

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 (0 or 1) for each strain. The tree is rooted at the top and branches outwards, with the outermost branches representing the individual strains. The tree is presented in a circular format, with the branches radiating from the center. The labels for each strain are placed at the end of the branches, and the number of copies is indicated by a small circle (0) or a square (1).