

A circular phylogenetic tree showing the relationships between various bacterial strains. The tree is divided into several major clades, with labels for each. The legend indicates that the tree is based on 16S rDNA gene cluster data, with 0 copies and 1 copy of the gene cluster. The tree is rooted at the top, and the branches are labeled with strain names. The tree is presented in a circular format, with the root at the top and the branches extending outwards.