Description of Additional Supplementary Files

File name: Supplementary Data 1

Description: Voay mtDNA alignment nexus file

File name: Supplementary Data 2

Description: Voay RAXML BEAST trees text file

File name: Supplementary Data 3

Description: RAxML maximum likelihood analyses, PAUP* parsimony analyses, and BEAST Bayesian tip-dating analyses used to test the stability of phylogenetic results. Trees with support scores can be found in Supplementary Data 2.