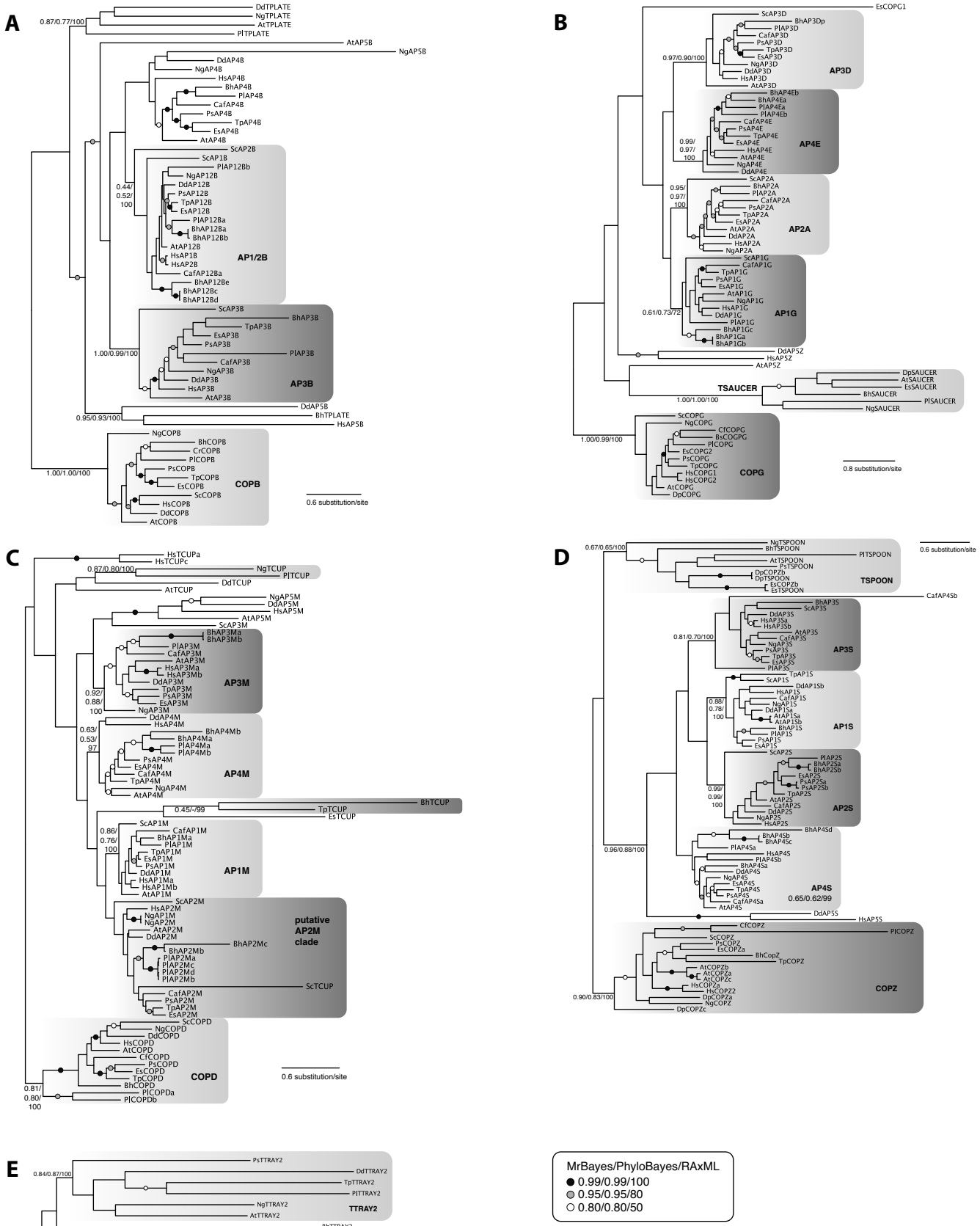

Data S2. Phylogenetic analyses of heterotetrameric adaptor coat complex subunits, Related to STAR methods, Figure S3. (**A**) large beta, (**B**) large EGADZ, (**C**) medium, (**D**) small, and (**E**) beta propeller-alpha solenoid. In all figures, MrBayes topology is shown, onto which support values from PhyloBayes and RAxML analyses are overlayed. Support values are shown for critical nodes, otherwise replaced as inset.