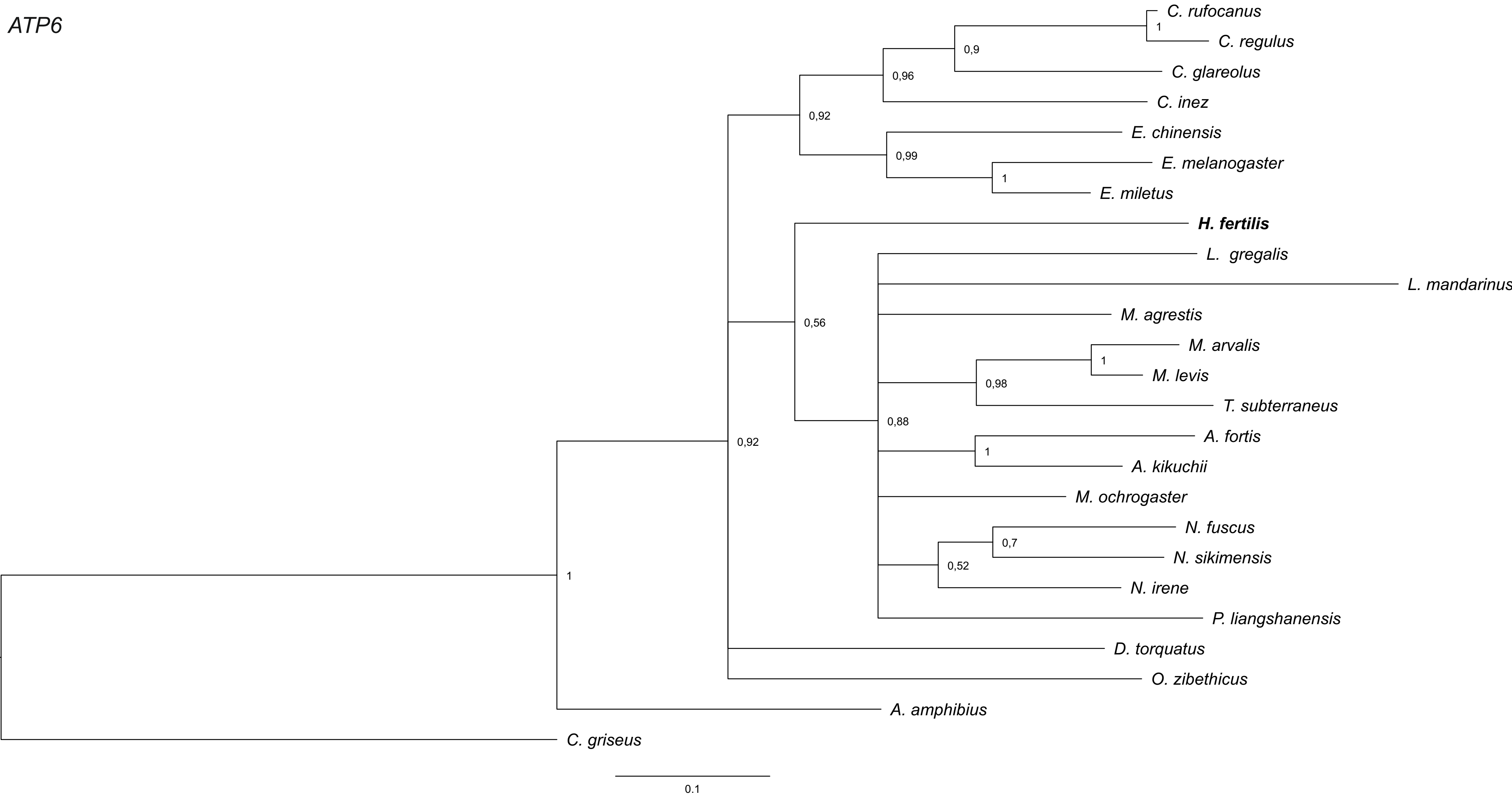
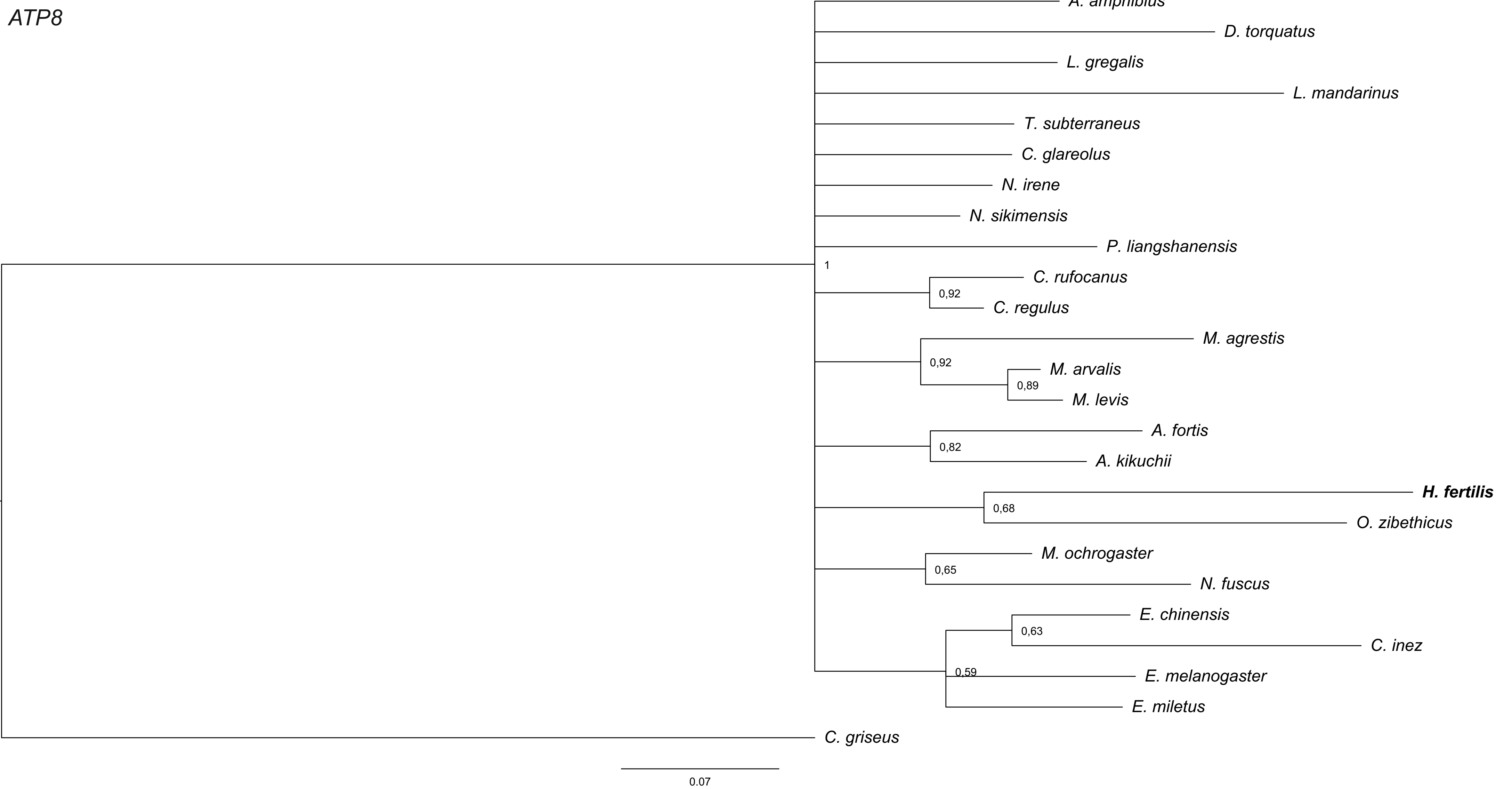


Figure S4. Bayesian phylogenetic reconstruction of tribes Arvicolini and Clethrionomyini inferred separately from alignments of 13 mitochondrial protein-coding genes.
Bayesian trees inferred from separate PCGs partitioned by codon position.
Node labels display BI posterior probabilities (PP).

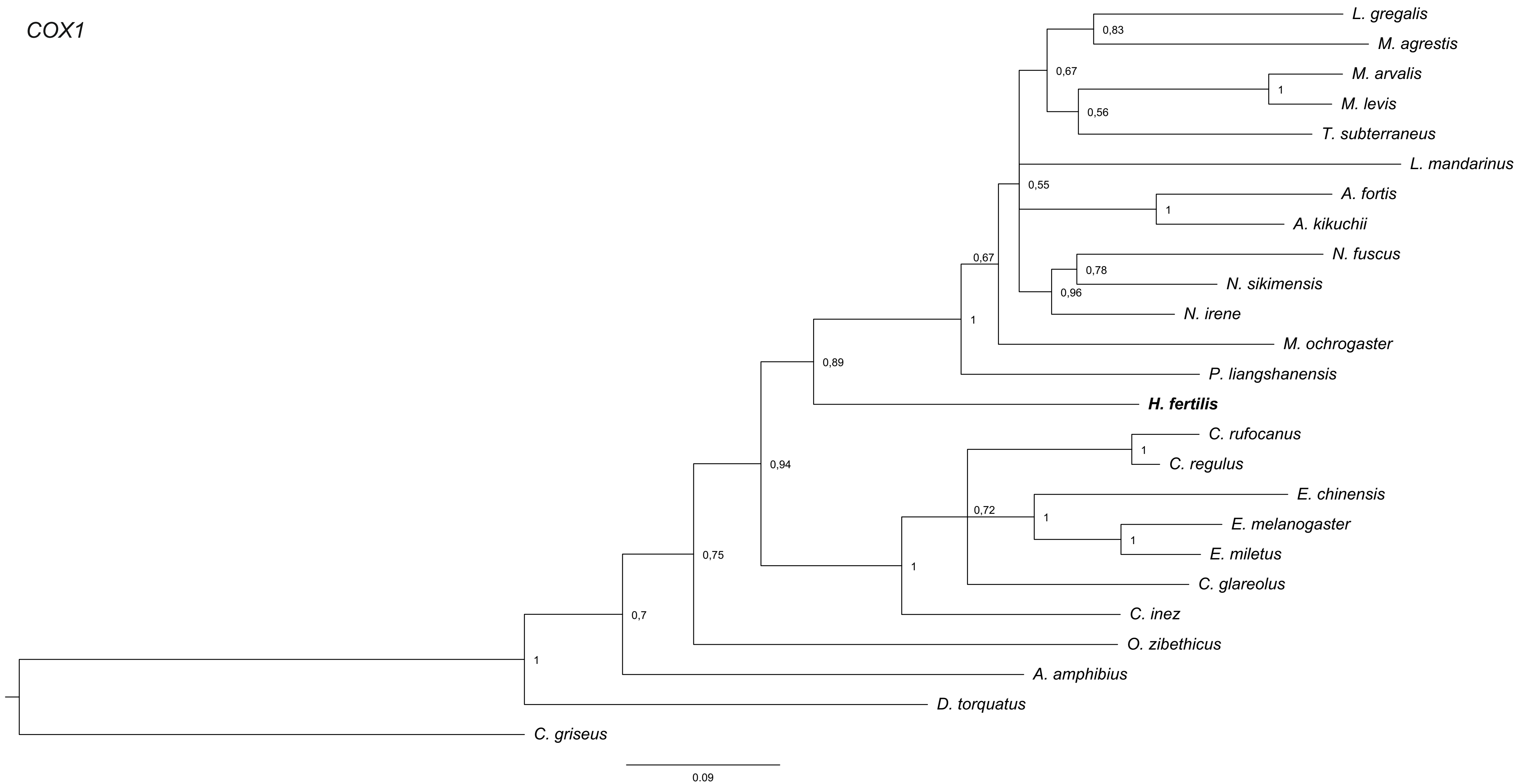
ATP6



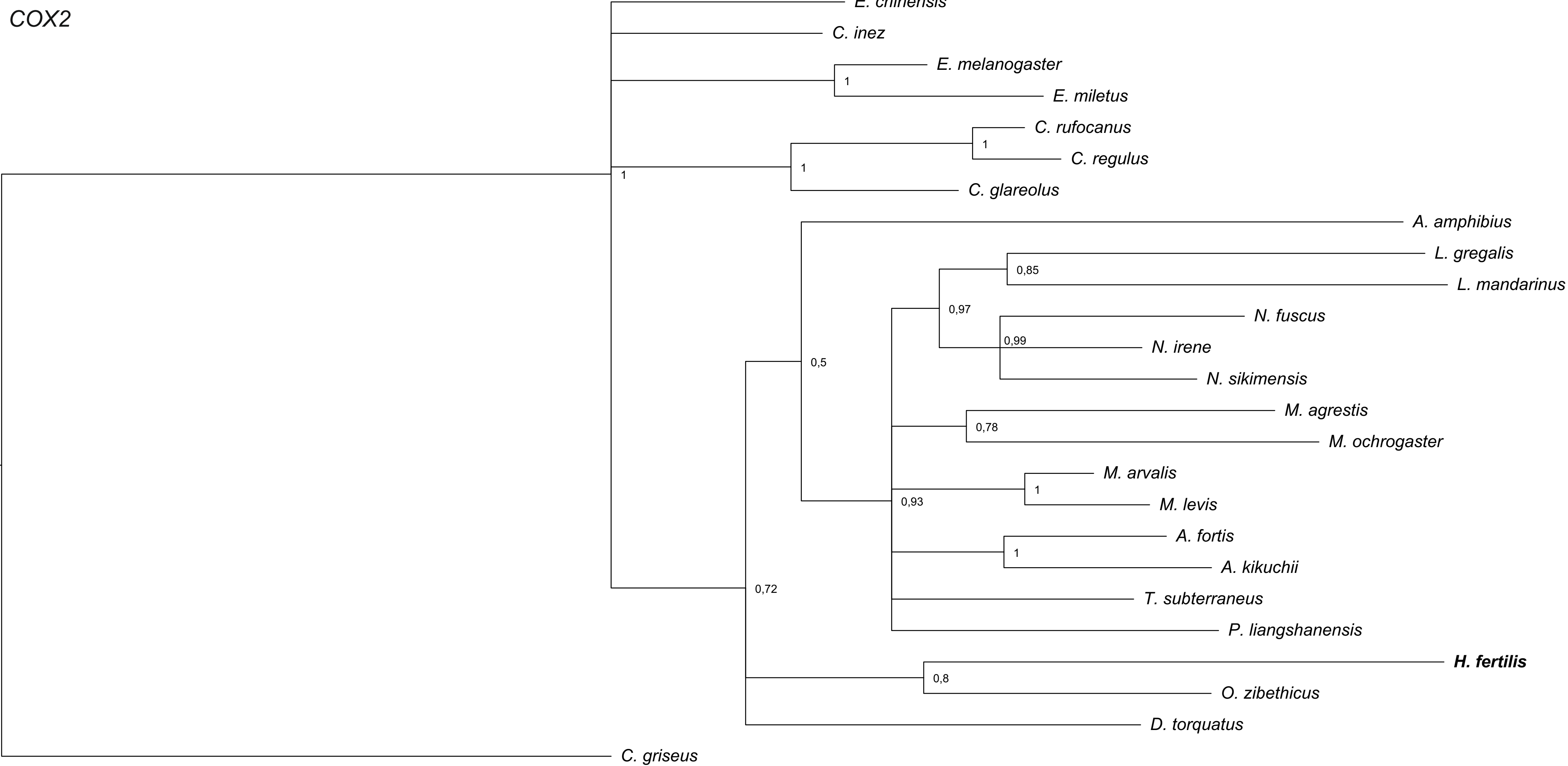
ATP8



COX1

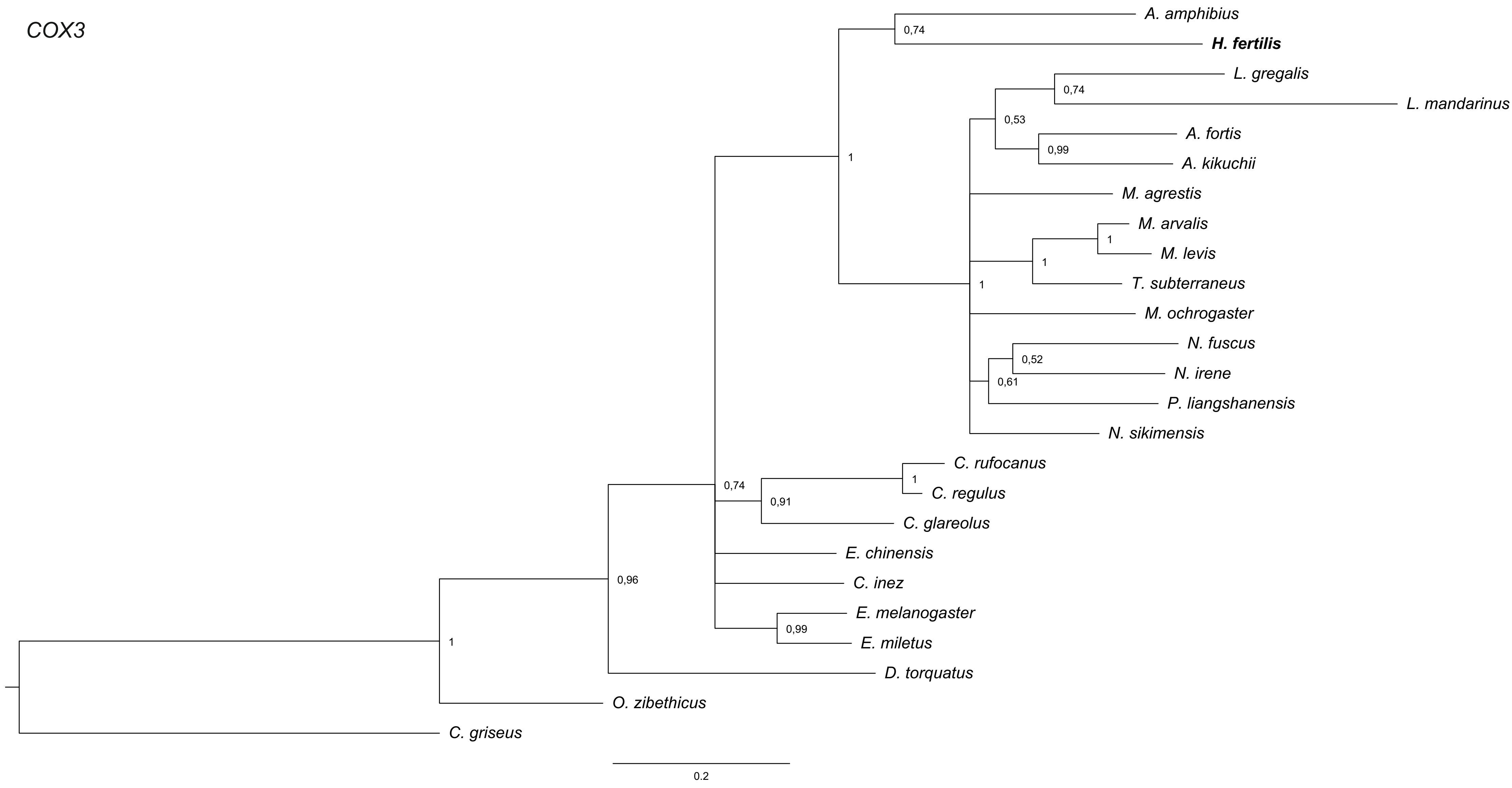


COX2



0.09

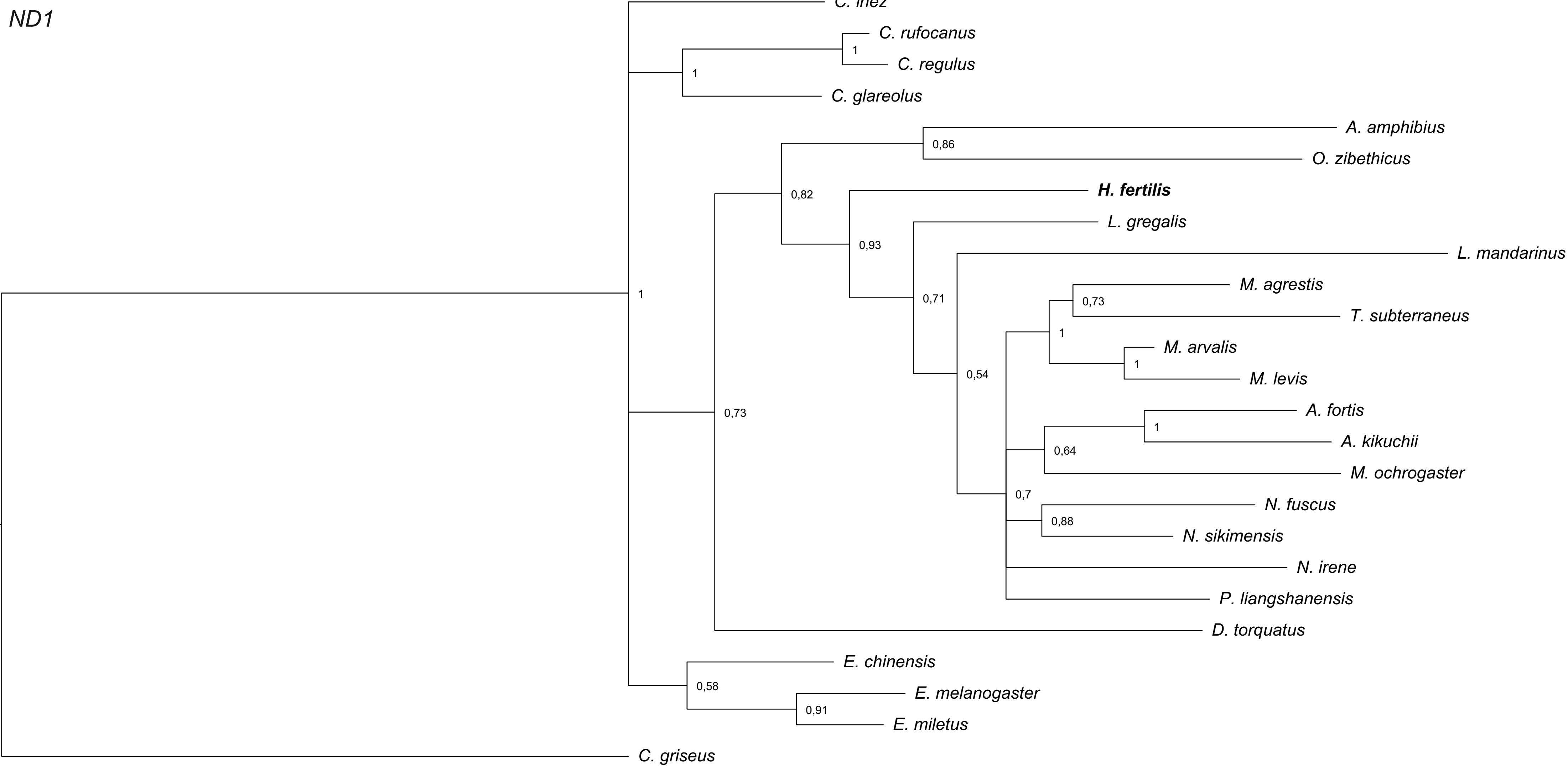
COX3





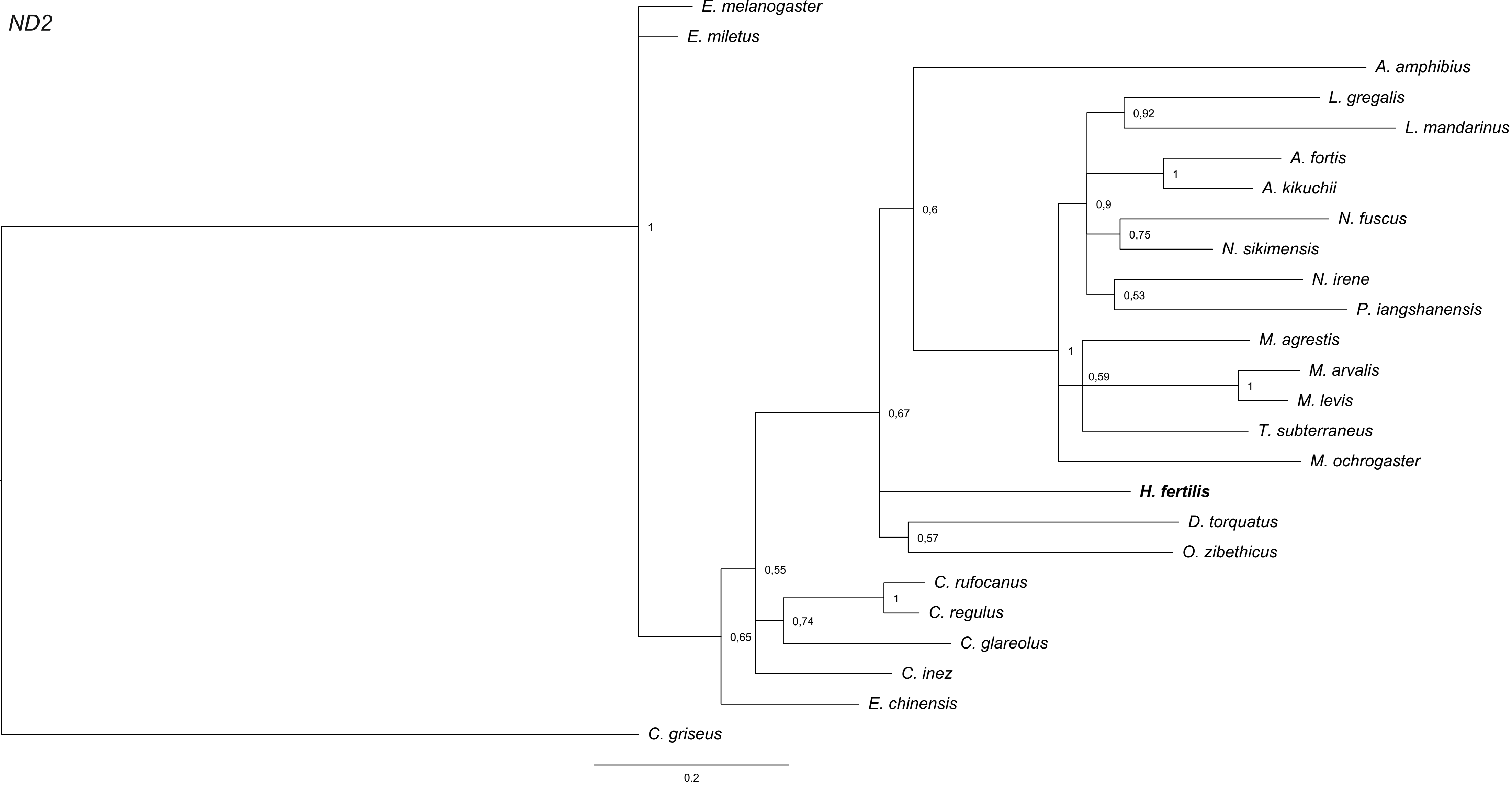
Additional cytochrome *b* sequences downloaded from Genbank are labelled with asterisk.

ND1

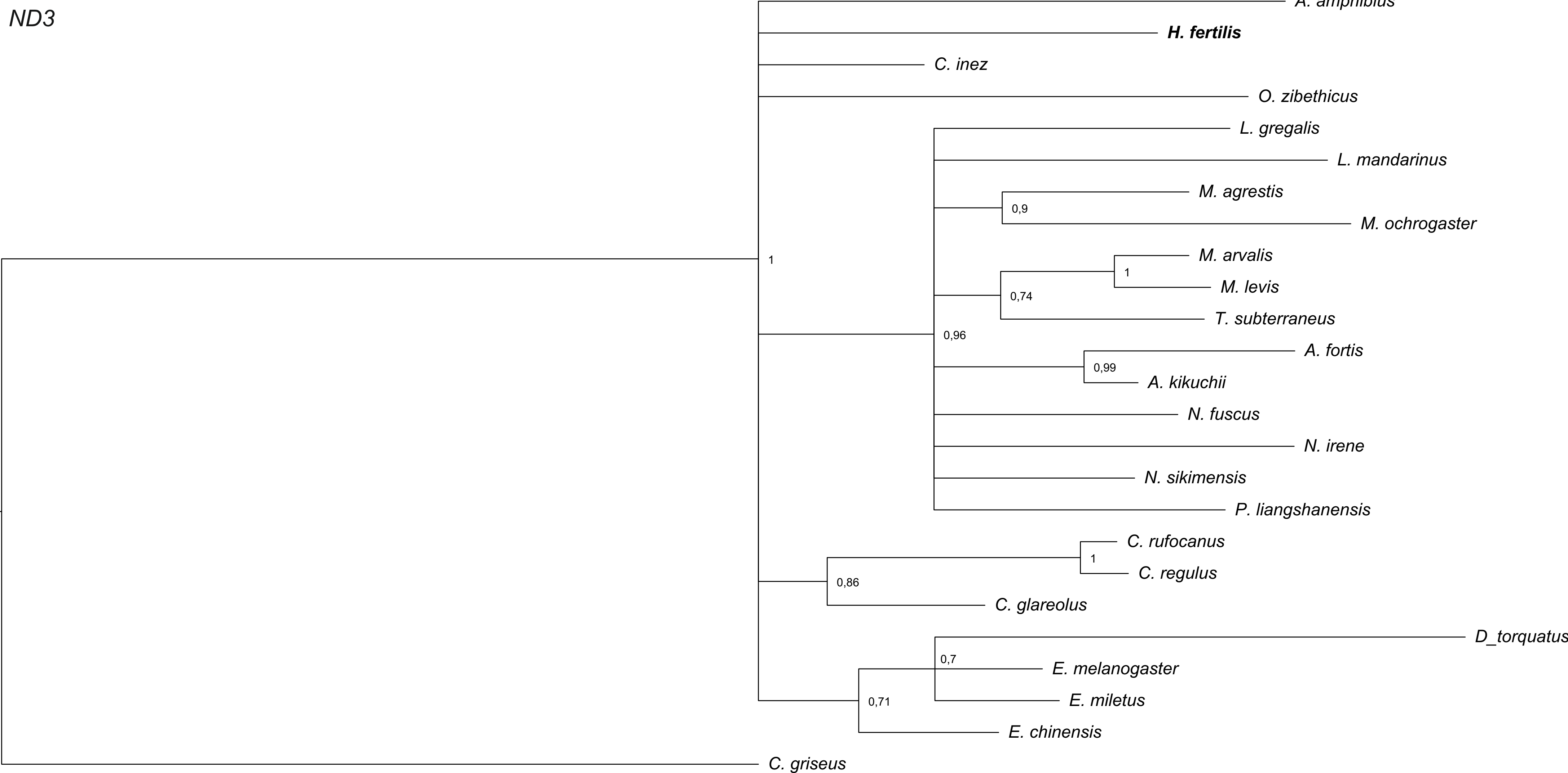


0.2

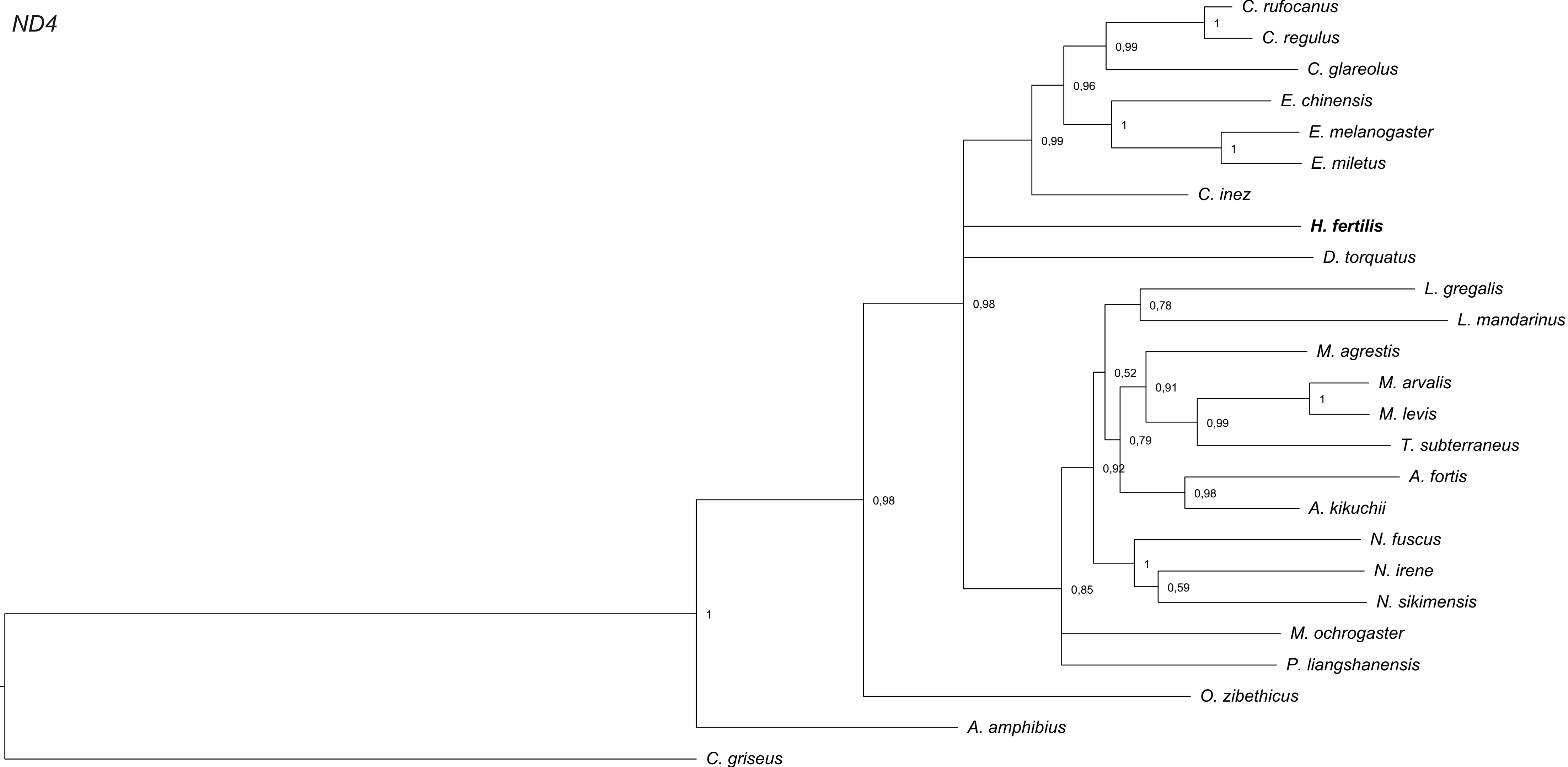
ND2



ND3

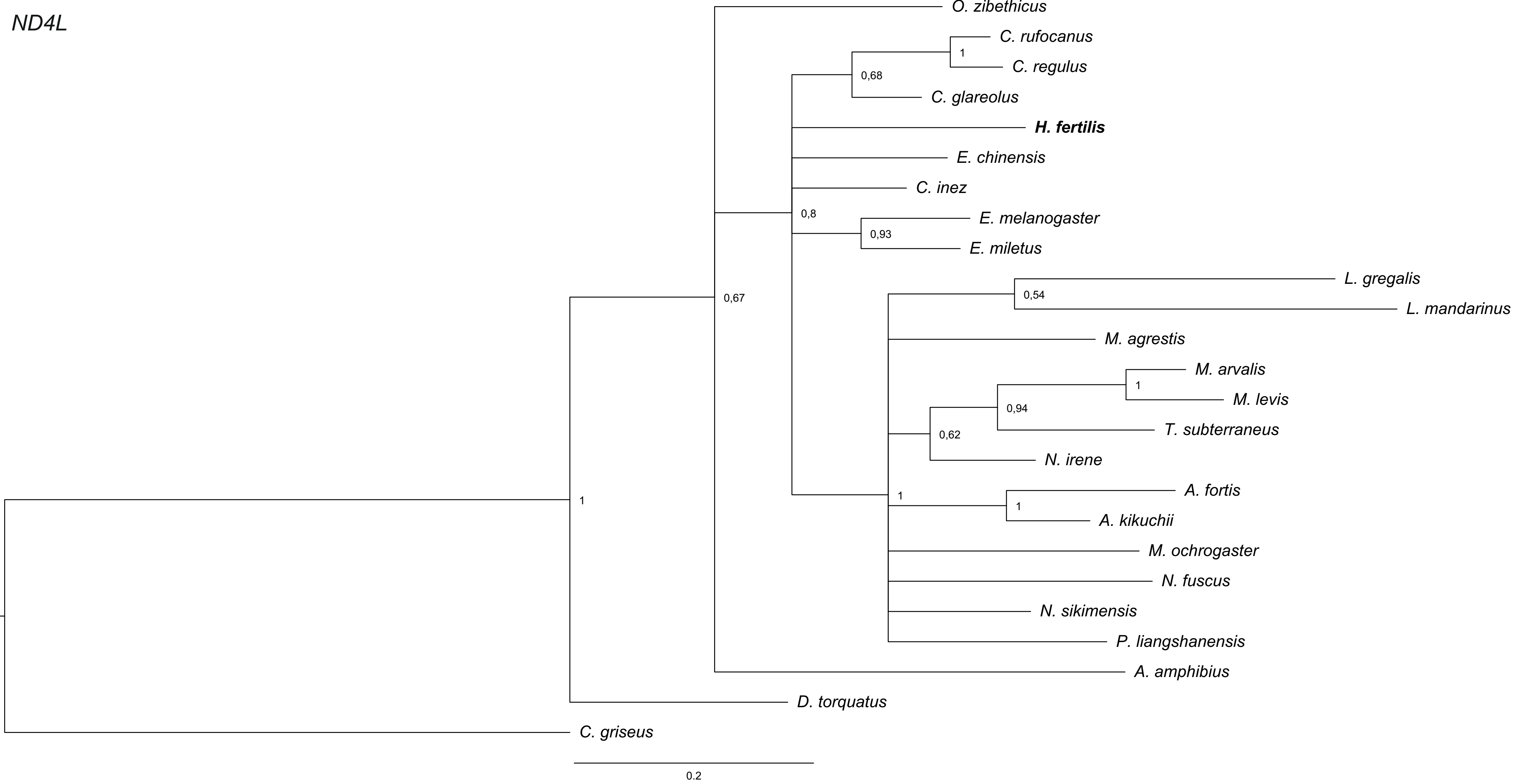


ND4

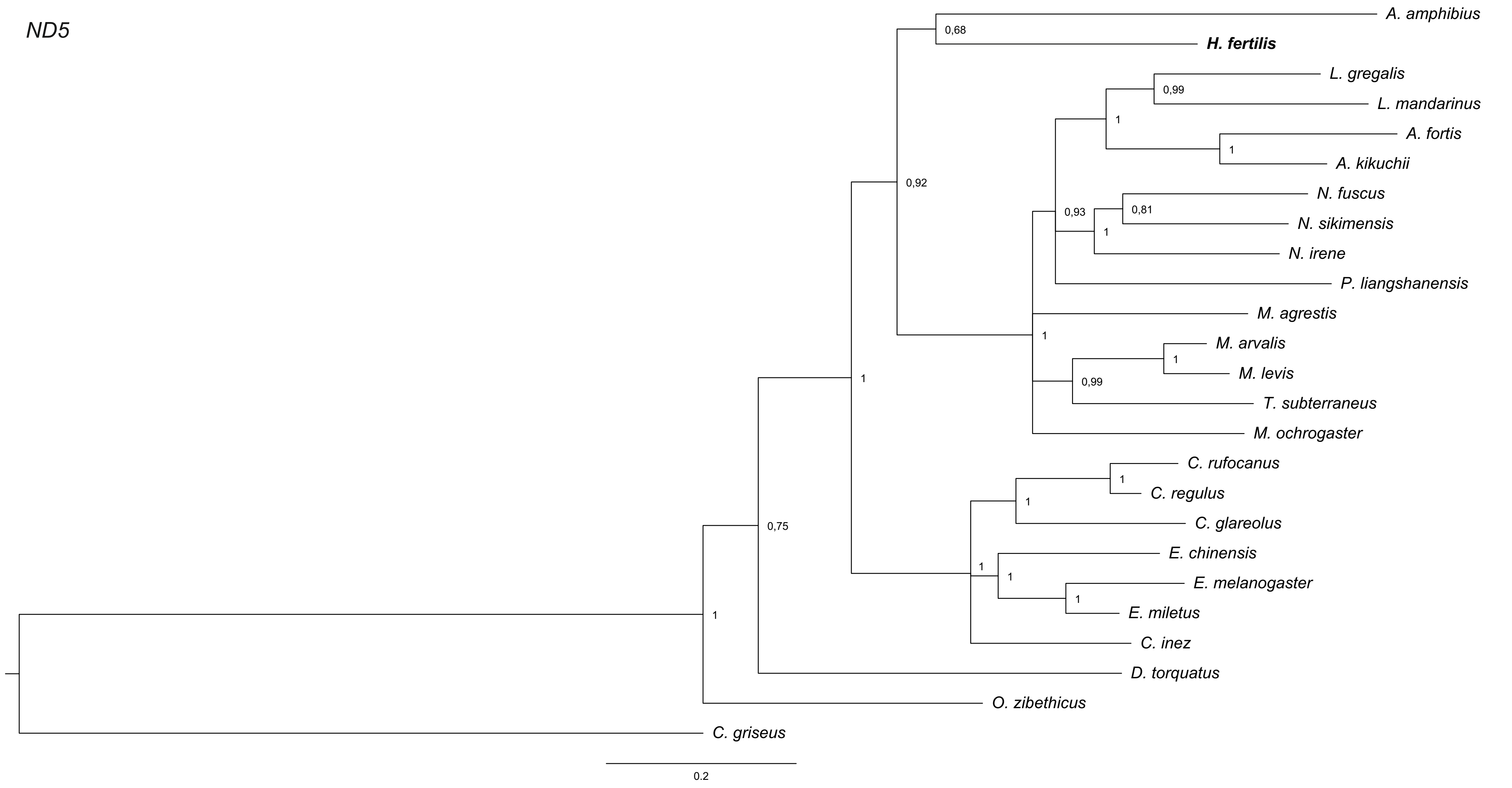


0.2

ND4L



ND5



ND6

