Descriptions of Additional Supplementary Files

Supplementary Data 1

Description: Vibrio cholerae O1 isolates and genomes used in this study. The following are shown: name, year and country of isolation, source, epidemiological information, biotype, serotype, ctxB and VSP-II alleles, genomic wave, sublineage, antibiotic resistance phenotype, antibiotic resistance gene and element, other typing results, identification as an outlier according to LSD2, GenBank and EBI-ENA accession numbers.