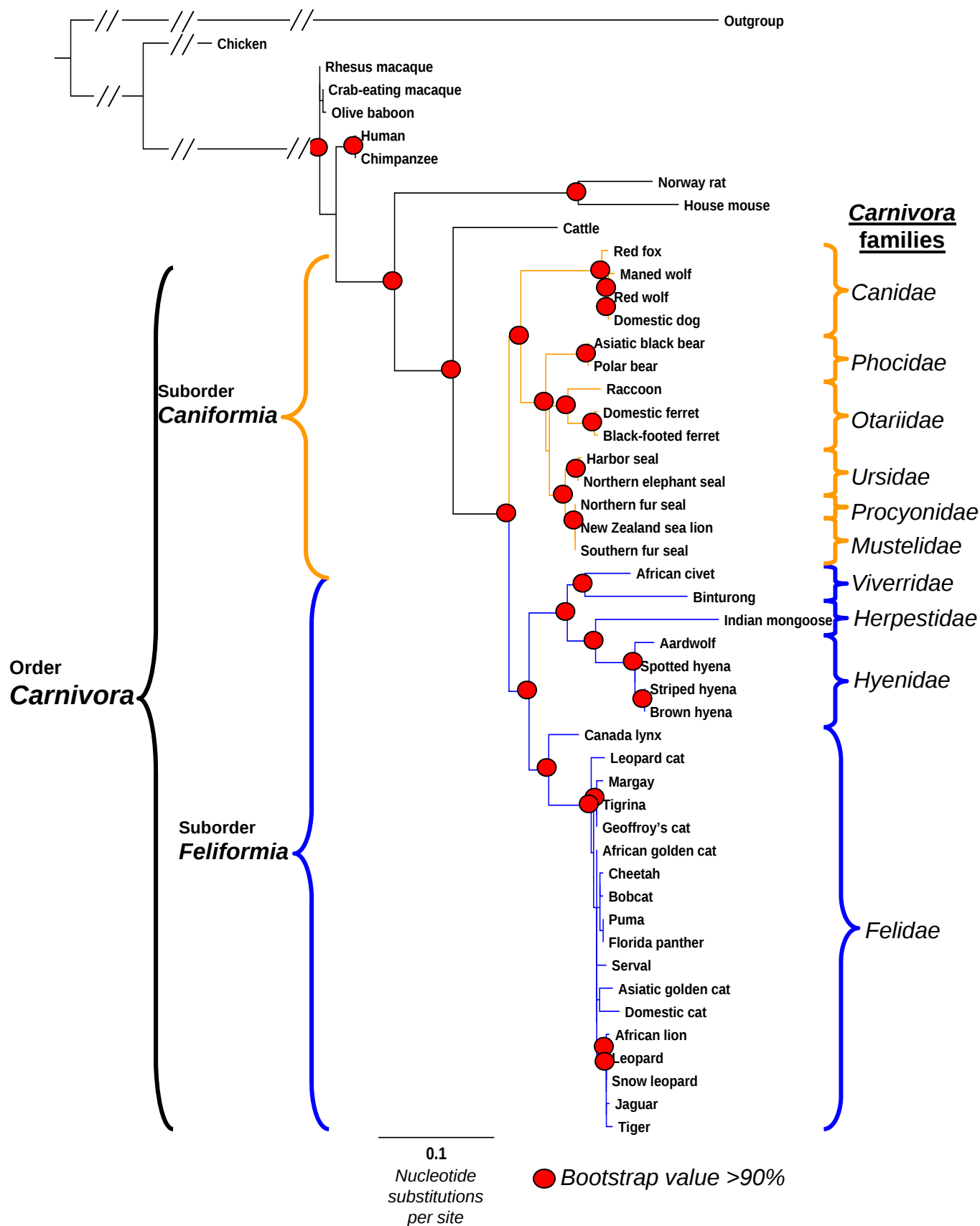


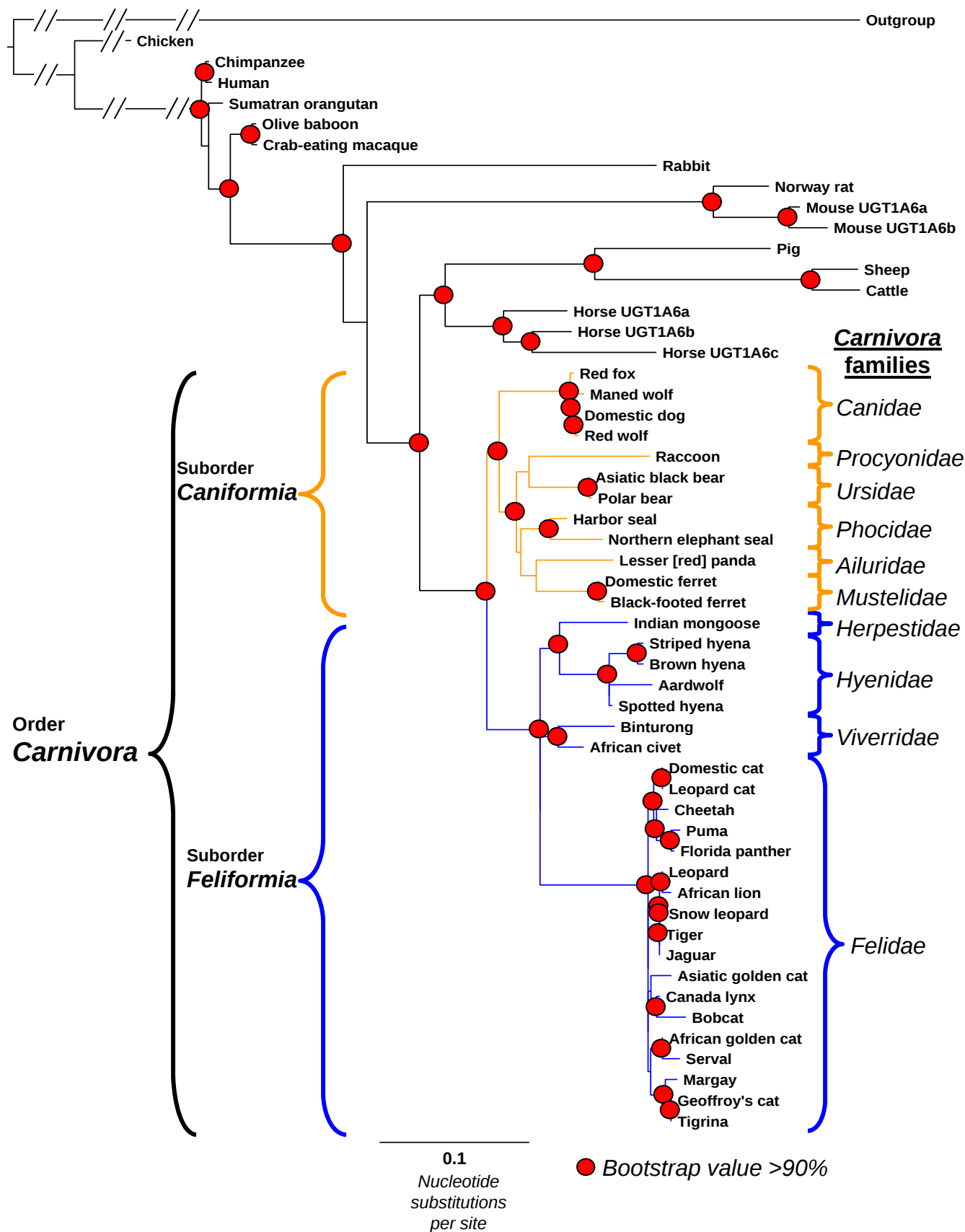
A

UGT1A1 Maximum Likelihood phylogenetic tree



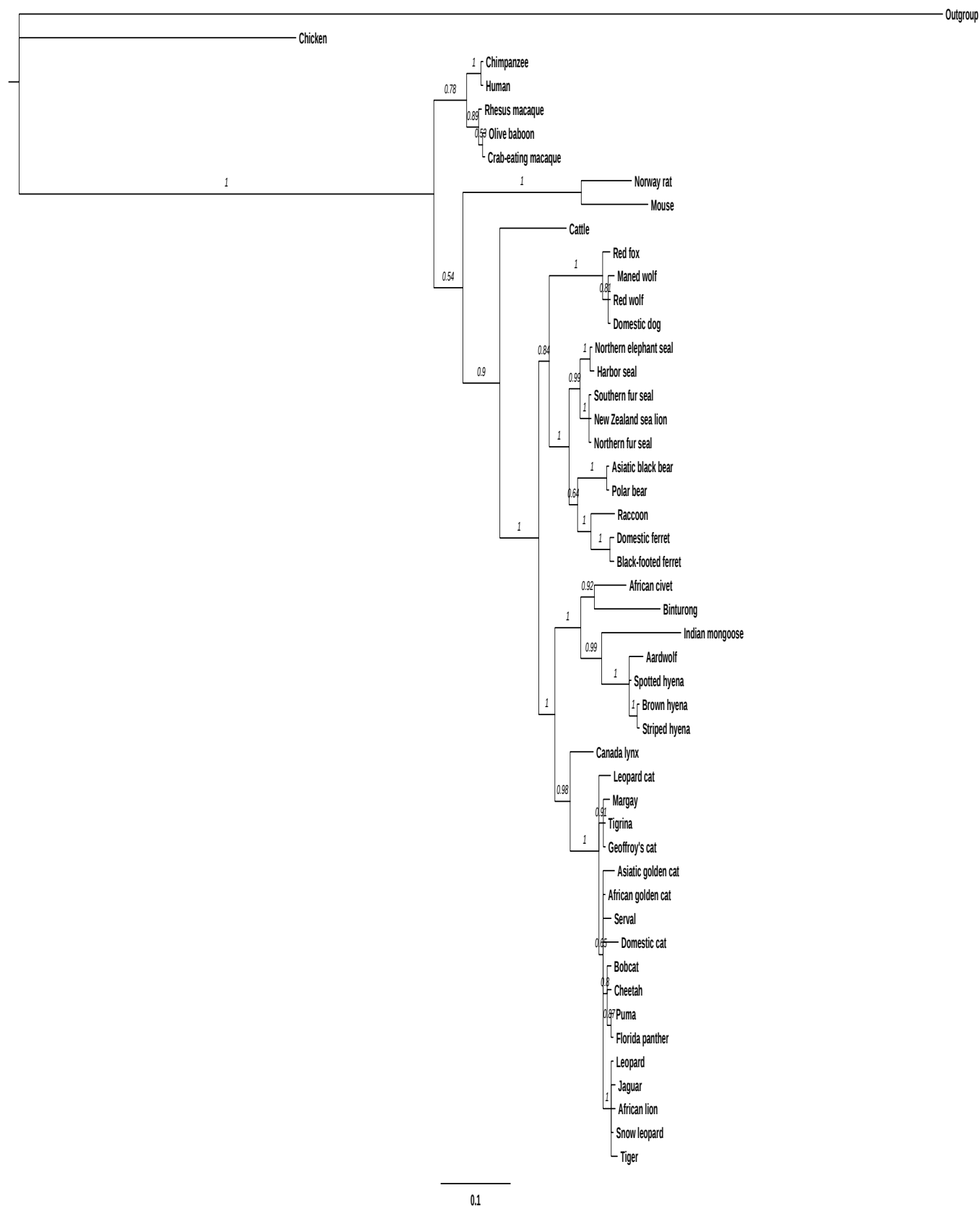
B

UGT1A6 Maximum Likelihood phylogenetic tree



C

UGT1A1 Bayesian phylogenetic tree



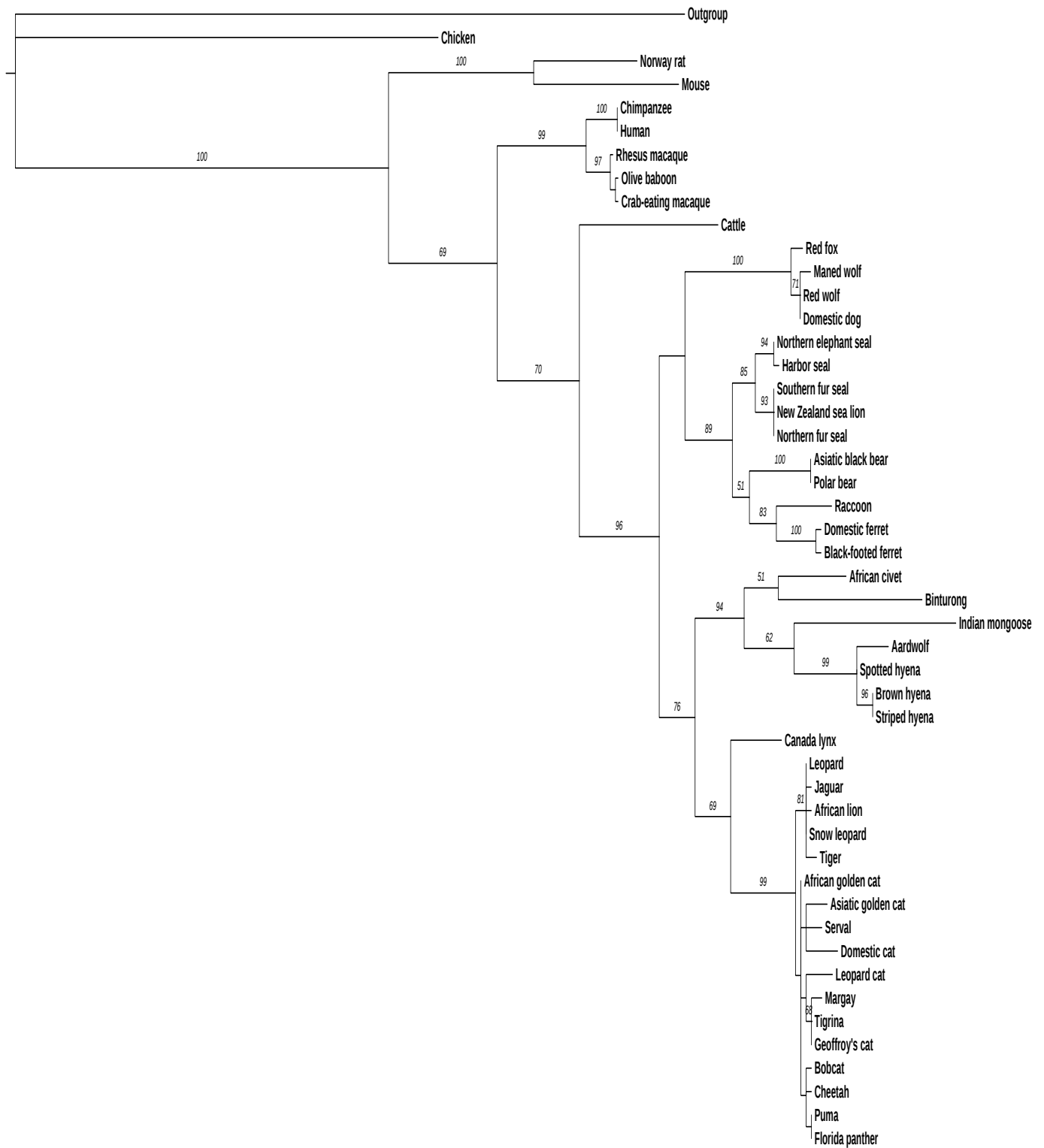
D

UGT1A6 Bayesian phylogenetic tree



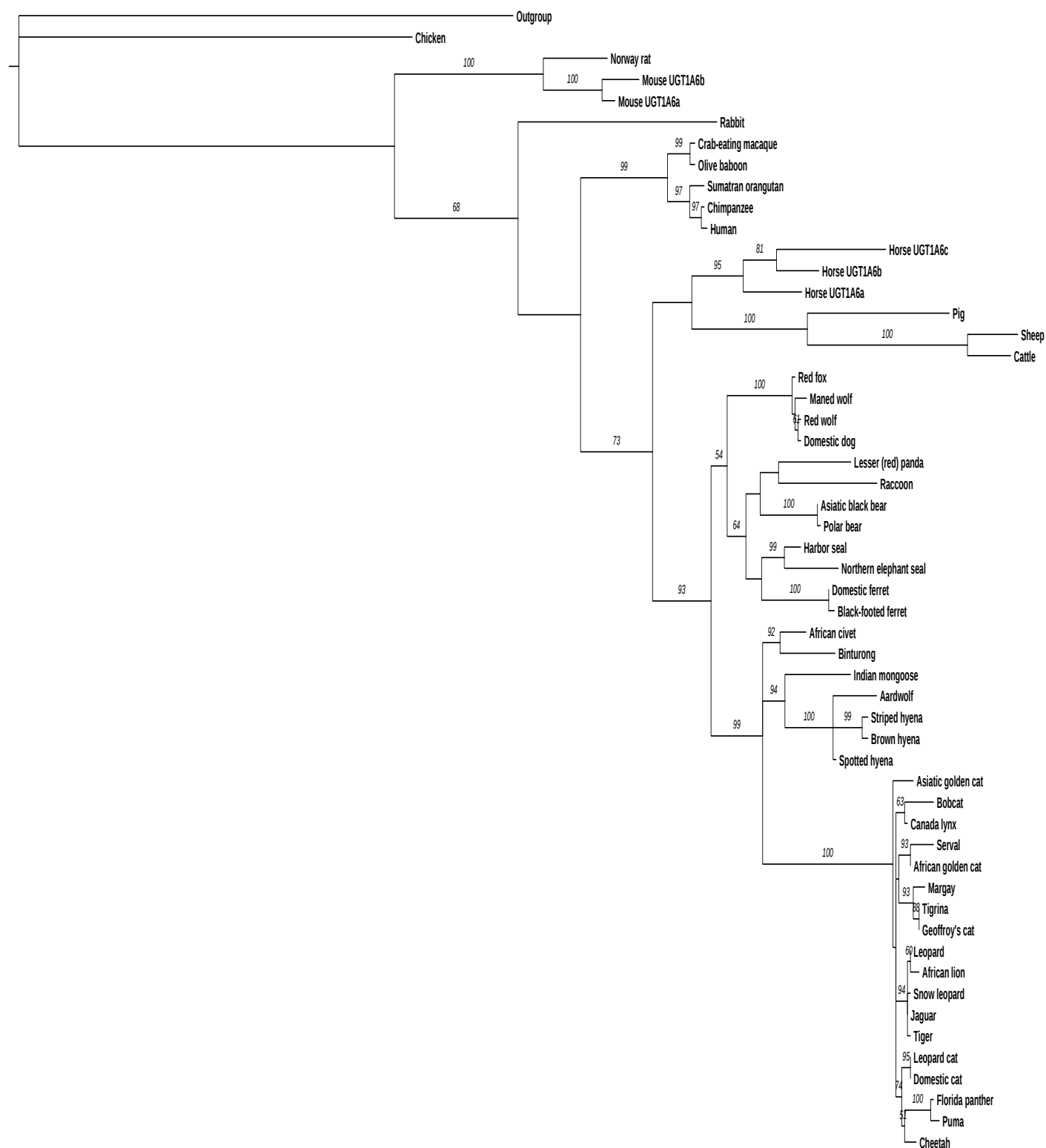
E

UGT1A1 Maximum Parsimony phylogenetic tree



F

UGT1A6 Maximum Parsimony phylogenetic tree



0.05