

A circular phylogenetic tree showing the relationships between various bacterial strains based on their 16S rDNA sequences. The tree is divided into several major clades, with labels indicating the number of copies of the gene for each strain (0 copies or 1 copy). The tree is rooted at the top, and the branches radiate outwards, forming a circular structure. The labels for the strains are placed along the branches, and the number of copies is indicated by a small circle next to the label.