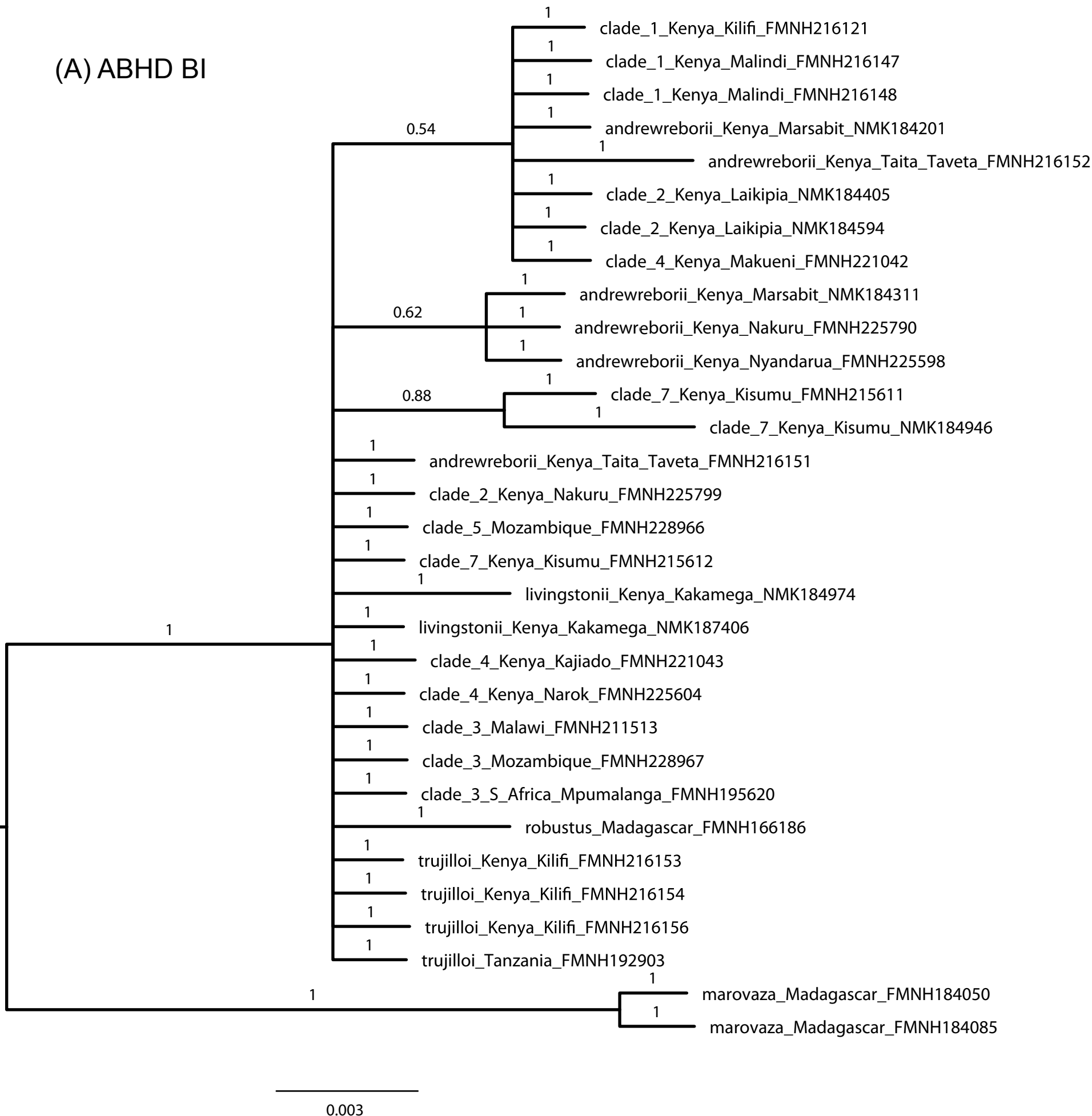
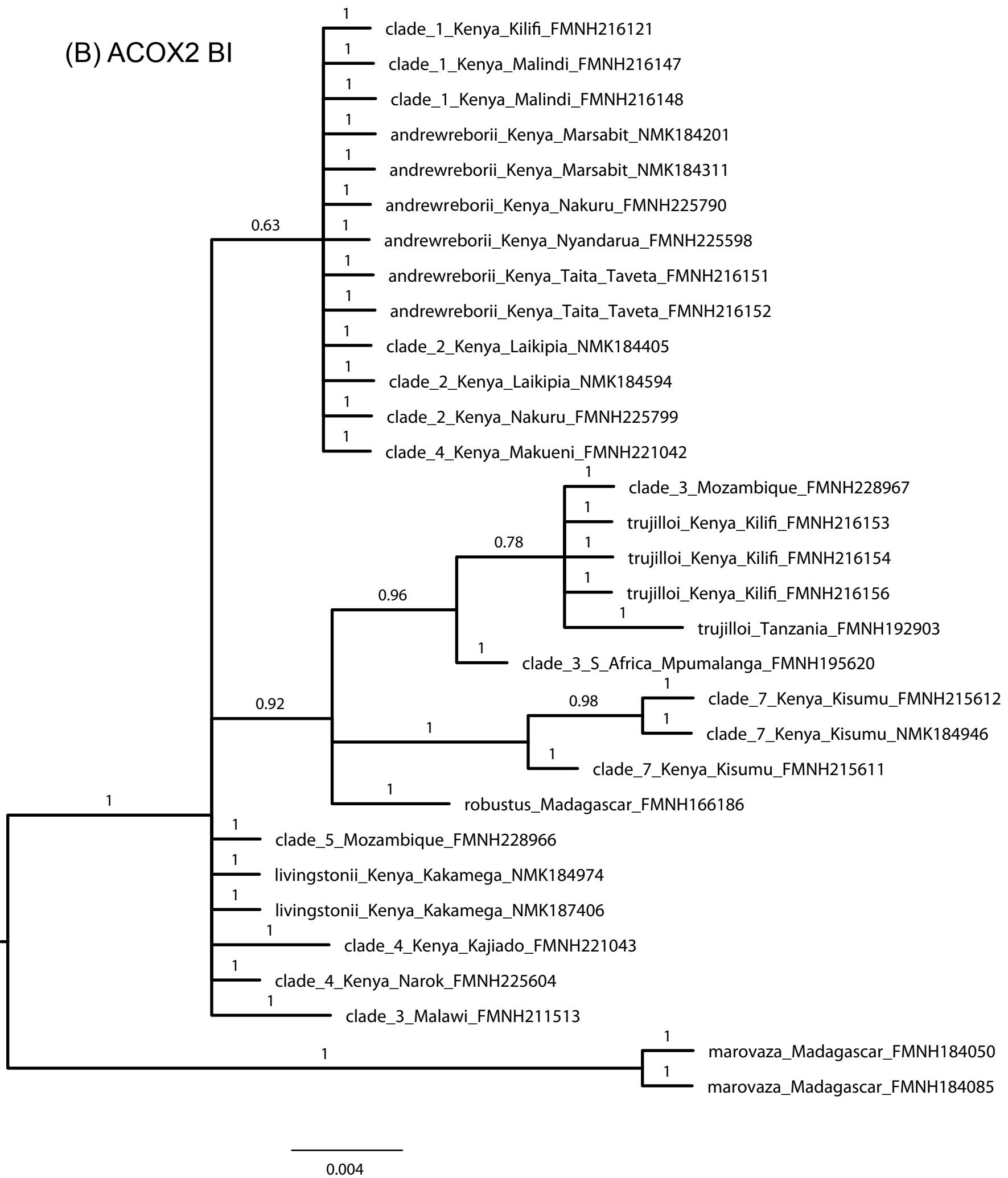


Figure 1. Supplemental Material. Bayesian gene trees inferred for nuclear introns using MrBayes (A–F), and maximum likelihood gene trees inferred for nuclear introns using RAxML (G–L). Node support is indicated above branches.

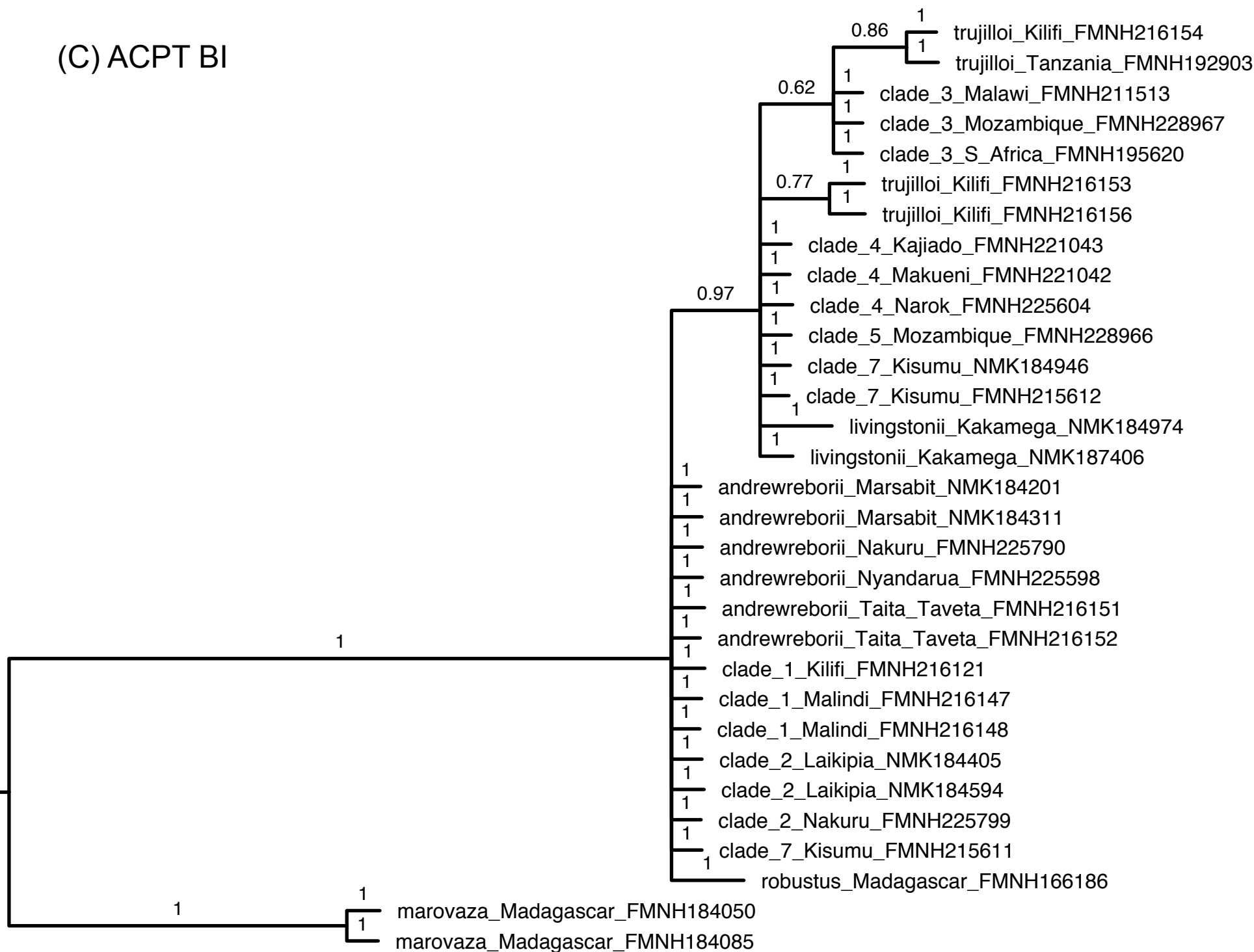
(A) ABHD BI



(B) ACOX2 BI

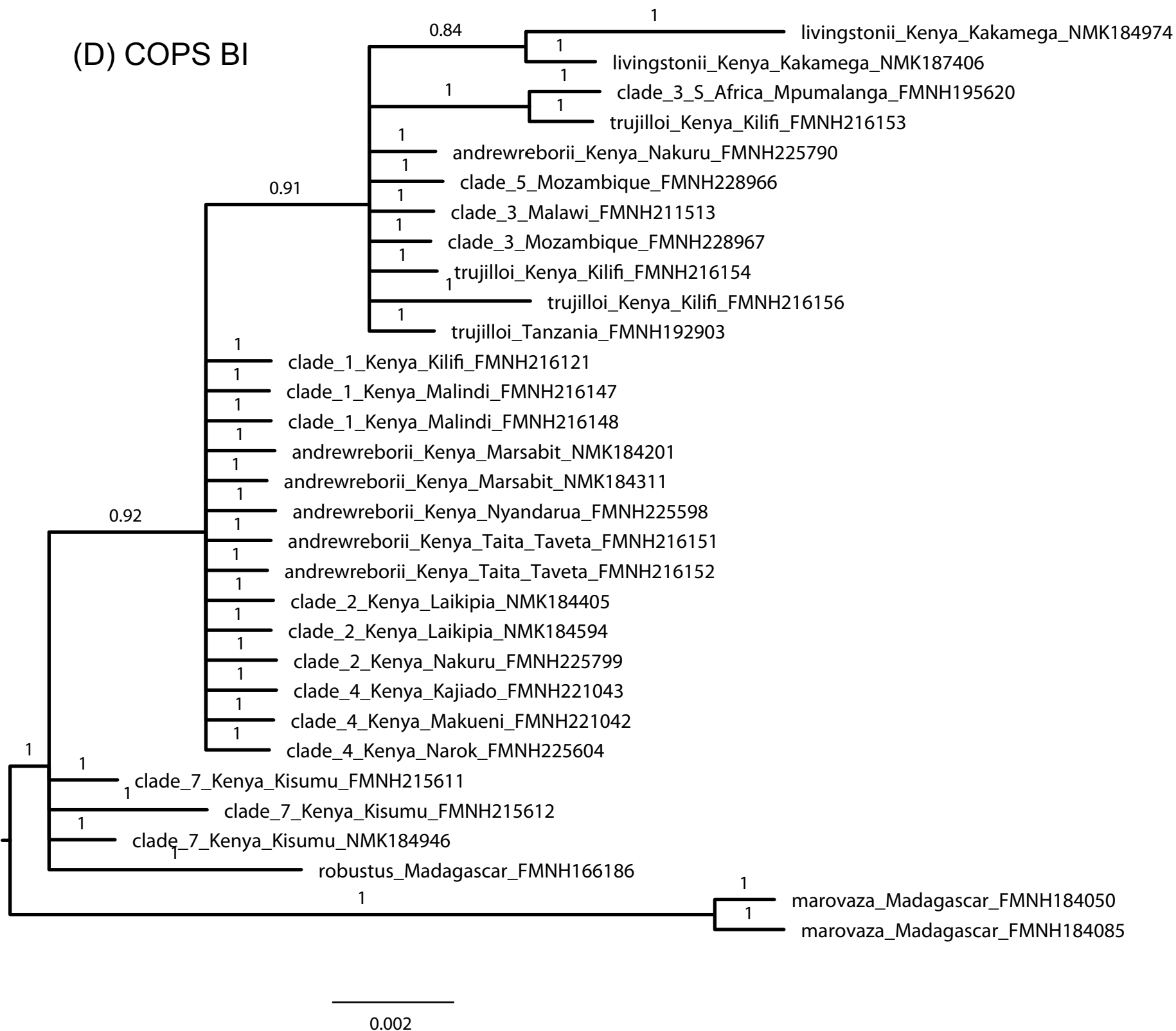


(C) ACPT BI

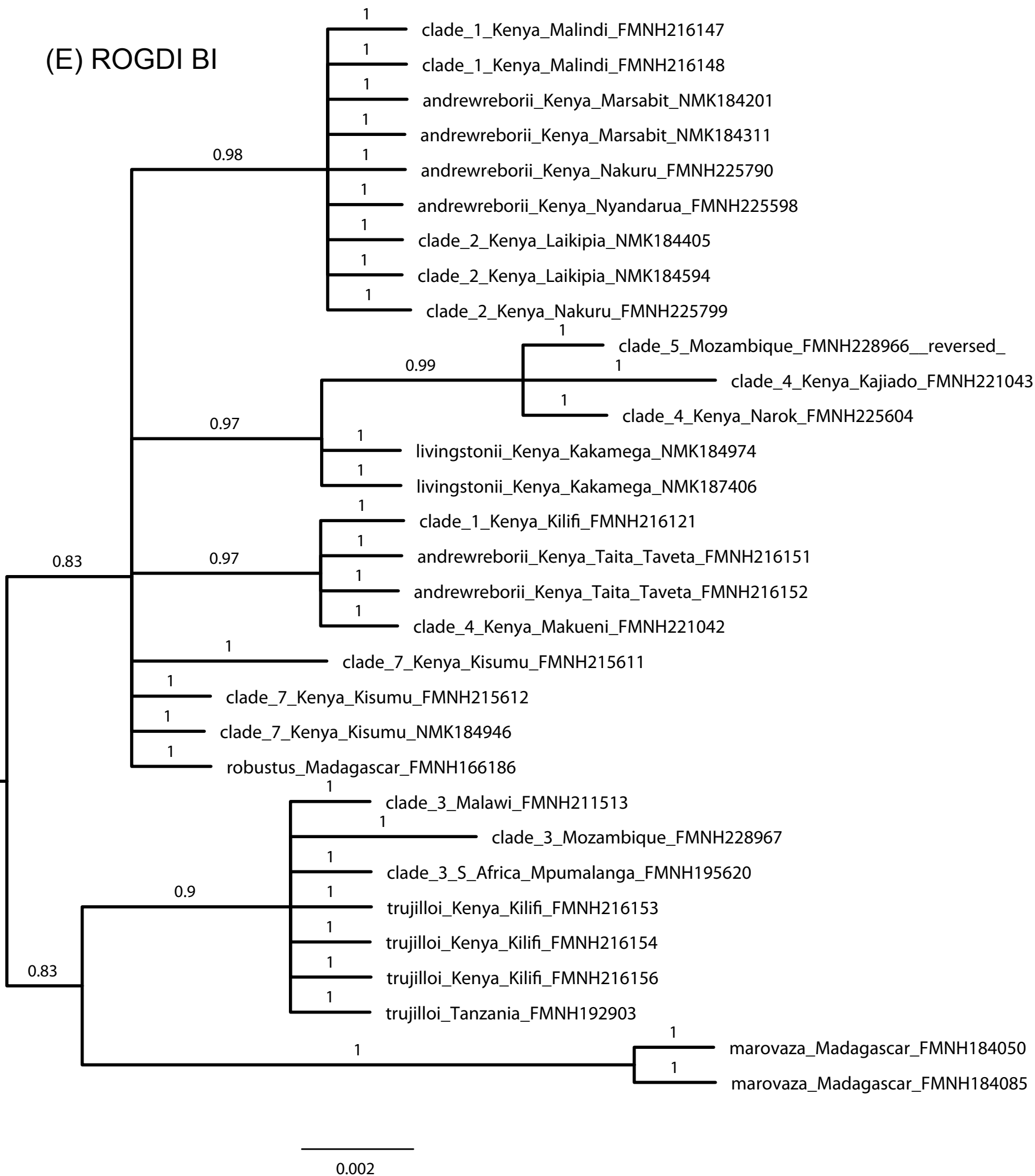


0.005

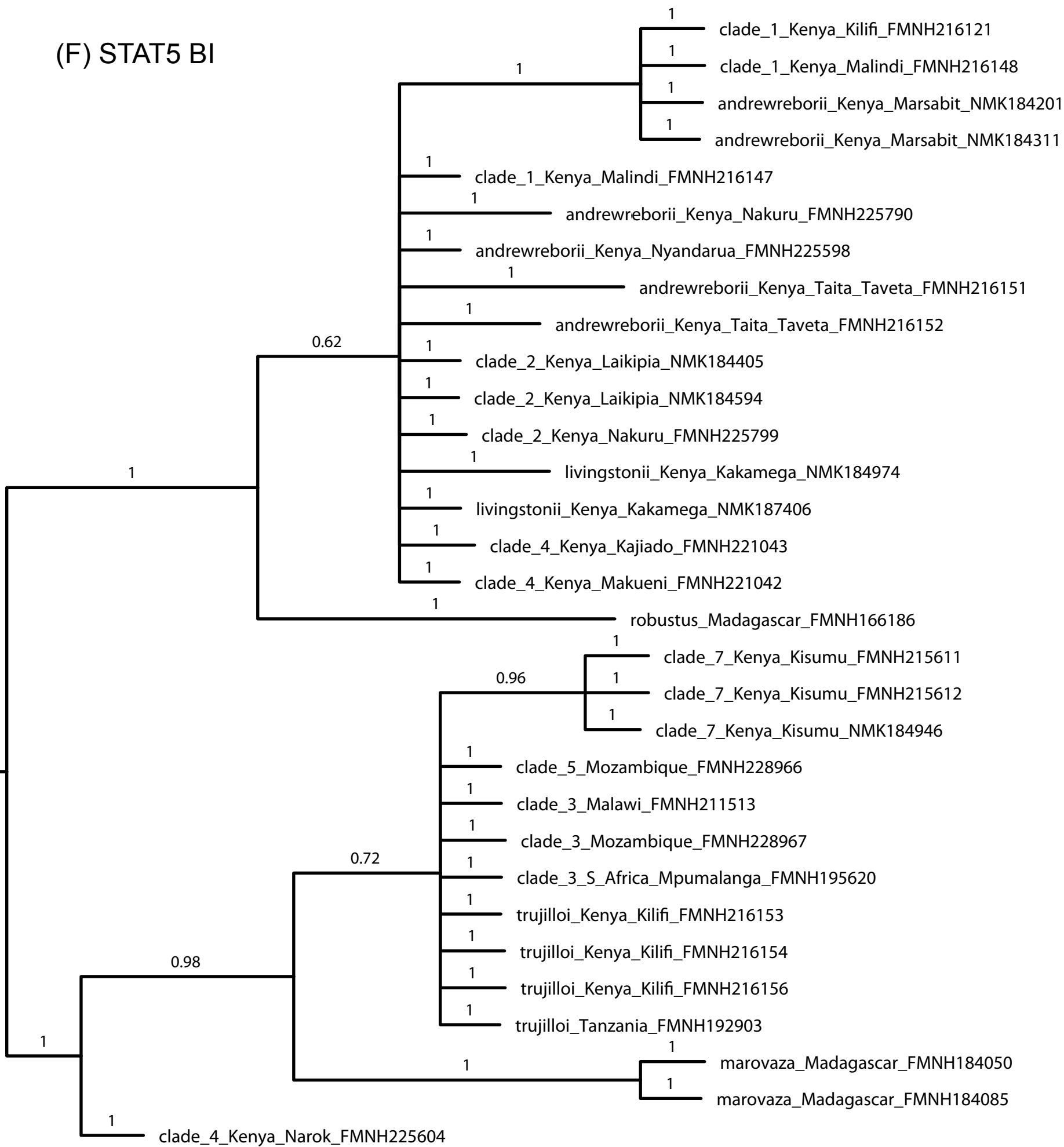
(D) COPS BI



(E) ROGDI BI

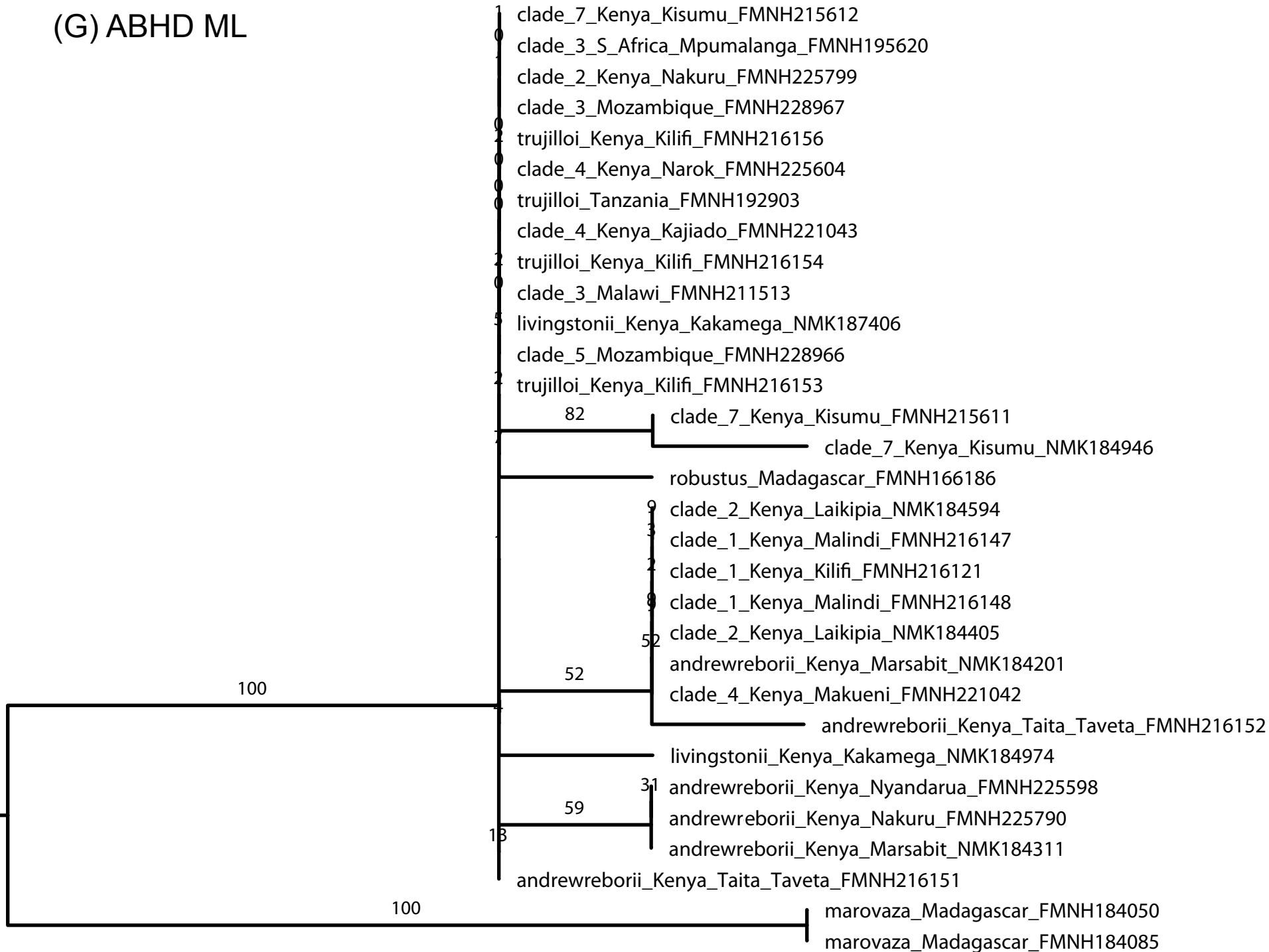


(F) STAT5 BI



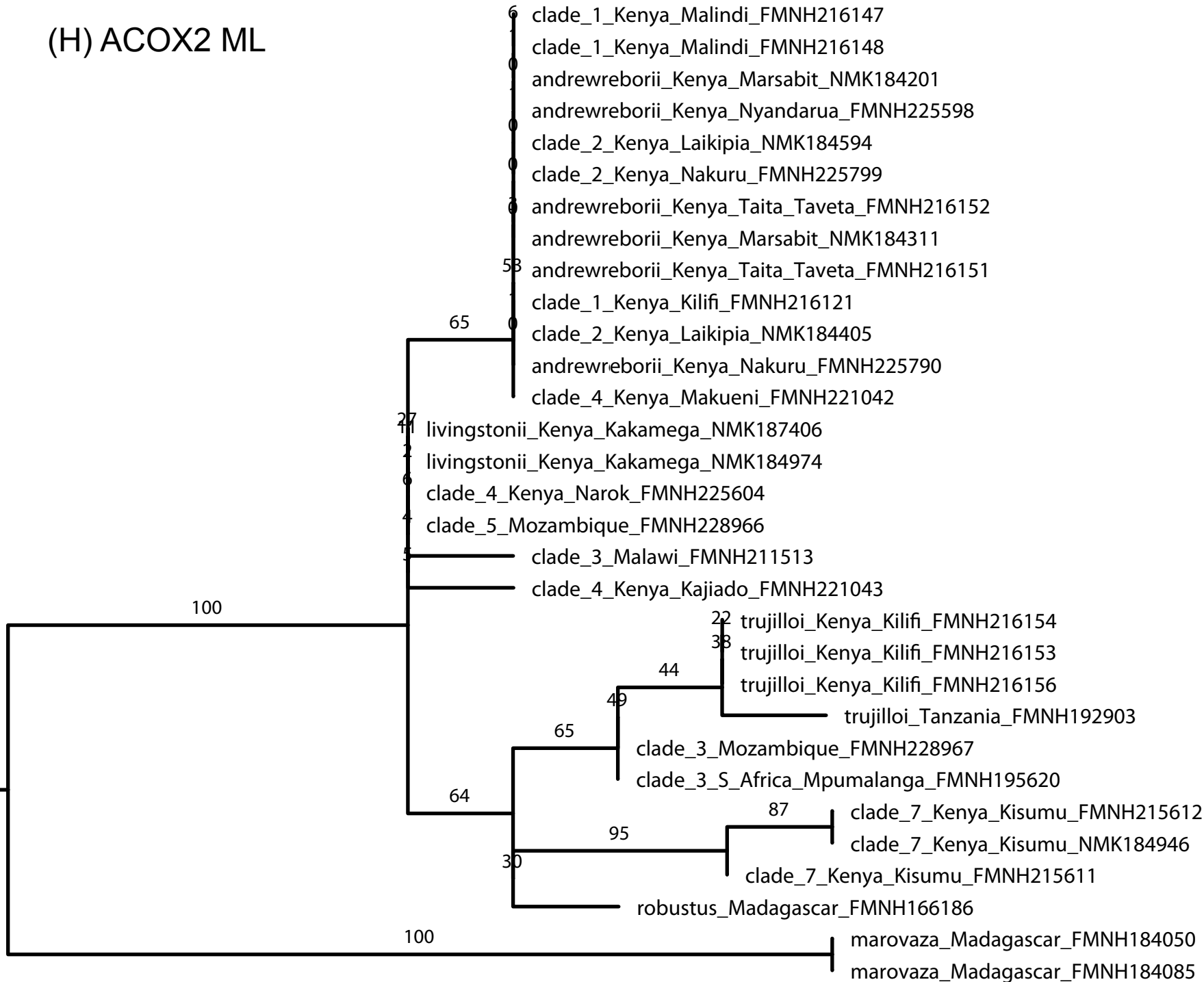
0.004

(G) ABHD ML



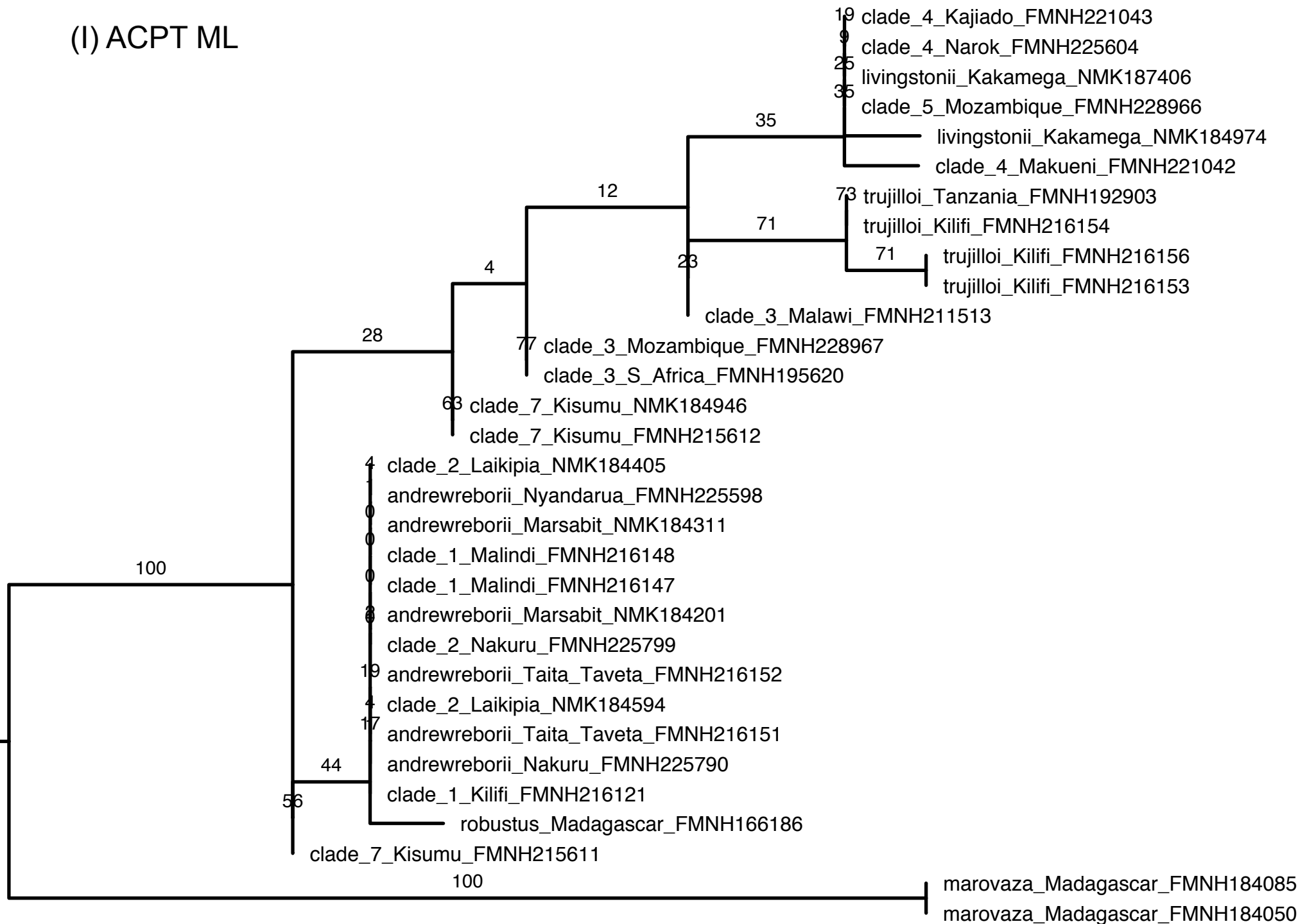
0.003

(H) ACOX2 ML



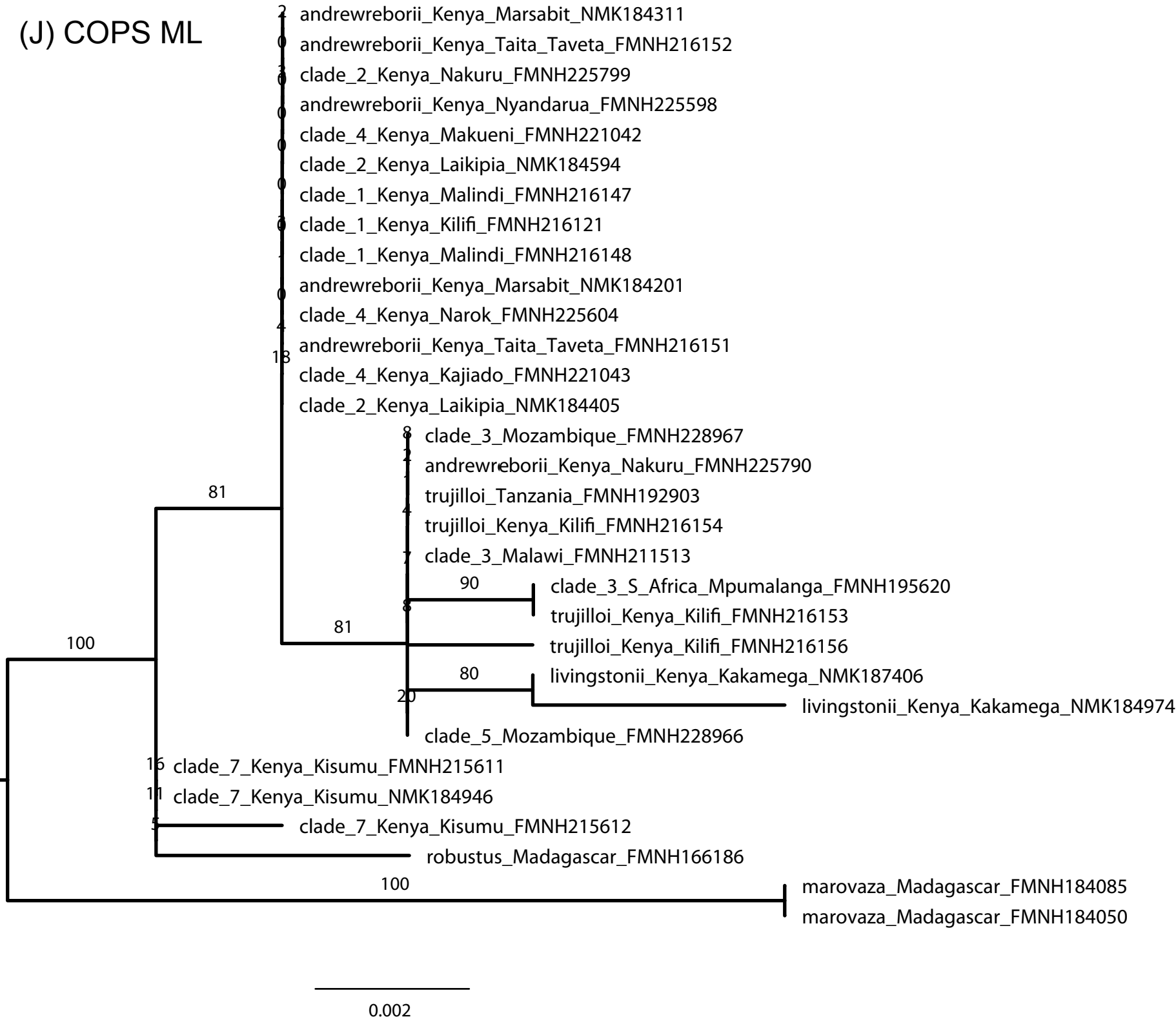
0.004

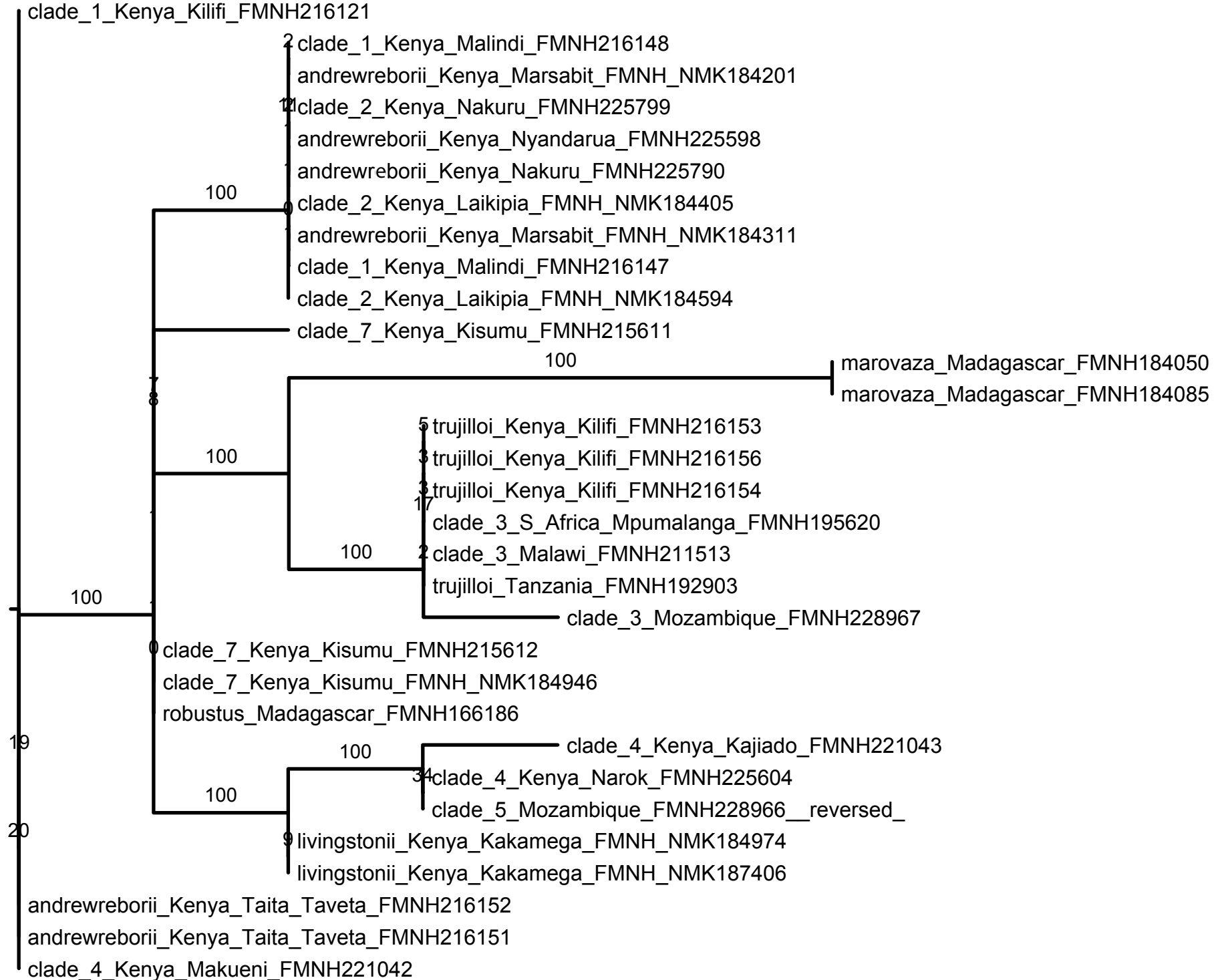
(I) ACPT ML



0.009

(J) COPS ML





K) ROGDI ML

0.002

(L) STAT5 ML

