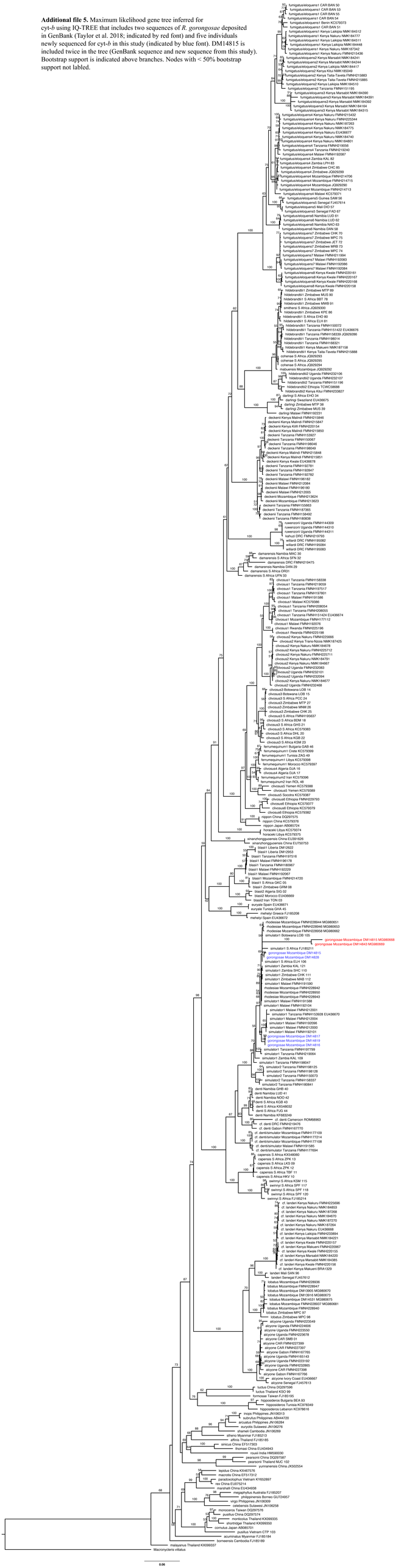Additional file 5. Maximum likelihood gene tree inferred for cyt-*b* using IQ-TREE that includes two sequences of *R. gorongosae* deposited in GenBank (Taylor et al. 2018; indicated by red font) and five individuals newly sequenced for cyt-*b* in this study (indicated by blue font). DM14815 is included twice in the tree (GenBank sequence and new sequence from this study). Bootstrap support is indicated above branches. Nodes with < 50% bootstrap support not labled.