

A circular phylogenetic tree showing the relationships between various bacterial strains. The tree is divided into several major clades, each represented by a different color. The strains are labeled with their accession numbers and names, such as *Escherichia coli*, *Salmonella enterica*, and *Staphylococcus aureus*. The tree is rooted at the top, and the branches represent the evolutionary distance between the strains. A legend in the top right corner indicates that the color of the branches corresponds to the number of copies of the 16S rDNA gene cluster: 0 copies (black) and 1 copy (white).