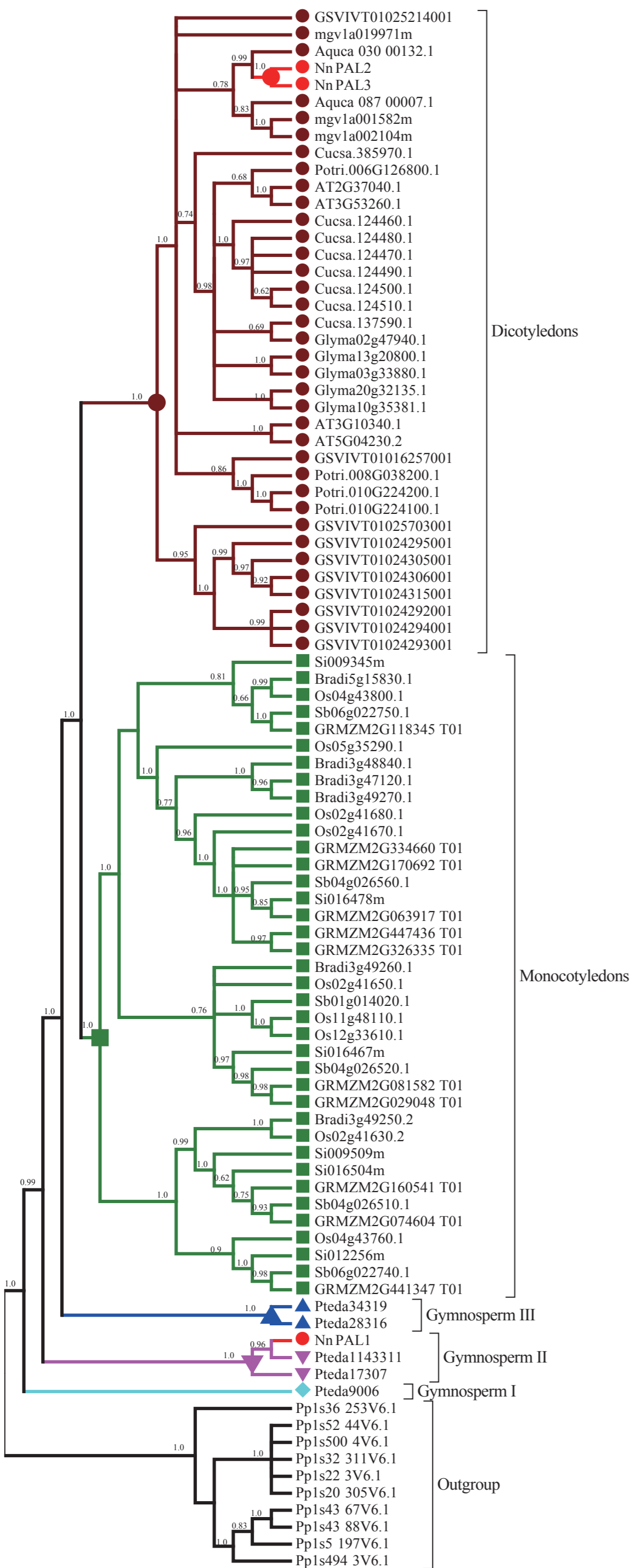


Figure S5. Phylogenetic trees of the phenylalanine ammonia lyase gene family constructed by BI method (a) and NJ method (b). Posterior probability and bootstrap values demonstrated on each branch of BI tree and NJ tree, respectively.

(a)



(b)

