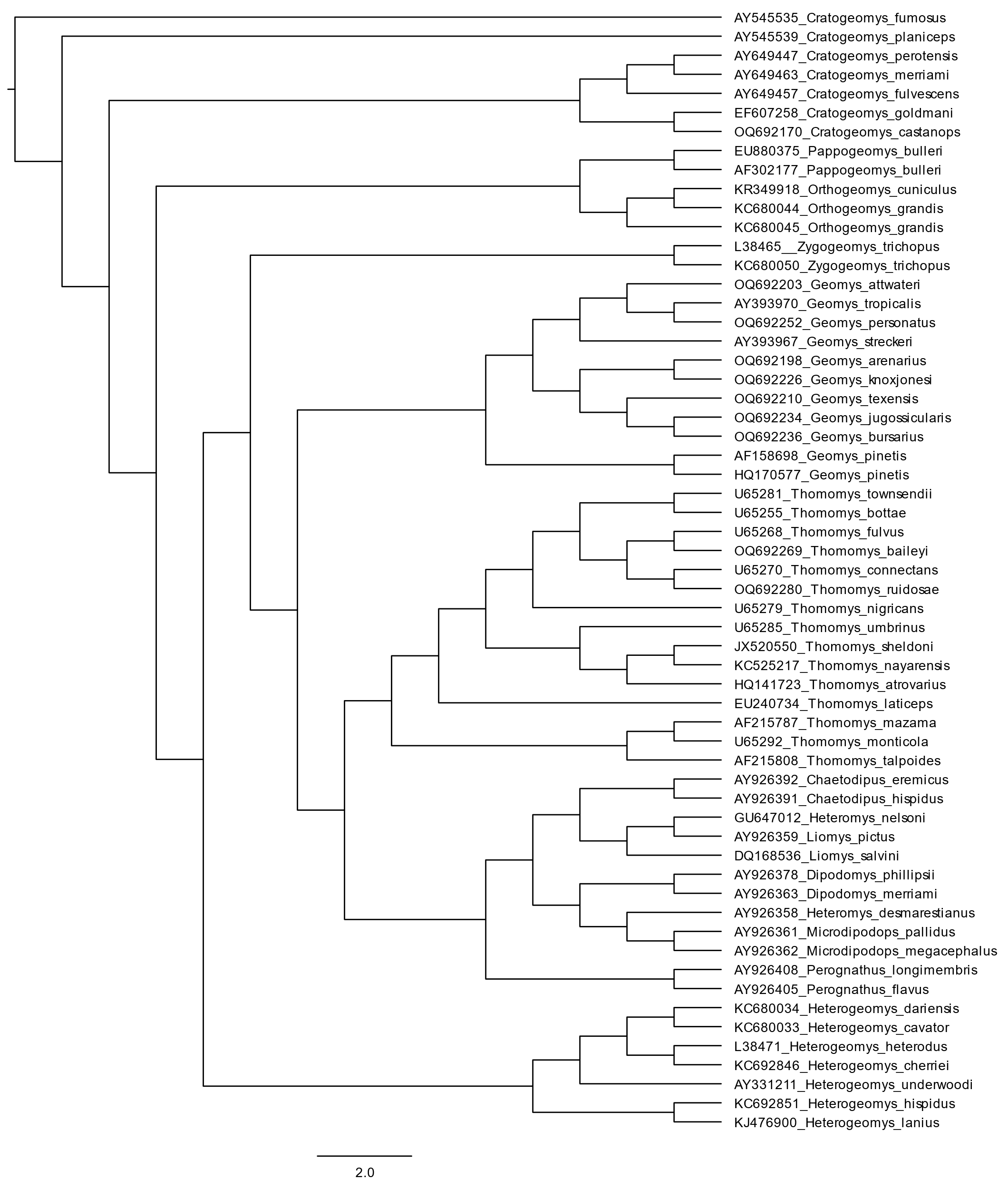
Álvarez-Castañeda, S. T., and C. A. Segura-Trujillo*.* 2025. Genus-level review of pocket gophers in the family Geomyidae. Therya. DOI:10.12933/therya-25-6153

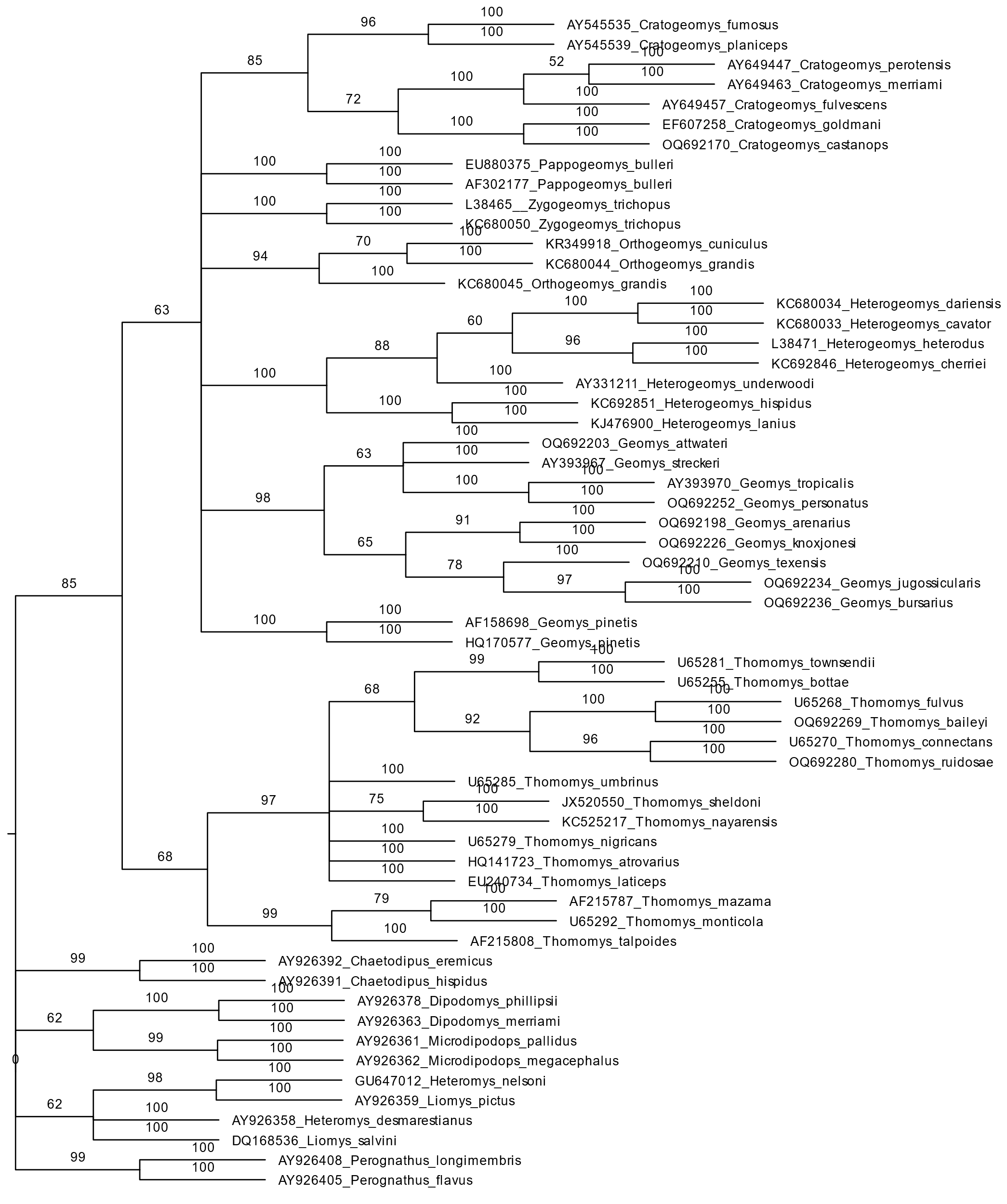
**Supplementary material 3**

Phylogenetic inference using neighbor-joining, unweighted pair group method with arithmetic mean (UPGMA), maximum-parsimony, maximum-likelihood, and Bayesian inference for the cytochrome b and cytochrome oxidase subunit 1 sequences.

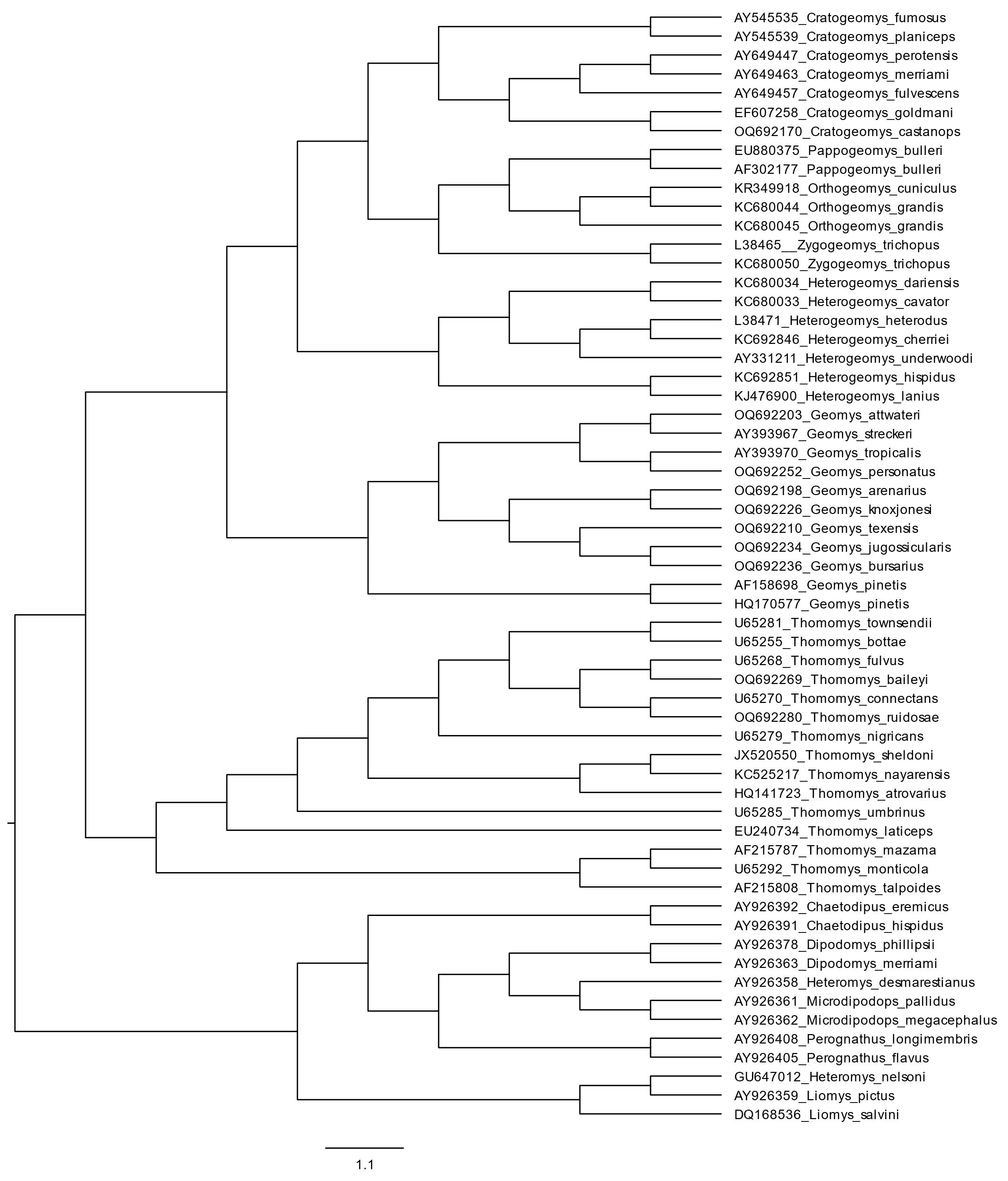
Cytochrome b neighbor-joining.



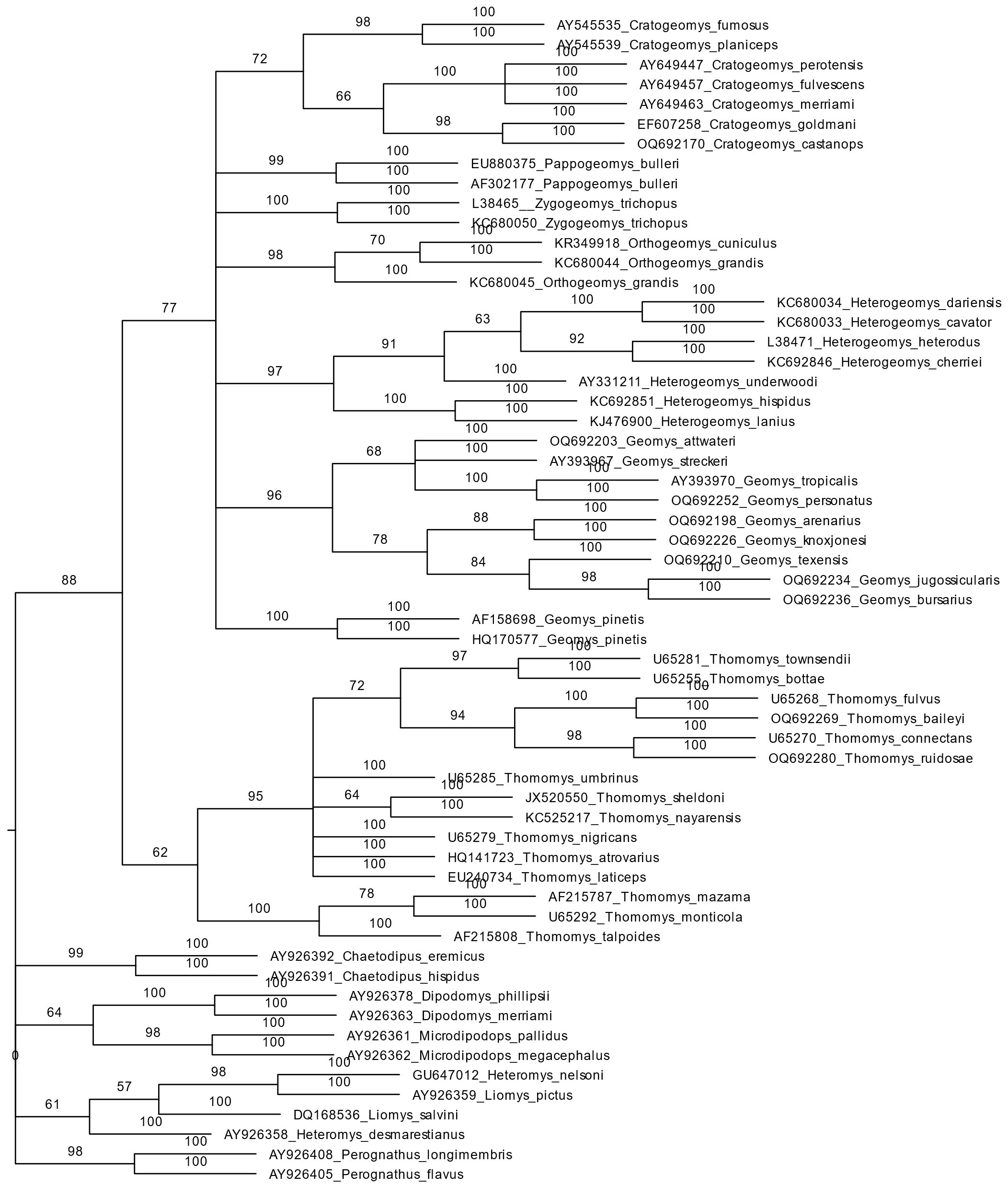
Cytochrome b neighbor-joining bootstrap consensus tree generating 100 pseudoreplicates.



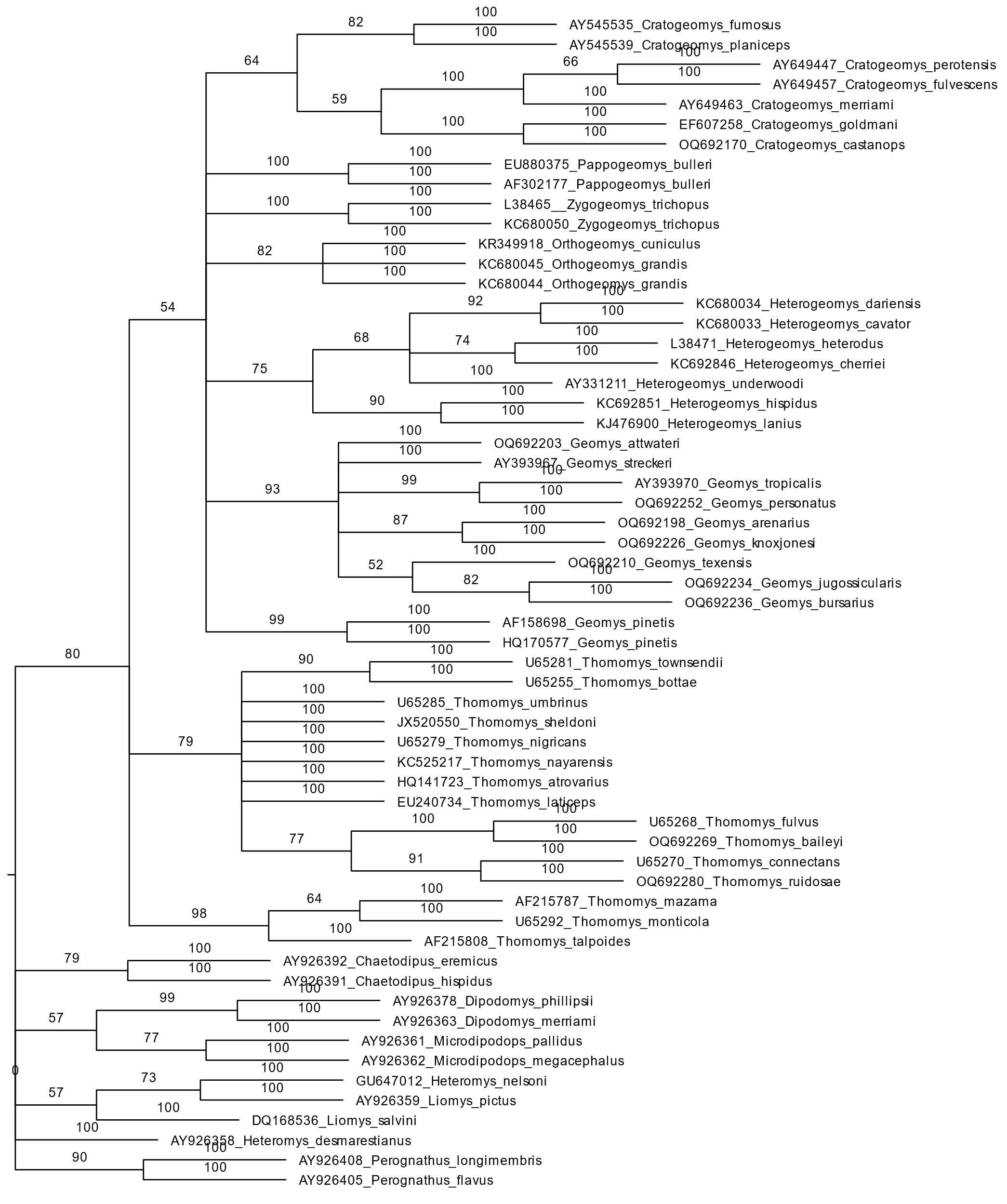
Cytochrome b unweighted pair group method with arithmetic mean.



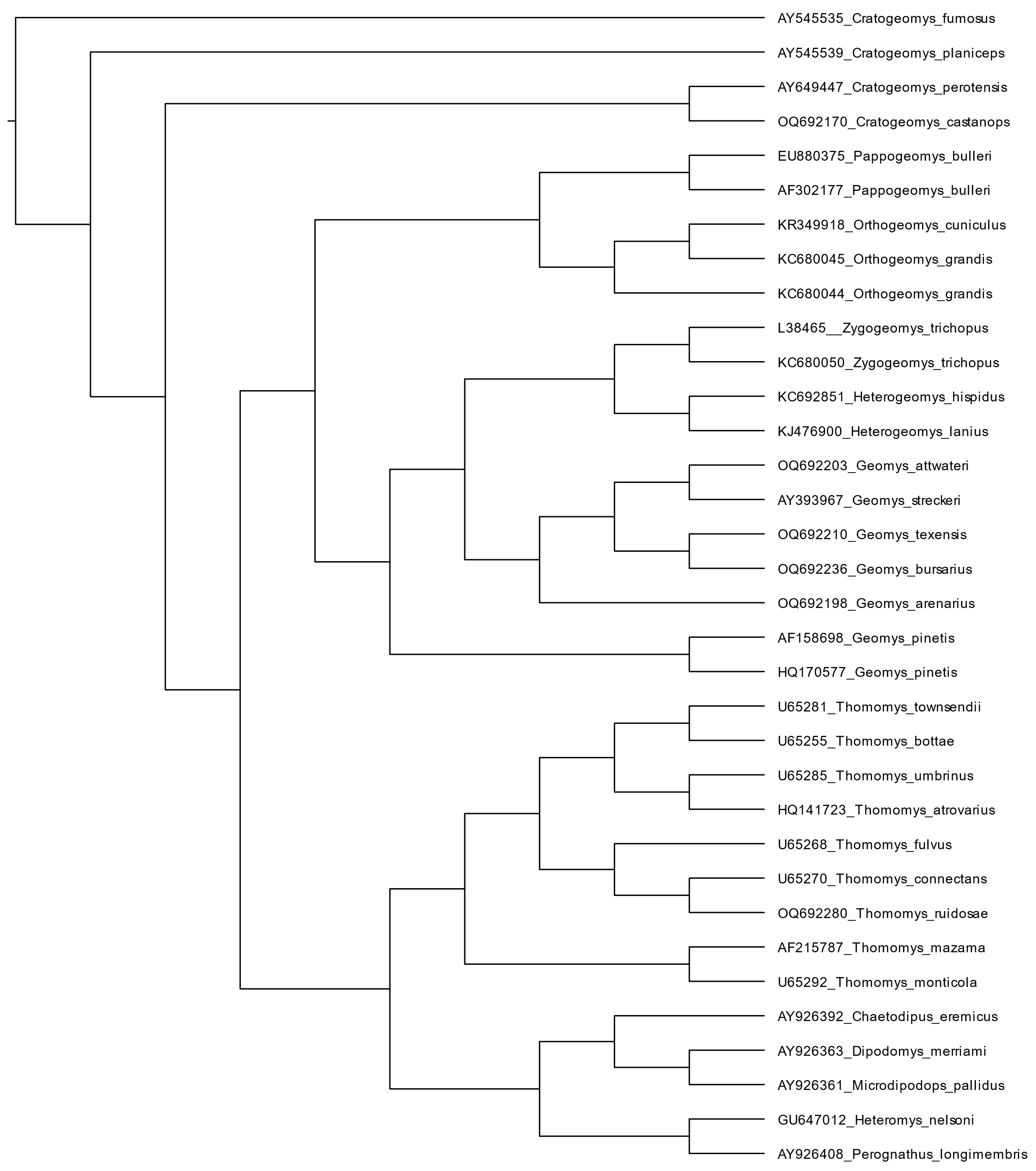
Cytochrome b unweighted pair group method with arithmetic mean bootstrap consensus tree generating 100 pseudoreplicates.



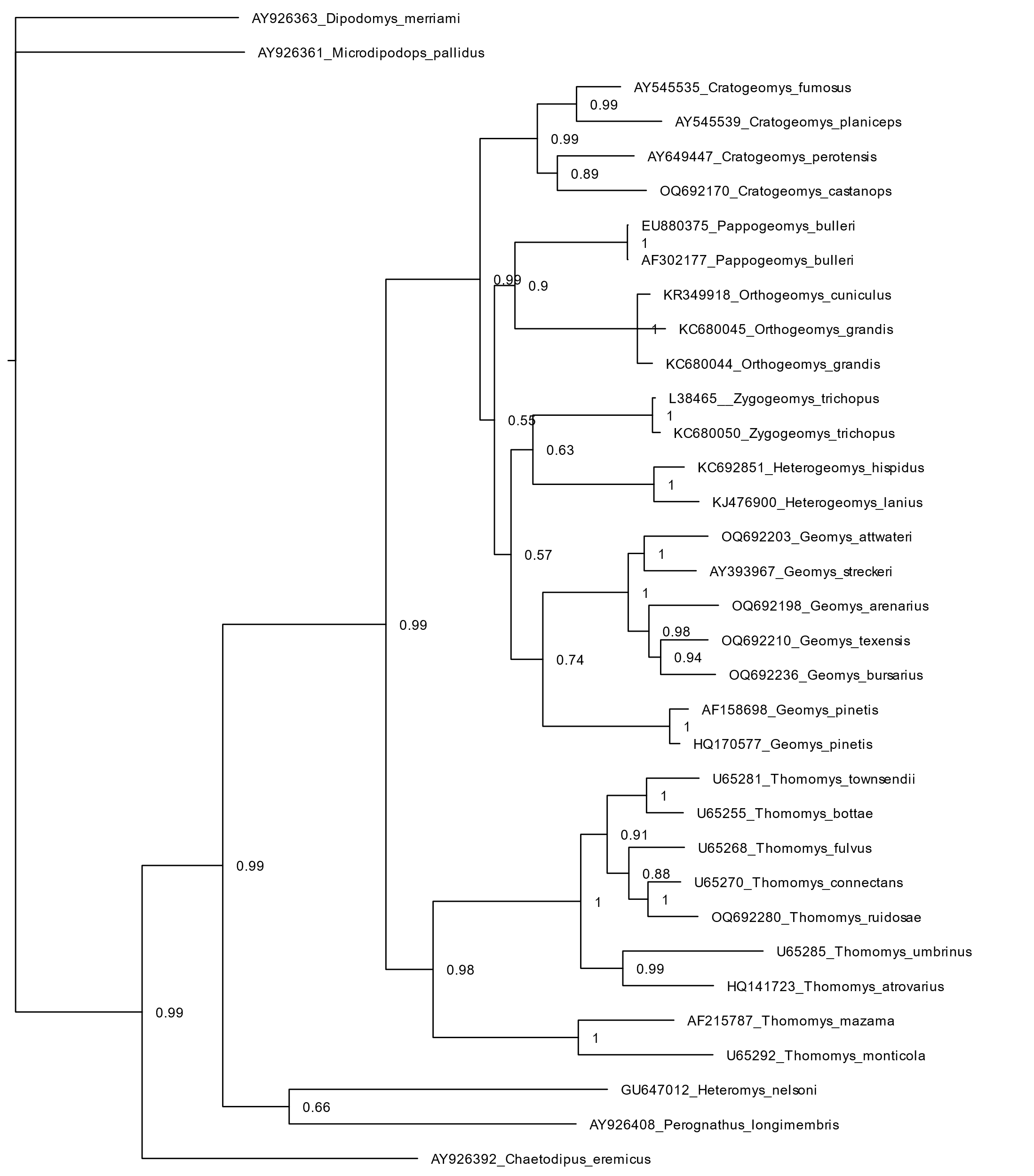
Cytochrome b maximum-parsimony.



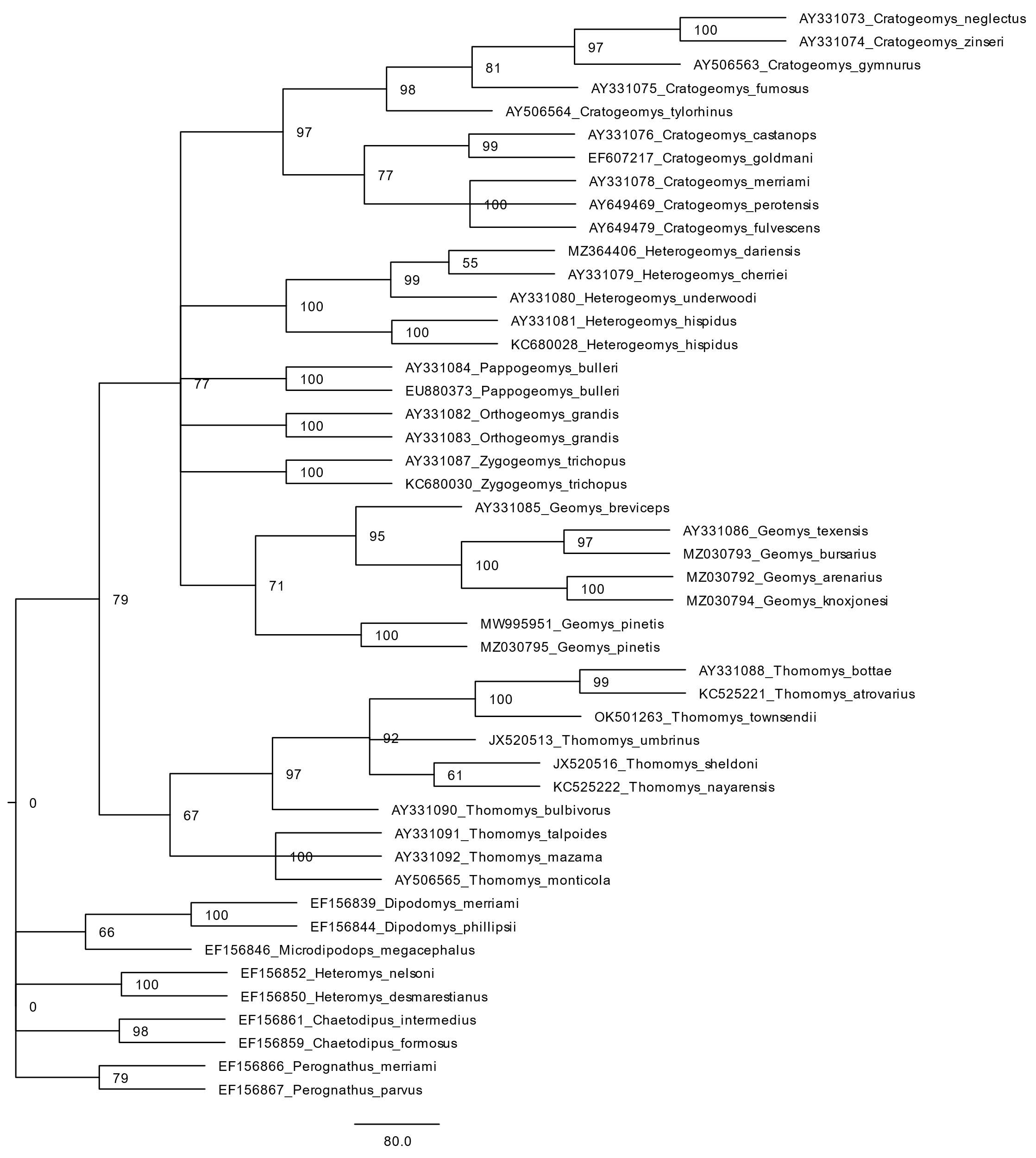
Cytochrome b maximum Likelihood method.



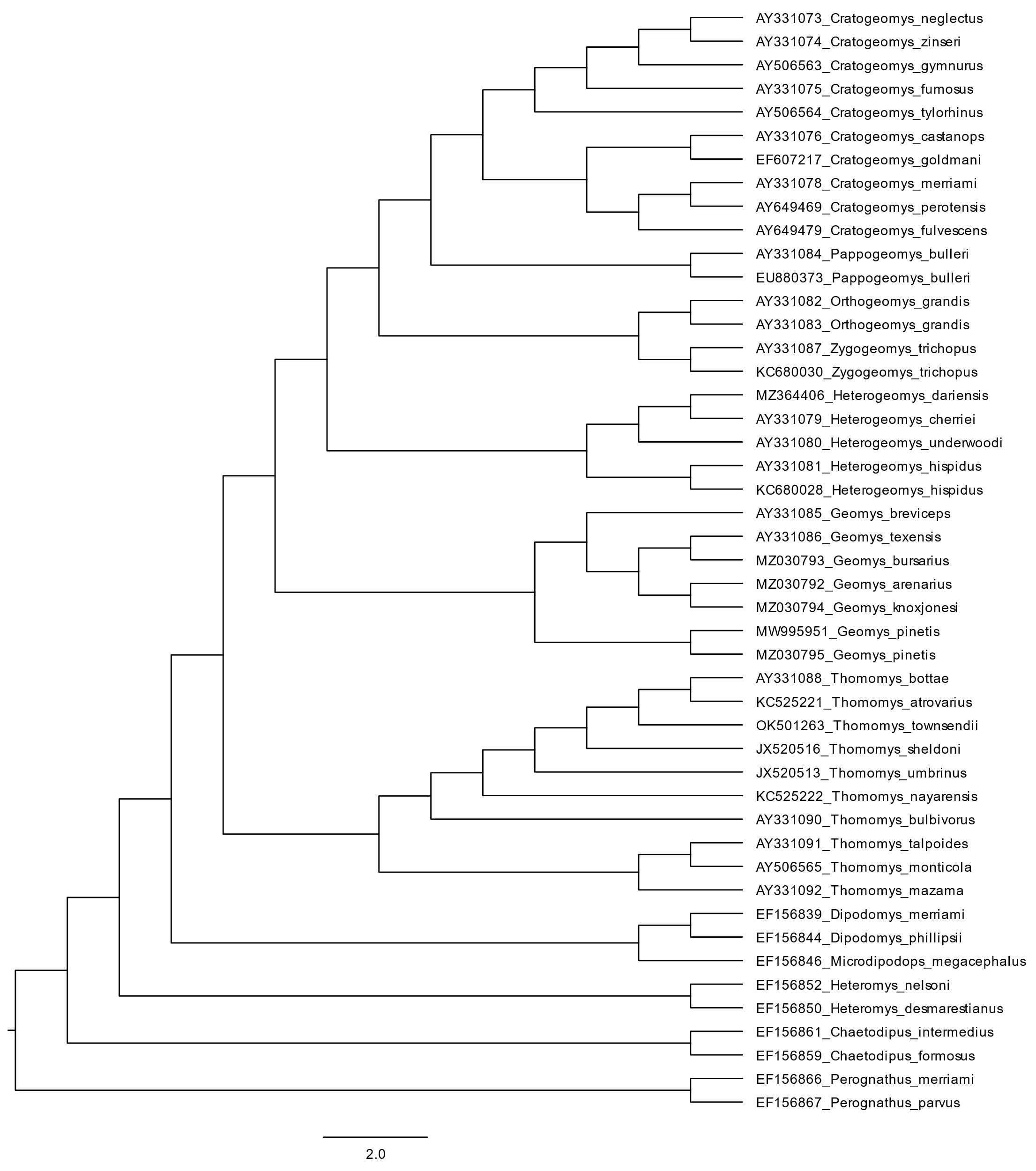
Cytochrome b Bayesian maxima probability.



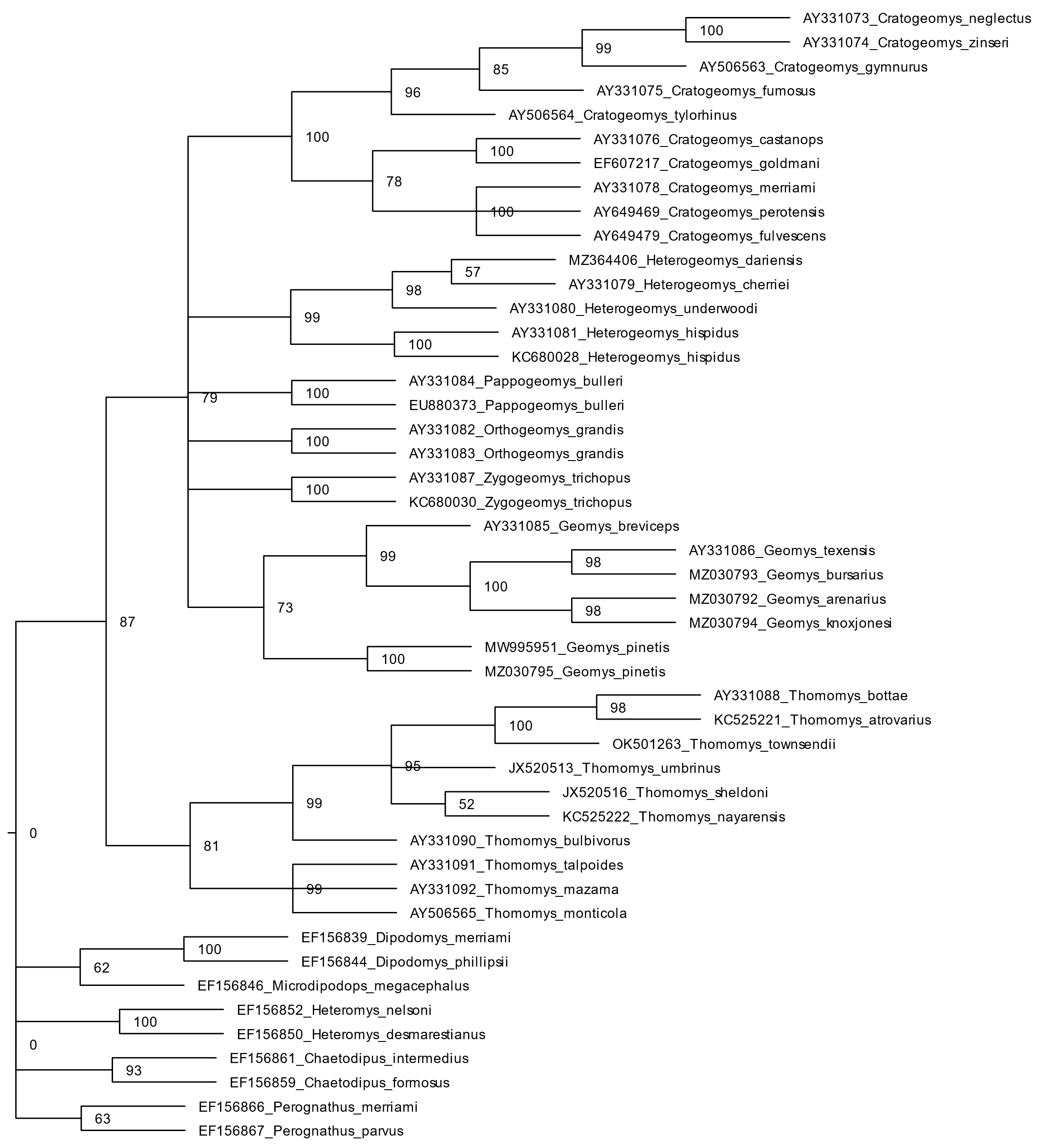
Cytochrome oxidase subunit 1 neighbor-joining bootstrap consensus tree generating 100 pseudoreplicates.



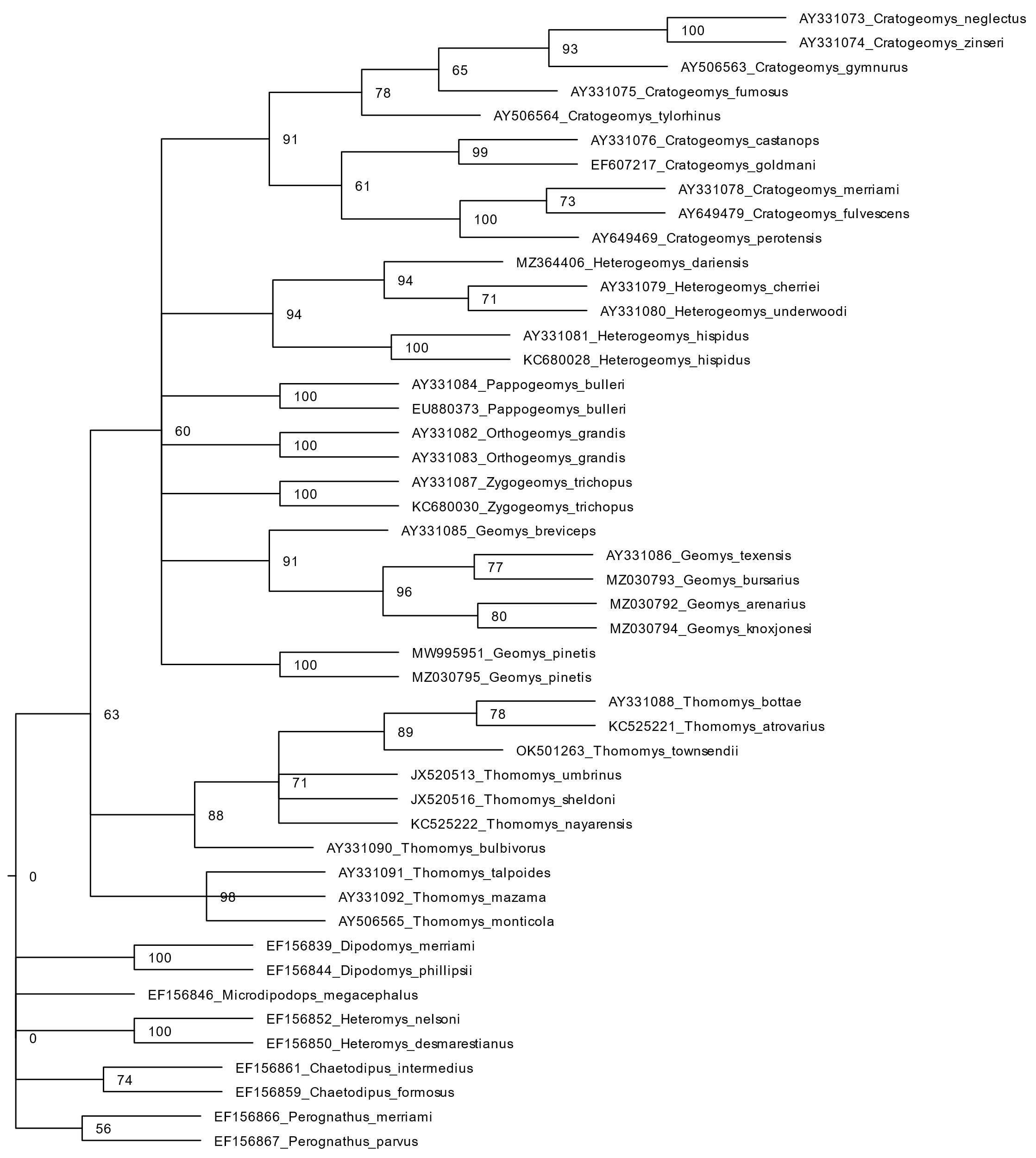
Cytochrome oxidase subunit 1 unweighted pair group method with arithmetic mean.



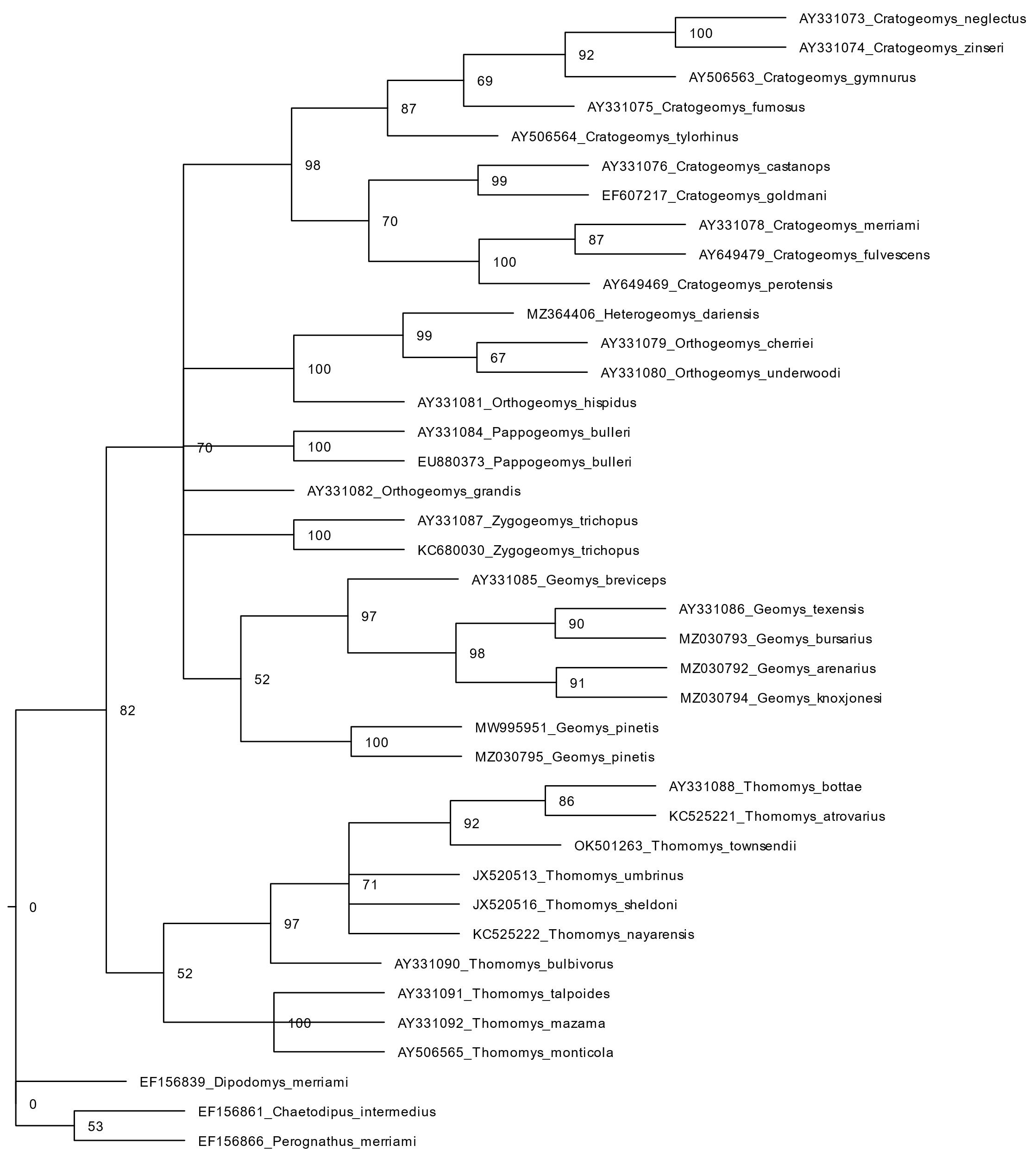
Cytochrome oxidase subunit 1 unweighted pair group method with arithmetic mean bootstrap consensus tree generating 100 pseudoreplicates.



Cytochrome oxidase subunit 1 maximum-parsimony.



Cytochrome oxidase subunit 1 maximum Likelihood method.



Cytochrome oxidase subunit 1 Bayesian maxima probability.

