

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 background is light gray, and the tree is drawn in black lines. The labels for the strains are small and placed at the ends of the branches.