

Supplementary Material 4: Neighbor-joining tree for Genetic relatedness targeting 18s rRNA gene of *Cryptosporidium* isolates;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 for *C. parvum* and *C. hominis* isolates were selected from GenBank, using the accession numbers indicated in the text.

