

A circular phylogenetic tree showing the relationships between 16S rDNA gene clusters. The tree is rooted in the center and branches outwards. The branches are labeled with accession numbers. A legend in the top right corner indicates that grey shading represents 0 copies and white represents 1 copy. The tree is divided into several major clades, with some clades showing a high degree of similarity (e.g., the large clade on the right).