

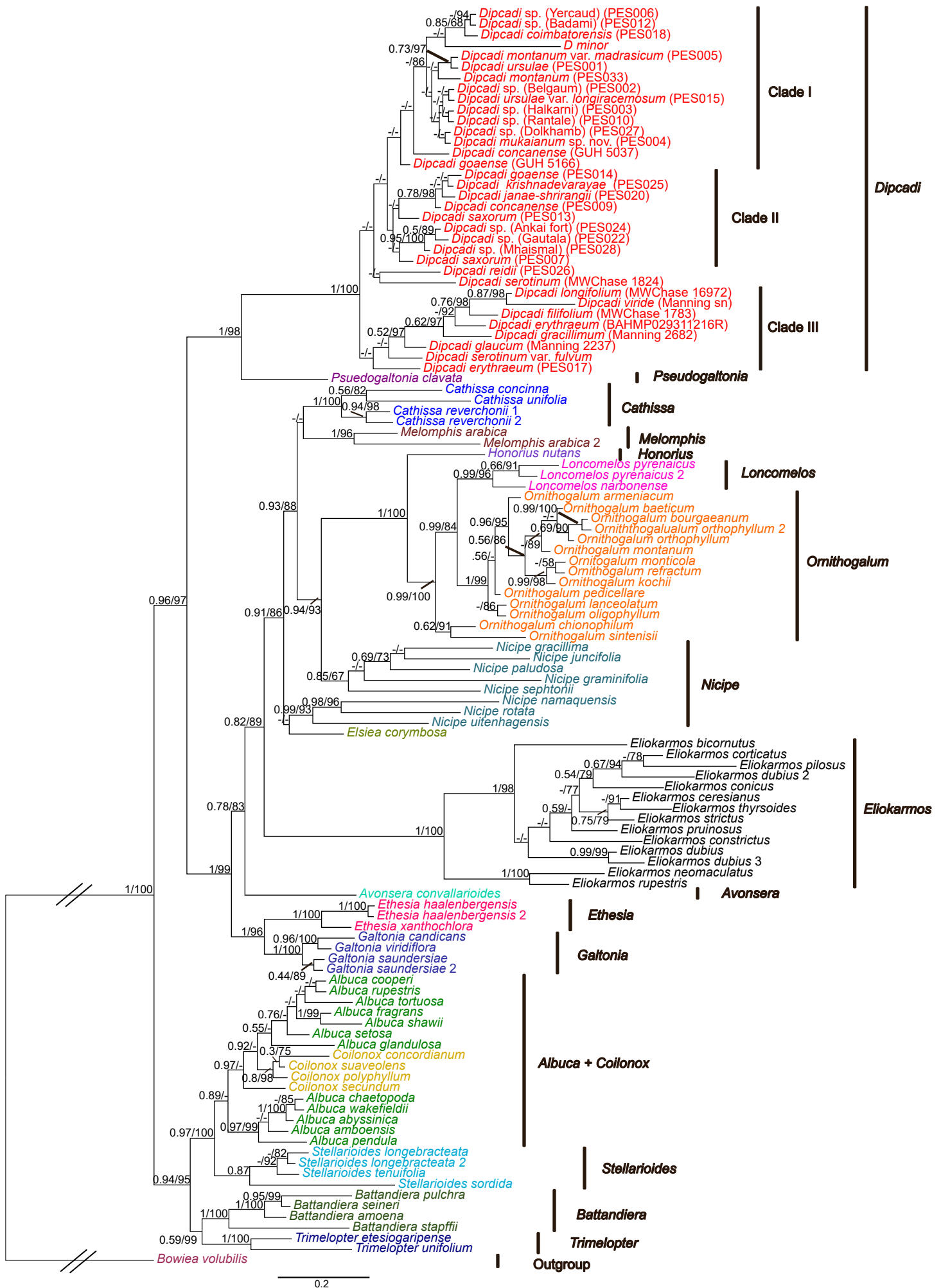


Fig. S1. Bayesian analysis of combined *matK* + *rbcL* + *trnL-F* sequences. Bayesian posterior probability values and maximum likelihood bootstrap values (BPP/ML BS) are provided above branches. Voucher ID of each *Dipcadi* taxa provided in parentheses. (-) indicates ML BS values < 50% and BI PP values < 0.5.