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2 **Figure S1. Comparison of the genomic region comprising *uvrY* in different *E. coli* strains.**

3 Analysis of the chromosomal region next to *uvrY* in the complete genome sequences of *pks*
 4 island-positive, but CPE-negative uropathogenic *E. coli* strain 536 with those of colibactin-
 5 producing newborn meningitis isolate IHE3034 as well as to K-12 laboratory strain MG1655
 6 indicated that *E. coli* 536 carries a 6.1-kb chromosomal deletion spanning the region between
 7 *uvrC* and *fliY* and thus lacks the *uvrY* gene.