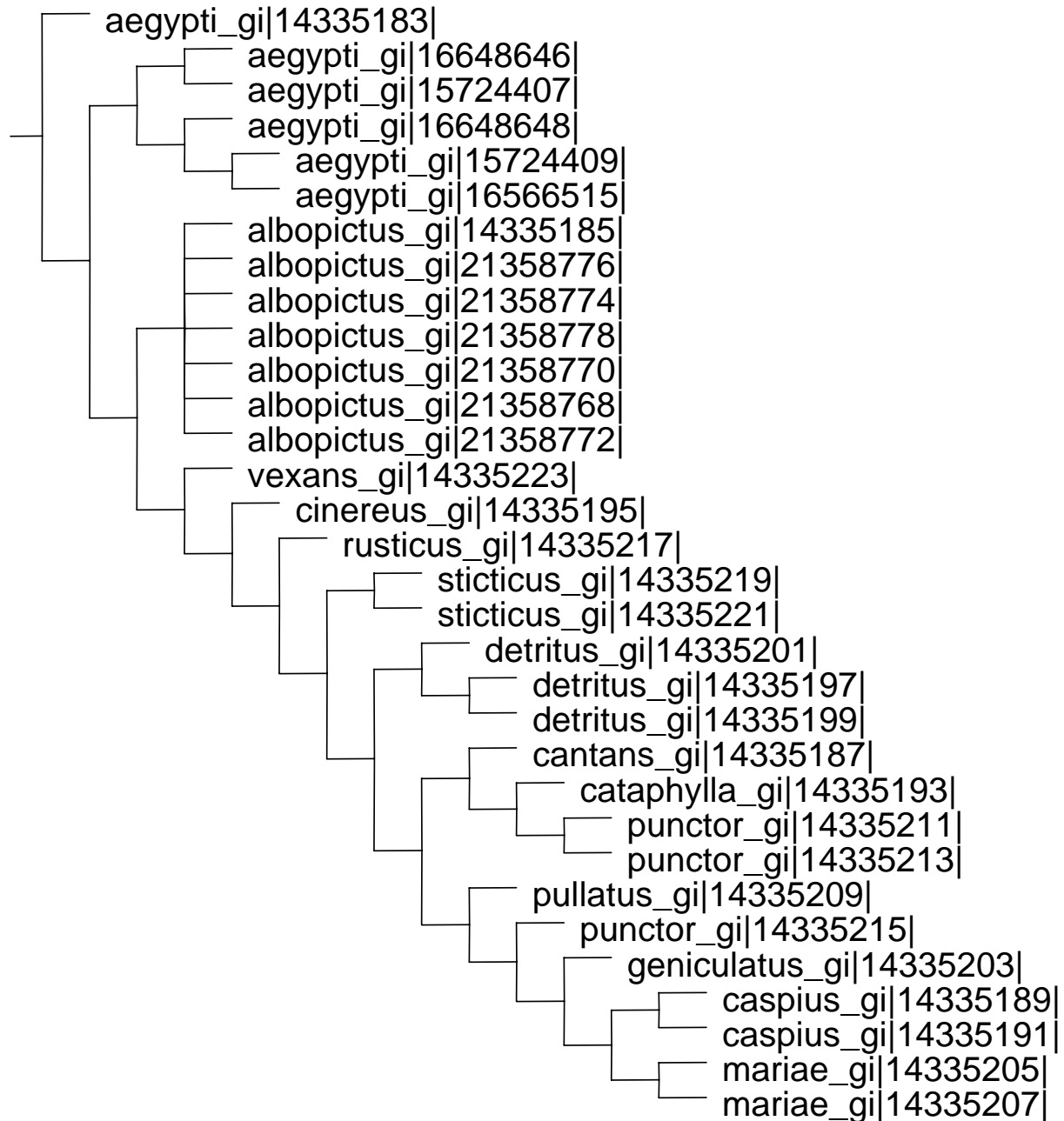


Aedes

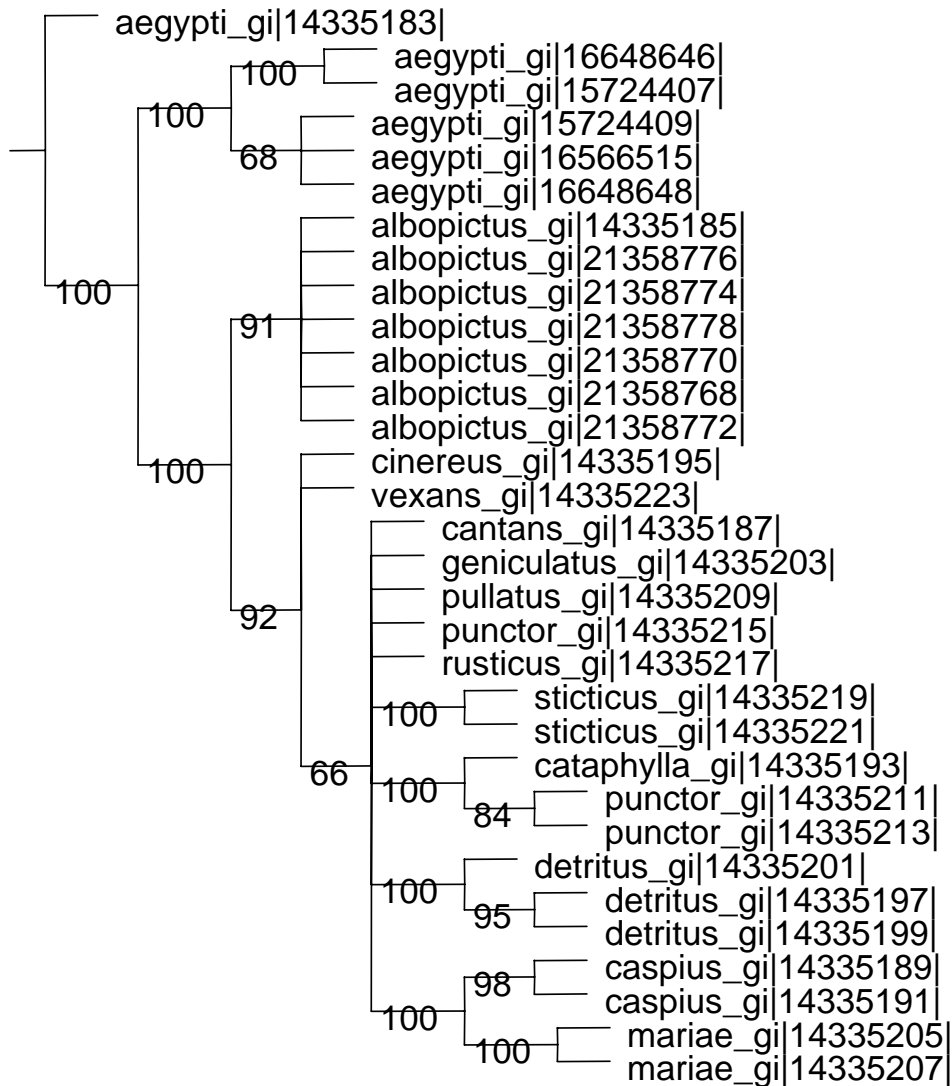
Strict consensus of 2 trees (0 taxa excluded)



Aedes

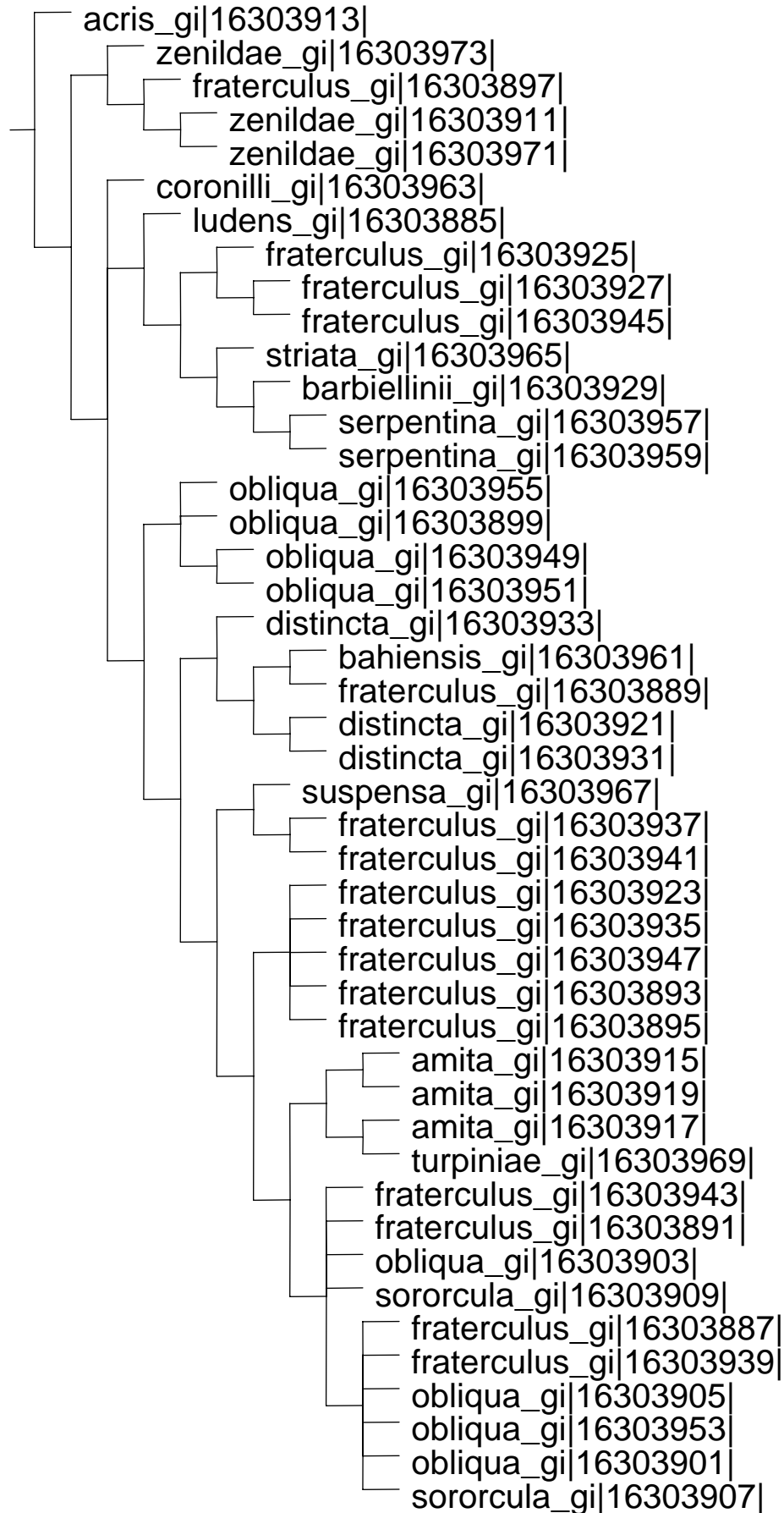
Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



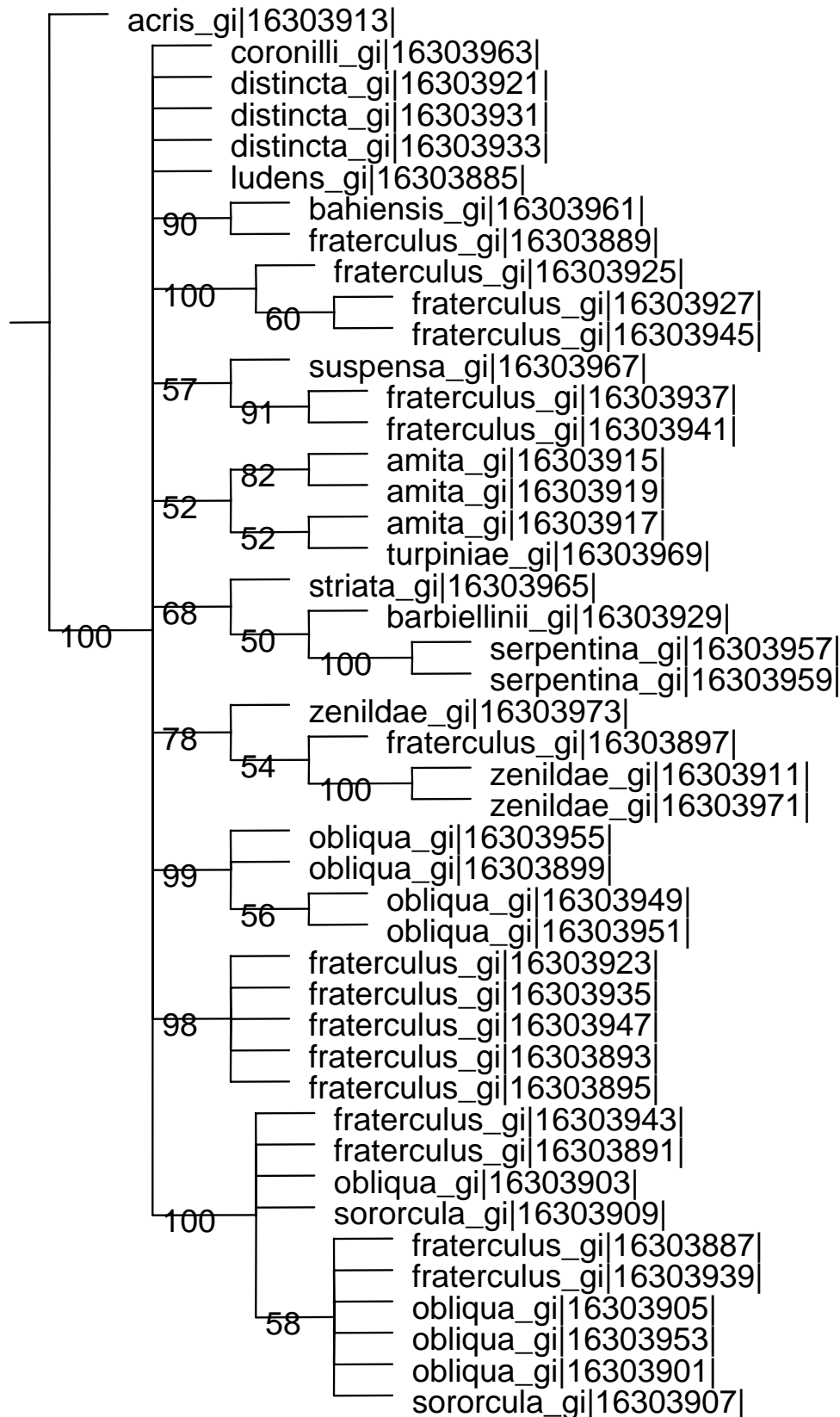
Anastrepha

Strict consensus of 2 trees (0 taxa excluded)



Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap

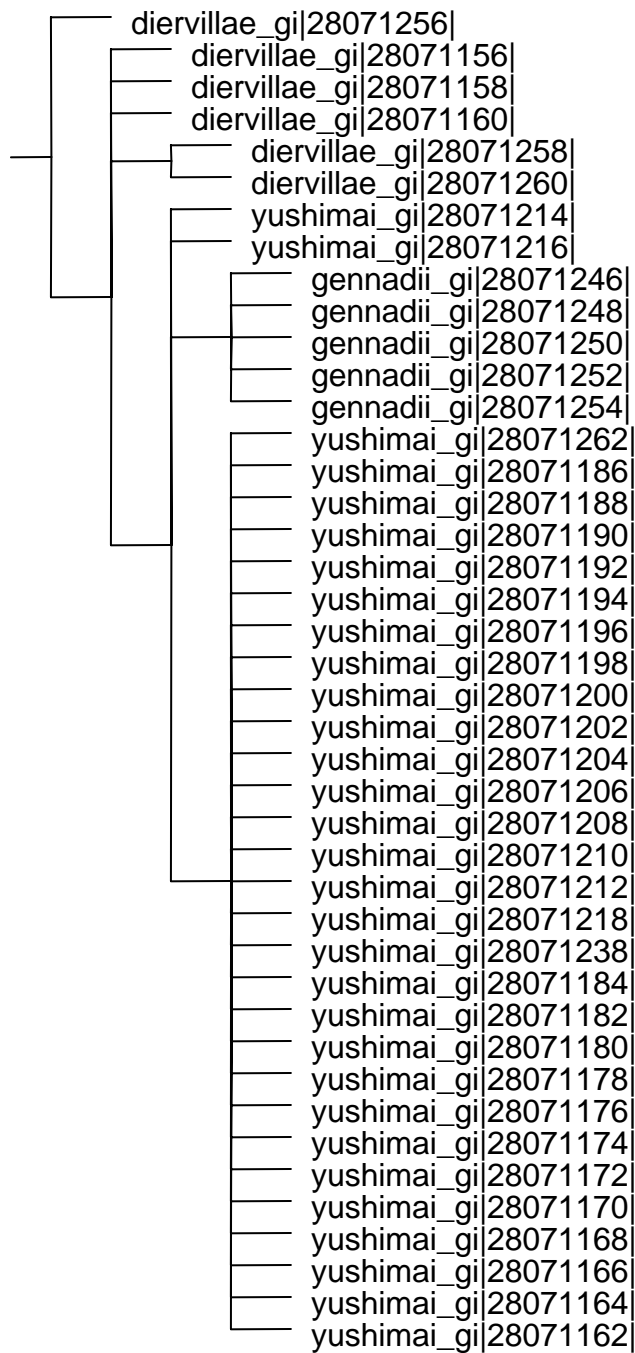


[illegible]

[illegible]

Asphondylia

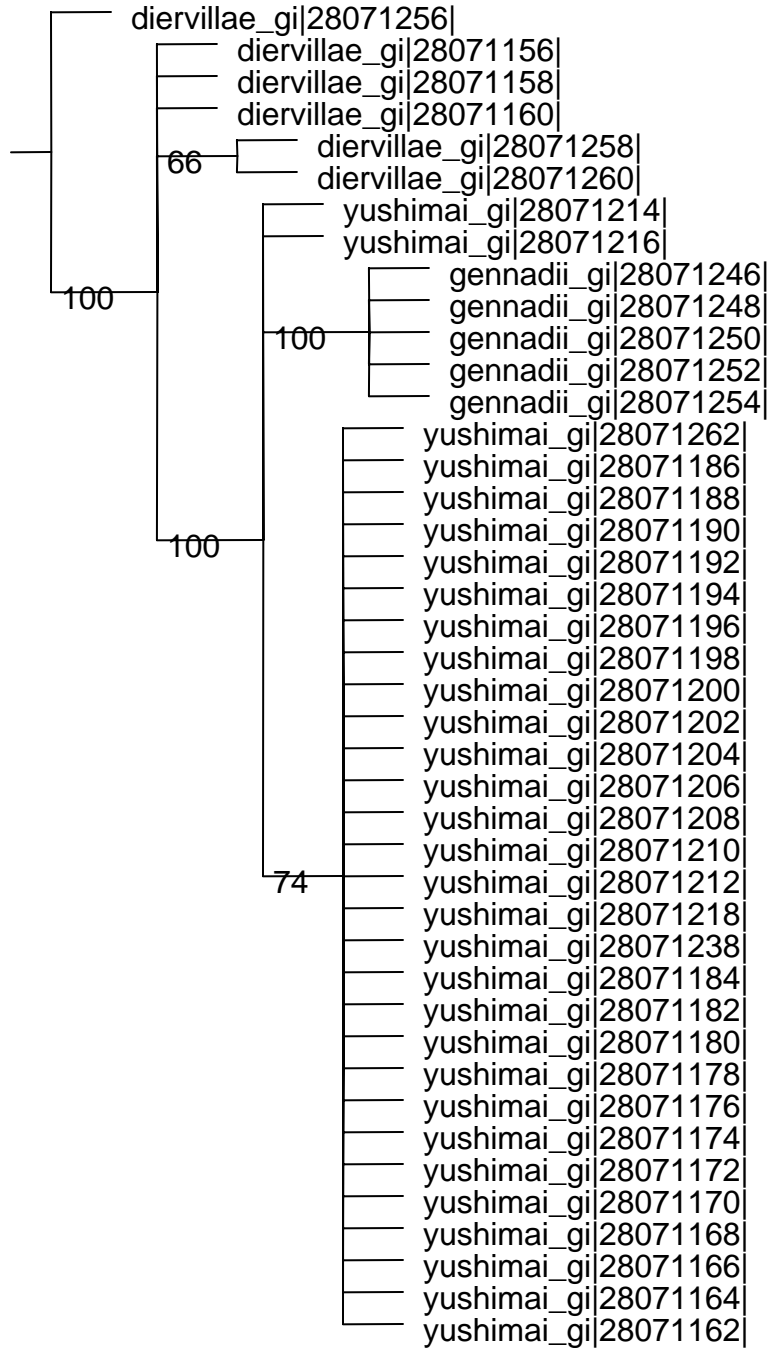
Strict consensus of 10 trees (0 taxa excluded)



Asphondylia

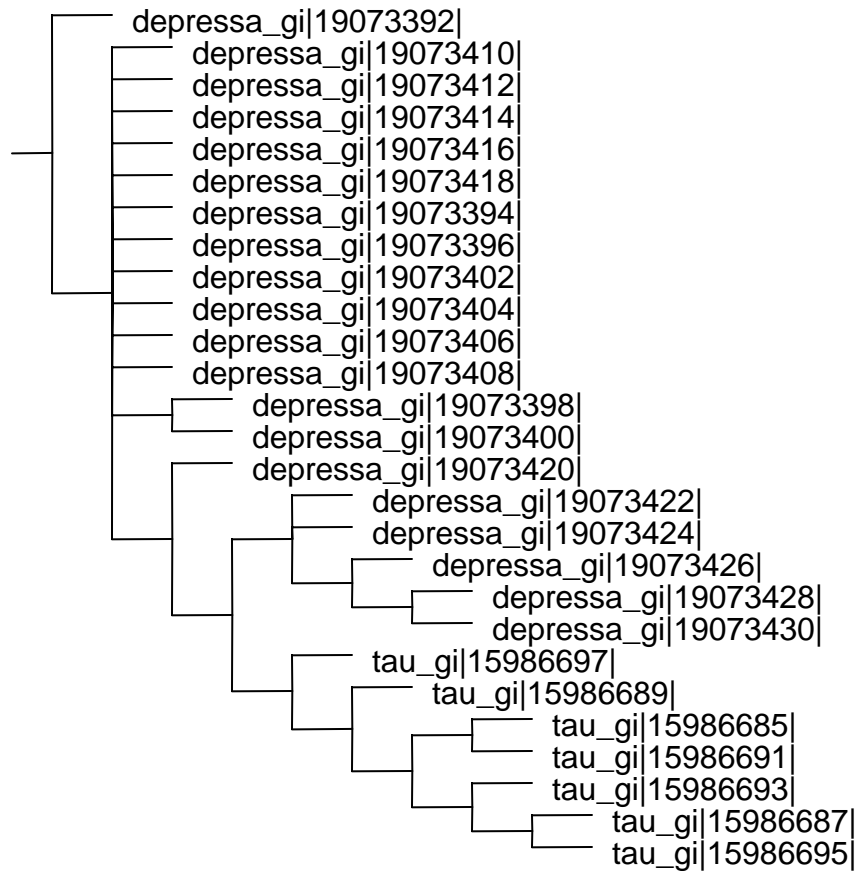
Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



Bactrocera

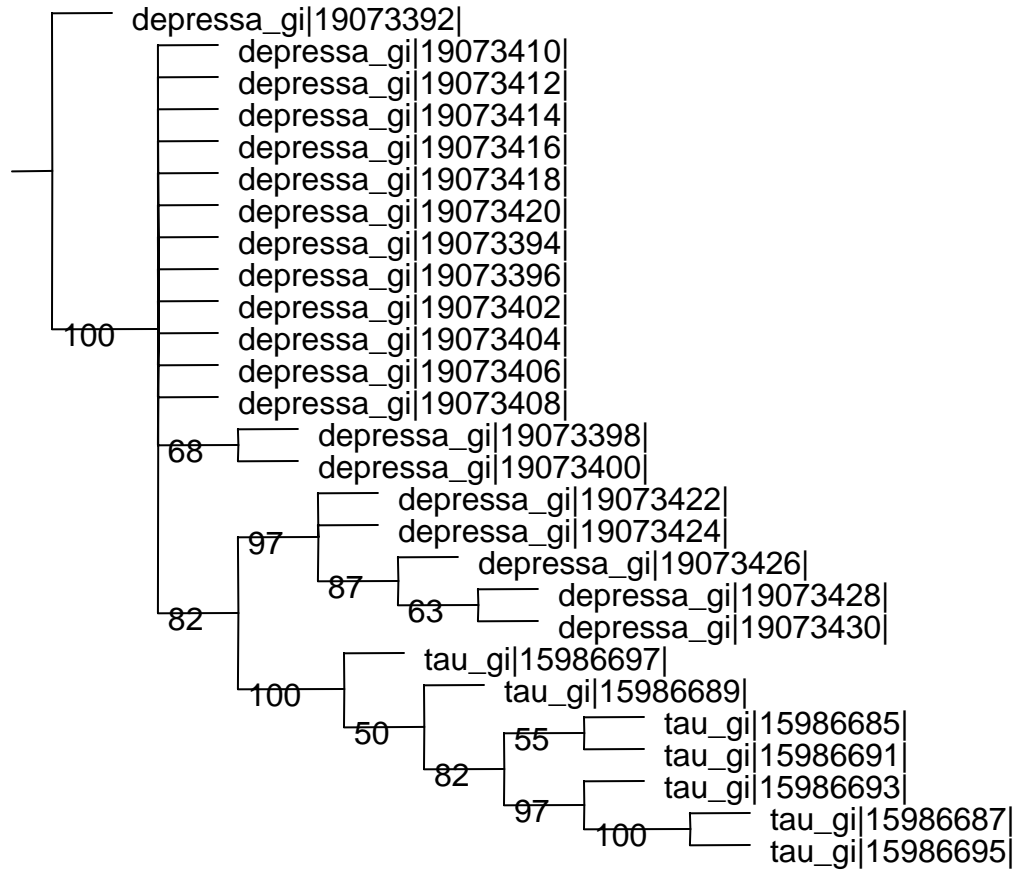
Strict consensus of 3 trees (0 taxa excluded)



Bactrocera

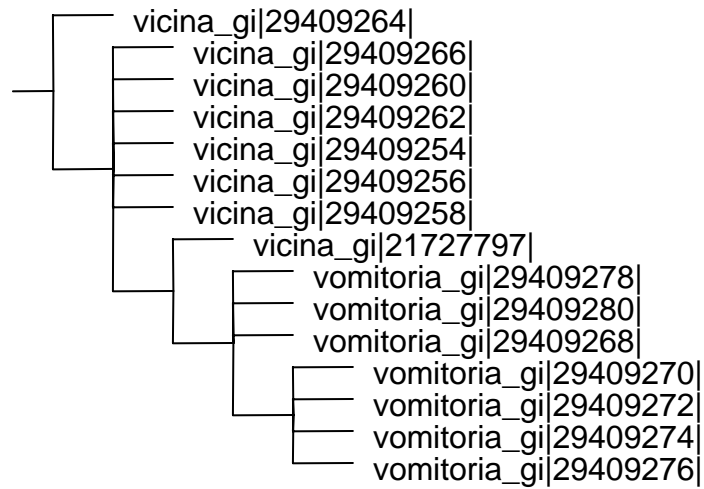
Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



Calliphora

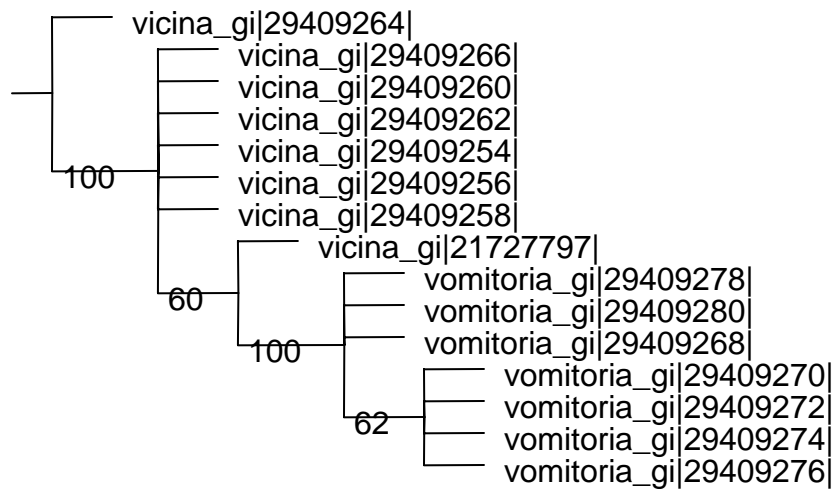
Strict consensus of 1 trees (0 taxa excluded)



Calliphora

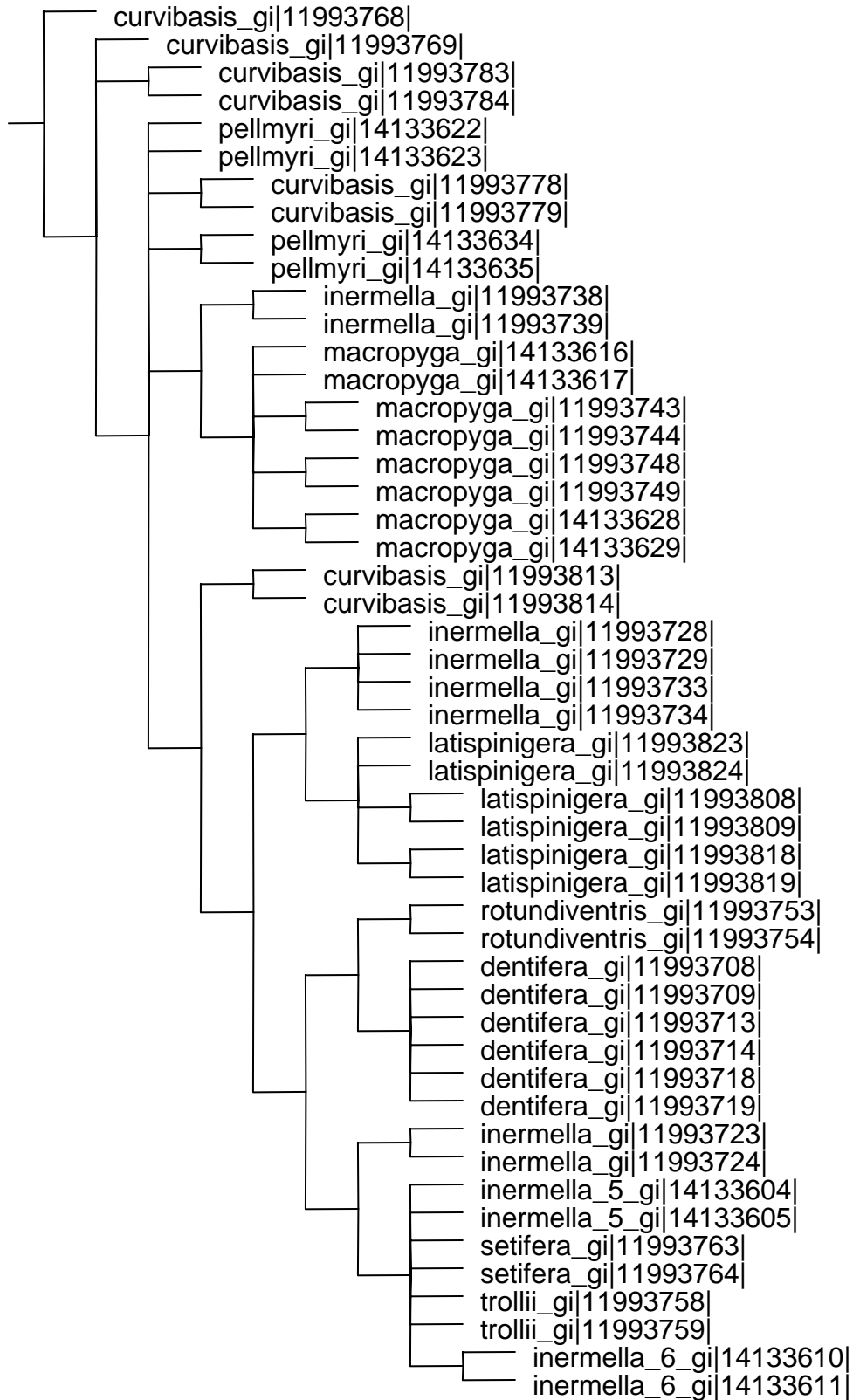
Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



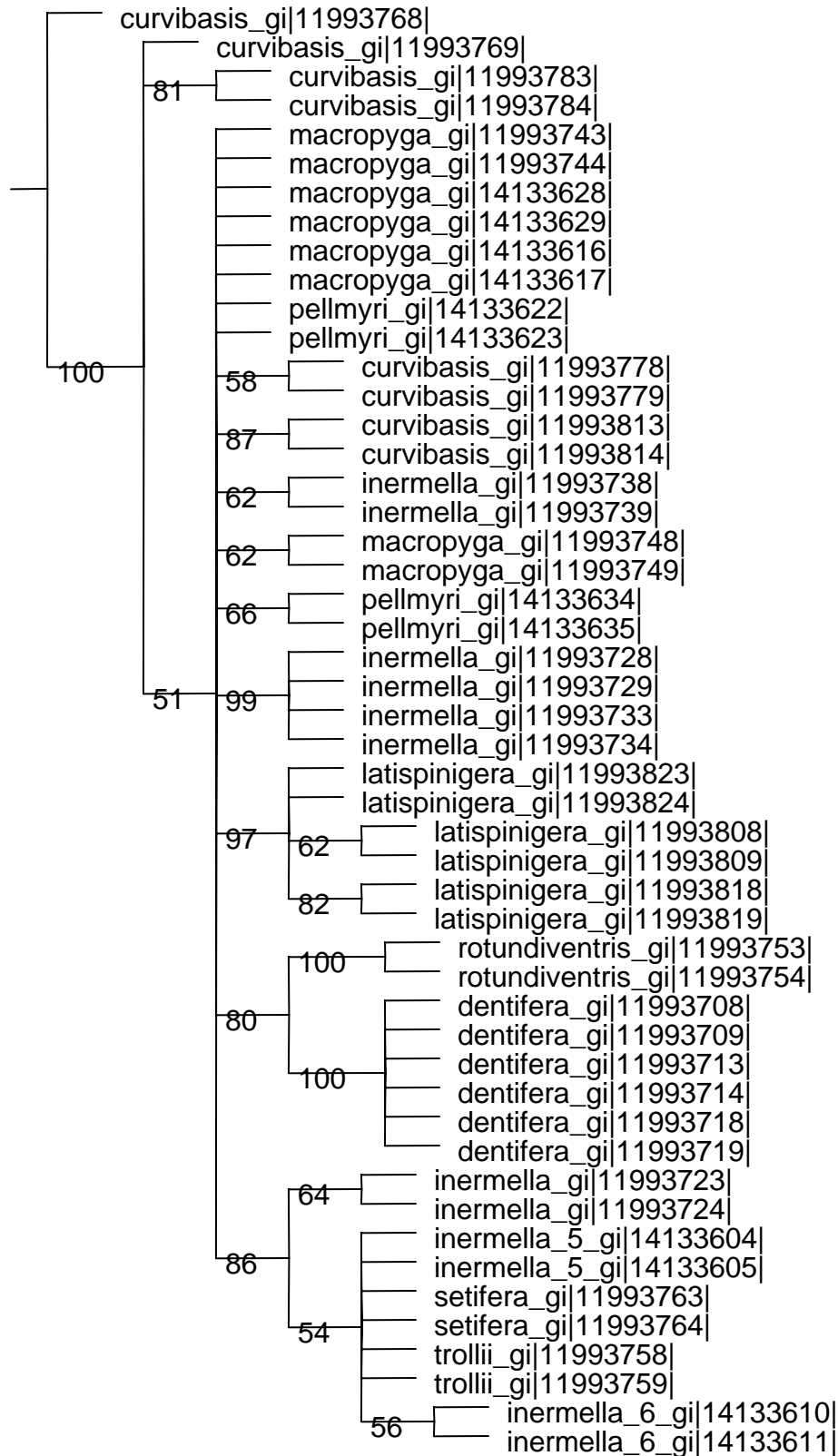
Chiastocheta

Strict consensus of 9 trees (0 taxa excluded)



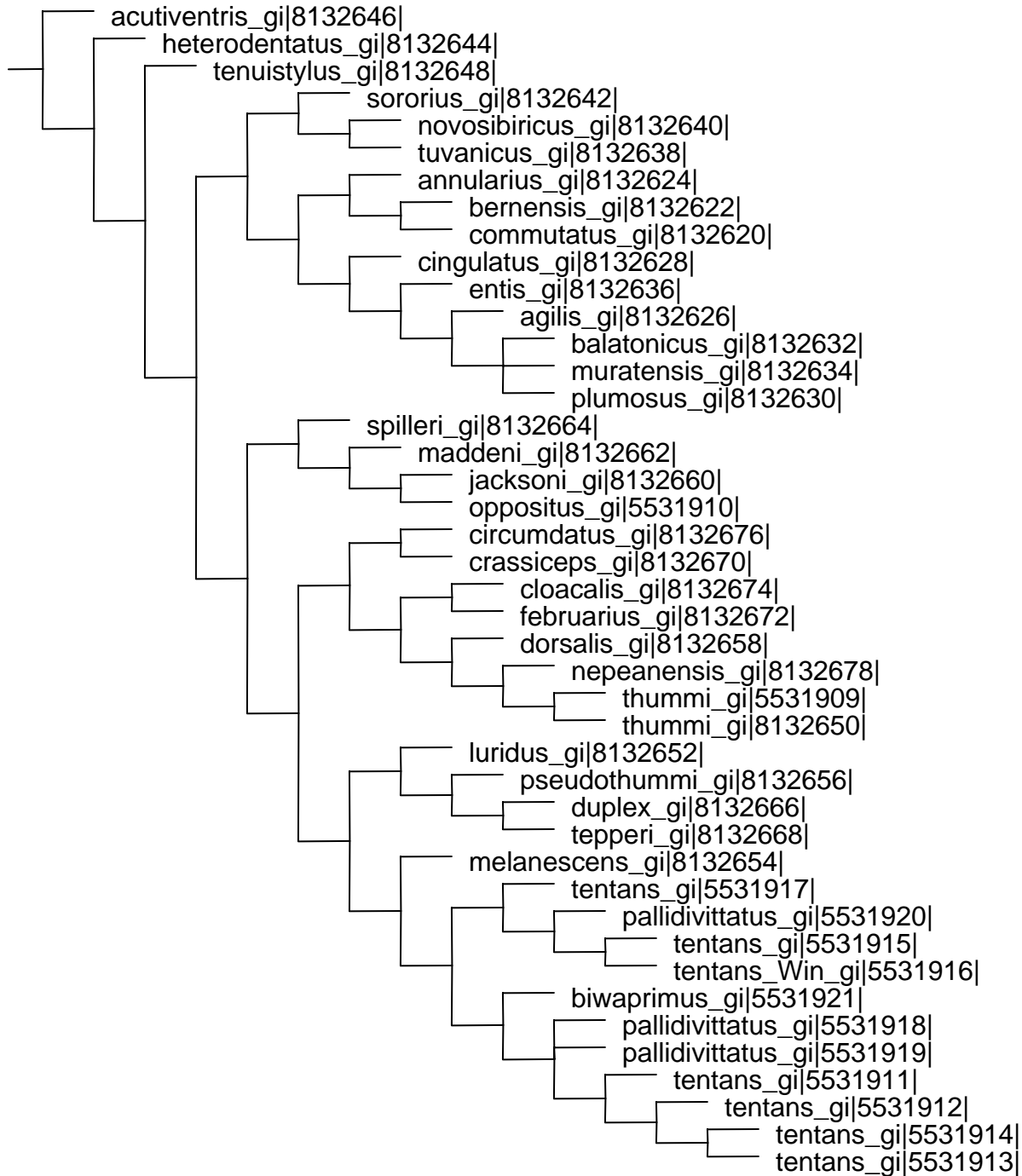
Chiastocheta

Bootstrap Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



Chironomus

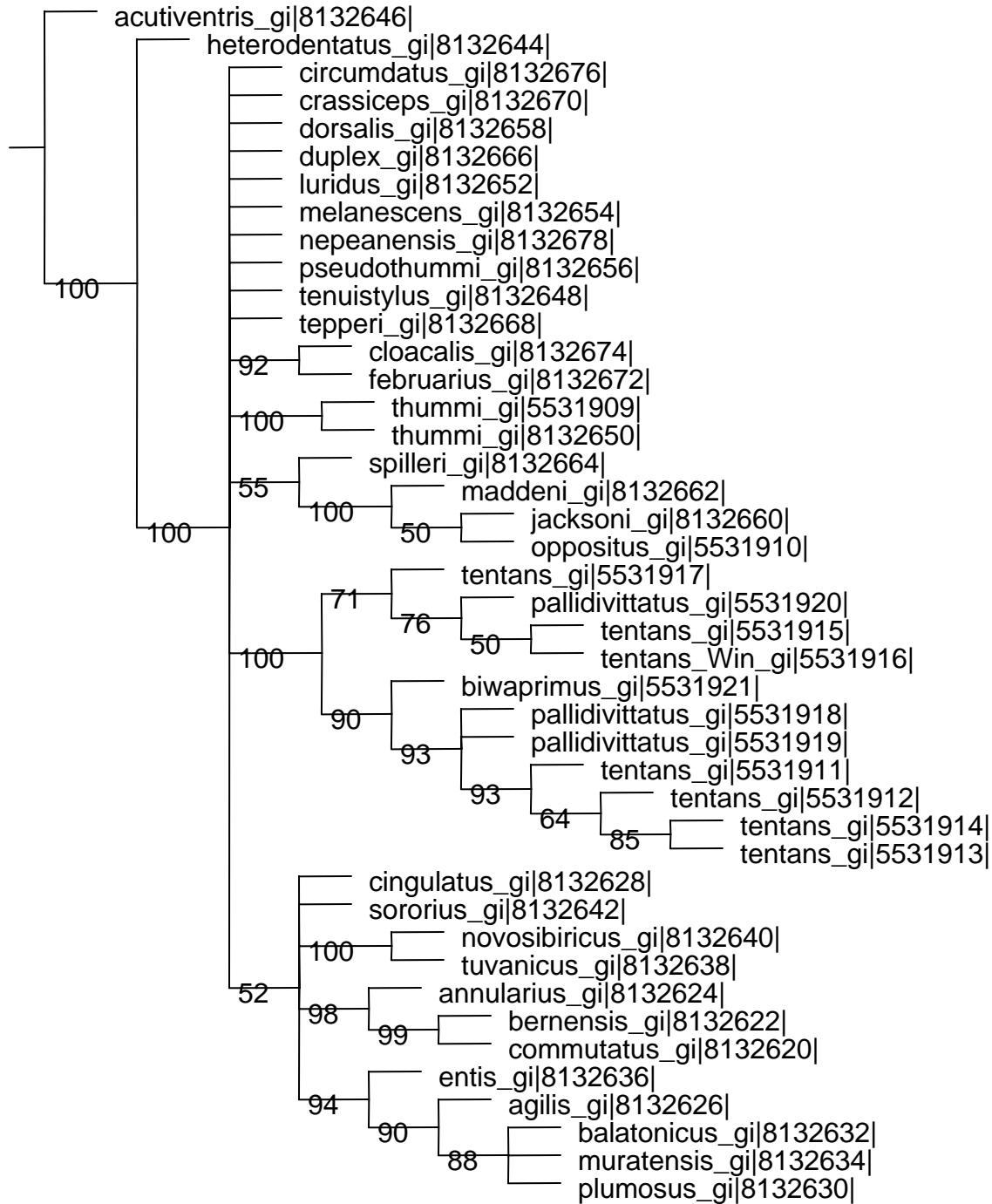
Strict consensus of 6 trees (0 taxa excluded)



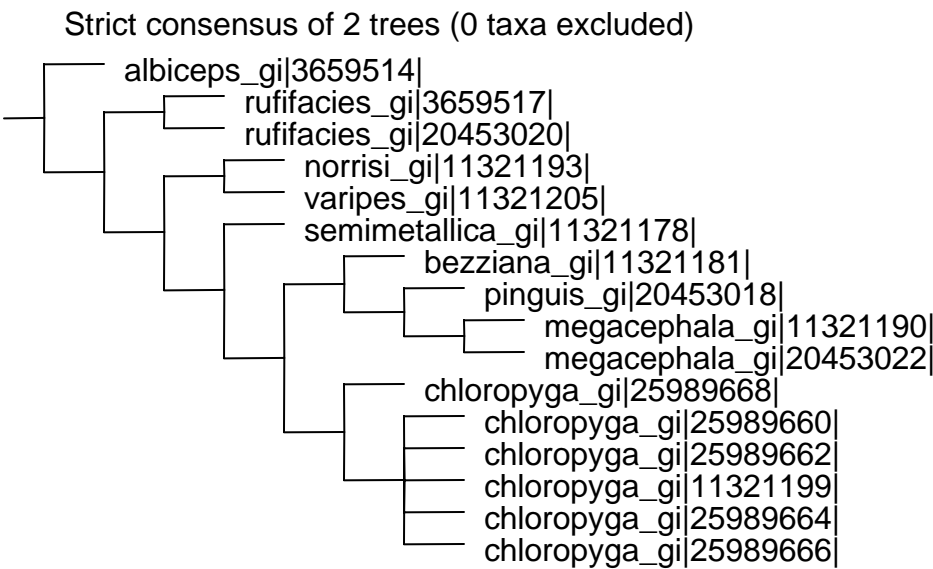
Chironomus

Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap

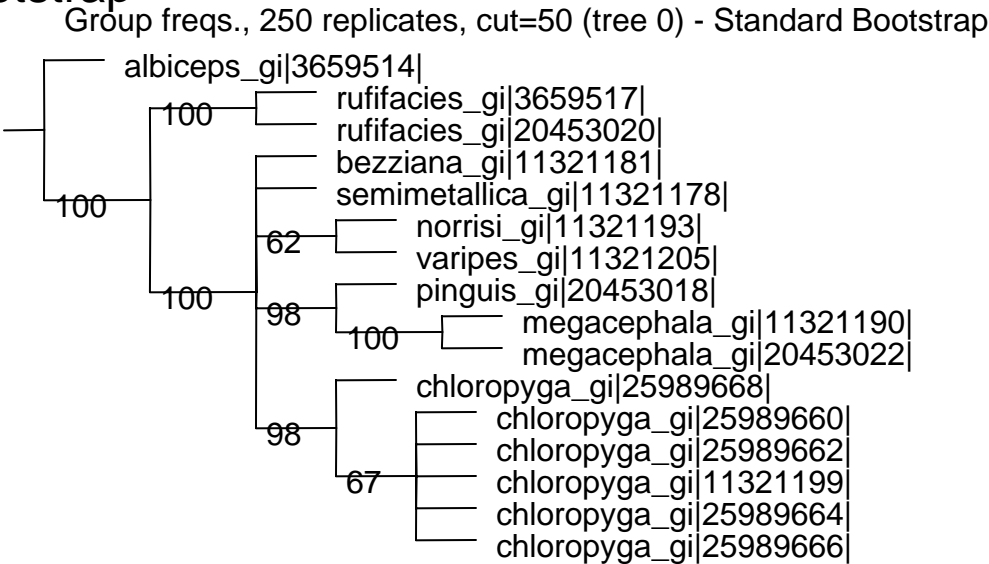


Chrysomya



Chrysomya

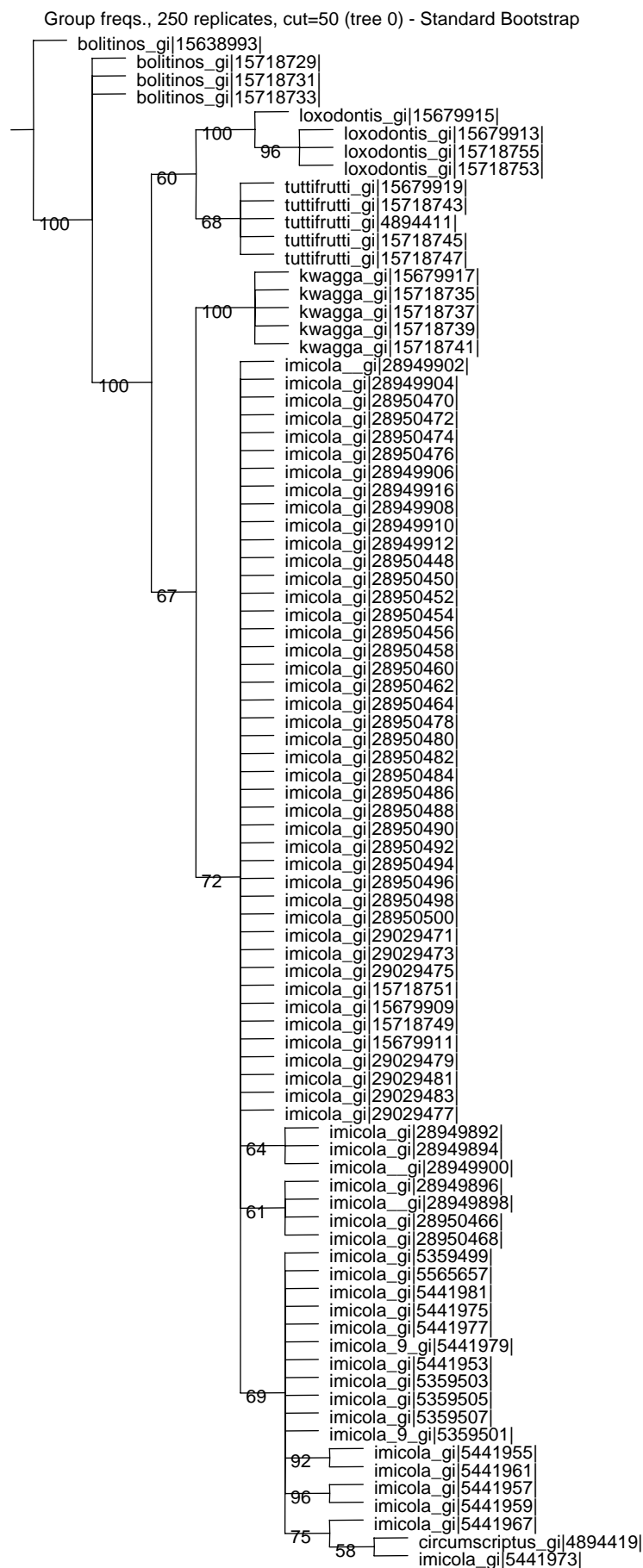
Bootstrap



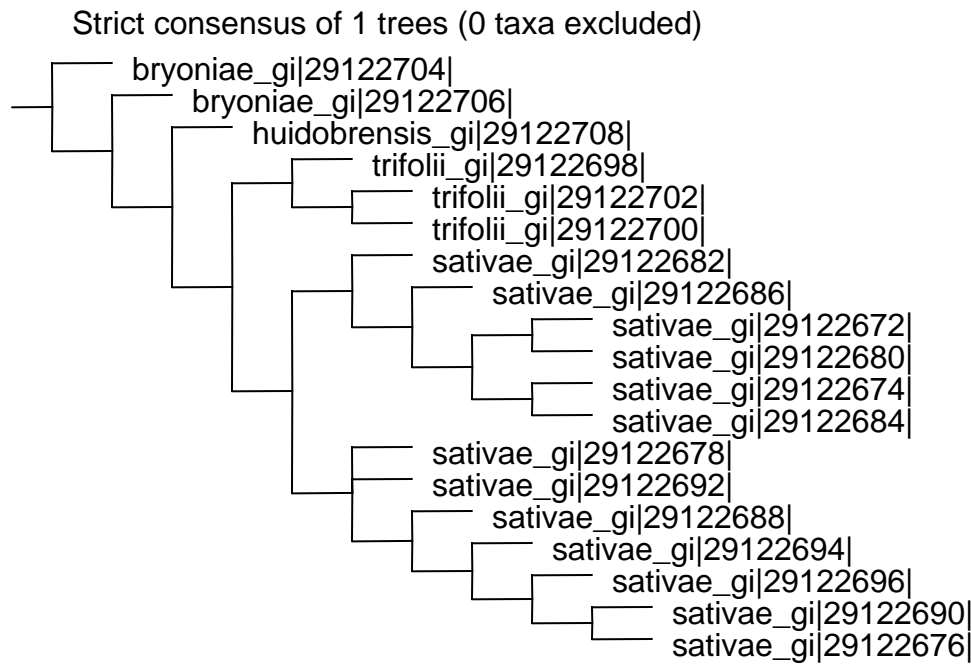
Culicoides



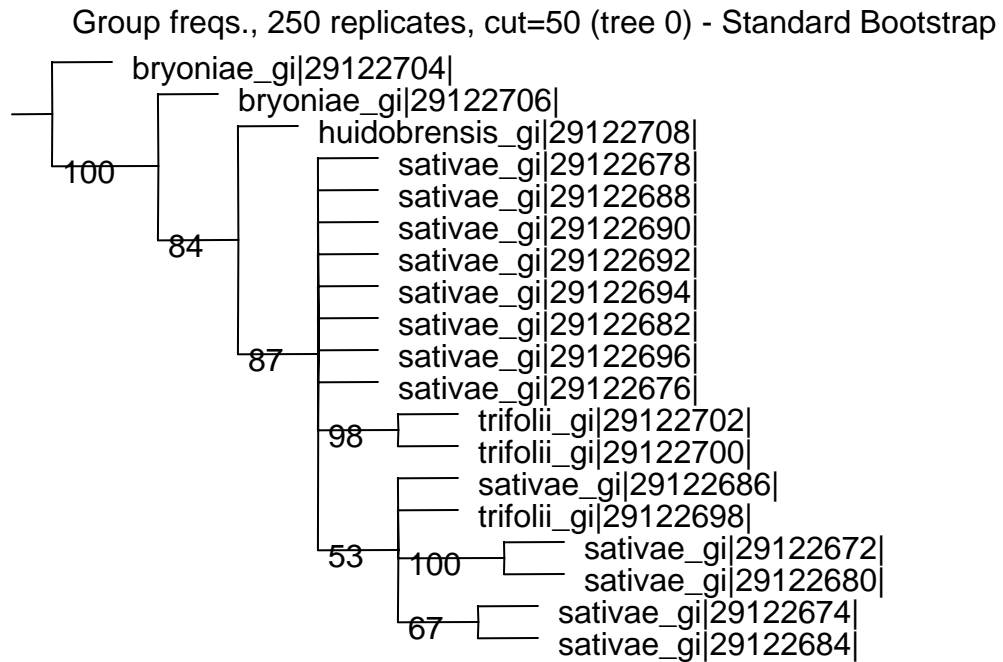
Culicoides Bootstrap



Liriomyza

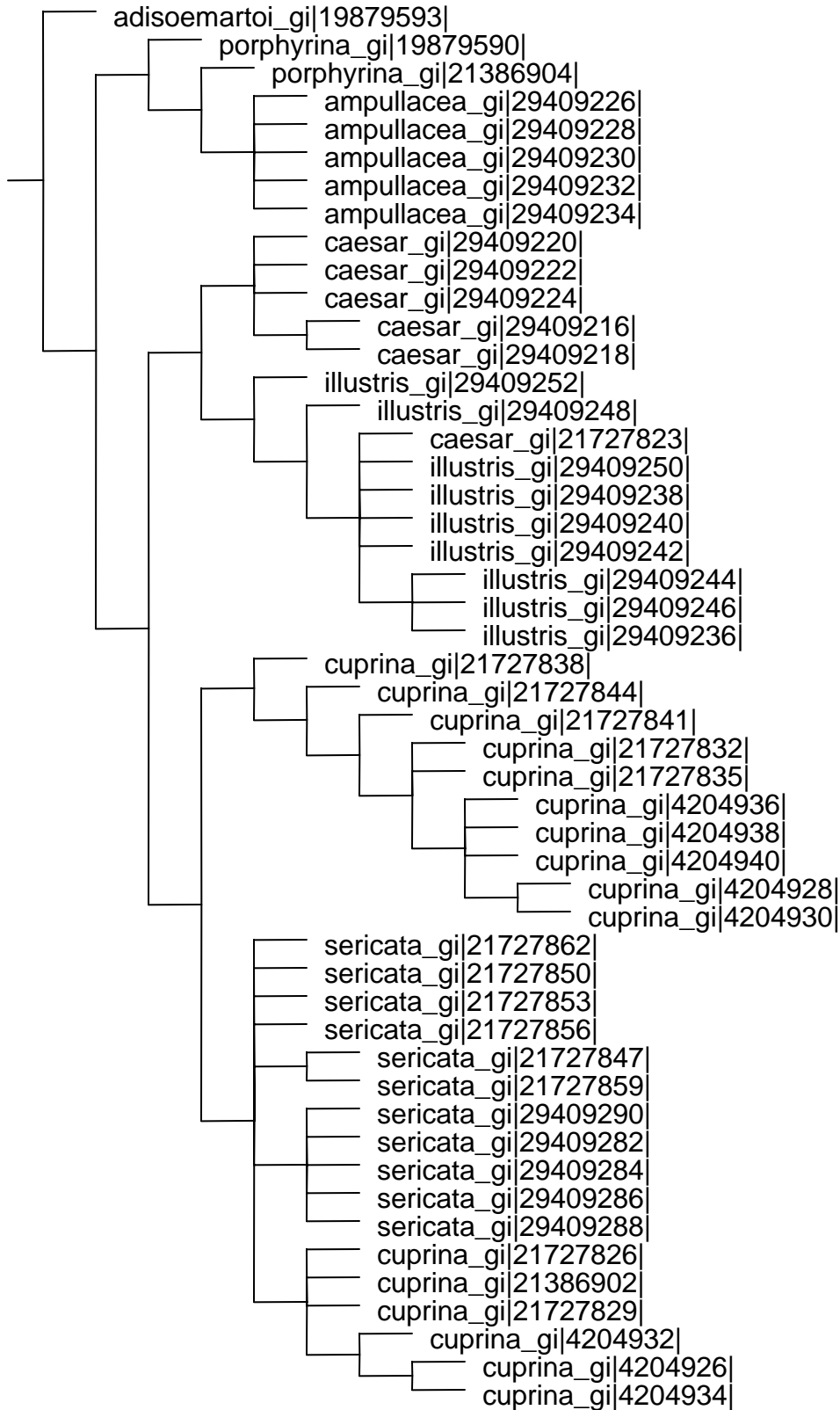


Liriomyza
Bootstrap



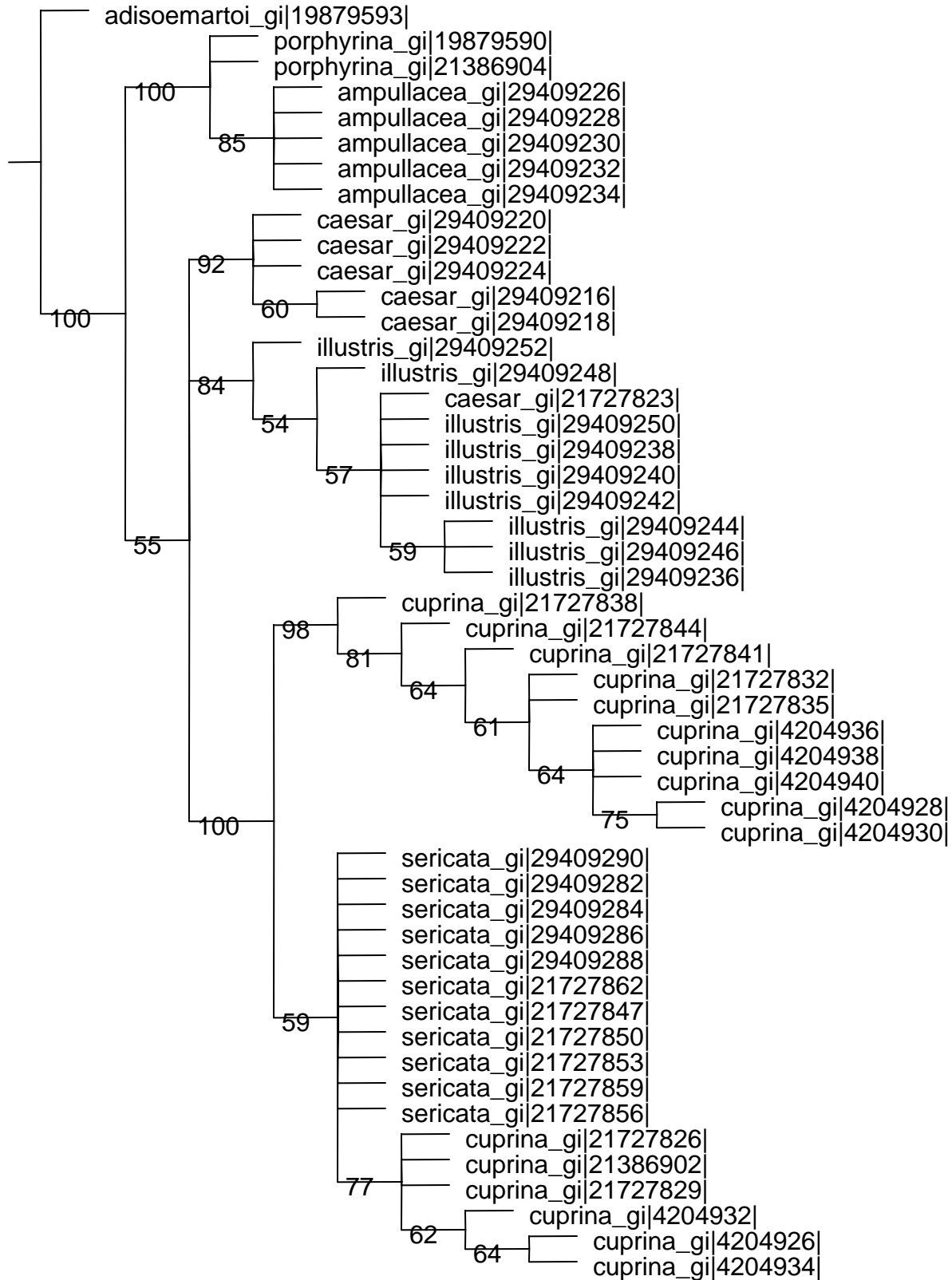
Lucilia

Strict consensus of 7 trees (0 taxa excluded)



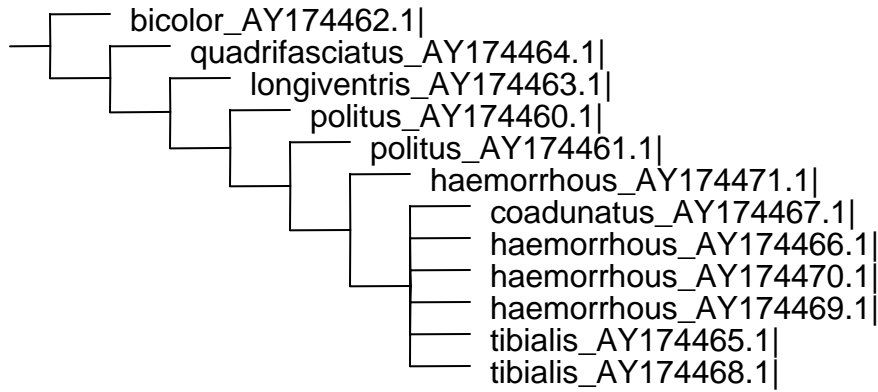
Lucilia Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



Paragus

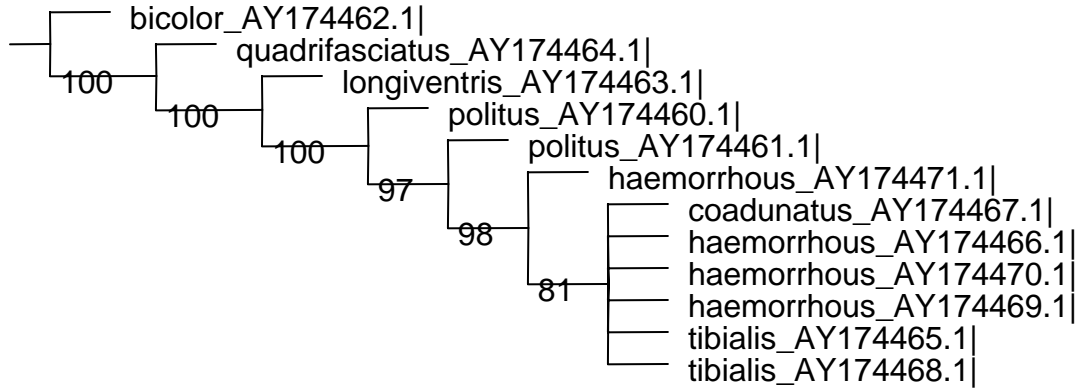
Strict consensus of 1 trees (0 taxa excluded)



Paragus

Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



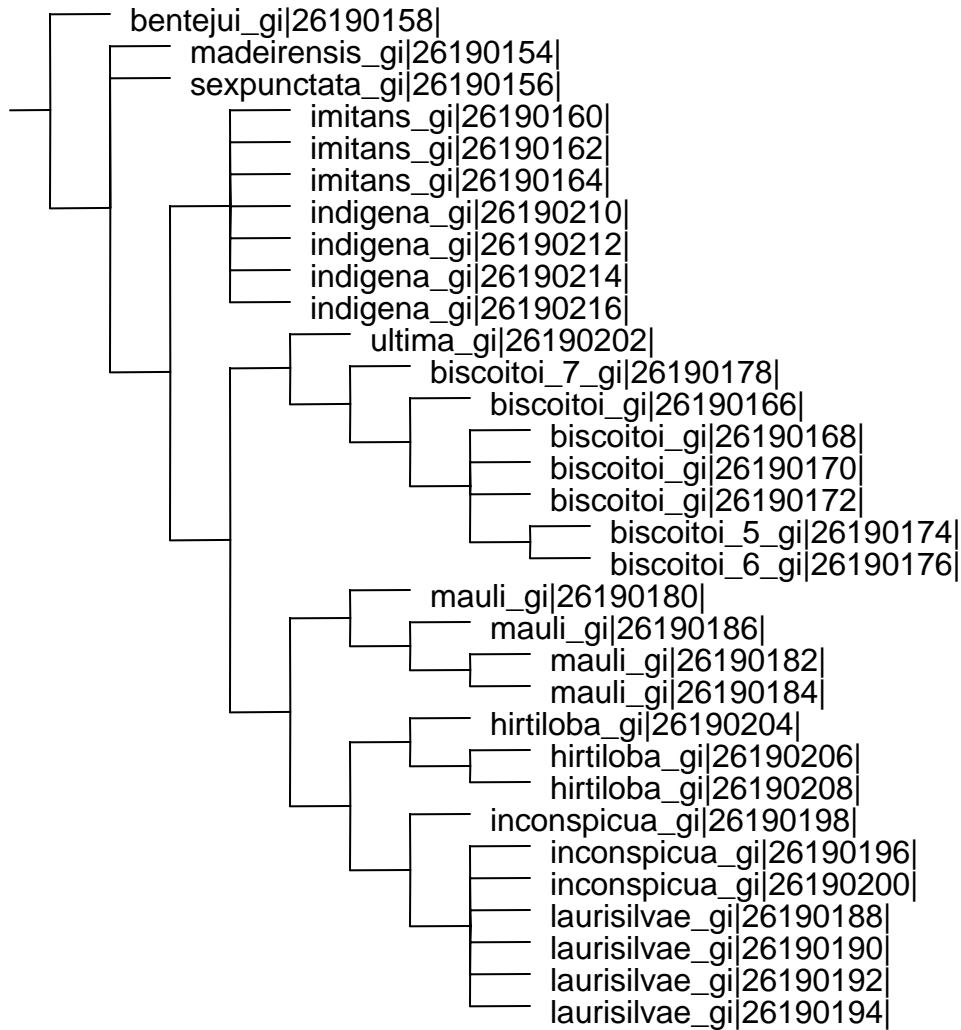
Phytomyza

Bootstrap

Phylogenetic tree of the genus *Plantago* based on ITS2 sequences. The tree shows relationships between various species, with bootstrap values indicated at the nodes. The species names are listed on the right, each followed by its GenBank accession number in brackets. The tree is rooted on the left and branches to the right. The species are: *aconiti_gi|9799490|*, *aquilegiovora_gi|9799496|*, *erigerophila_gi|9799505|*, *flavicornis_gi|9799487|*, *ilicis_gi|9799508|*, *Plantaginis_gi|9799493|*, *glabricola_gi|9799520|*, *glabricola_gi|9799514|*, *jucunda_gi|9799526|*, *vomitoriae_gi|9799532|*, *ditmani_gi|9799568|*, *nemopantheri_gi|9799559|*, *ilicicola_gi|9799586|*, *ilicicola_gi|9799589|*, *opacae_gi|9799607|*, *opacae_gi|9799604|*, *verticillatae_gi|9799538|*, *verticillatae_gi|9799553|*, and *verticillatae_gi|9799556|*. Bootstrap values are shown at the nodes: 100, 65, 100, 92, 54, 97, 100, 100, 100, 100.

Sapromyza

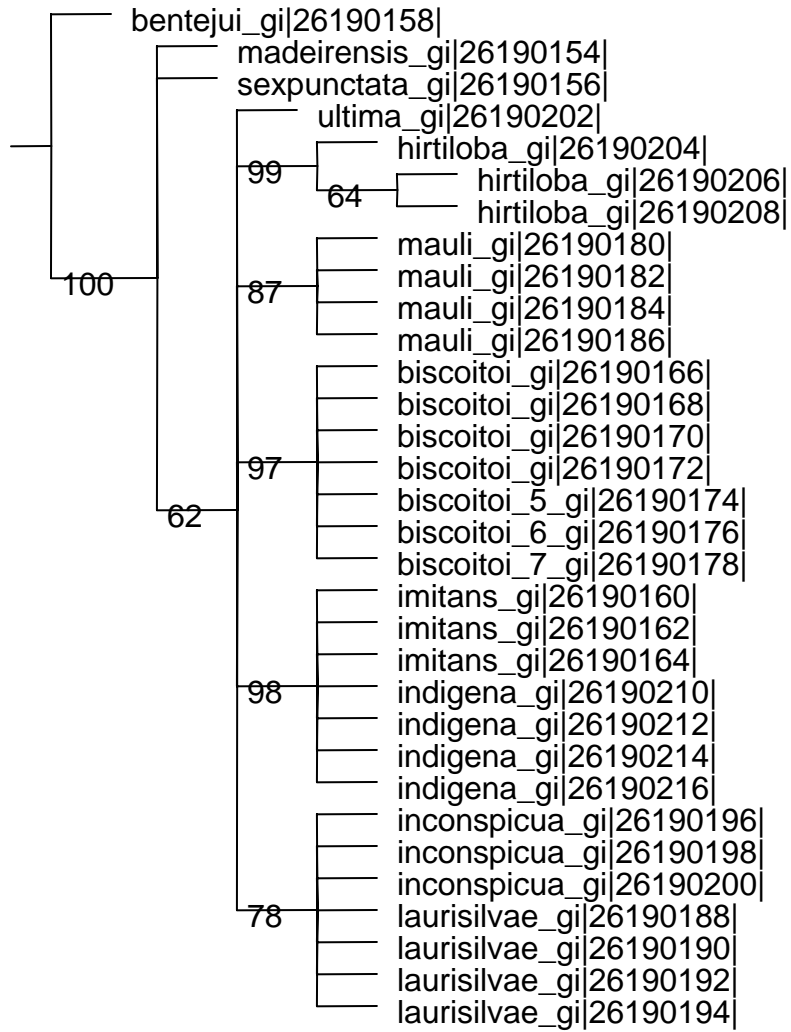
Strict consensus of 9 trees (0 taxa excluded)



Sapromyza

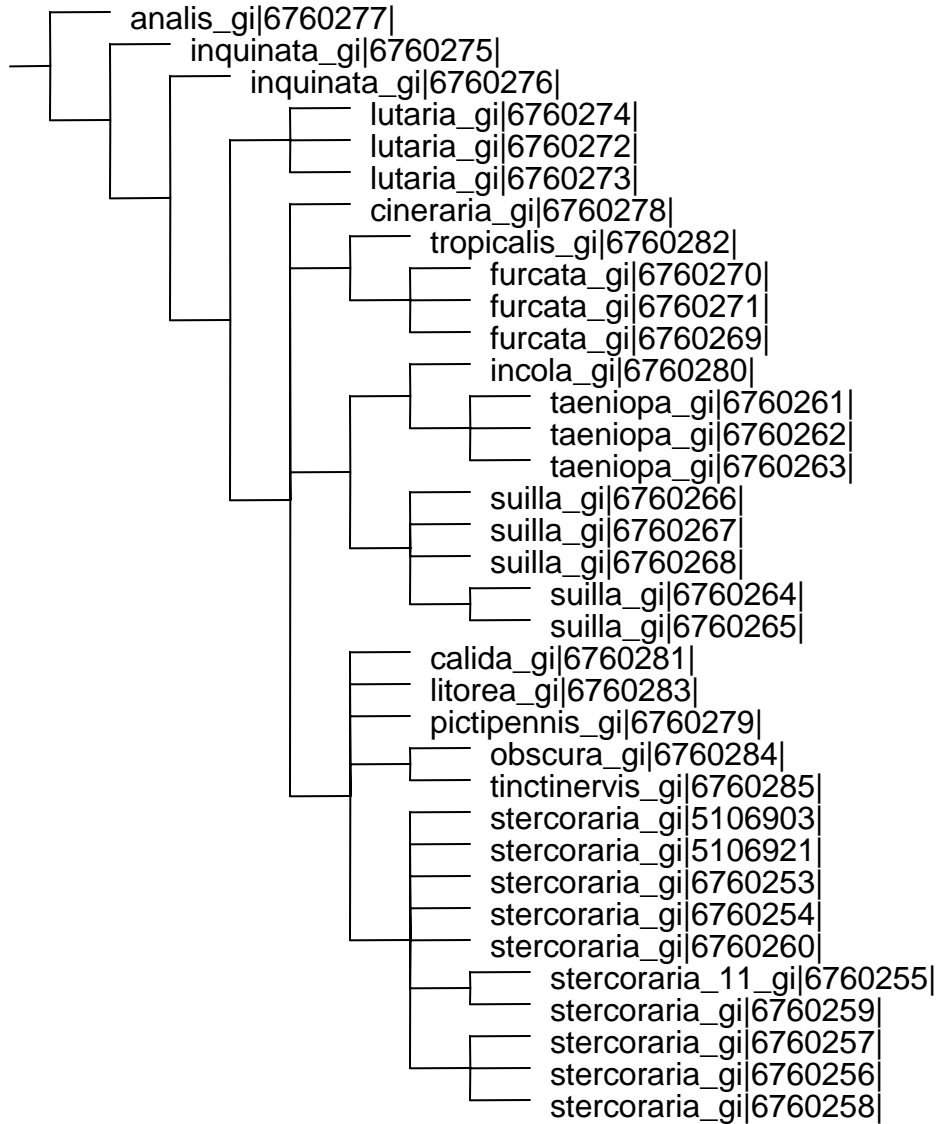
Bootstrap

Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



Scathophaga

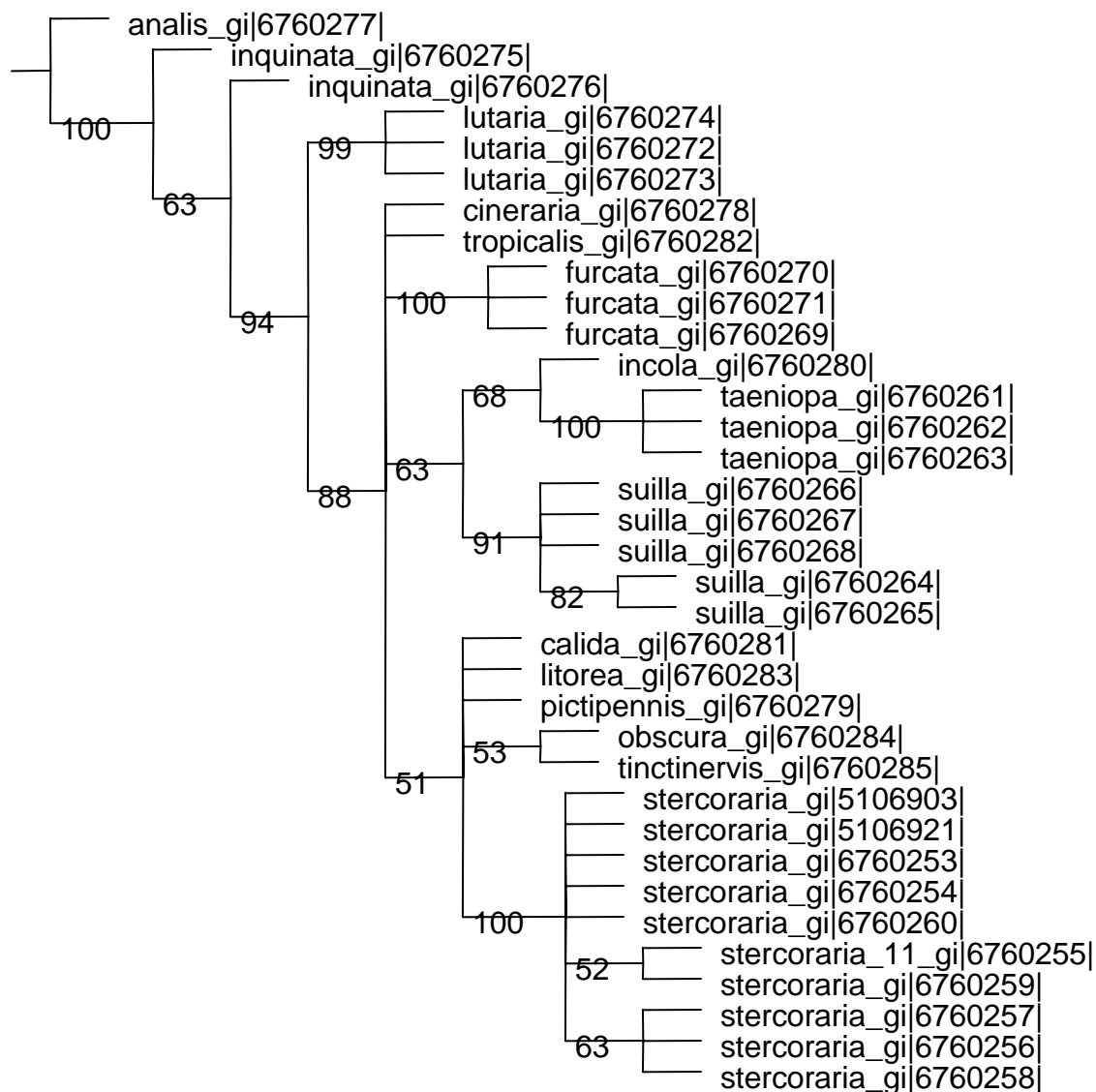
Strict consensus of 5 trees (0 taxa excluded)



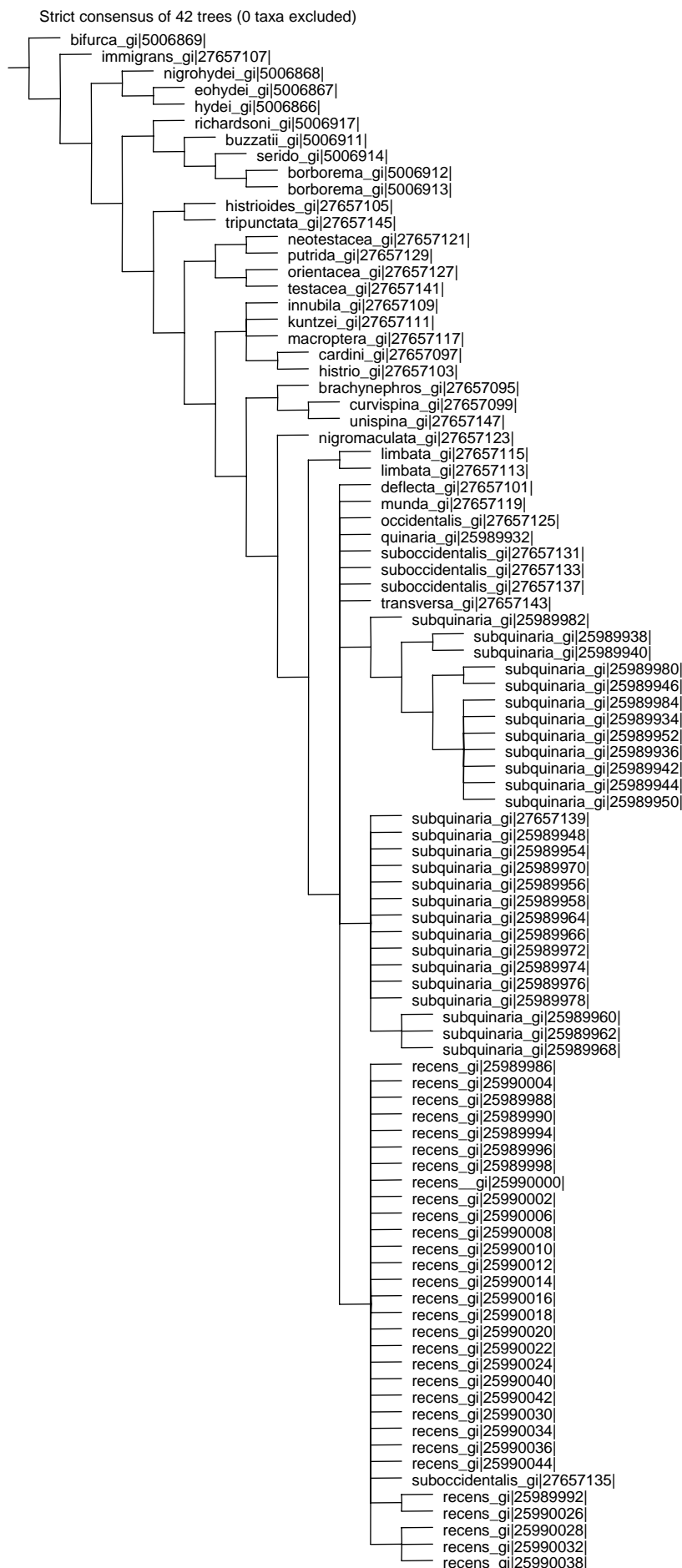
Scathophaga

Bootstrap

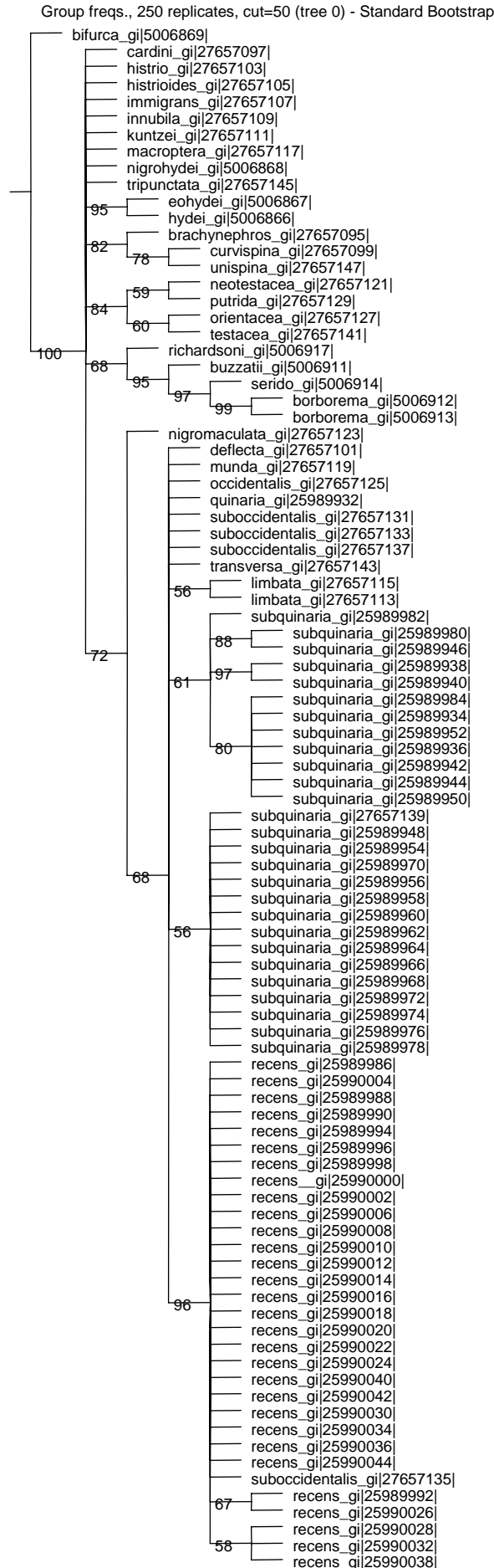
Group freqs., 250 replicates, cut=50 (tree 0) - Standard Bootstrap



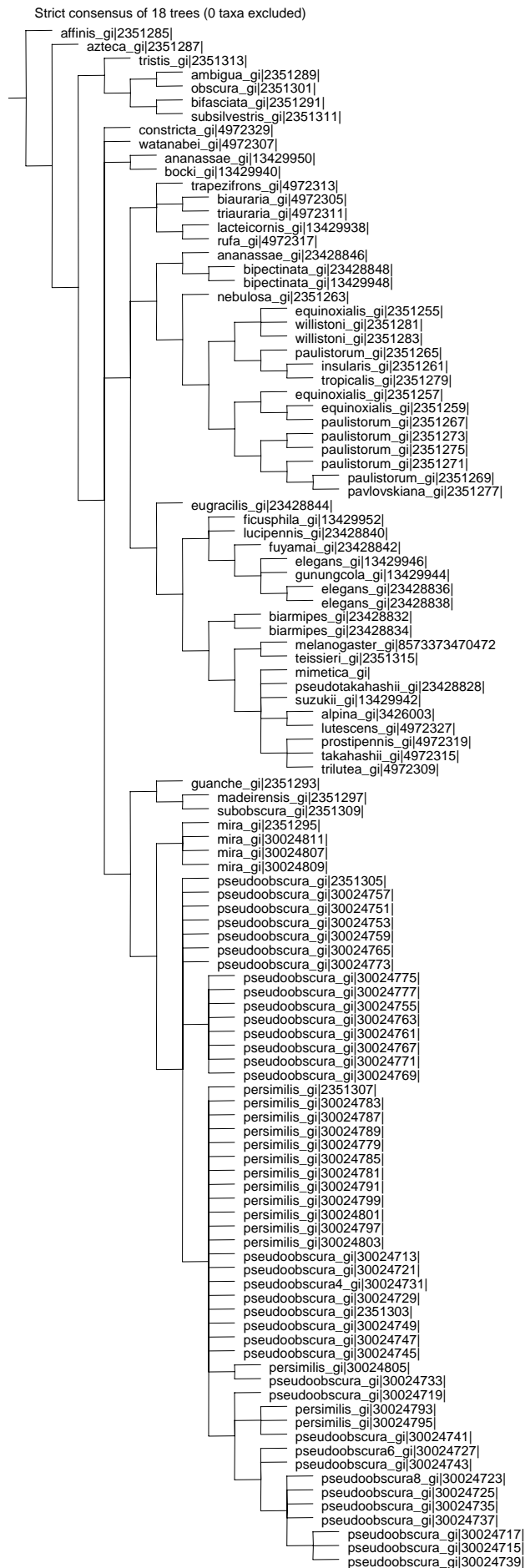
Subgenus *Drosophila*



Subgenus *Drosophila* Bootstrap



Subgenus *Sophophora*



Subgenus *Sophophora* Bootstrap

