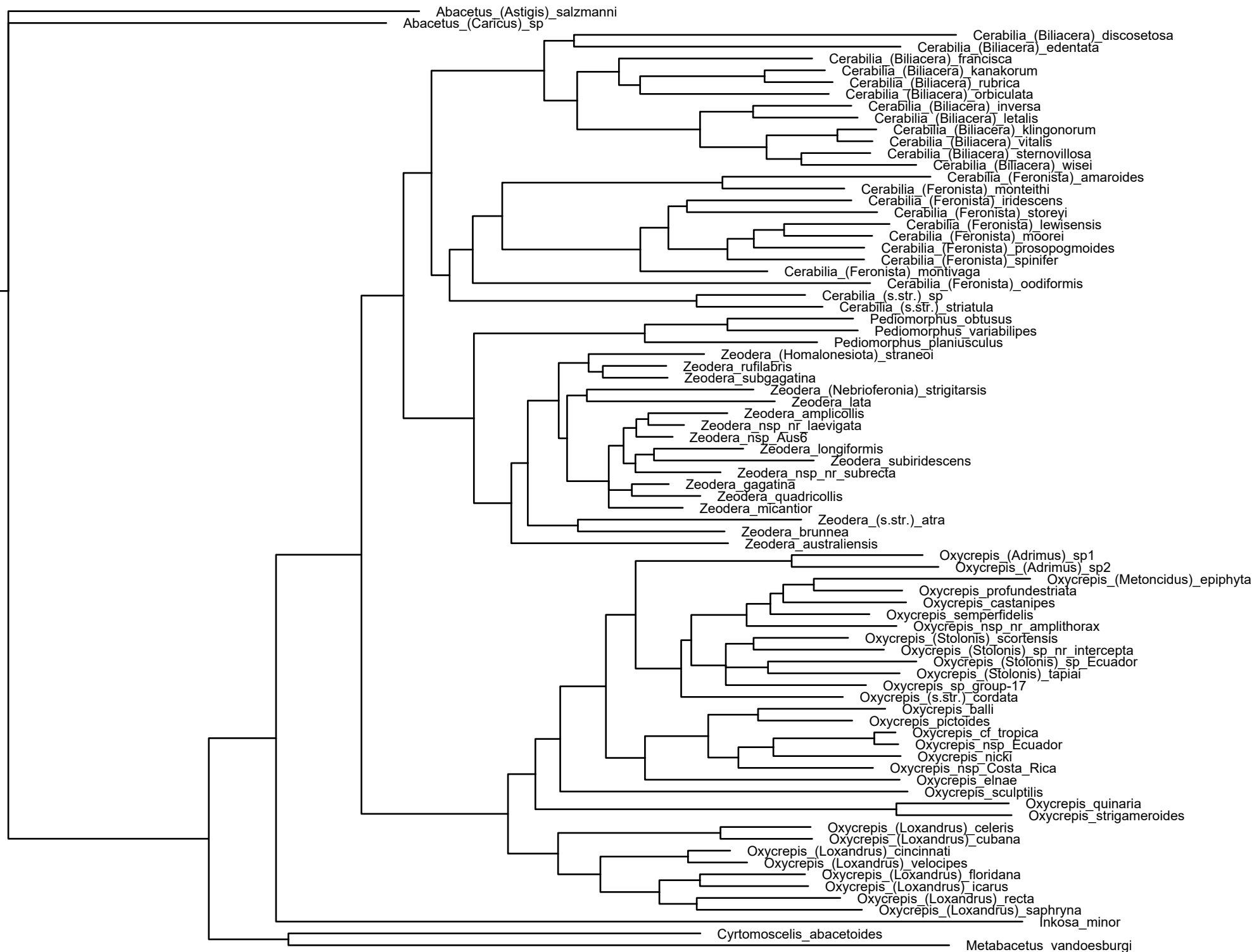
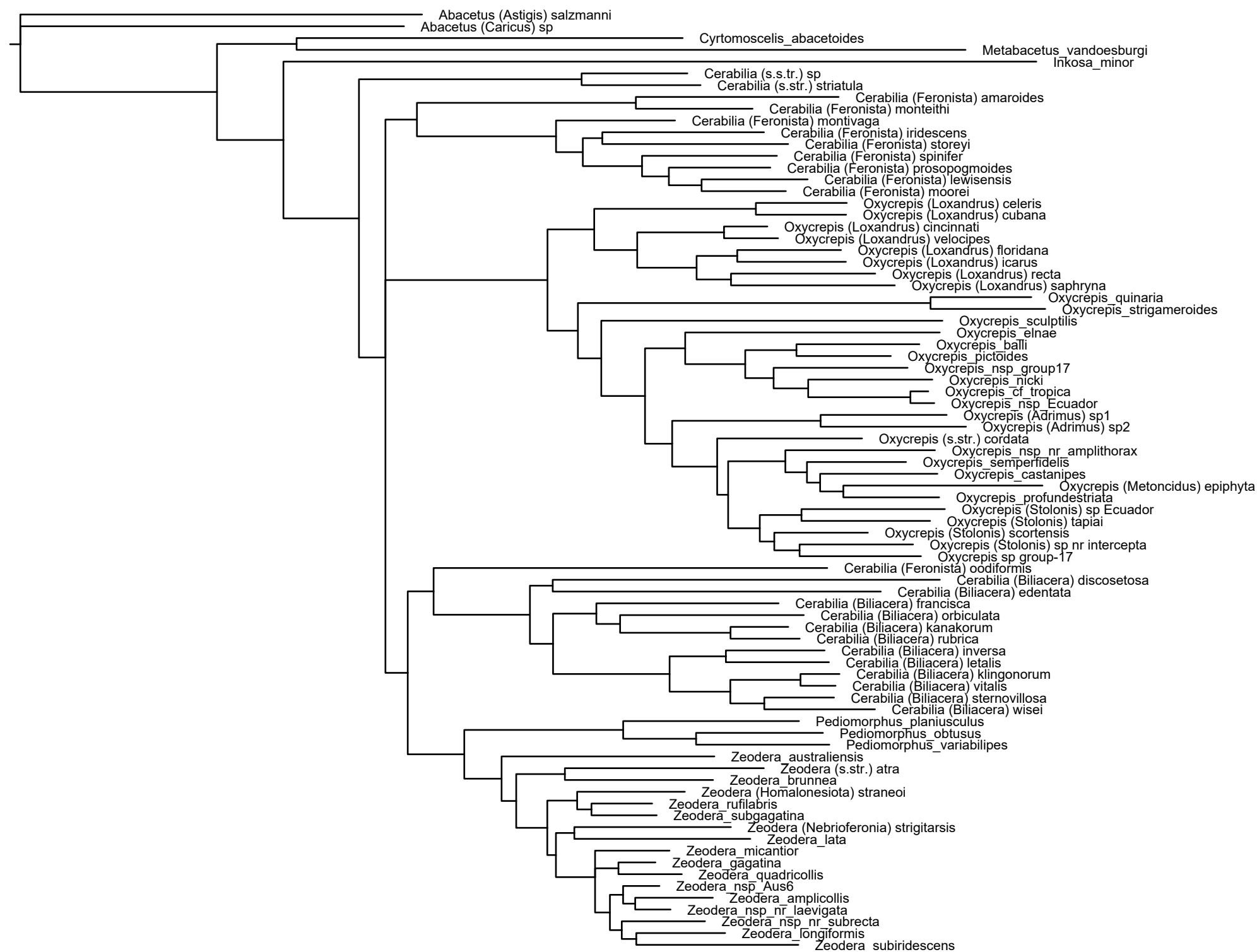


Tree from MrBayes analysis of all data for 80 OTUs



0.03

Tree from MrBayes analysis of 80 OTUs using DNA sequence data only



0.03

Consensus tree from TNT parsimony analysis of morphological data

