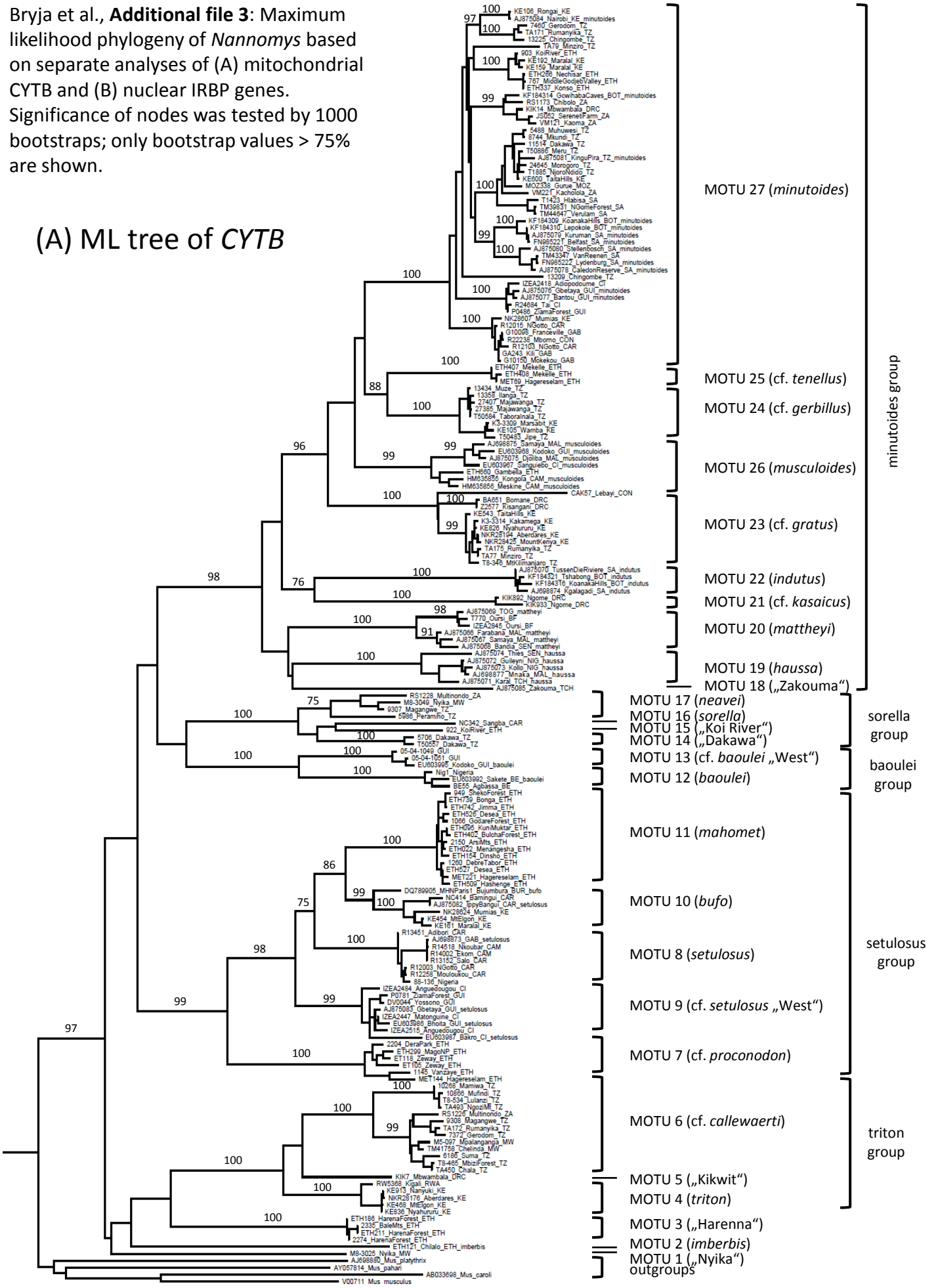
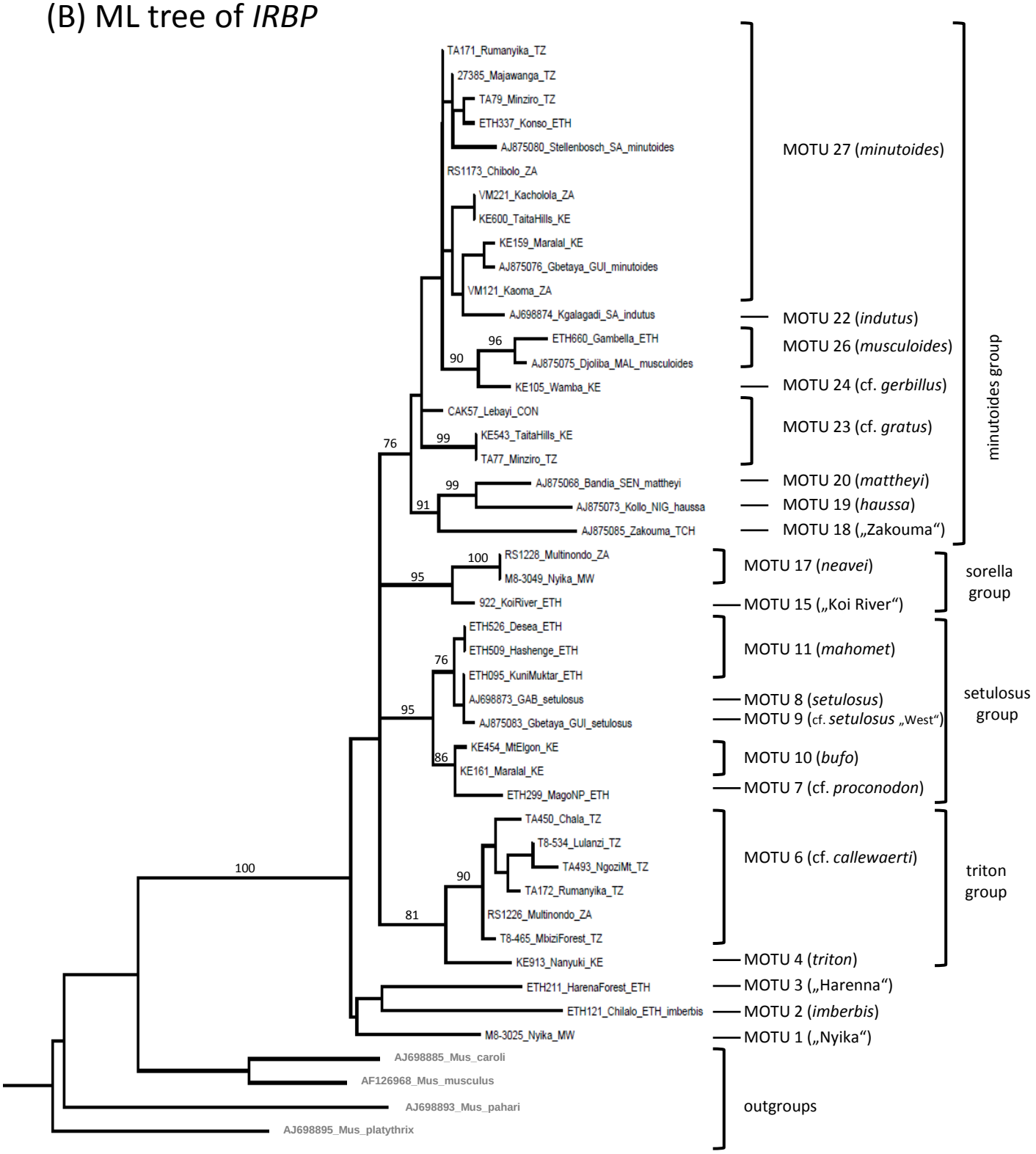
Bryja et al., **Additional file 3**: Maximum likelihood phylogeny of *Nannomys* based on separate analyses of (A) mitochondrial CYTB and (B) nuclear IRBP genes. Significance of nodes was tested by 1000 bootstraps; only bootstrap values > 75% are shown.

(A) ML tree of *CYTB*



(B) ML tree of *IRBP*