



SUPPLEMENTAL FIG. 1A. Eighty percent majority rule consensus of the Bayesian analysis of the ITS sequences from Fig. 1. Numbers above branches indicate Bayesian posterior probabilities (B-PP), bootstrap values (>50% are shown) for maximum likelihood (BS-ML) and maximum parsimony (BS-MP) analyses (B-PP/BS-ML/BS-MP). African species are indicated with an arrow.

