

Phylogenetic tree of the 16S rDNA sequences of the bacterial strains. The tree is rooted on the left and branches out to the right. The strains are labeled with their IDs and bootstrap values. The tree shows two main clusters: one on the left (strains 337.2, 3C26.1, 3B21.6, 2388) and one on the right (strains 4235.6, 430.1, 133.1, 36.04, 3B120.7, 3B21.6, 3B21.7, 3B3.1, 4389.2, 14.1, 3B21.1, alpha.1, 13.1, 3B11.2, 3B15.6, 3B17.6, 139.18, alpha1). A scale bar of 0.004 is shown at the bottom.