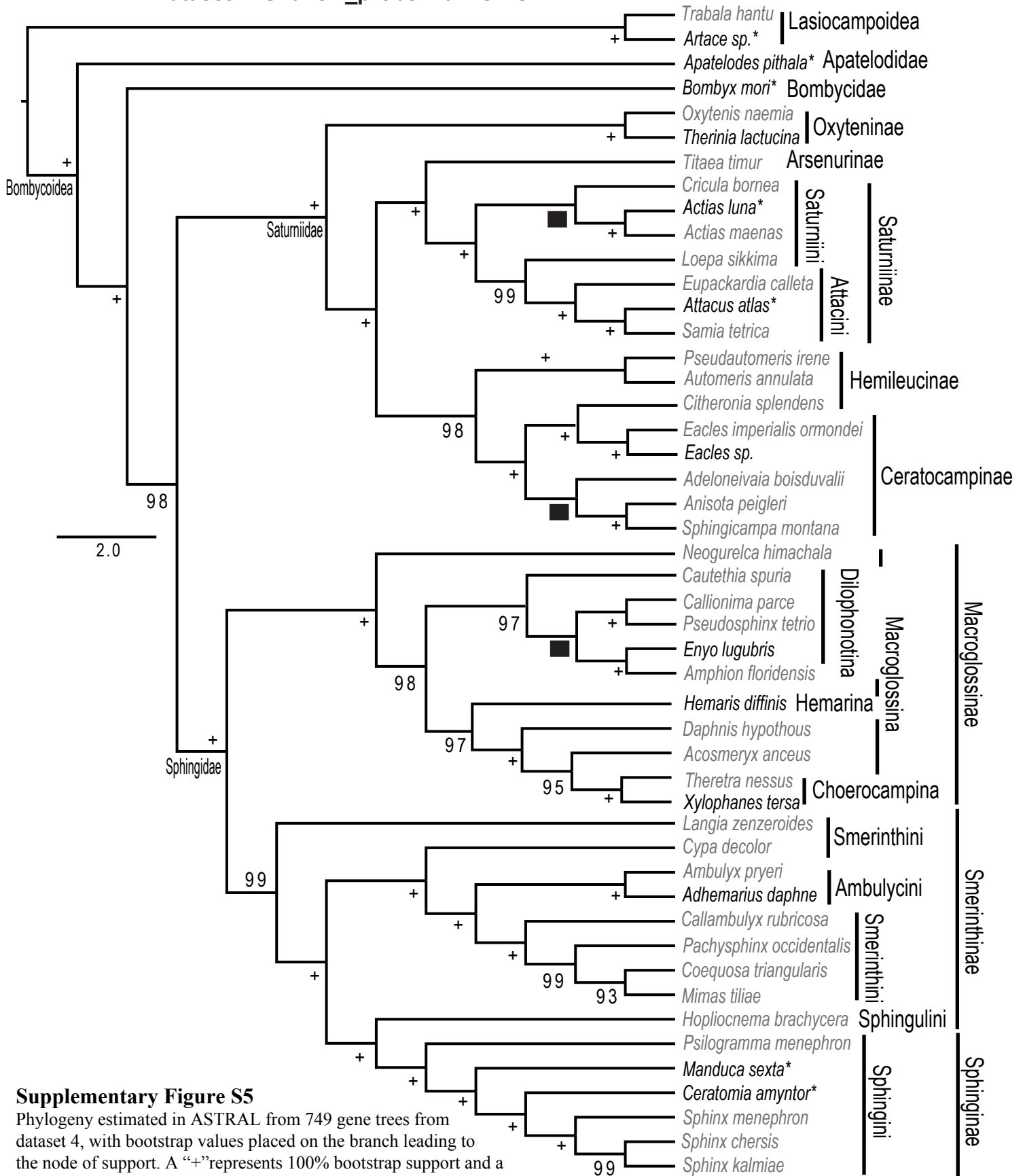


Dataset 4: Shallow_probe+flanks ASTRAL



Supplementary Figure S5

Phylogeny estimated in ASTRAL from 749 gene trees from dataset 4, with bootstrap values placed on the branch leading to the node of support. A "+" represents 100% bootstrap support and a black square represents support under 70%. Taxa in grey are sequenced anchored phylogenomic data and taxa in black are data from ESTs, transcriptomes, or genomes. An asterisk following a taxon name indicates a reference taxon used to make the Lep1 probe set.