

## 5. Supplementary figures S1–S2

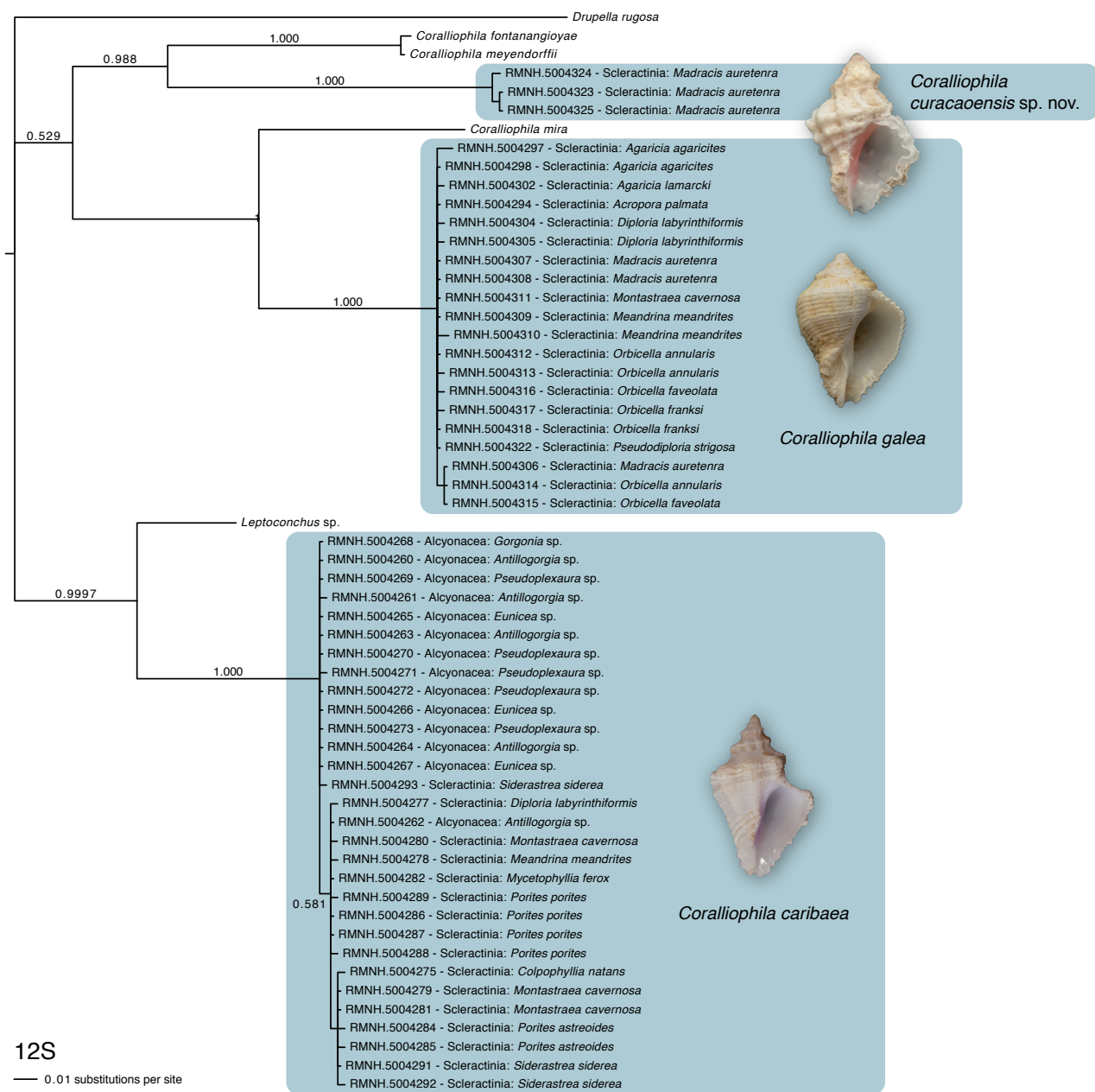


Fig. S1. Phylogenetic tree based on the 12S rRNA (12S) marker of a selection of the collected *Coralliophila* spp. specimens. The three shaded clades represent the three species of *Coralliophila* found in the present study. Within the shaded clusters, tip labels represent the host order and host species with which the snail was associated. Specimens outside the shaded cluster are previously published sequences. Branch labels are posterior probabilities (PP). Intraspecific PP-values are not shown (except for the main clade within *C. caribaea*). Scale bar: 0.01 substitutions per site.

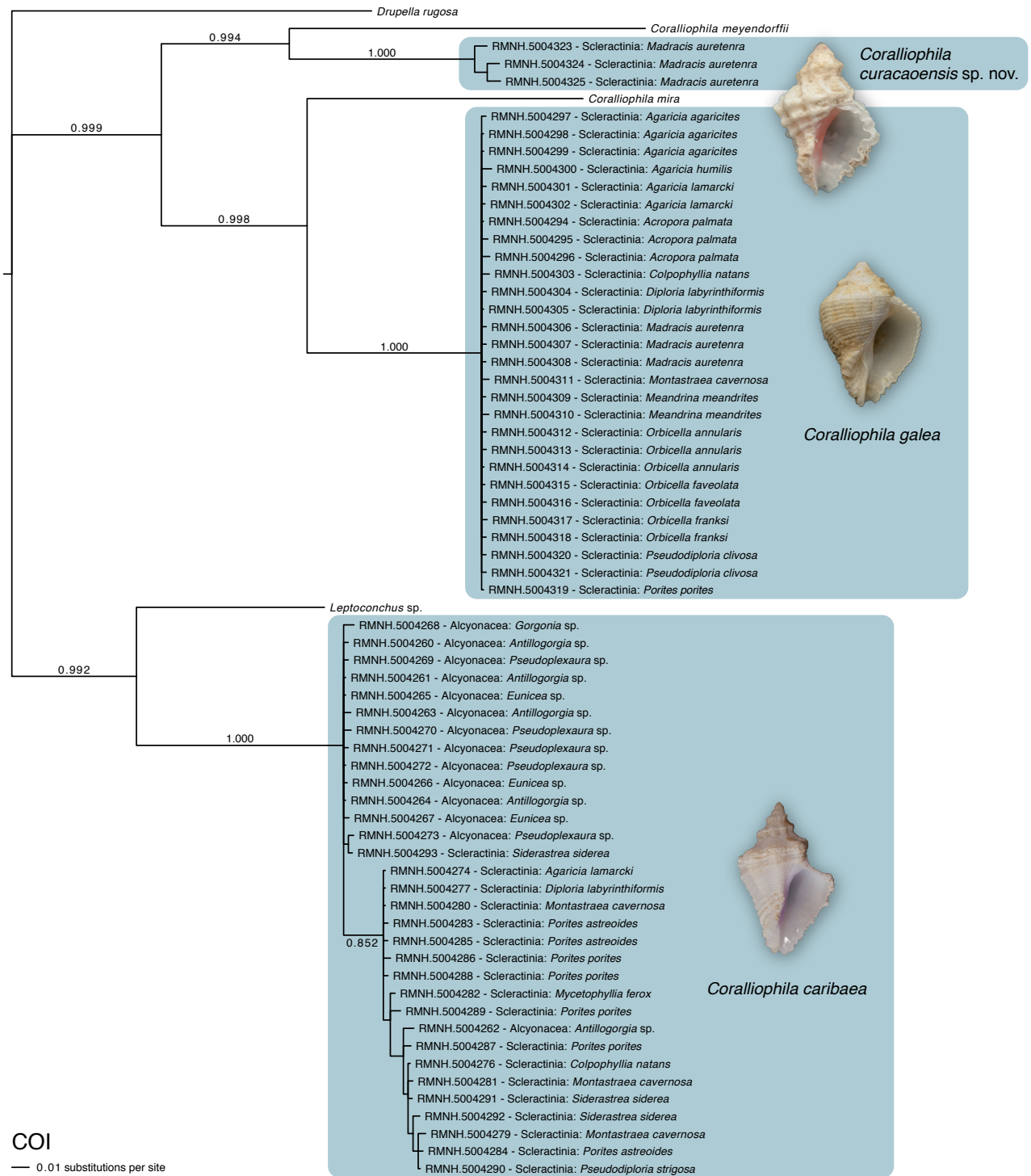


Fig. S2. Phylogenetic tree based on the cytochrome *c* oxidase subunit I (COI) marker of a selection of the collected *Coralliophila* spp. specimens. The three shaded clades represent the three species of *Coralliophila* found in the present study. Within the shaded clusters, tip labels represent the host order and host species with which the snail was associated. Specimens outside the shaded cluster are previously published sequences. Branch labels are posterior probabilities (PP). Intraspecific PP-values are not shown (except for the main clade within *C. caribaea*). Scale bar: 0.01 substitutions per site.