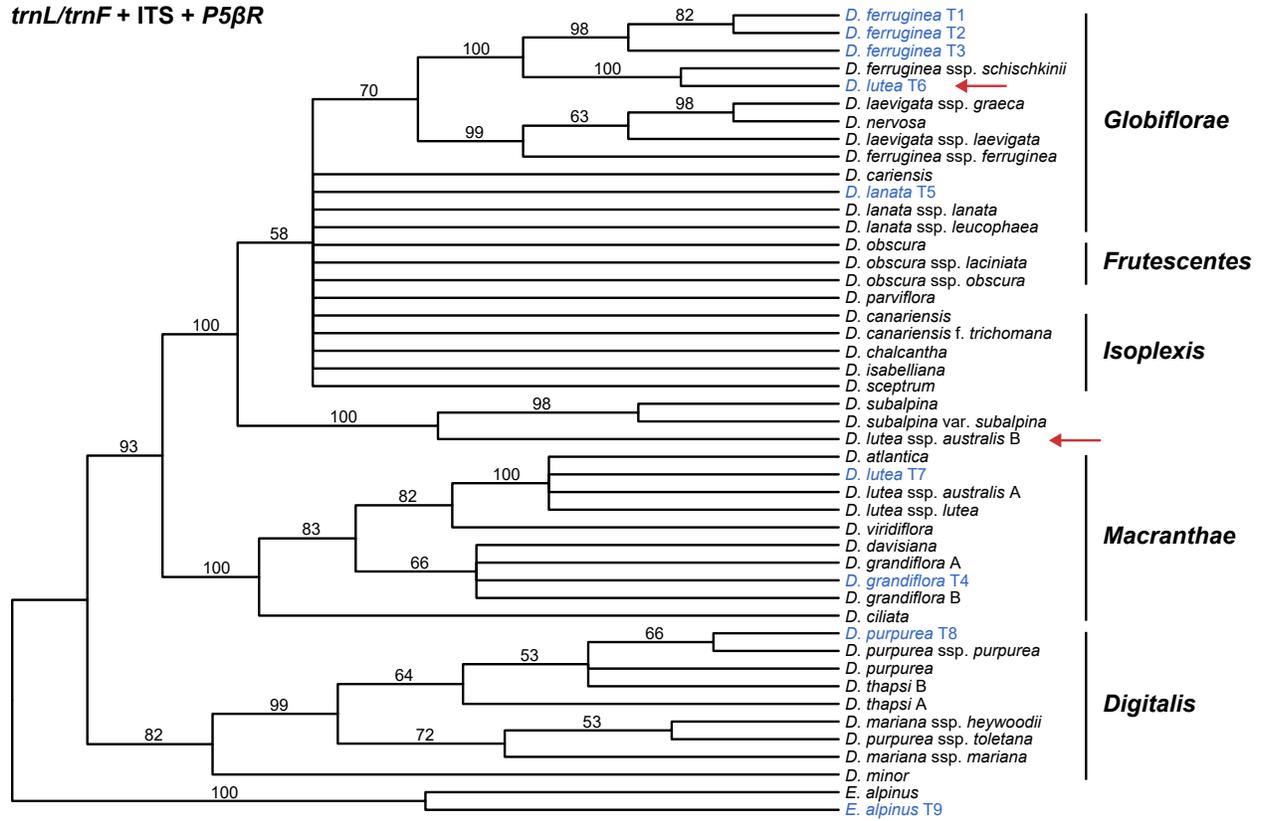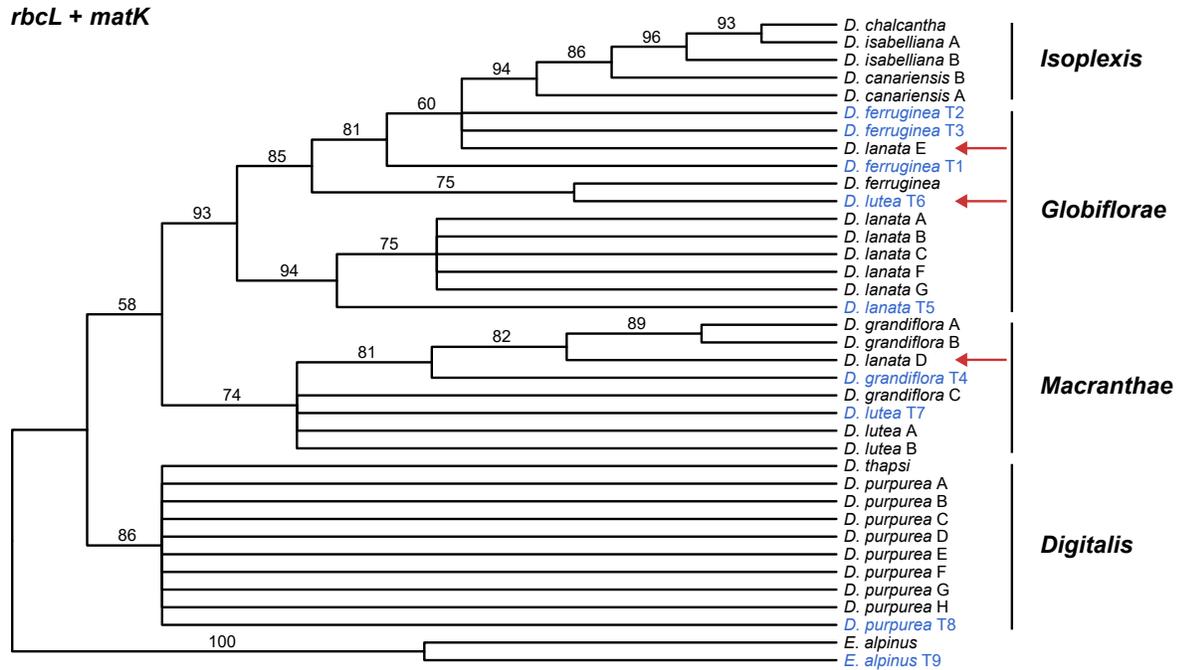Figure S1. Phylogenetic analysis of marker data sets. Top: 3-locus data set containing plastid *trnL* + *trnF*, nuclear ITS1 + 5.8S + ITS2, and nuclear *P5βR*. Bottom: 2-locus data set containing plastid *matK* + *rbcL*. Taxa newly assembled in this study (labeled T1–T9) are shown in blue. Taxa grouped in unexpected positions are marked with a red arrow. Branches with bootstrap values <50% were collapsed. Subgenera are shown at right. Trees were rooted on *Erinus*.