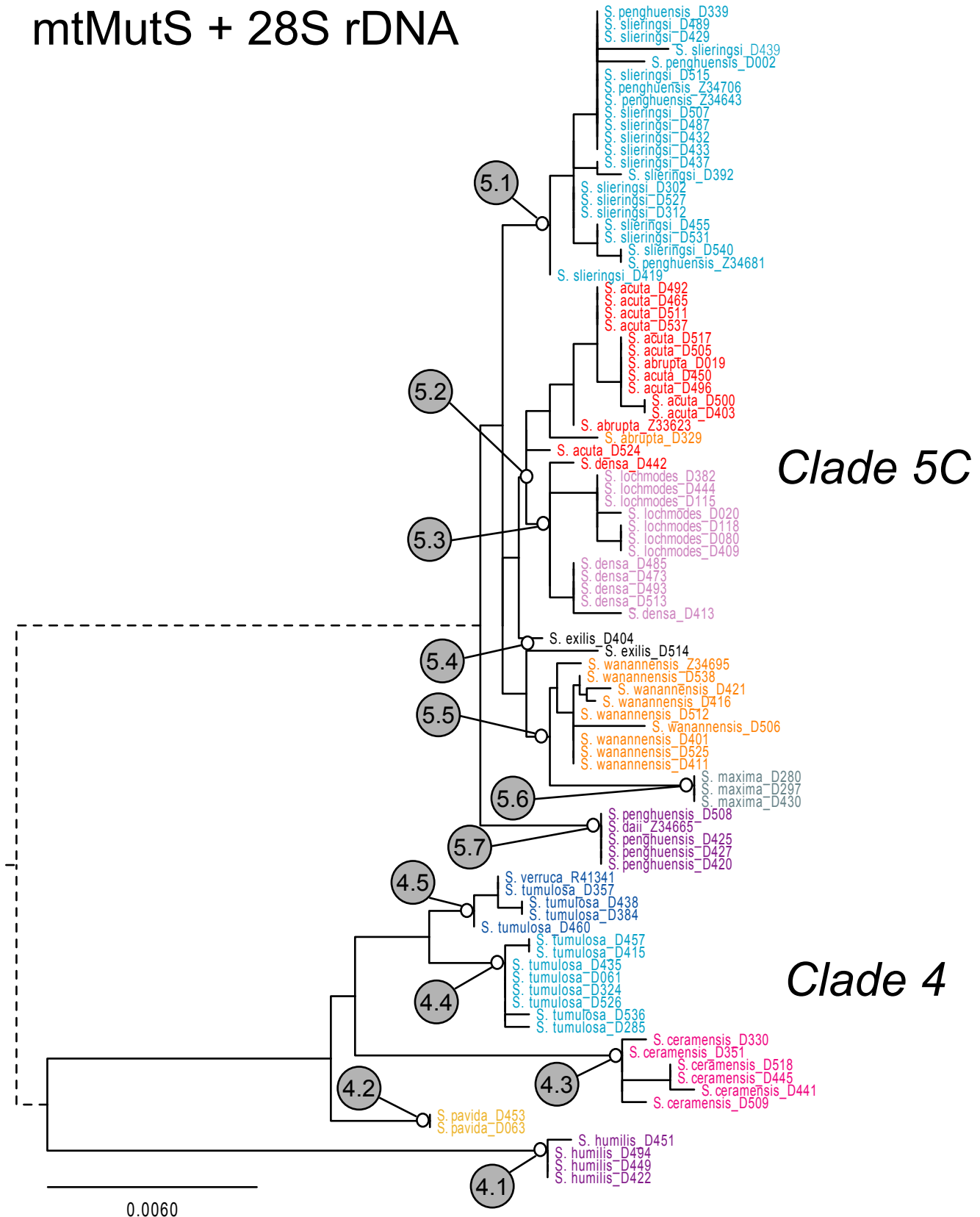


mtMutS + 28S rDNA



Maximum likelihood tree for *Sinularia* clades 4 and 5C from combined analysis of concatenated *mtMutS* and 28S *rDNA* barcode sequences. Numbered circles indicate molecular operational taxonomic units (MOTUs) delimited using a 0.3% genetic distance threshold. Colors of taxon label corresponds to MOTUs. Tree is midpoint-rooted with the branch connecting clades 4 and 5C pruned to facilitate readability.