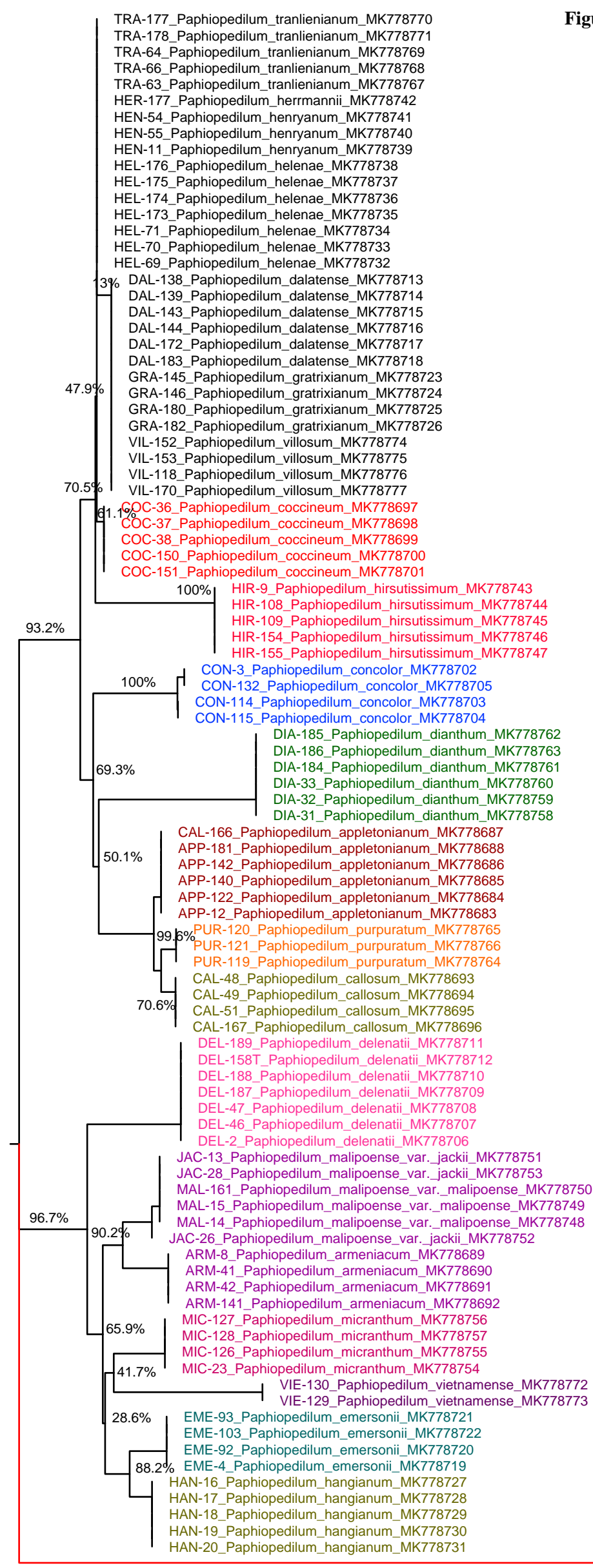
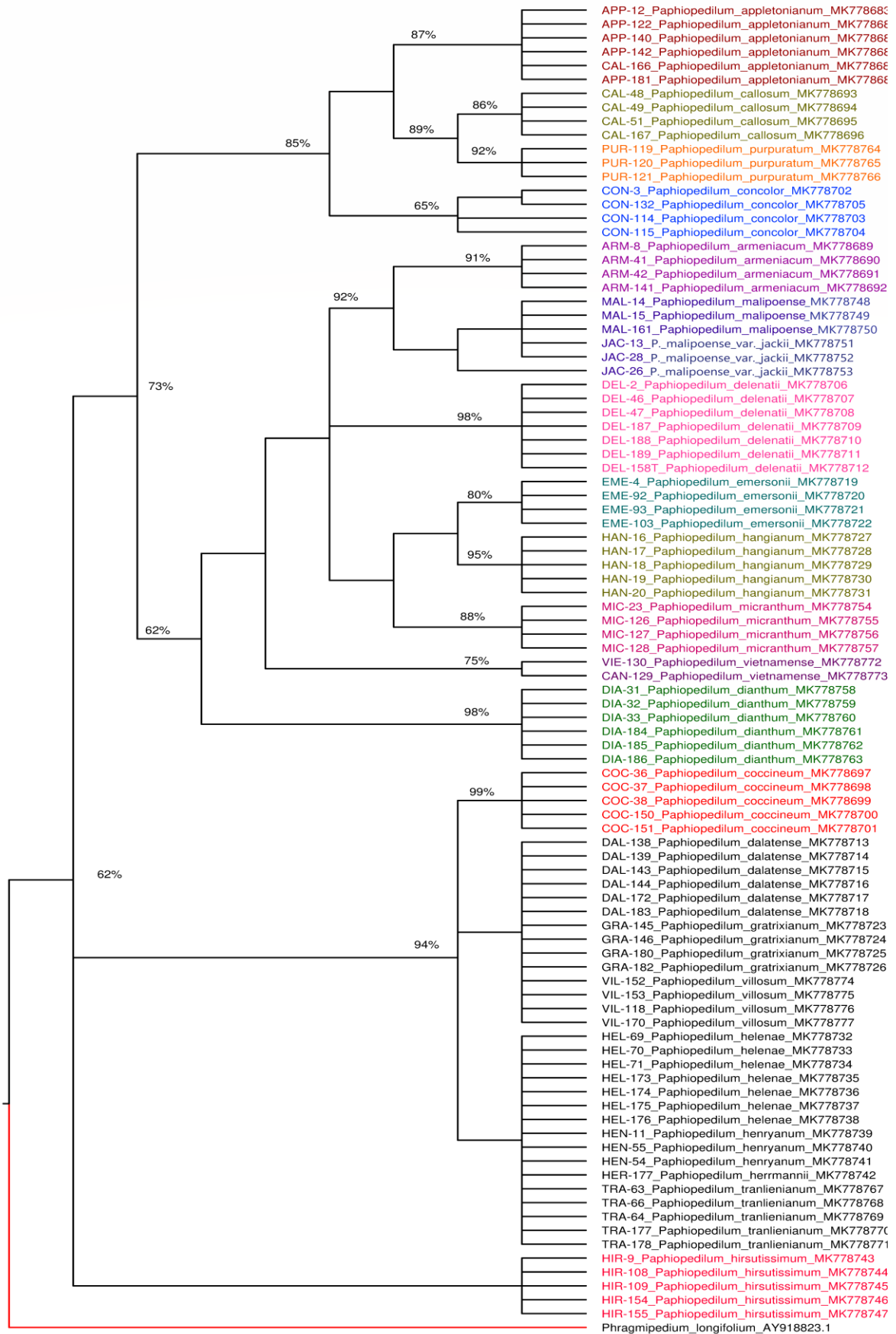


Figure S3. Phylogenetic trees of 21 Vietnamese *Paphiopedilum* species based on ITS region using different methods.

- (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)

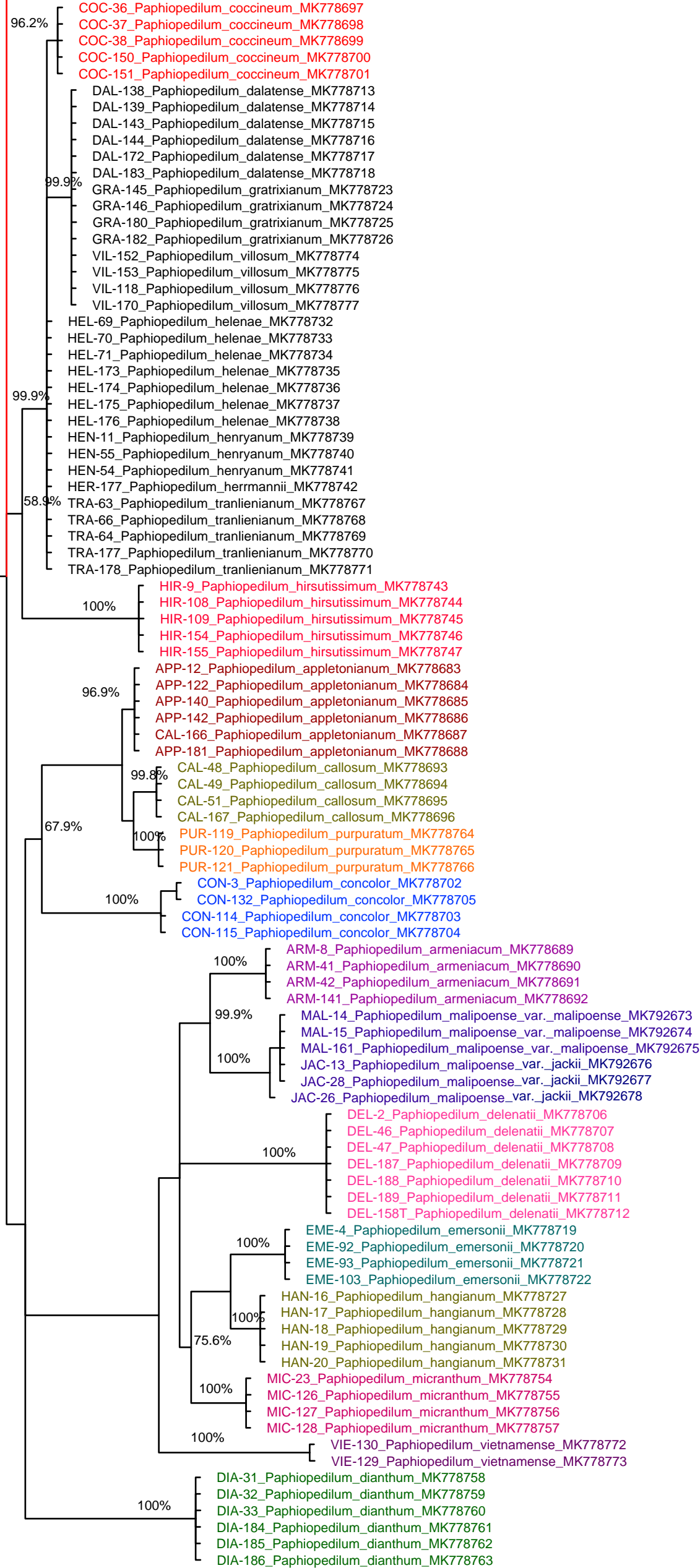


Phragmipedium_longifolium_AY918823.1

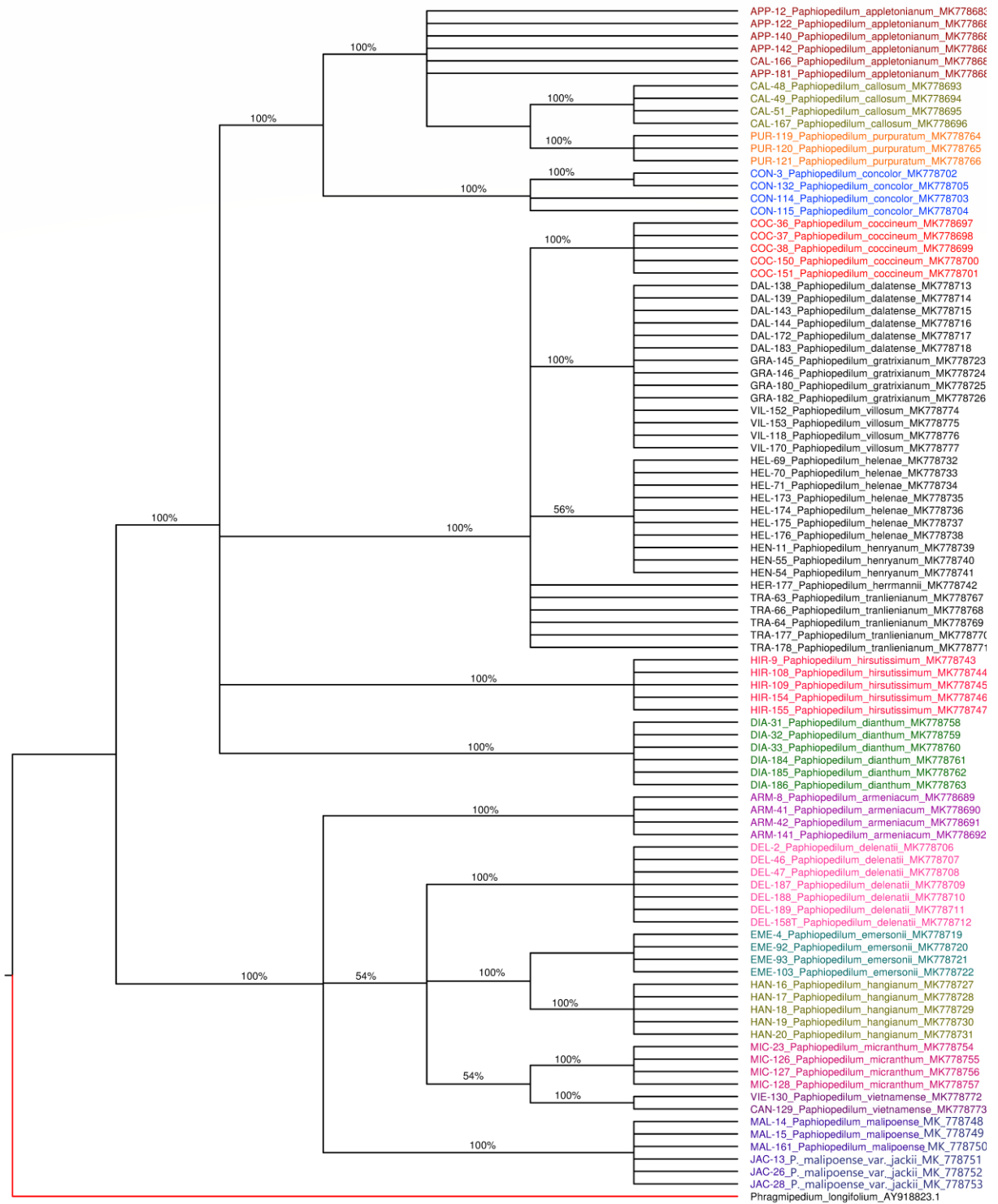


1.0

B. Maximum Likelihood method in PAUP*

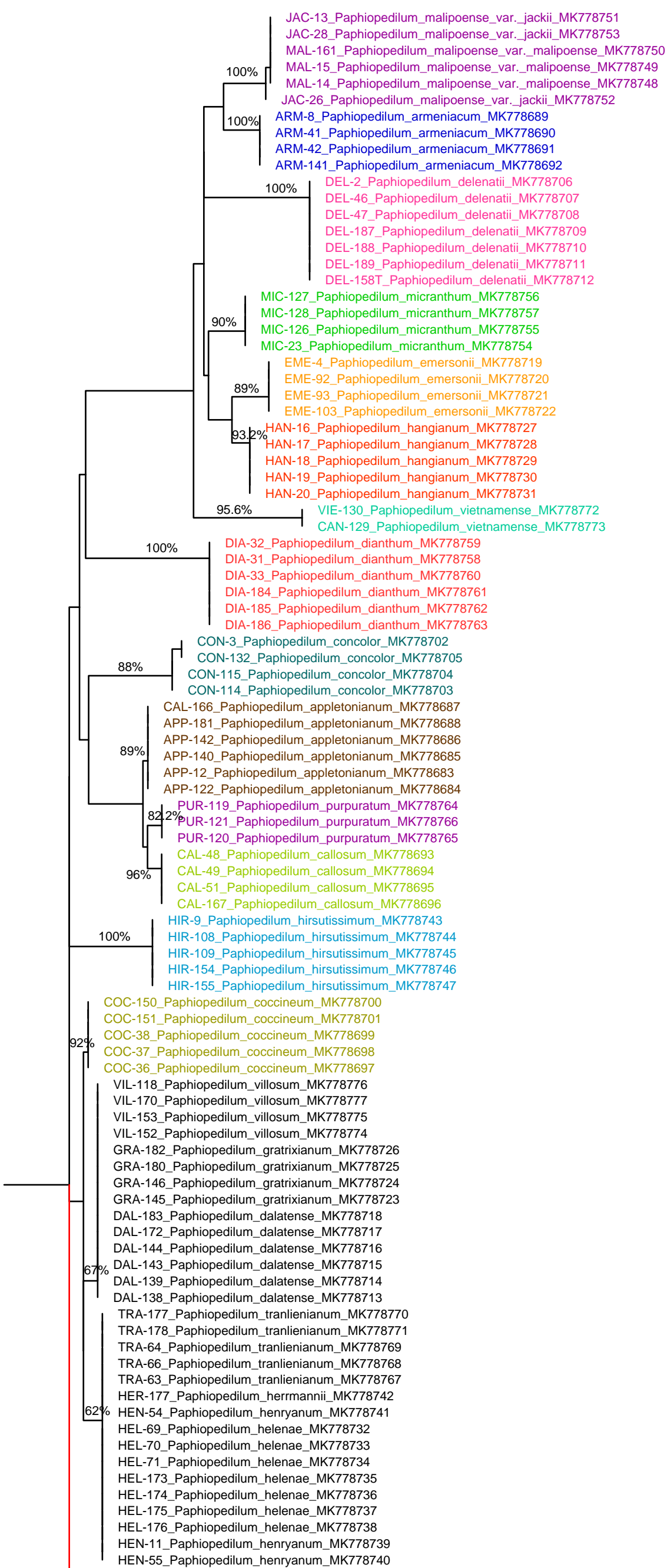


0.02



0.8

D. Maximum Parsimony method in PAUP*



Phragmipedium_longifolium_AY918823.1

E. Maximum Likelihood method in MEGA