

Supplementary Material 2: Neighbor-joining tree for Genetic relatedness of pathogenic *E. coli* strains identified in this study;The scale indicates the number of nucleotide substitutions per position. Numbers above the branches indicate bootstrap percentages (those above 50%) based on 1,000 replicates. Reference strains of pathogenic *E. coli* were selected from GenBank, using the accession numbers indicated in the text

