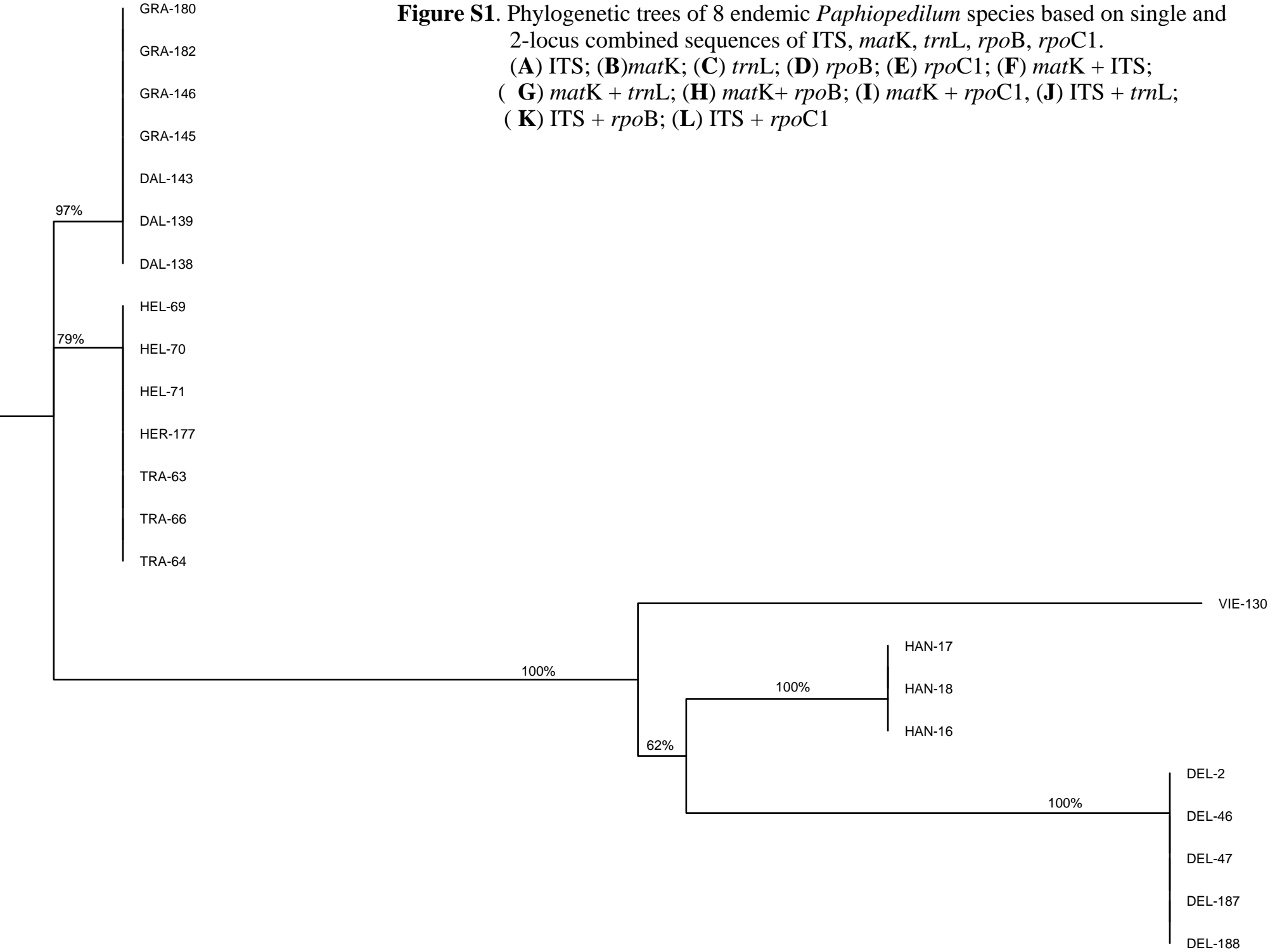
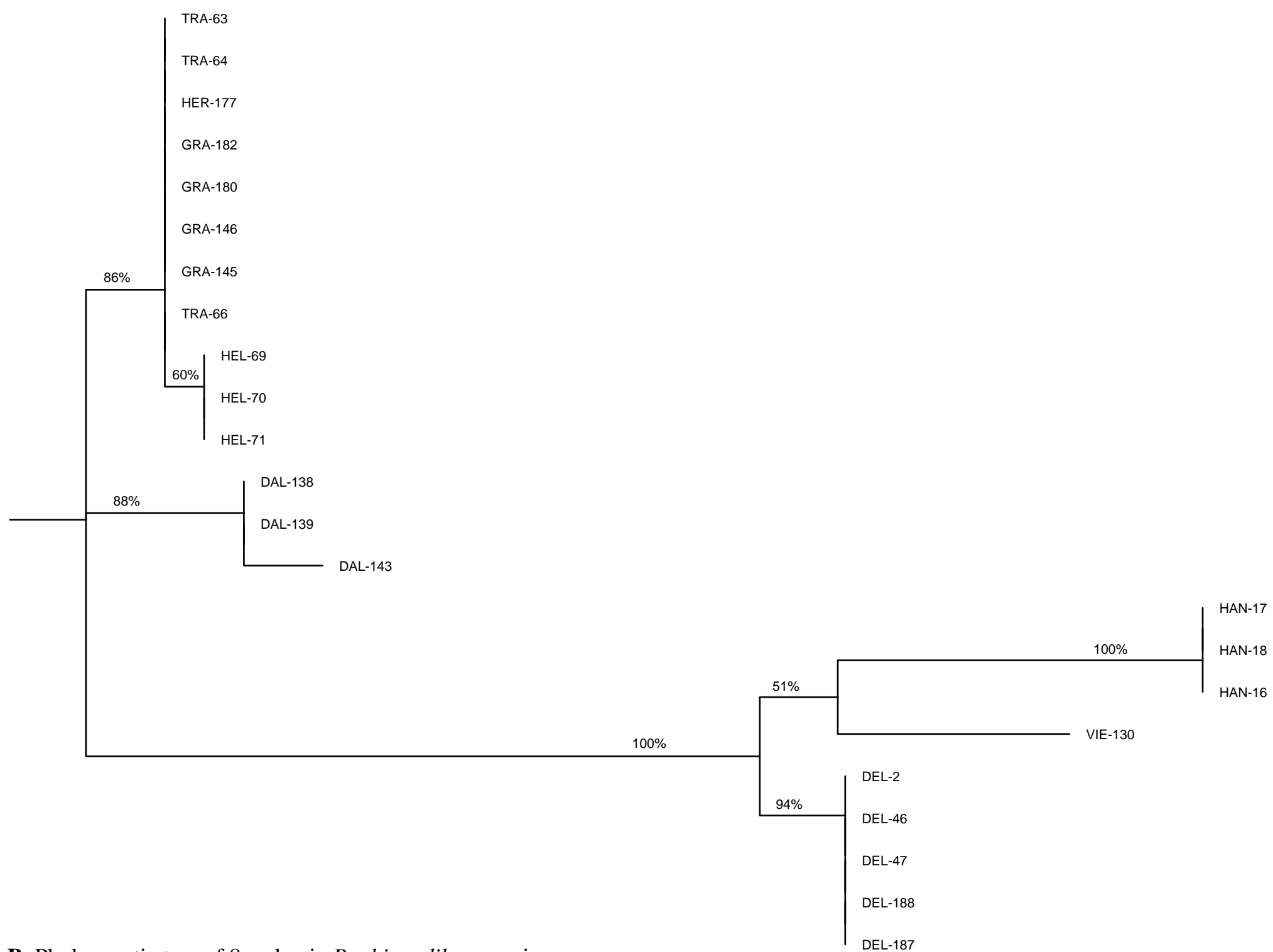


Figure S1. Phylogenetic trees of 8 endemic *Paphiopedilum* species based on single and 2-locus combined sequences of ITS, *matK*, *trnL*, *rpoB*, *rpoC1*.
(**A**) ITS; (**B**) *matK*; (**C**) *trnL*; (**D**) *rpoB*; (**E**) *rpoC1*; (**F**) *matK* + ITS;
(**G**) *matK* + *trnL*; (**H**) *matK*+ *rpoB*; (**I**) *matK* + *rpoC1*, (**J**) ITS + *trnL*;
(**K**) ITS + *rpoB*; (**L**) ITS + *rpoC1*



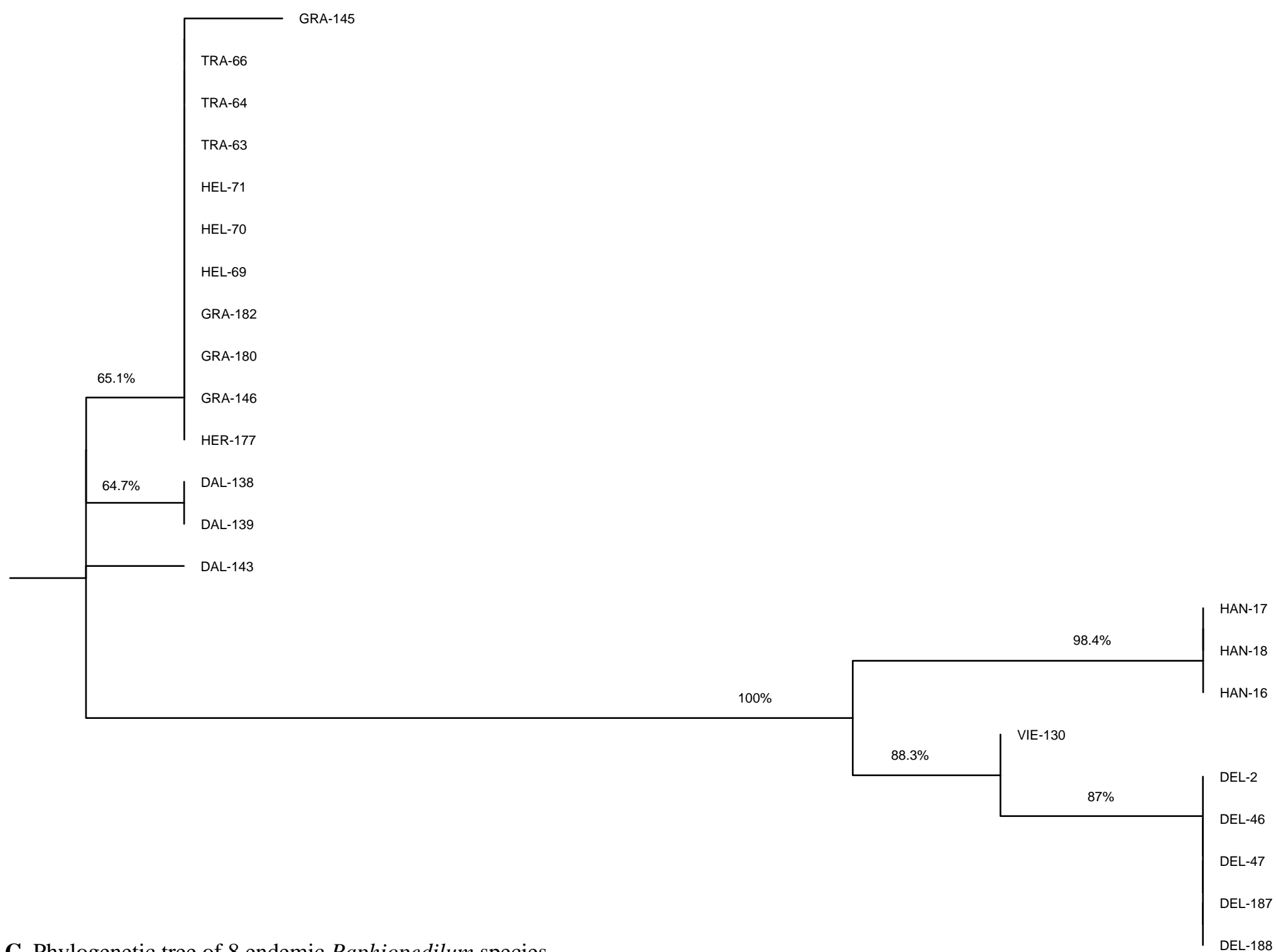
A. Phylogenetic tree of 8 endemic *Paphiopedilum* species
based on single ITS locus

0.008



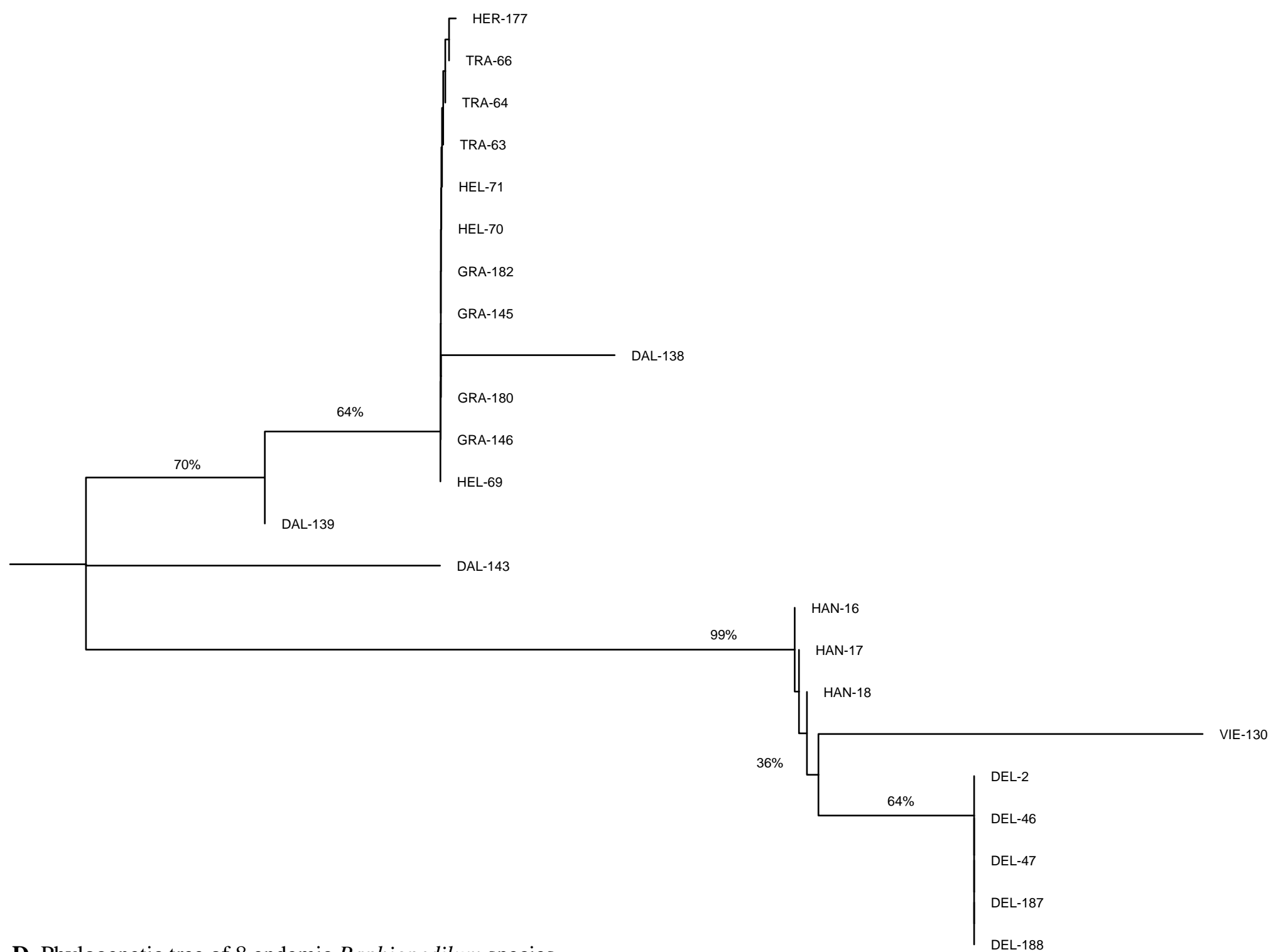
B. Phylogenetic tree of 8 endemic *Paphiopedilum* species
based on single *matK* locus

0.003



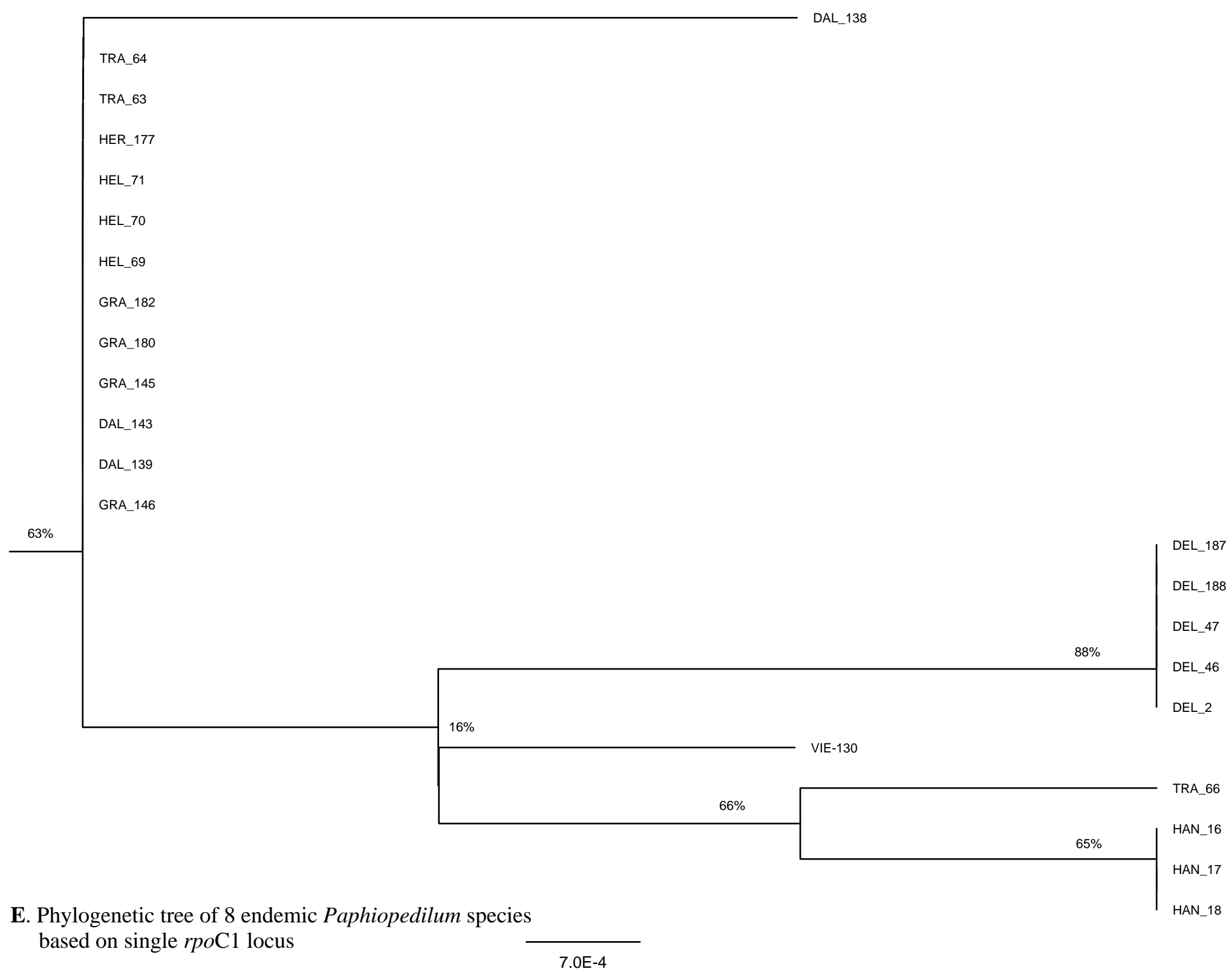
C. Phylogenetic tree of 8 endemic *Paphiopedilum* species
based on single *trnL* locus

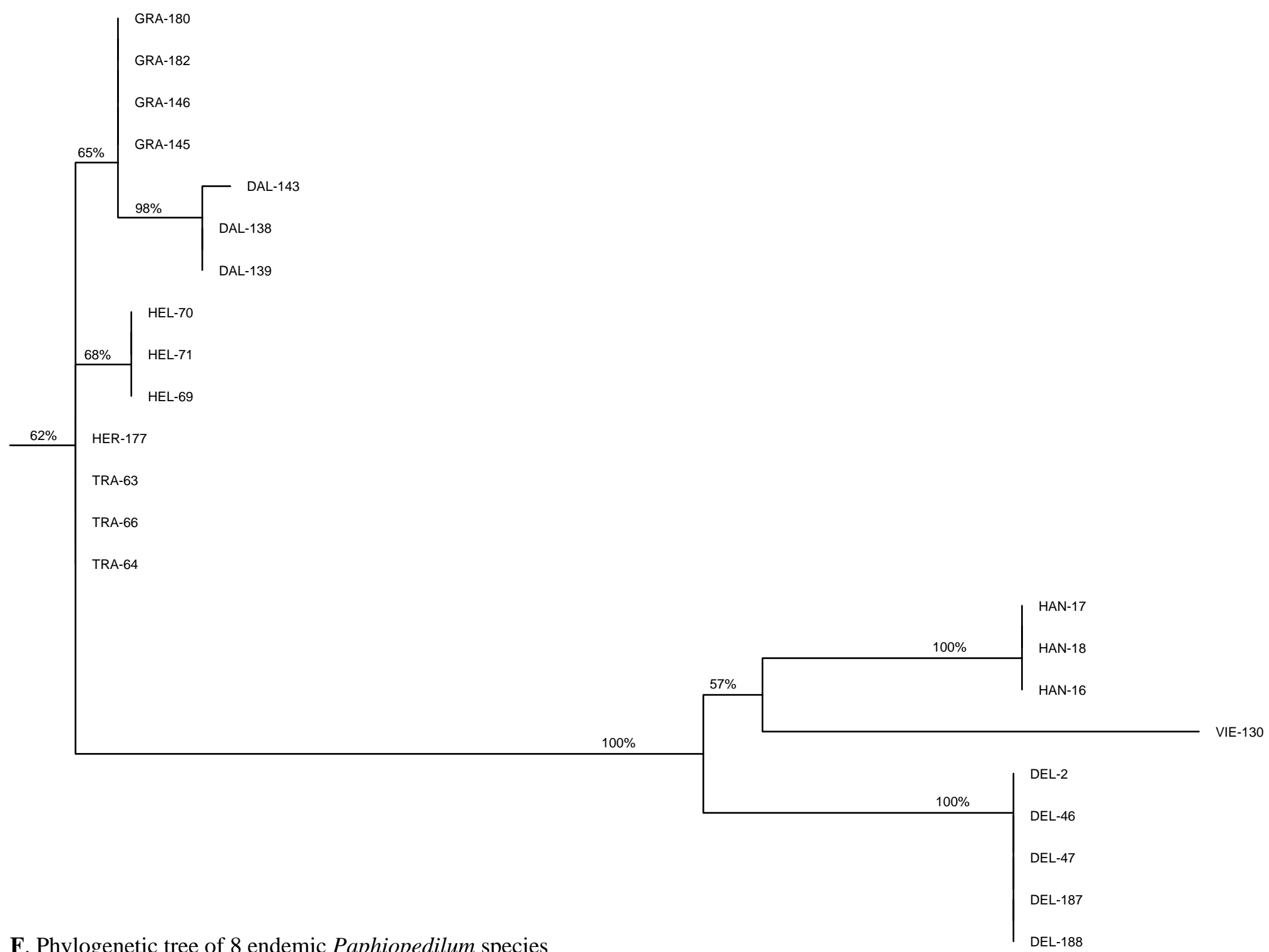
0.003



D. Phylogenetic tree of 8 endemic *Paphiopedilum* species
based on single *rpoB* locus

0.002



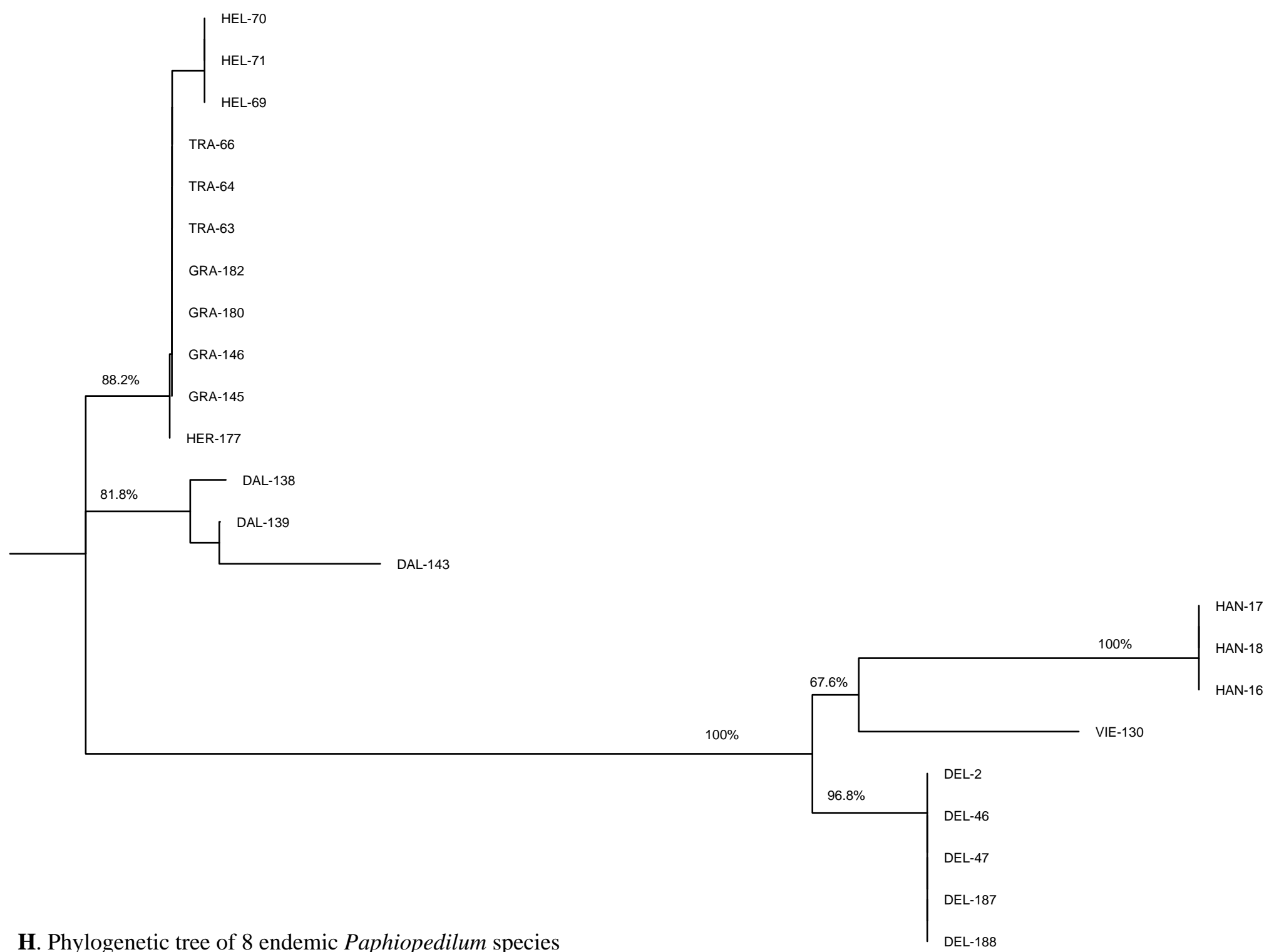


F. Phylogenetic tree of 8 endemic *Paphiopedilum* species based on combined *matK* + ITS locus



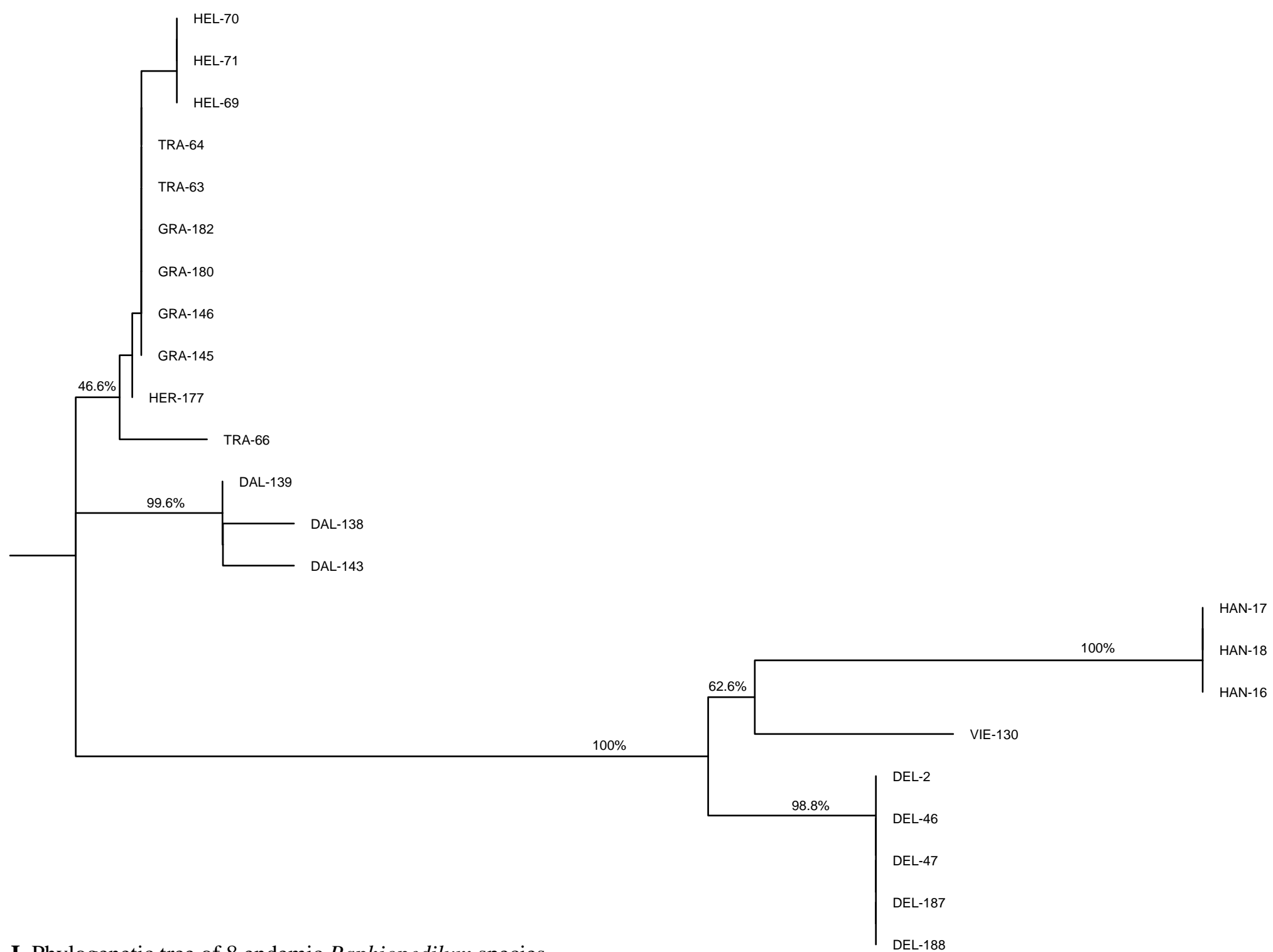
G. Phylogenetic tree of 8 endemic *Paphiopedilum* species
based on combined *matK* + *trnL* locus

0.003



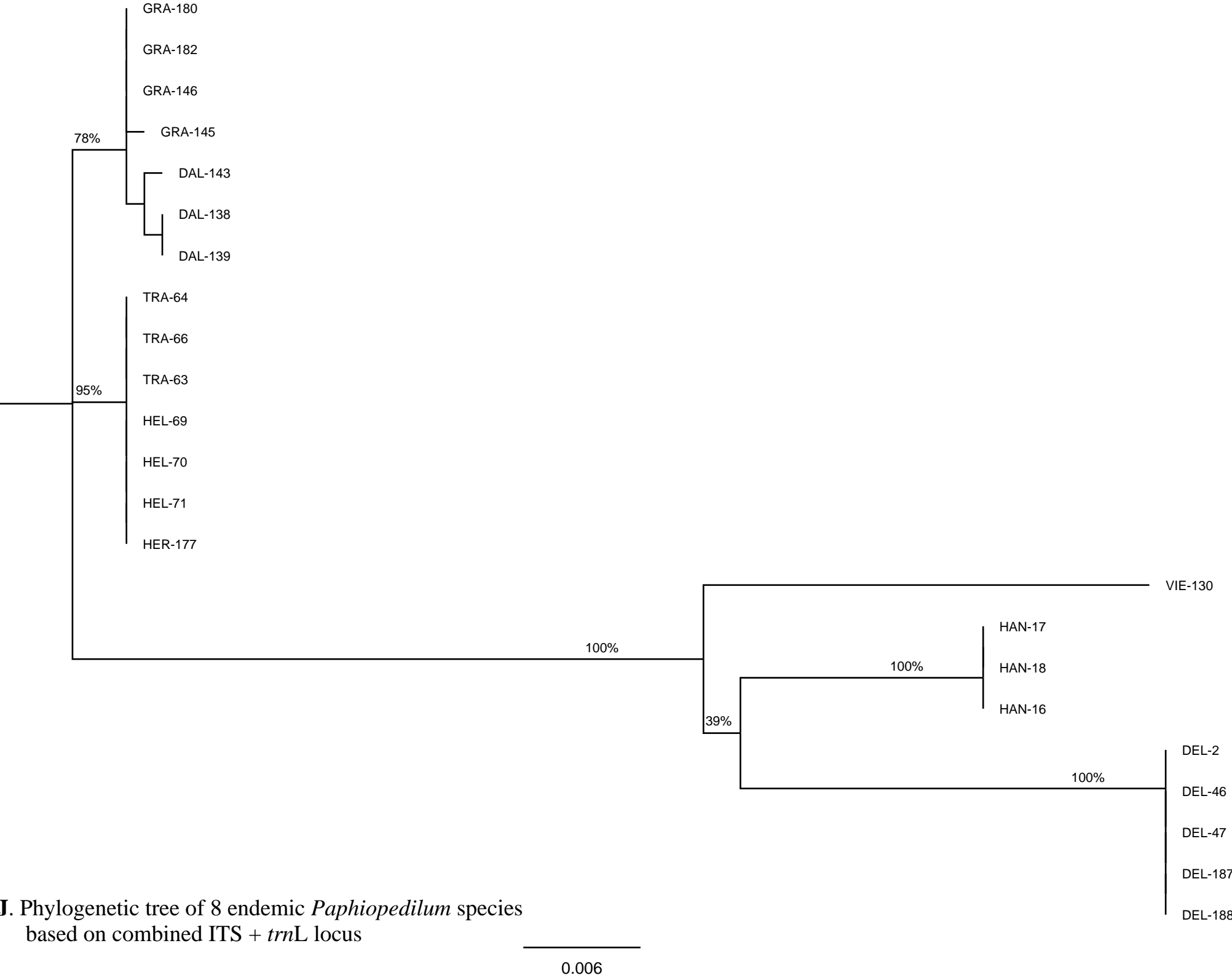
H. Phylogenetic tree of 8 endemic *Paphiopedilum* species based on combined *matK* + *rpoB* locus

0.003

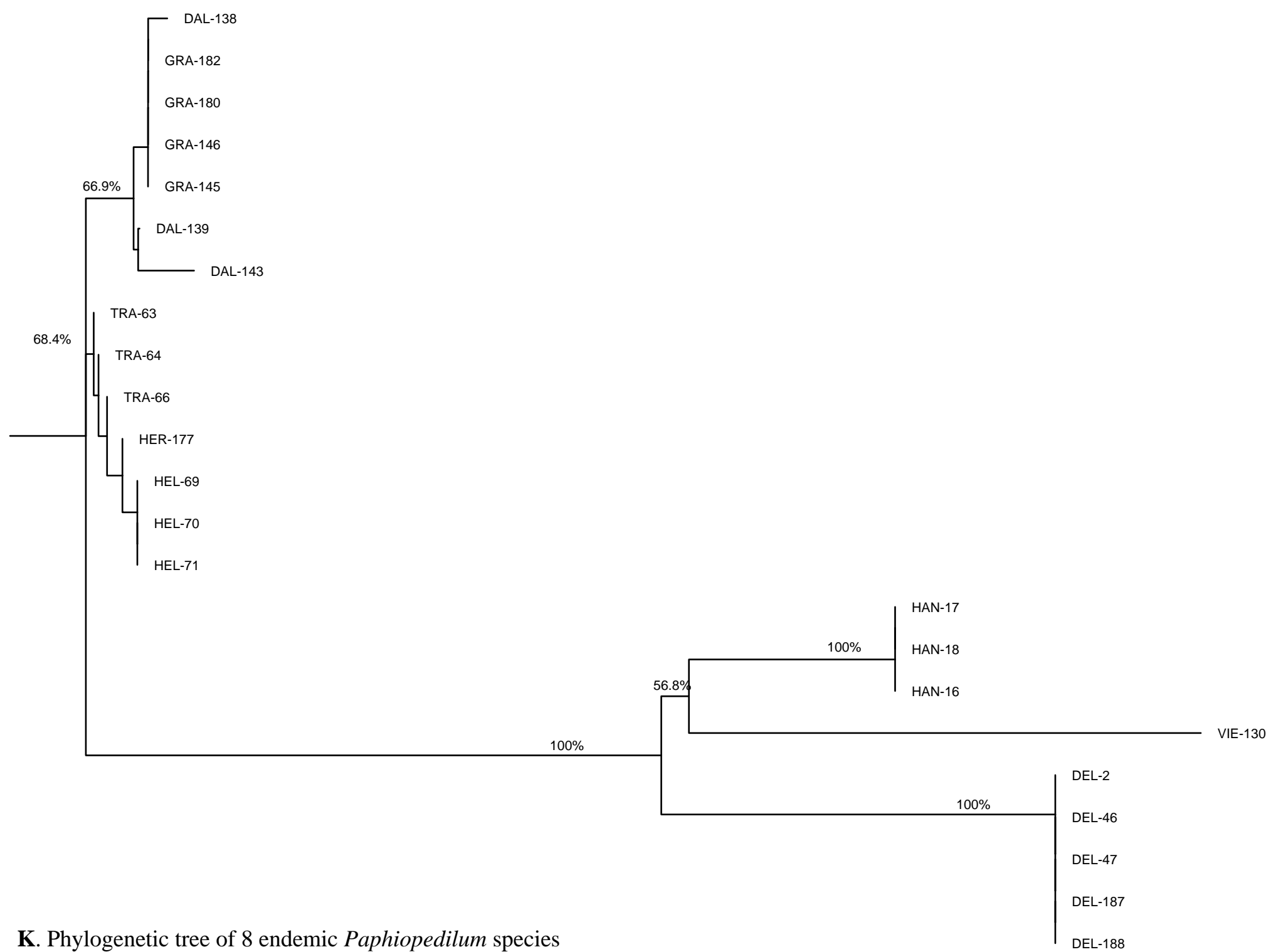


I. Phylogenetic tree of 8 endemic *Paphiopedilum* species
based on combined *matK* + *rpoC1* locus

0.003

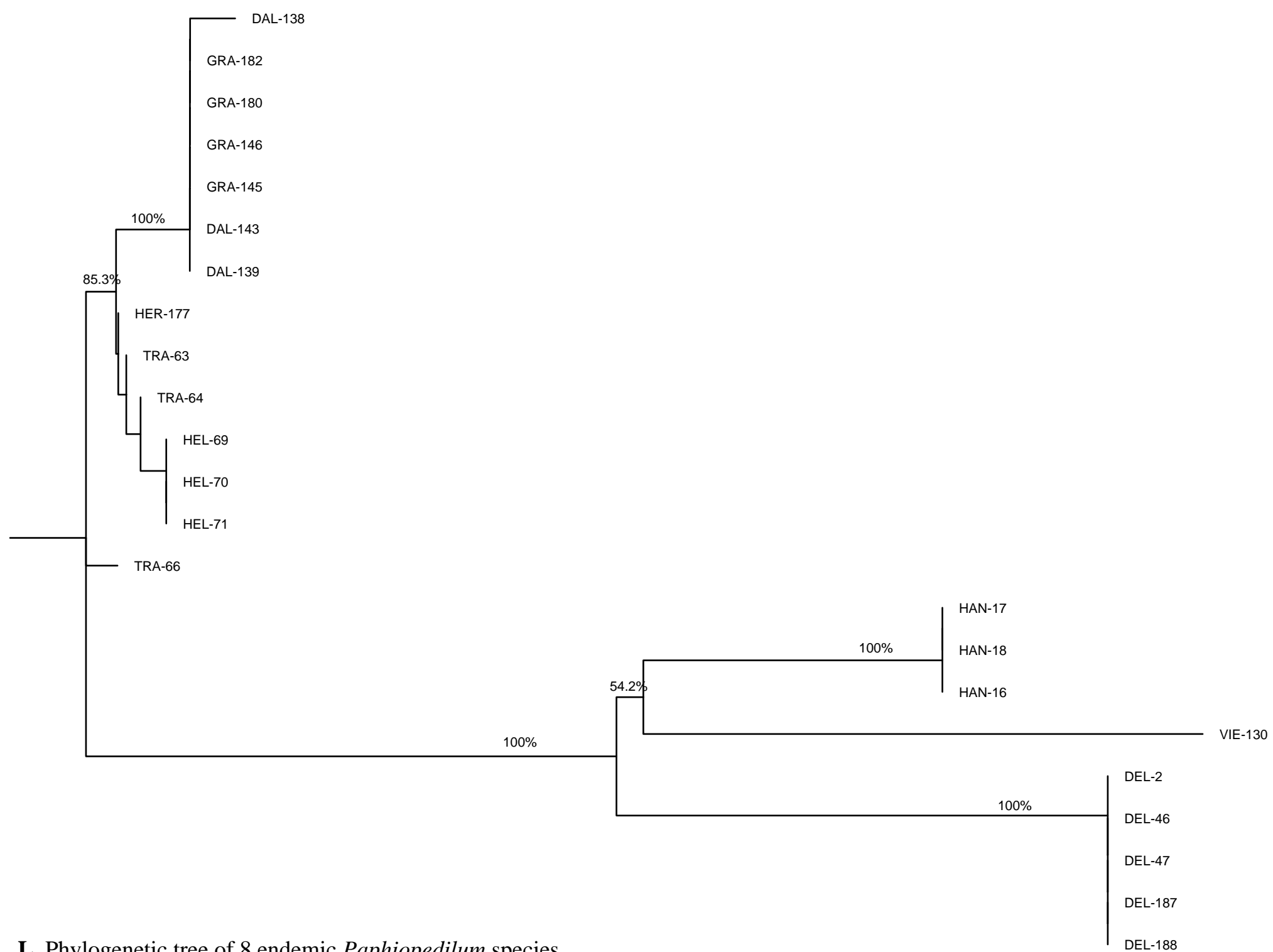


J. Phylogenetic tree of 8 endemic *Paphiopedilum* species based on combined ITS + *trnL* locus



K. Phylogenetic tree of 8 endemic *Paphiopedilum* species based on combined ITS + *rpoB* locus

0.006



L. Phylogenetic tree of 8 endemic *Paphiopedilum* species based on combined ITS + *rpoC1* locus

0.005