

Electronic Supplementary Material (Coral Reefs)
Evolution and biogeography of the *Zanclus-Scleractinia* symbiosis

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Fig. S1. Single-locus Bayesian inference (BI) and maximum likelihood (ML) phylogenetic hypotheses of coral-associated *Zanlea*. Numbers at nodes represent Bayesian posterior probabilities in BI analyses and bootstrap values in ML analyses. Each recovered clade is highlighted with a different code, as detailed in the ‘Results’ section.

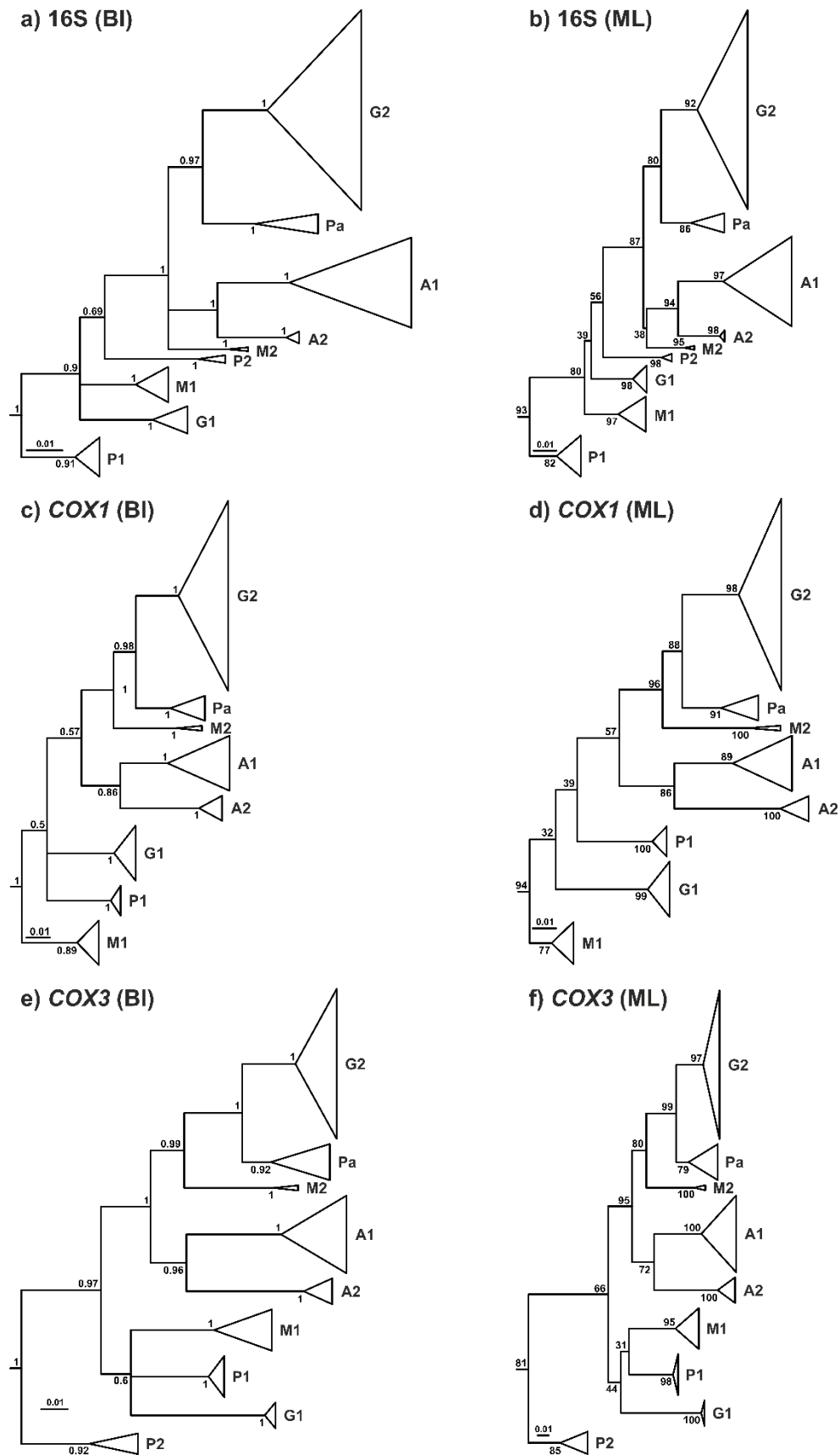


Fig. S3. Medium-joining networks of *Zancklea* clades based on a) *COX1* and b) *COX3* regions

