

Figure S4

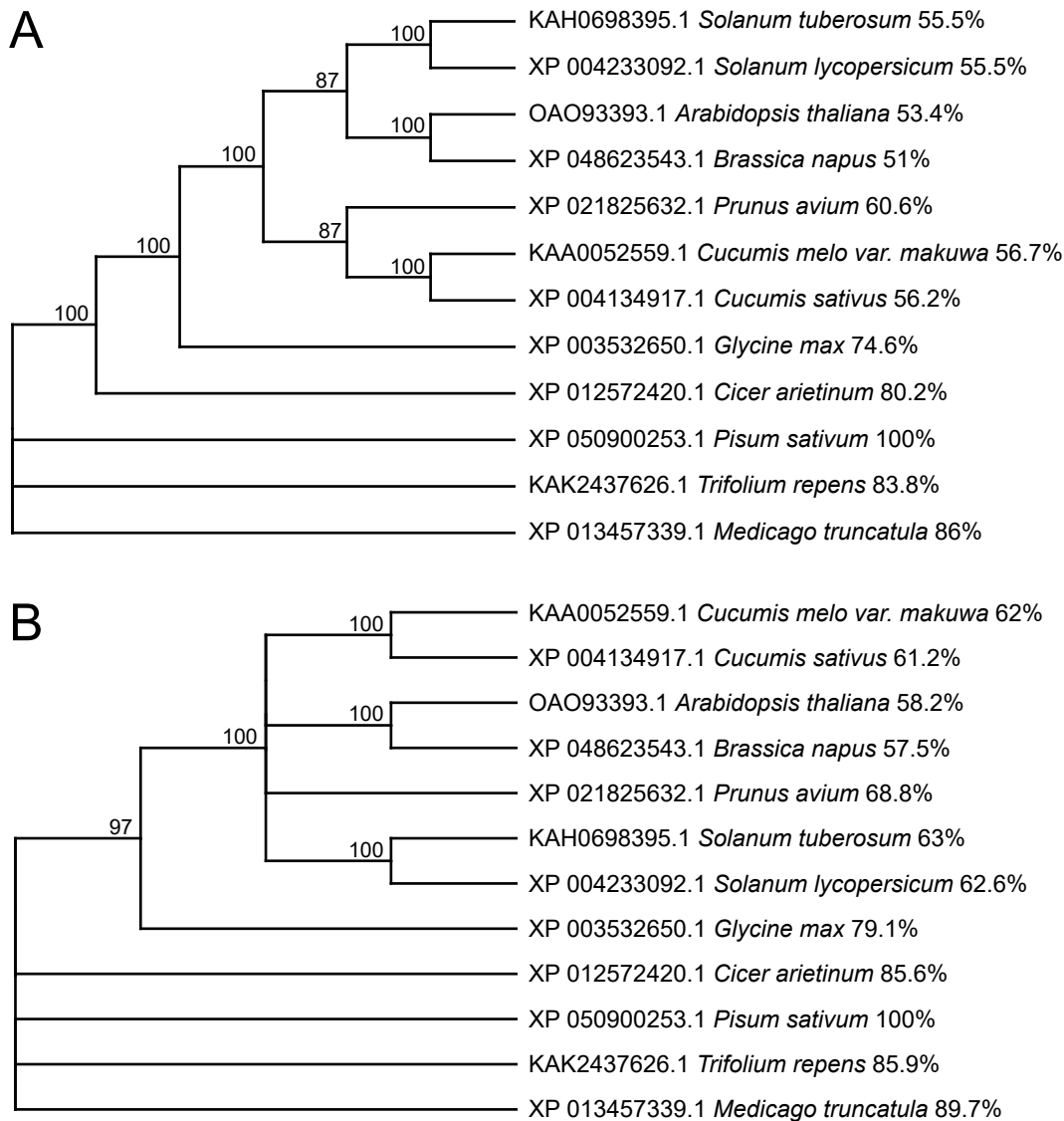


Figure S4. Phylogenies of the entire protein sequence of Psat7g091800.1 (XP 050900253.1) from 'PI180693' (A) and FLS2-like domain encoding sequence parts (B) reflect taxonomic distance of represented plant families. Homologs in other plant families have been identified with BLAST searches in the NCBI protein database, selected for best hits and maximum likelihood trees were constructed using IQ-TREE. Percent identity compared to query from *P. sativum* are indicated for every accession.