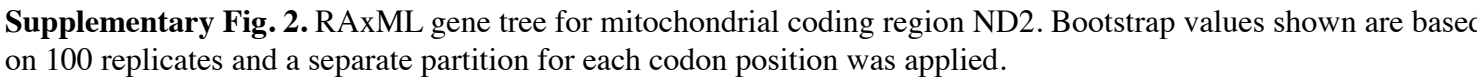
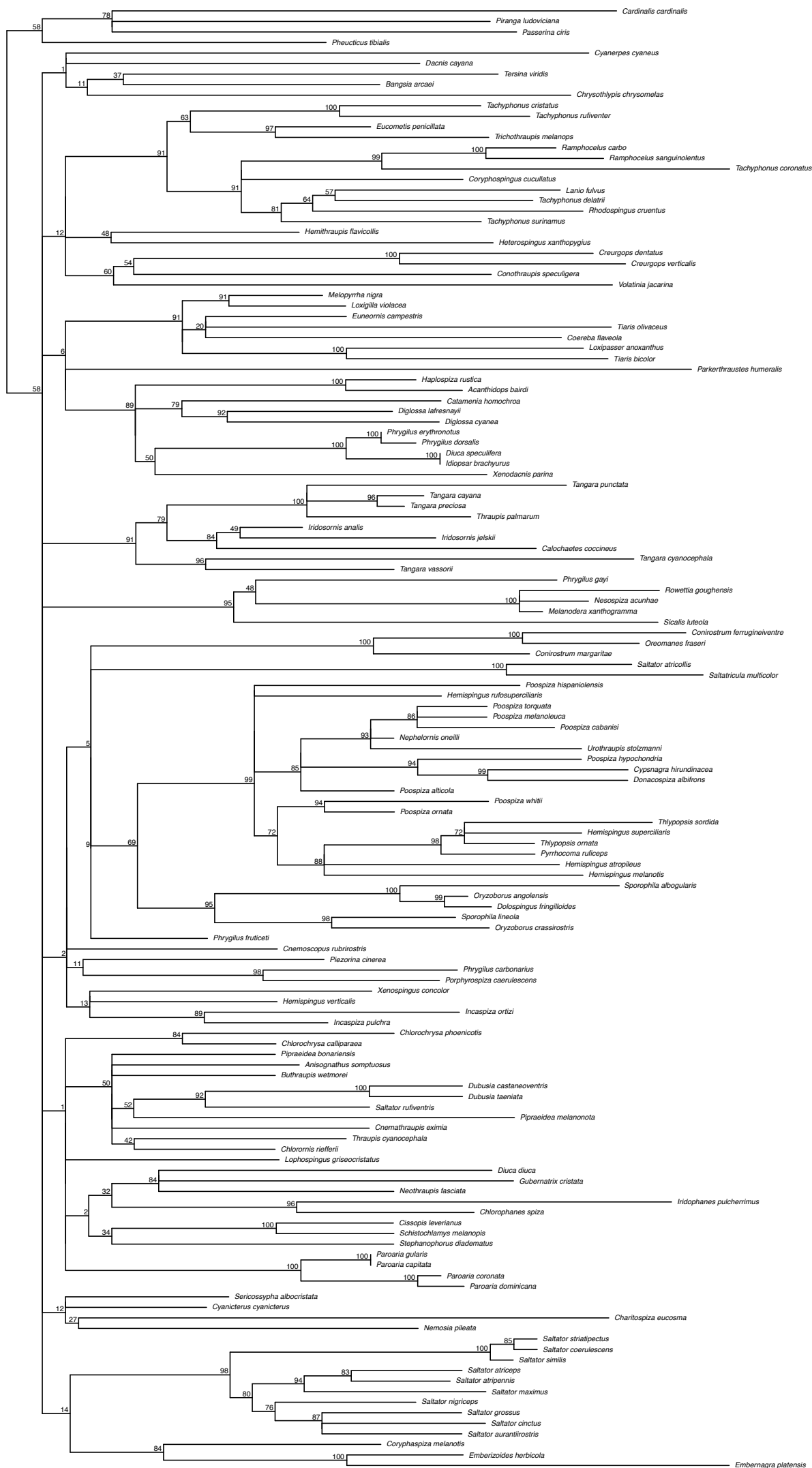




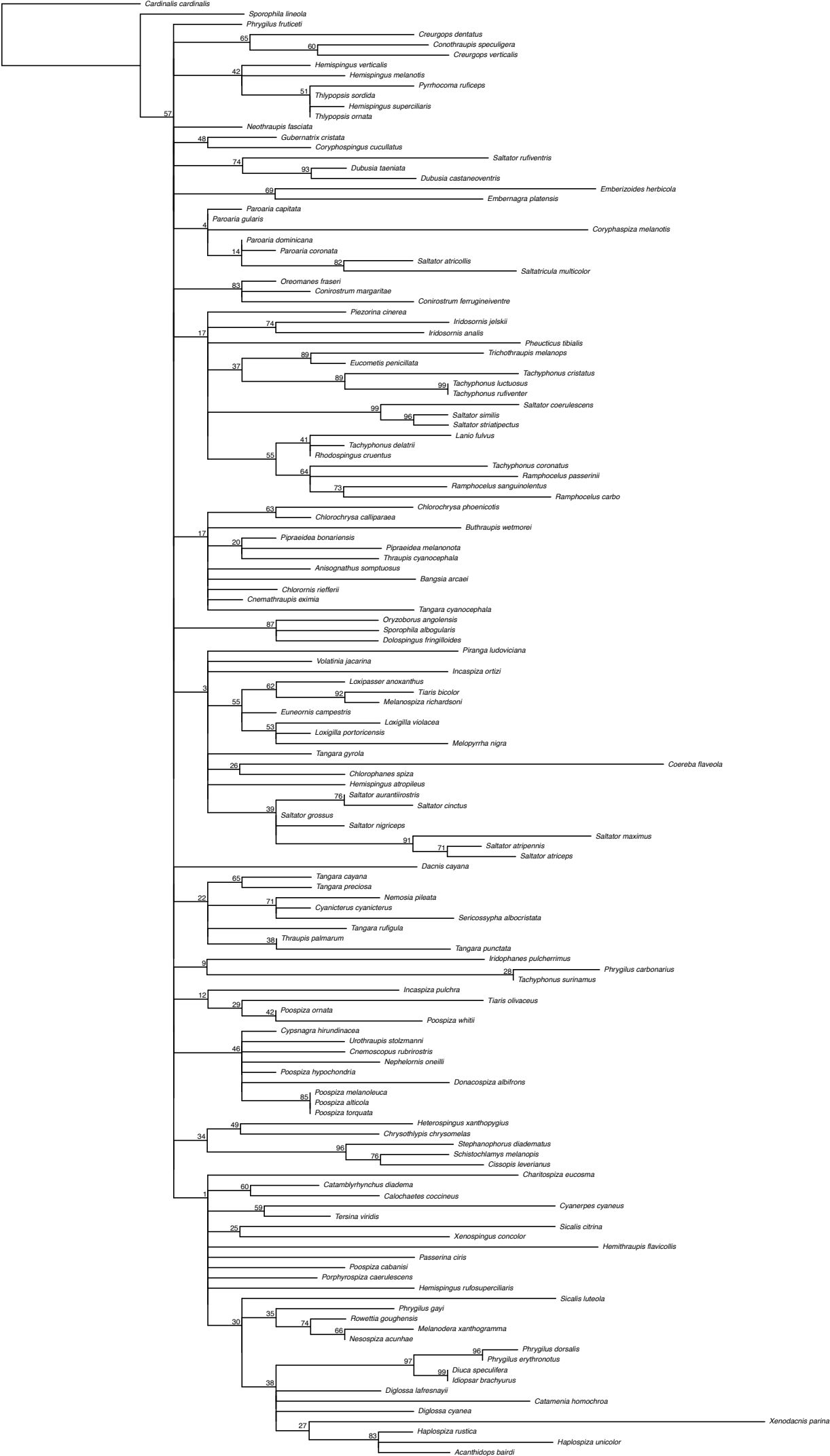
Supplementary Fig. 1. RAxML gene tree for mitochondrial coding region cyt *b*. Bootstrap values shown are based on 100 replicates and a separate partition for each codon position was applied.



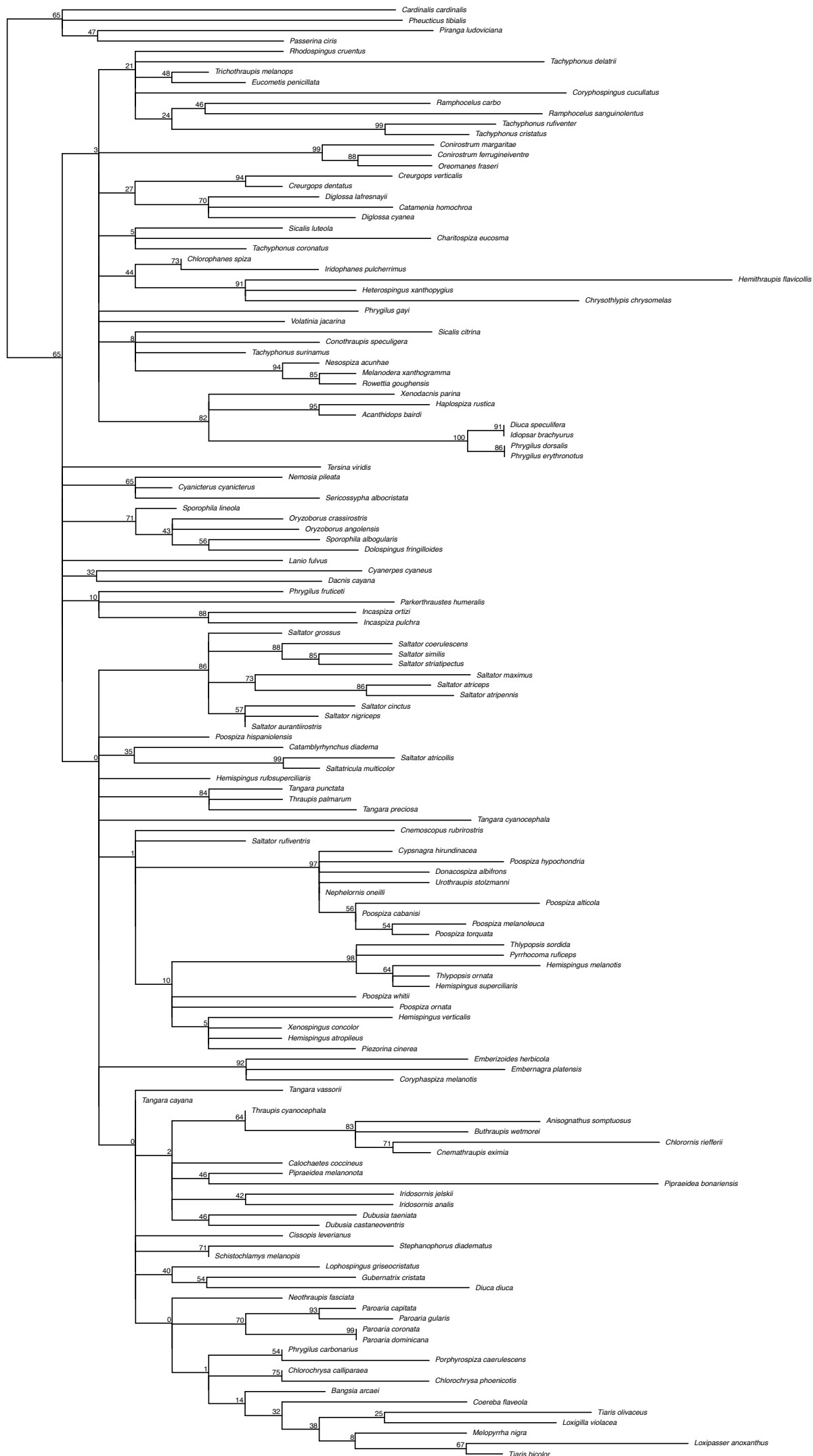
Supplementary Fig. 2. RAxML gene tree for mitochondrial coding region ND2. Bootstrap values shown are based on 100 replicates and a separate partition for each codon position was applied.



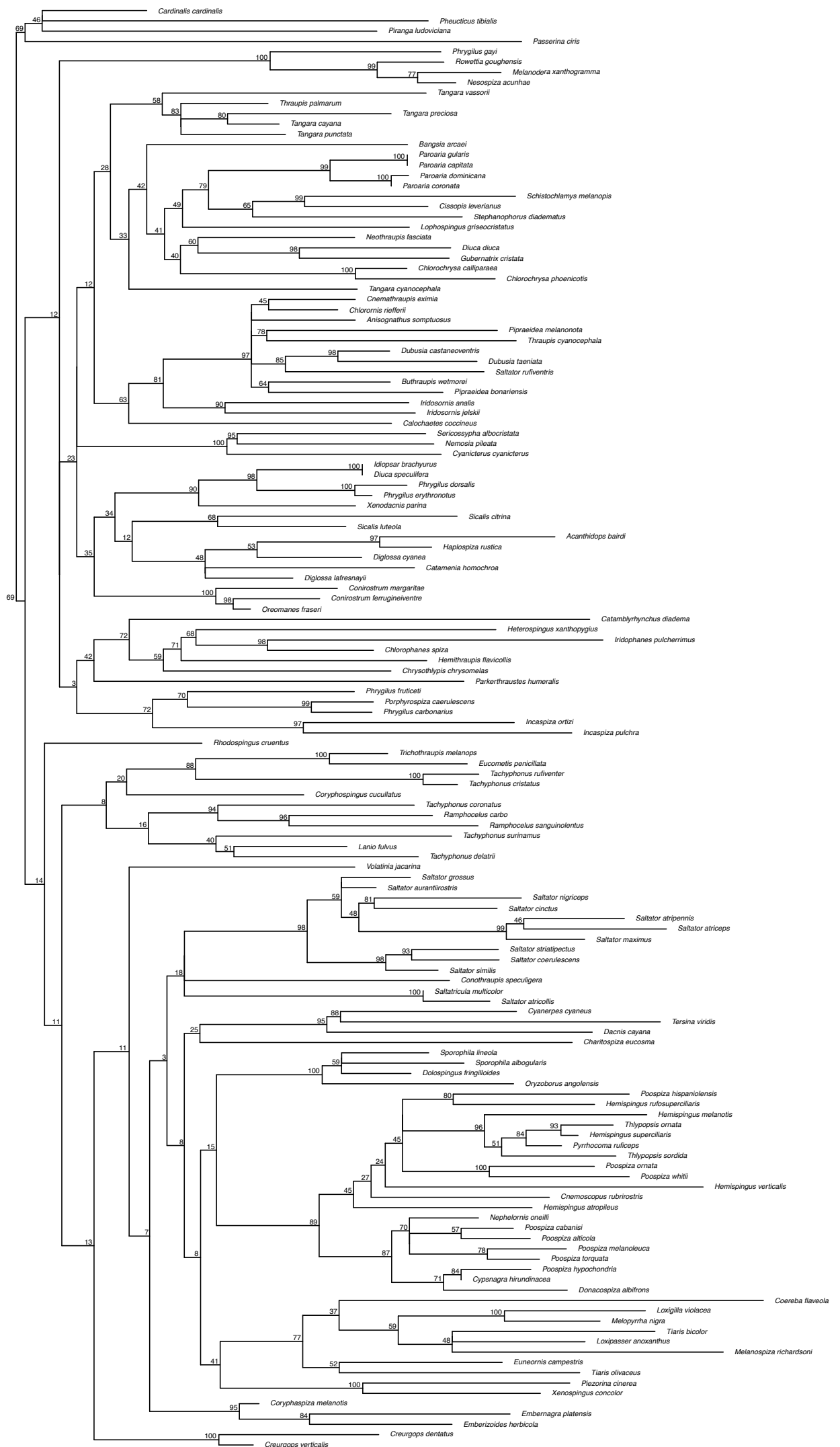
Supplementary Fig. 3. RAxML gene tree for nuclear intron ACO1-I9. Bootstrap values shown are based on 100 replicates and a single partition was applied.



Supplementary Fig. 4. RAxML gene tree for nuclear intron FGB-I5. Bootstrap values shown are based on 100 replicates and a single partition was applied.



Supplementary Fig. 5. RAxML gene tree for nuclear intron MB-I2. Bootstrap values shown are based on 100 replicates and a single partition was applied.



Supplementary Fig. 6. RAXML gene tree for mitochondrial nuclear exon RAG1. Bootstrap values shown are based on 100 replicates and a separate partition for each codon position was applied.