

SUPPLEMENTARY DATA

FIG. S2-1. Consensus phylogram obtained from 119935 bayesian trees with higher posterior probability from *matK* sequence

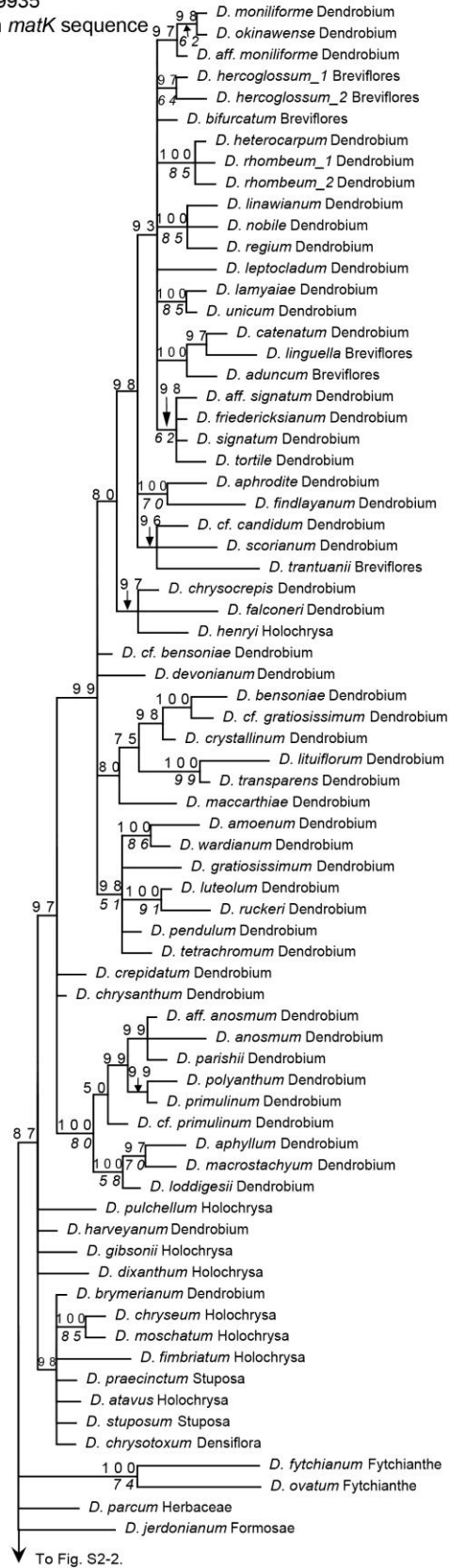


FIG. S2-2.

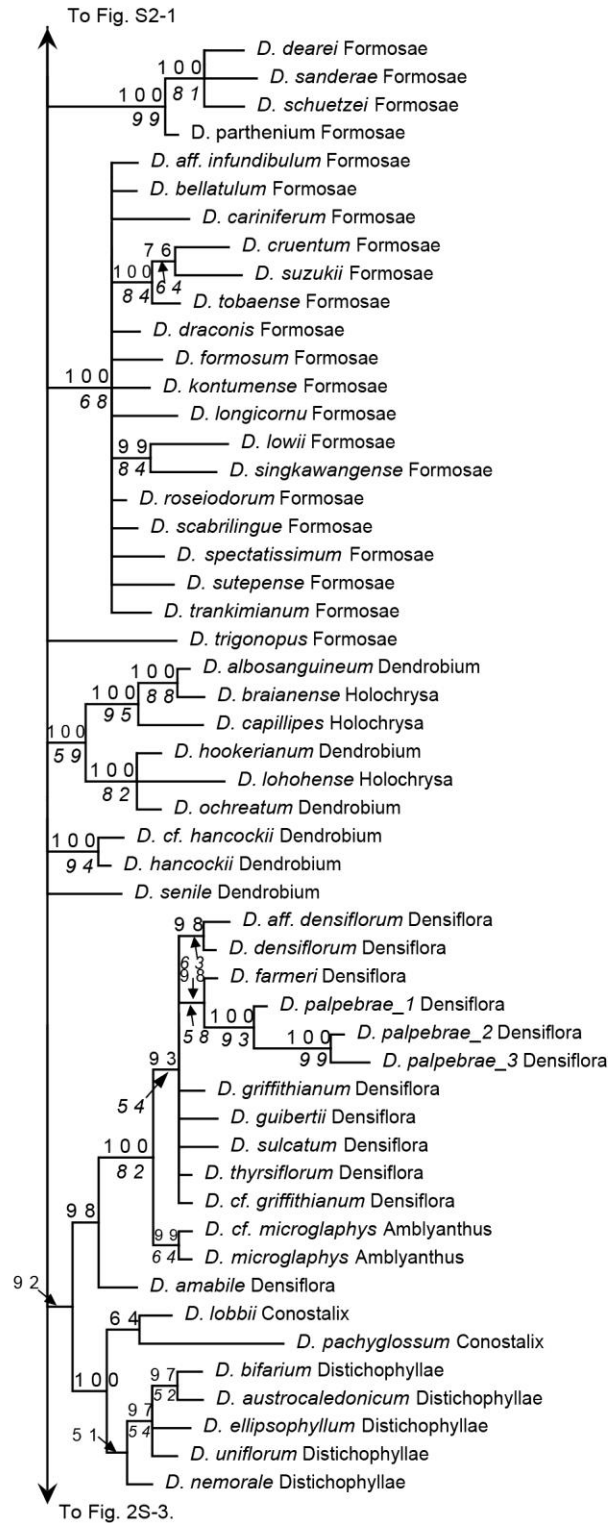


FIG. S2-3.

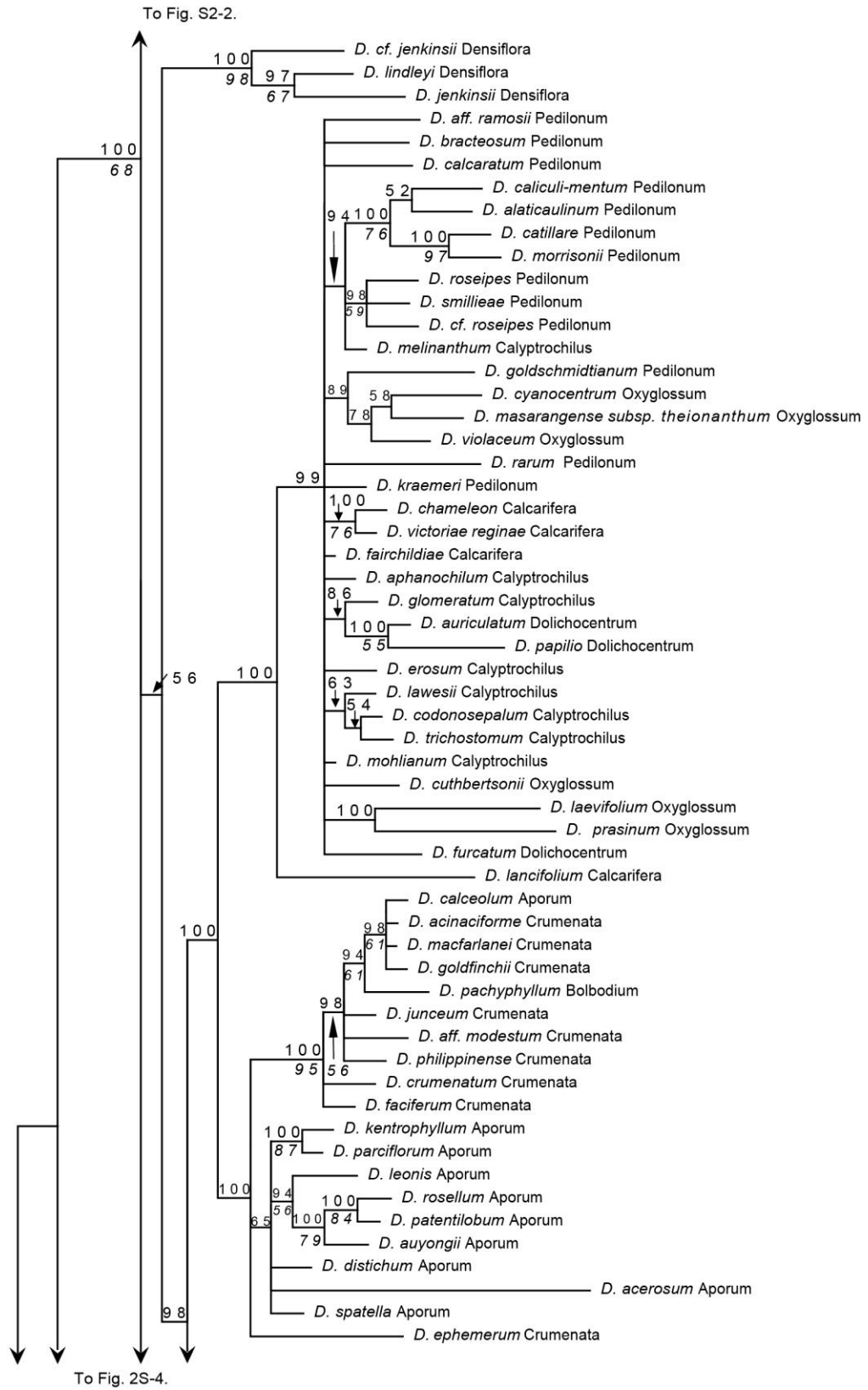


FIG. S2-4.

