCATCTAGGTTGATGGTACAAAGTGTCCGTGCC  
TGGTAAACTGATGGCAATCTGCACTGAAGCTGCATAATAATGTAACAC



**Figure S2.** The three dimensional structure prediction of CYCB protein