

**Supplemental Figure S1. Phylogenetic tree (UPGMA) of EAEC strain *aggR* genes isolated in Kagoshima**

The *aggR*-PCR amplicons of all EAEC strains isolated in Kagoshima were sequenced.

Representative strains are shown. The EAEC 042 strain was included in a phylogenetic representation. One SNP with a non-synonymous substitution was observed between a and b.