

PCA plot of 16S rDNA sequences from 16 bacterial strains. The x-axis is PC1 (12.2%) and the y-axis is PC2 (10.1%). Points are labeled with strain names and accession numbers. A color scale indicates the number of SNPs, ranging from 0 (black) to 16 (red). A dendrogram on the right shows hierarchical clustering of the strains based on their genetic distance.

● NC_060925.1	● NC_060937.1
● NC_060926.1	● NC_060938.1
● NC_060927.1	● NC_060940.1
● NC_060928.1	● NC_060941.1
● NC_060930.1	● NC_060942.1
● NC_060931.1	● NC_060943.1
● NC_060932.1	● NC_060944.1
● NC_060933.1	● NC_060945.1
● NC_060934.1	● NC_060946.1
● NC_060935.1	● NC_060947.1
● NC_060936.1	● NC_060948.1