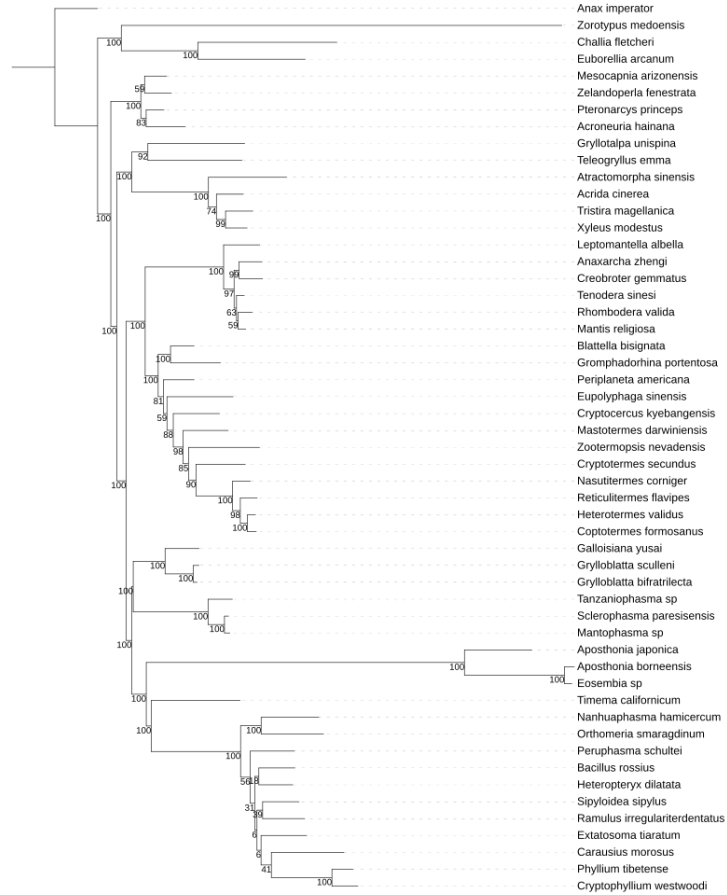
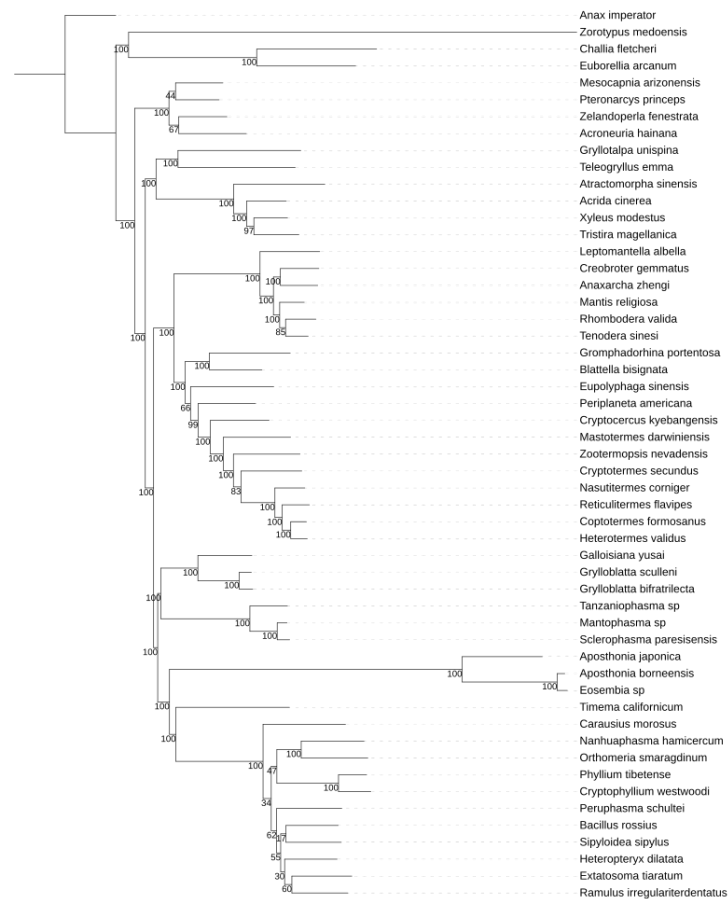


Tree scale: 1

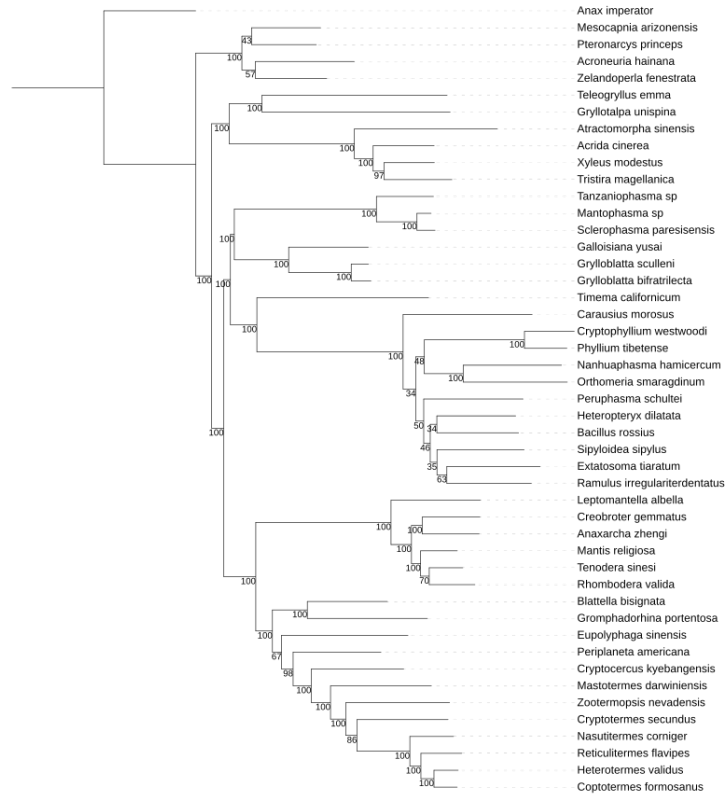


Maximum Likelihood
phylogeny with amino acids
alignment and all species

Tree scale: 1



Tree scale: 1



Maximum Likelihood
phylogeny with nucleotides
alignment and without LBA
groups