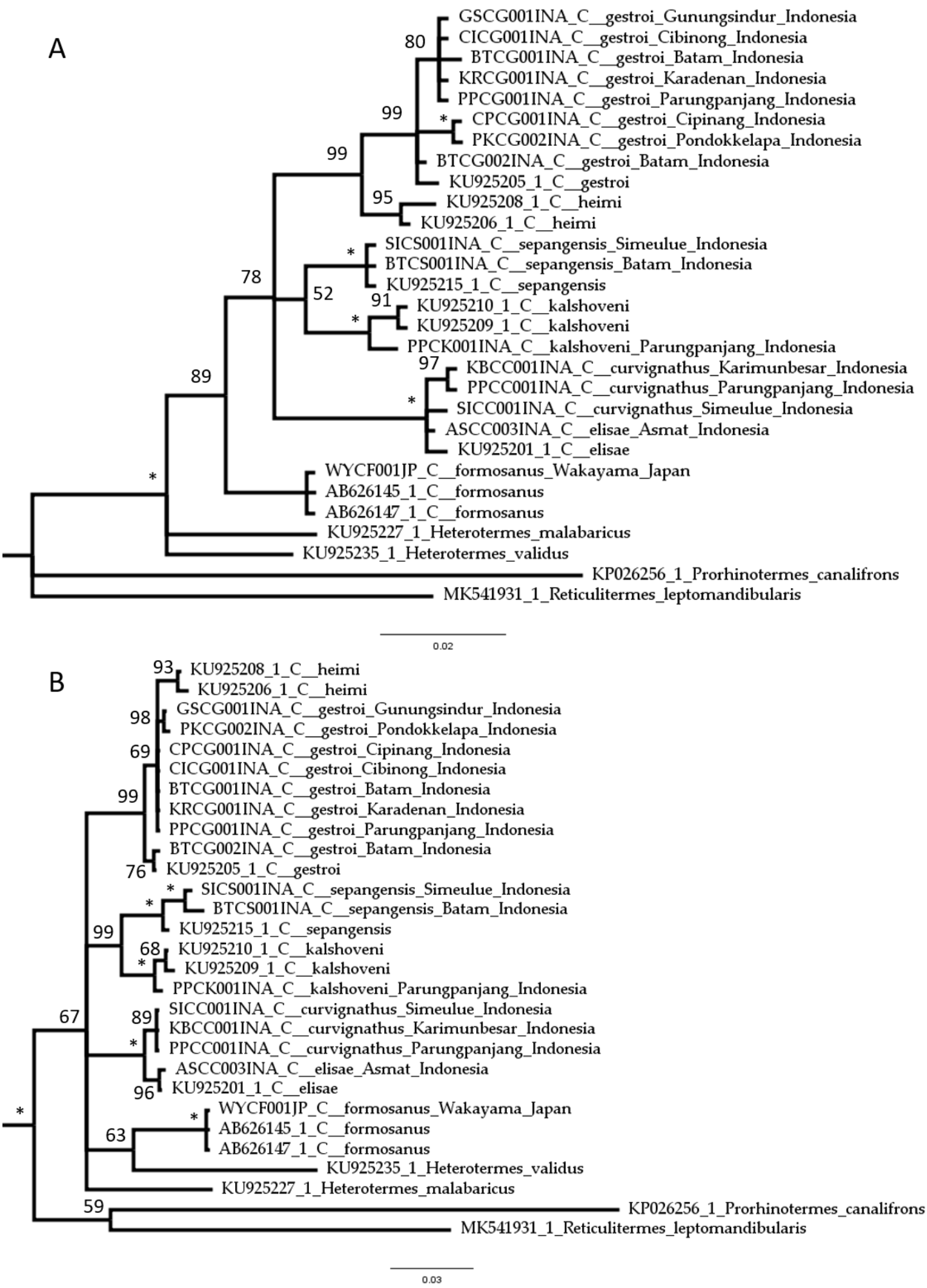

Figure S1. Tree inferred from 12S (A) and 16S (B) rRNA gene fragment by using MrBayes 3.2.7. N-generation was 10 x 10^6 with 3 x 10^5 burn-in. Substitution model was GTR_G for each gene. Sequences with sample code were taken from NCBI database. A hundred percent probability was represented by asterice (*) at the node.