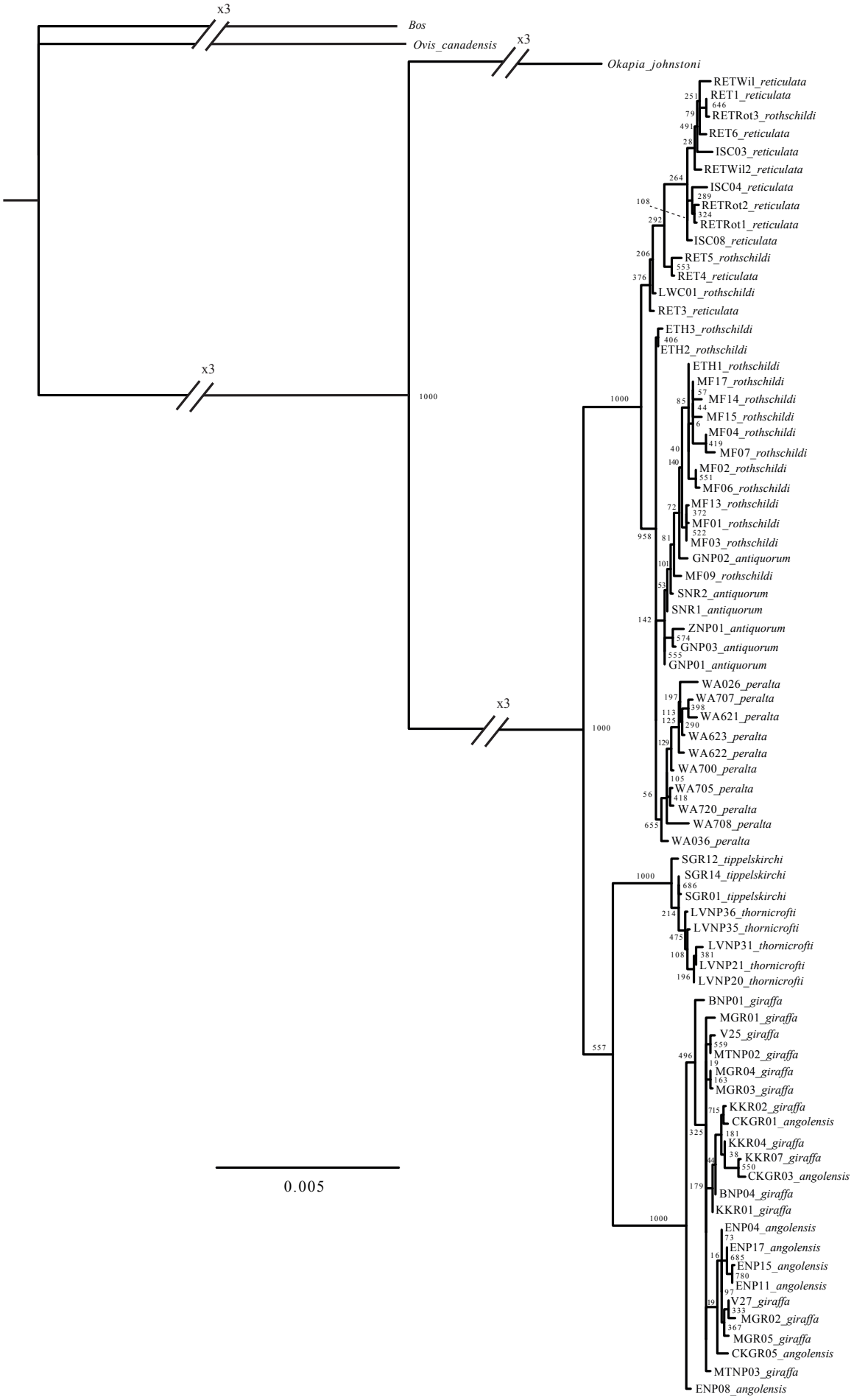

Supplementary file 7. PhyML tree from the Maximum Likelihood analysis of the concatenated alignment of 21 nuclear markers (17.275 bp) for 75 giraffes and three outgroup taxa. The bootstrap values are indicated for each node.