

A circular phylogenetic tree showing the relationships between various bacterial strains based on their 16S rDNA sequences. The tree is organized into several major clades, with branches radiating from a central point. The branches are labeled with strain identifiers, and the tree is color-coded to represent different copy numbers: 0 copies (light gray) and 1 copy (dark gray). The tree is set against a light gray background with a white circular border.