

Figure S1. Bayesian inference phylogenetic tree topology inferred from the concatenated dataset (18S rDNA, ITS-1, 28S rDNA and COI). Numbers at nodes indicate support values (posterior probability).

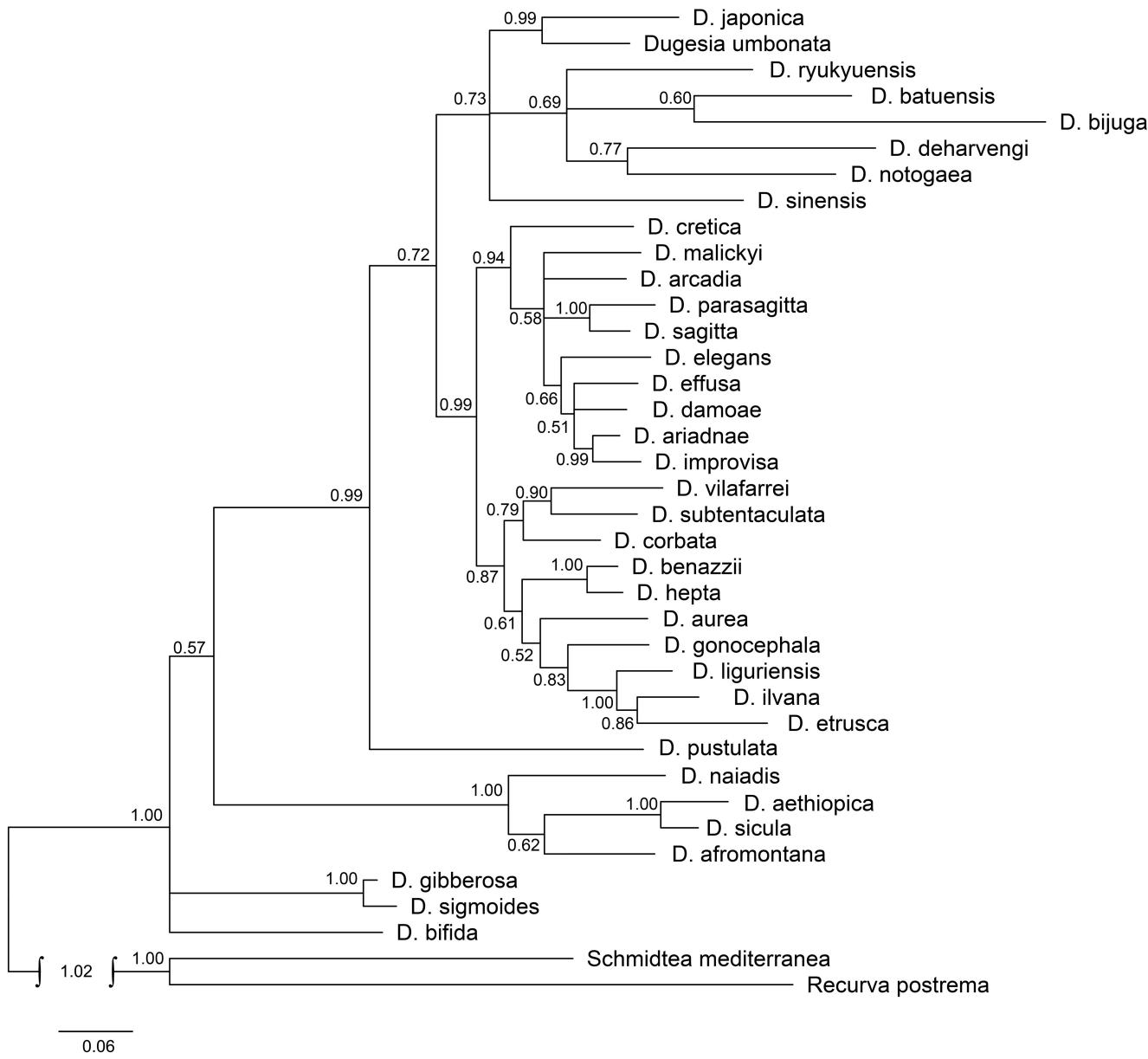


Figure S2. Bayesian inference phylogenetic tree based on COI dataset. Numbers at nodes indicate support values (posterior probability).

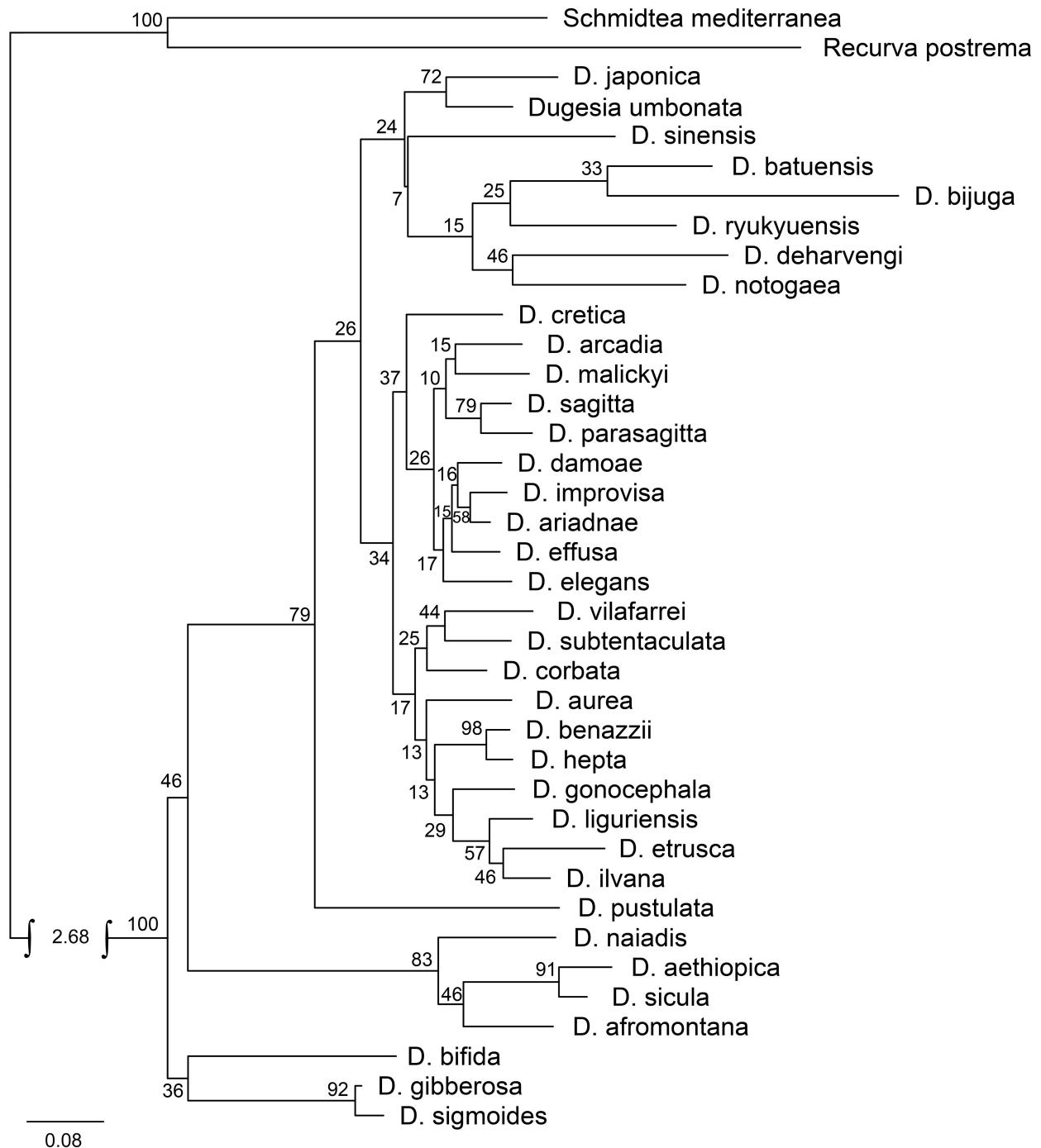


Figure S3. Maximum likelihood phylogenetic tree based on COI dataset. Numbers at nodes indicate support values (bootstrap).

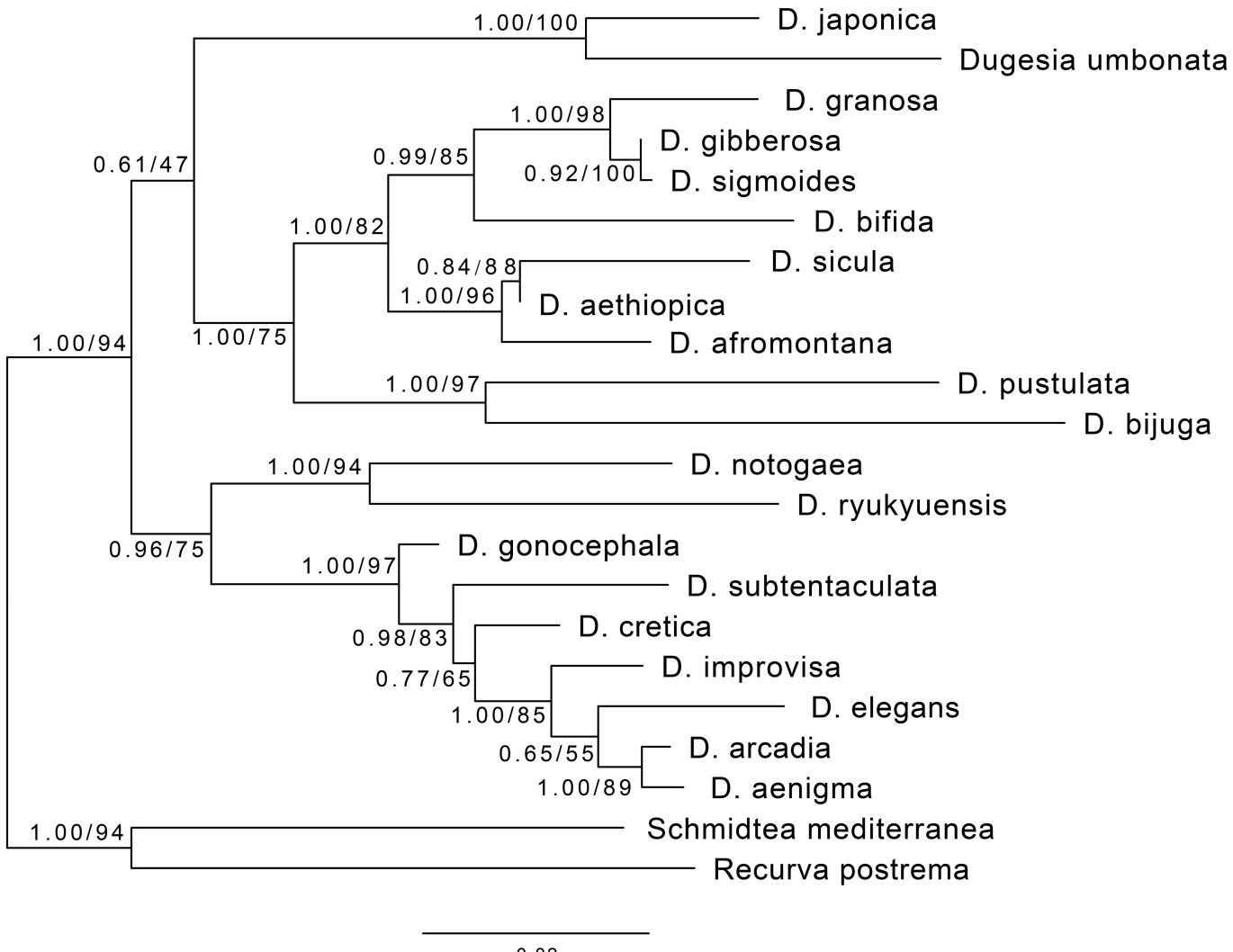


Figure S4. Maximum likelihood phylogenetic tree topology based on 18S rDNA dataset. Numbers at nodes indicate support values (posterior probability/bootstrap).

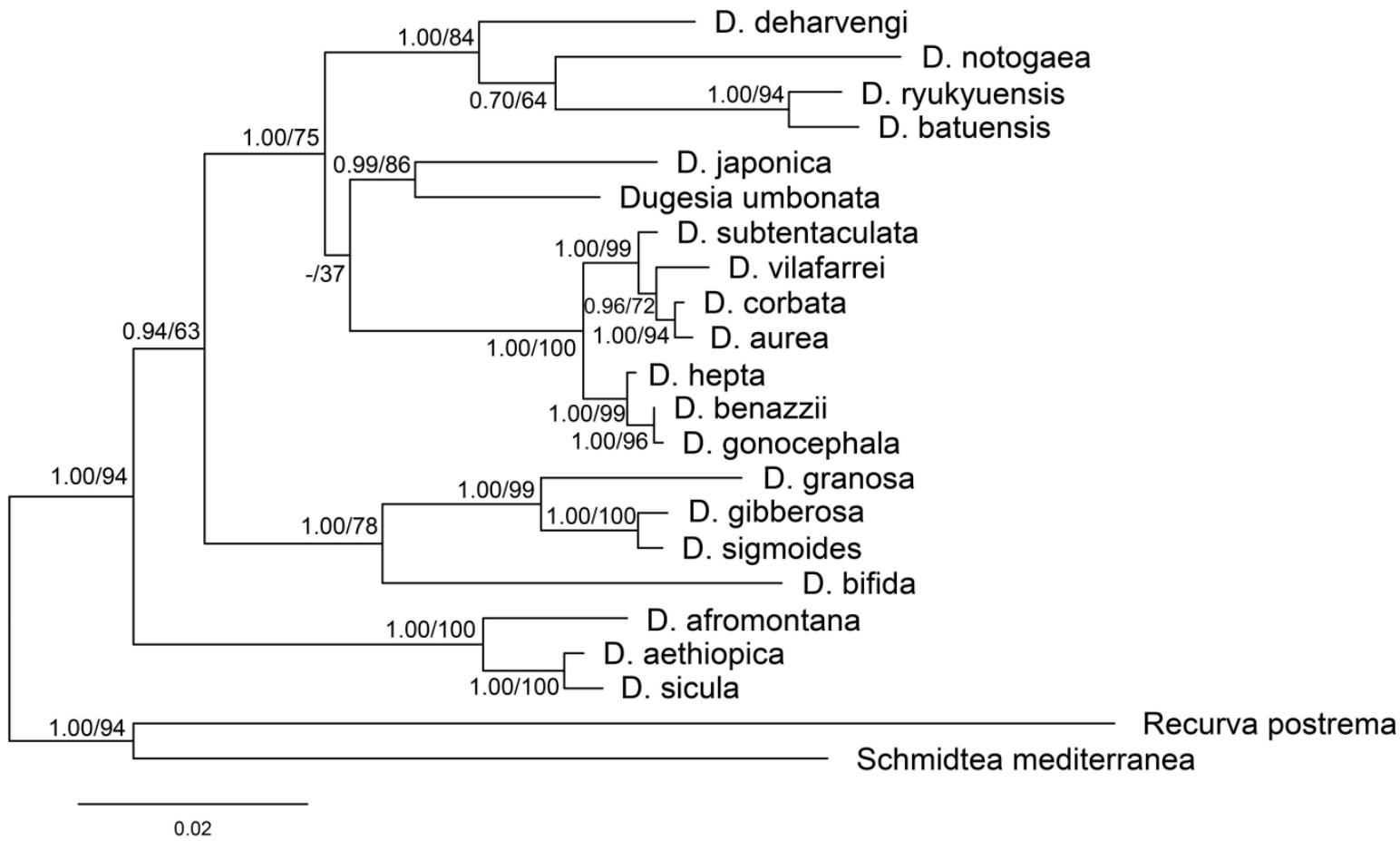


Figure S5. Maximum likelihood phylogenetic tree topology based on 28S rDNA dataset. Numbers at nodes indicate support values (posterior probability/bootstrap).

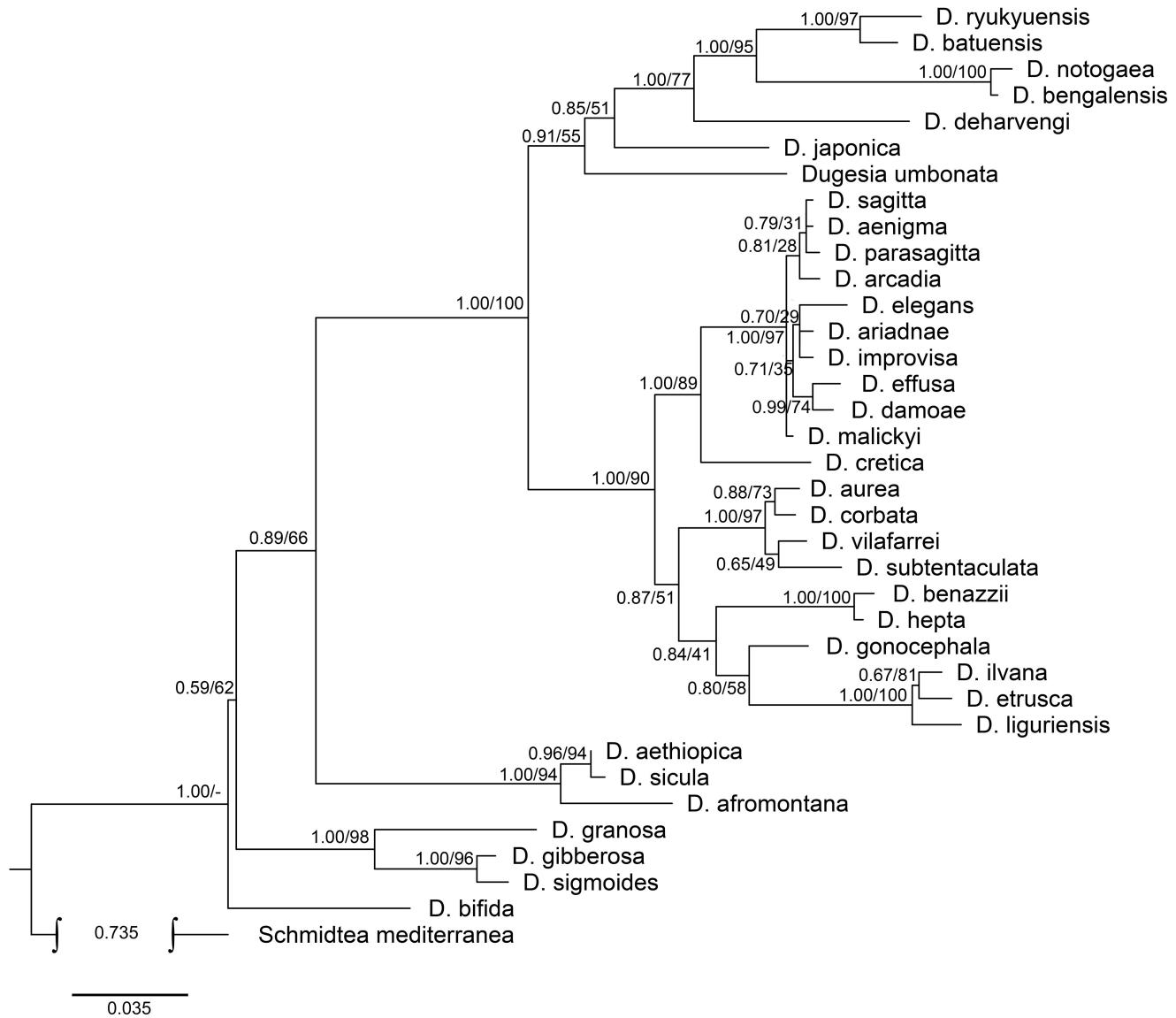


Figure S6. Maximum likelihood phylogenetic tree based on ITS-1 dataset. Numbers at nodes indicate support values (posterior probability/bootstrap).