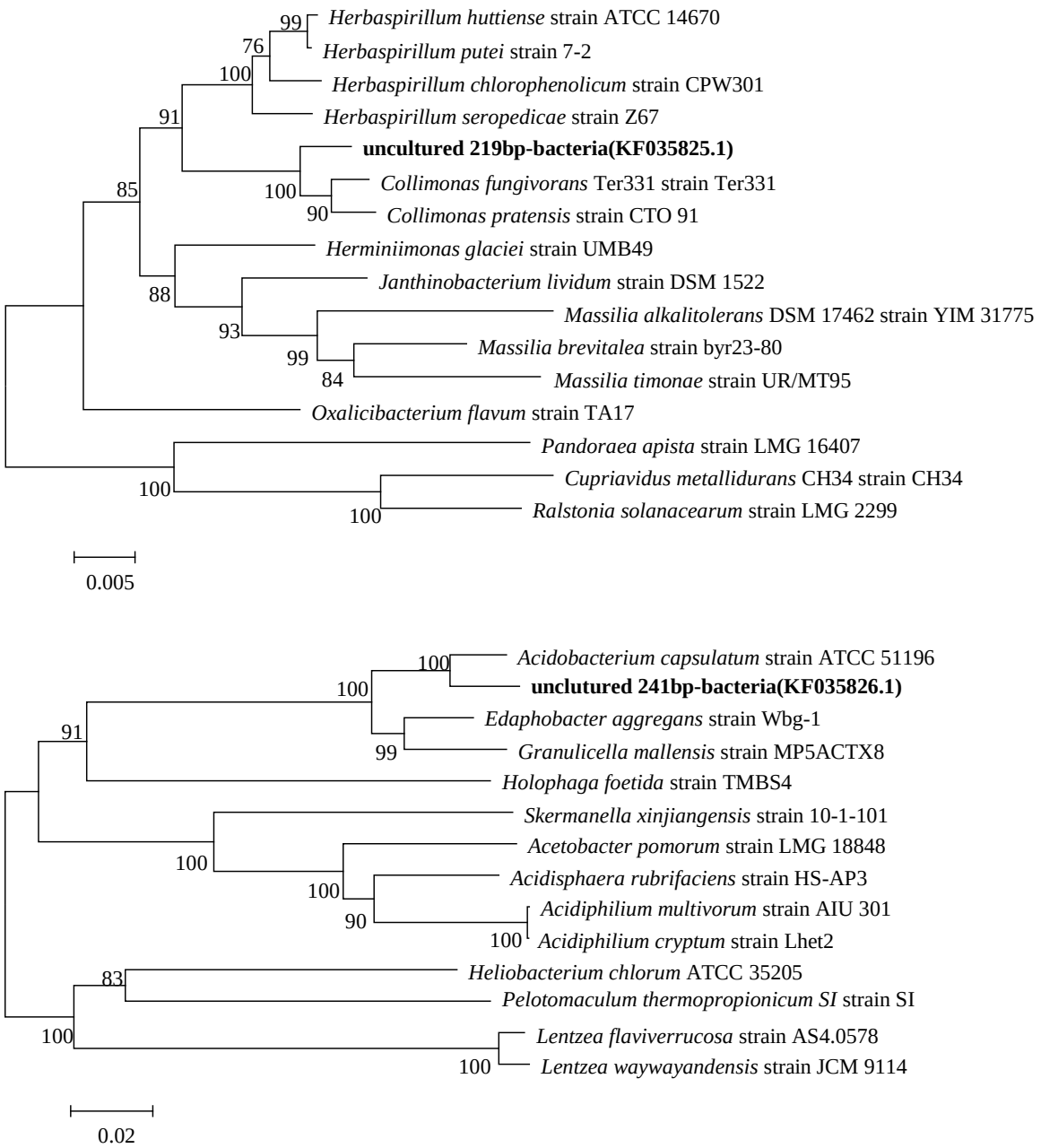

S1 Fig. Phylogenetic tree of the unclutured degraders.

Phylogenetic tree for the taxonomic location of the bacteria corresponding 219- and 241-bp TRFs. The tree is based on 16S rDNA sequence and produced by MEGA 5.0.