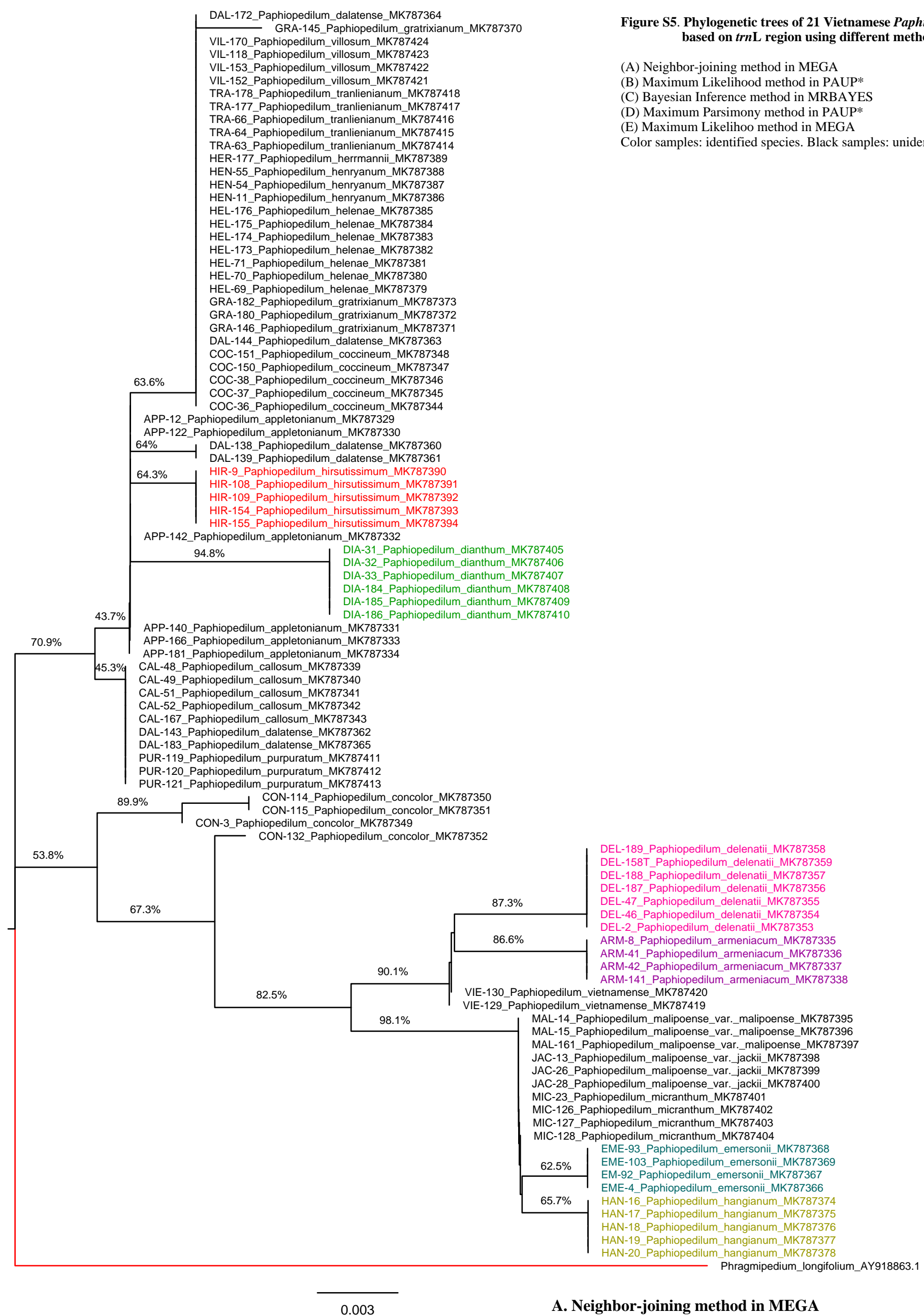
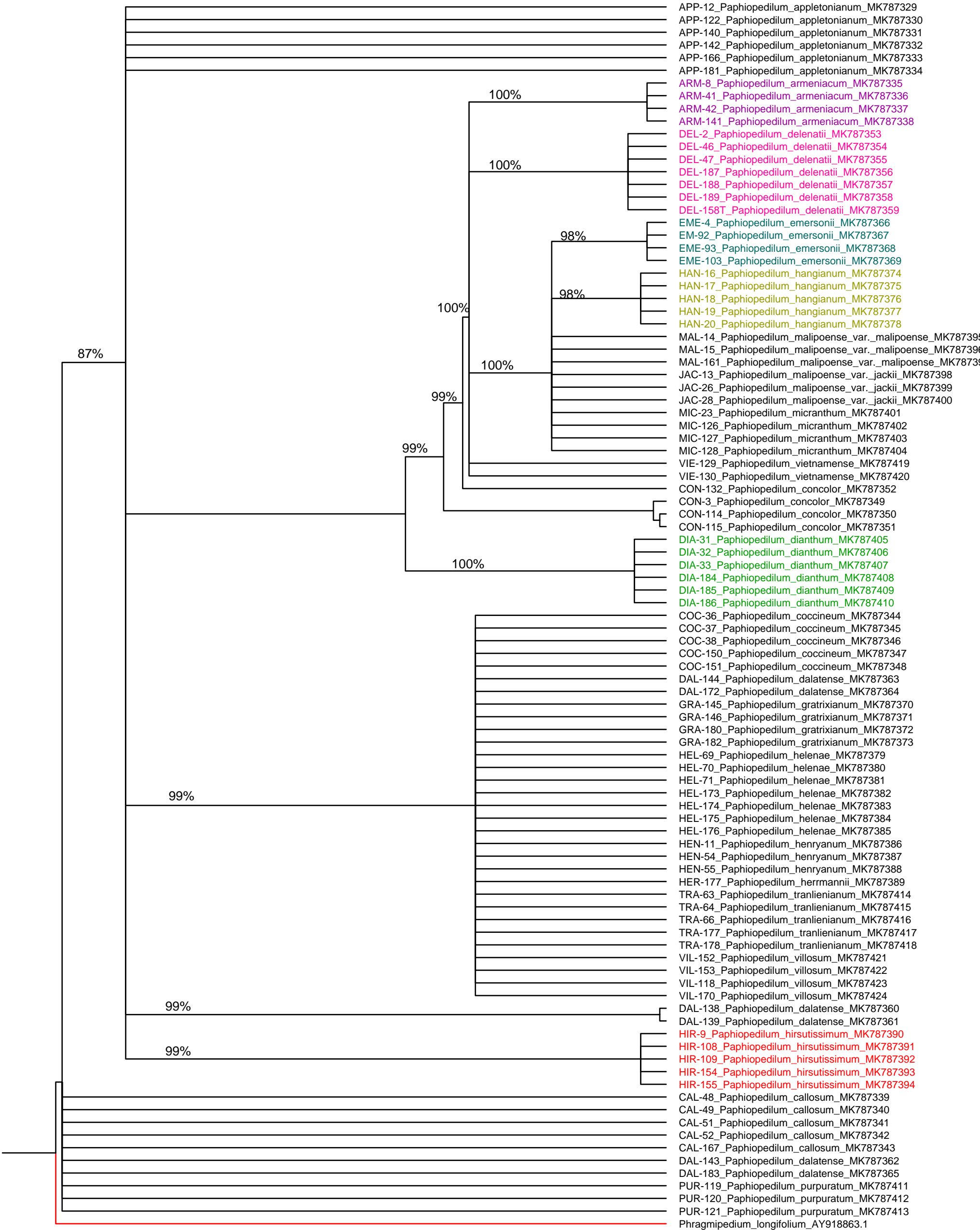
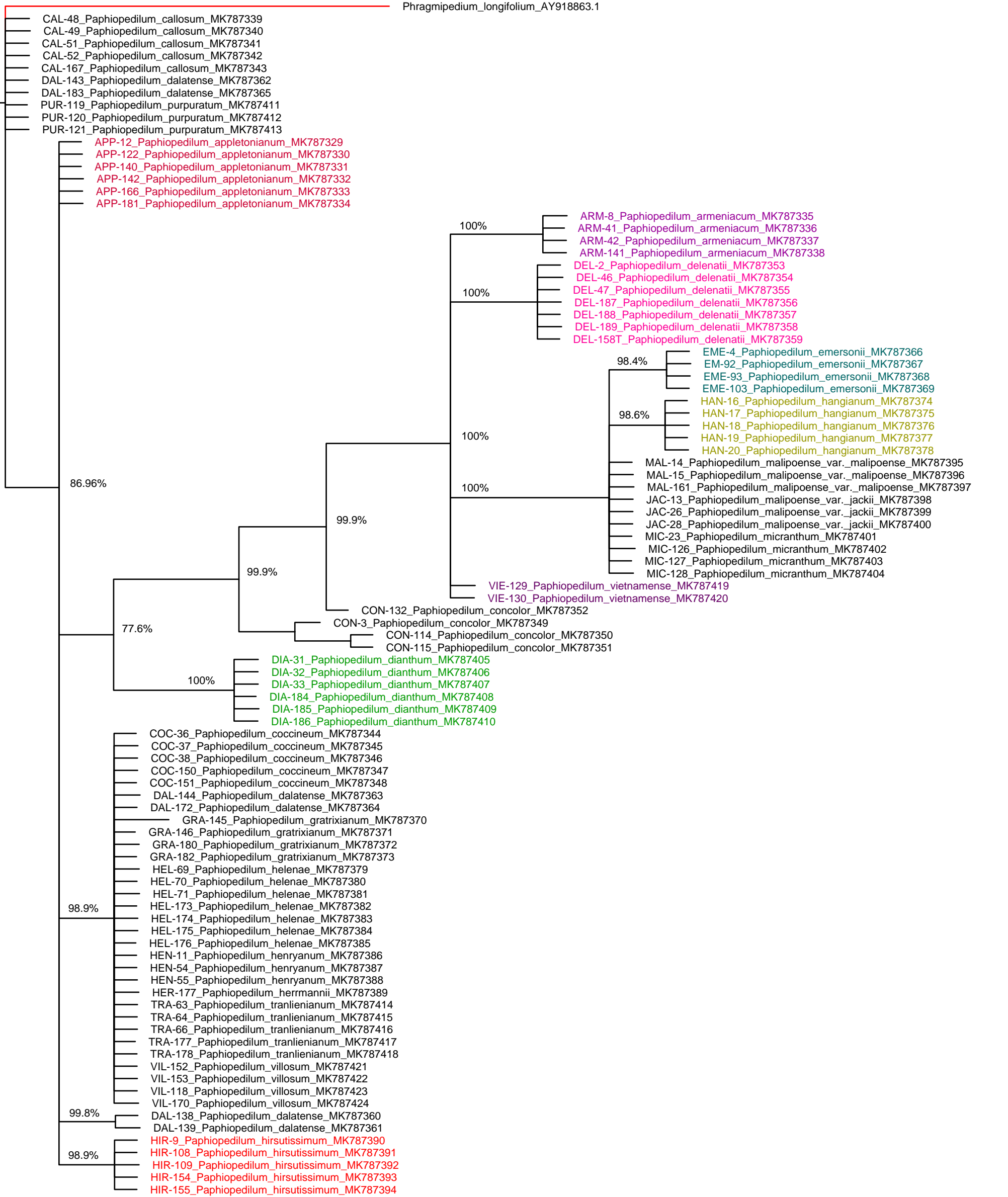


**Figure S5. Phylogenetic trees of 21 Vietnamese *Paphiopedilum* species based on *trnL* region using different methods.**

- (A) Neighbor-joining method in MEGA  
(B) Maximum Likelihood method in PAUP\*  
(C) Bayesian Inference method in MRBAYES  
(D) Maximum Parsimony method in PAUP\*  
(E) Maximum Likelihood method in MEGA  
Color samples: identified species. Black samples: unidentified species)

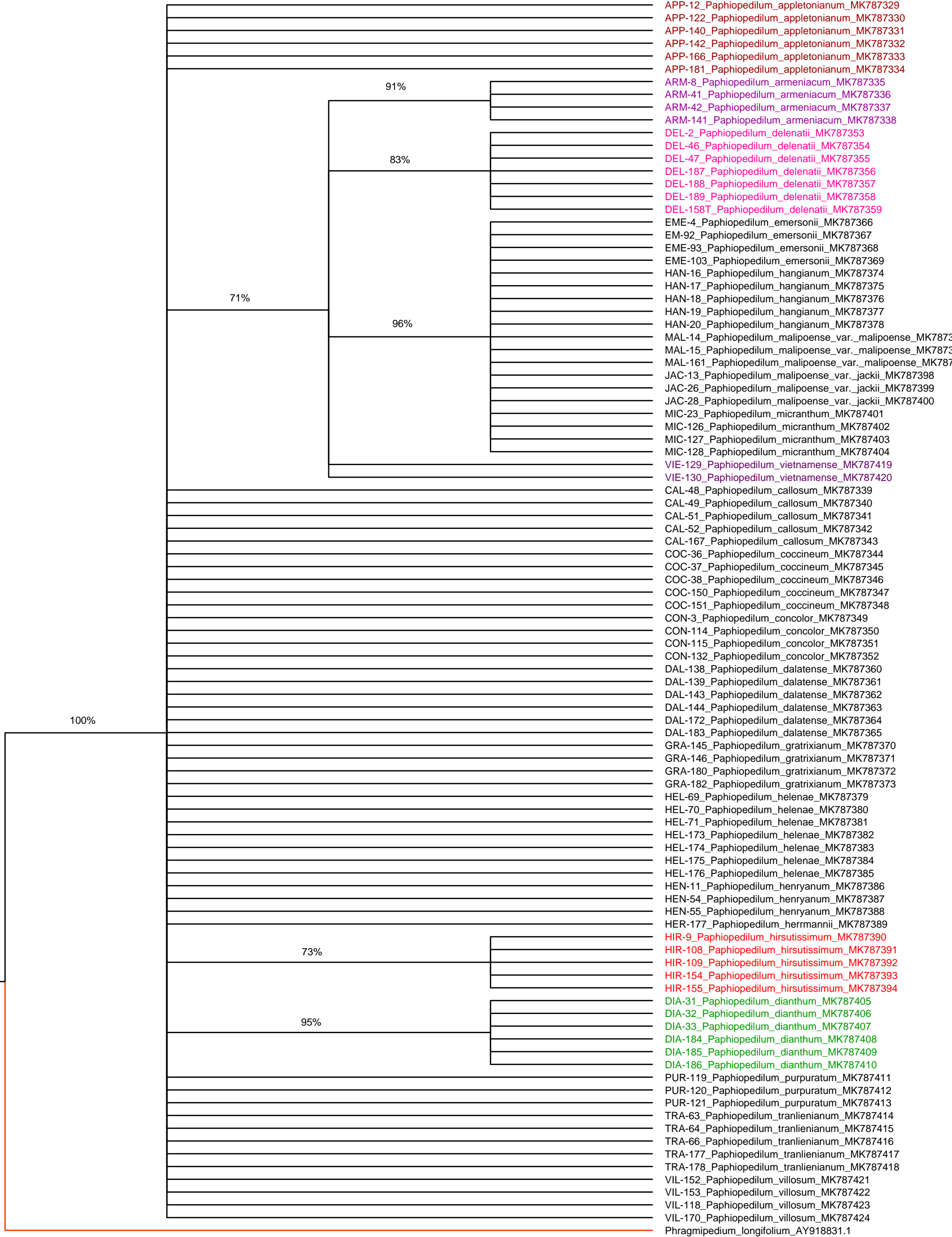






C. Bayesian Inference method in MRBAYES

9.0E-4



0.5

D. Maximum Parsimony in PAUP\*



