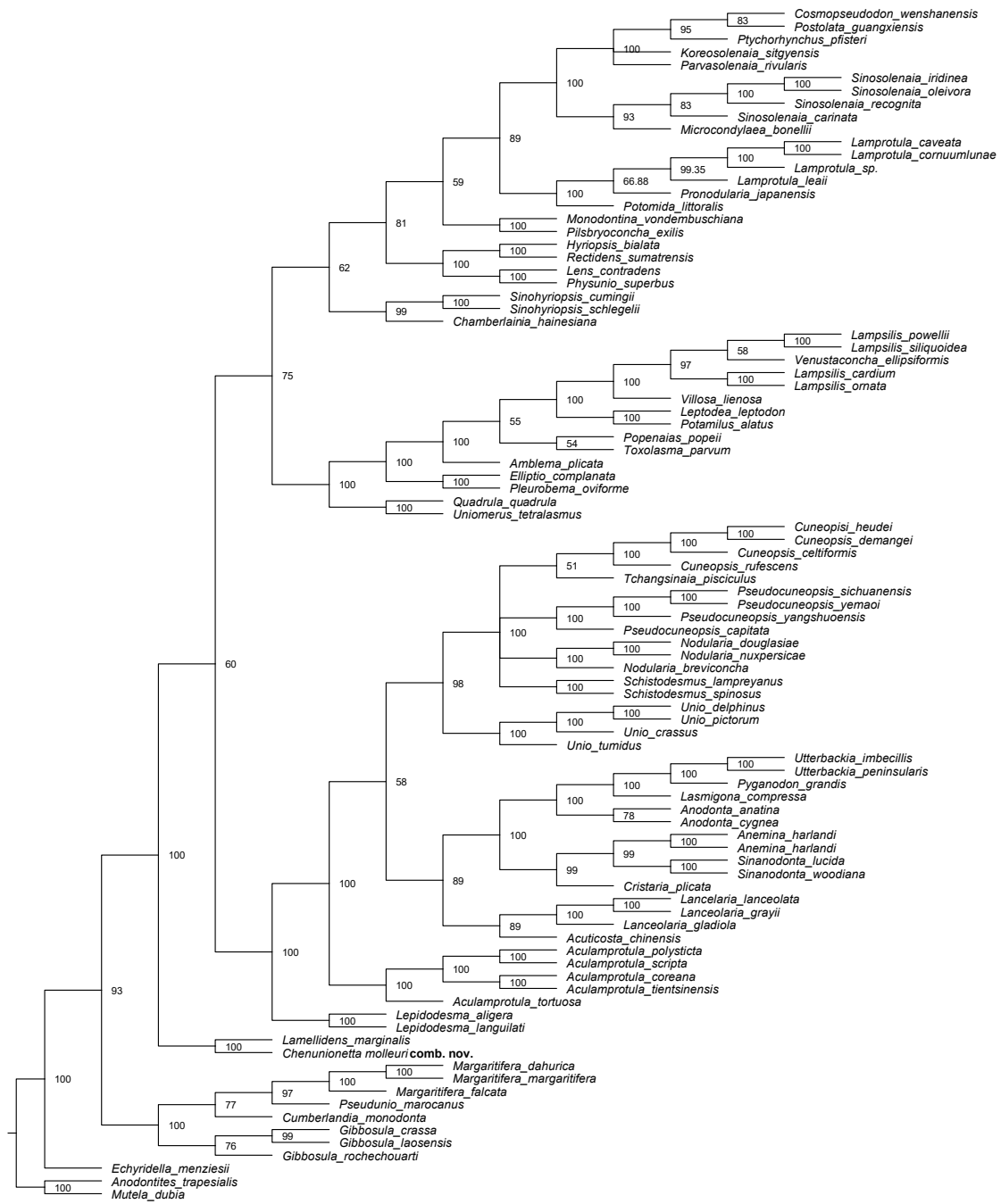




Supplementary Figure S7. Maximum parsimony phylogeny inferred from complete F-type mitogenome sequences. Bootstrap support values are indicated at nodes.