

Supplemental Table S1. Characteristics of *aggR*-positive EAEC strains tested and prevalences of virulence genes and the *bla<sub>CTX-M</sub>* genes.

No. Strain No	ST or ST complex (clx)	CG <sup>*</sup>	cluster	year	area	serotype	biofilm	Hep-2 aggR	EAEC virulences plasmid										chromosome			UPEC virulences				ESBL	
									aatA	aap	AAF/1	AAF/2	AAF/3	AAF/4	AAF/5	pet	astA	sepA	sat	pic	aaiC	sfa	pap	afa	aer	cnf1	CTX-M <sup>†</sup>
1 K7656	ST200 clx	CG200	C1	2009	K	O126:H27	0.64	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
2 K7658	ST200 clx	CG200	C1	2009	K	O126:HND	0.49	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
3 K7484	ST200 clx	CG200		2008	K	O127:H27	0.22	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	+	+	+	+	-
4 K7668	ST200 clx	CG200		2009	K	O127:H27	0.58	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
5 K7701	ST200 clx	CG200		2009	K	O127:H27	0.5	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
6 K7164	ST200 clx	CG200		2008	K	O127:H27	0.29	NA	+	+	+	-	-	-	-	-	+	+	-	-	+	+	-	-	-	-	-
7 K7225	ST200 clx	CG200		2008	K	O142:H5	0.28	AA	+	+	+	-	-	-	-	-	+	+	-	-	+	+	-	-	-	-	-
8 K4685	ST200 clx	CG200	C2	2005	K	OUT:H27	0.56	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
9 K4368	ST200 clx	CG200	C2	2005	K	O127a:H27	0.87	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
10 K601	ST200	CG200	C3	1998	K	O126:HND	0.48	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
11 K666	ST200	CG200	C3	1998	K	O126:HND	0.42	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
12 K1437	ST200	CG200	C3	2001	K	OUT:H27	1.88	AA	+	+	+	-	+	-	-	-	+	+	-	-	+	+	-	-	-	-	-
13 K2012	ST200	CG200	C3	2002	K	O126:H27	0.29	AA	+	-	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
14 K2013	ST200	CG200	C3	2002	K	O126:H27	0.54	AA	+	-	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
15 K2754	ST200	CG200	C3	2003	K	O126:HND	0.21	AA	+	-	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
16 K2832	ST200	CG200	C3	2003	K	O126:HND	0.45	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
17 K2855	ST200	CG200	C3	2003	K	OUT:H1	1.43	AA	+	-	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
18 K2858	ST200	CG200	C3	2003	K	O126:H27	0.61	AA	+	-	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
19 K3574	ST200	CG200	C3	2004	K	OUT:HND	0.68	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
20 K4610	ST200	CG200	C3	2005	K	O86:H27	0.55	NA	+	+	+	-	-	-	-	-	-	-	-	-	+	+	-	-	-	-	-
21 K5227	ST200	CG200	C3	2006	K	O126:HND	1.67	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	-
22 K6269	ST200	CG200	C3	2007	K	O126:H27	2.17	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
23 K7130	ST200	CG200	C3	2008	K	OUT:H27	0.68	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
24 K7132	ST200	CG200	C3	2008	K	O126:H27	1.50	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
25 K7240	ST200	CG200	C3	2008	K	OUT:H27	3.50	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
26 K7693	ST200	CG200	C3	2009	K	O126:H27	0.27	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
27 K7982	ST200	CG200	C3	2010	K	O126:HND	0.58	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
28 K9544	ST200	CG200	C3	2009	O	O126:H27	2.78	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
29 K2626	ST200 clx	CG200		2003	K	O126:H27	0.75	AA	+	+	+	-	+	-	-	-	+	-	-	-	+	+	-	-	-	-	-
30 K566	ST40	CG40	C4	1998	K	O111:H21	1.15	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	-
31 K2776	ST40	CG40	C4	2003	K	O111:HND	0.90	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	-
32 K7987	ST40	CG40	C4	2010	K	O127a:HNC	1.45	AA	+	+	+	-	-	-	-	-	-	-	-	-	+	+	-	-	-	-	-
33 K578	ST40	CG40	C5	1998	K	O111:H21	0.72	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	-
34 K708	ST40	CG40	C5	1998	K	O111:HND	1.02	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	-
35 K771	ST40	CG40	C5	1998	K	O111:H21	0.85	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	+
36 K878	ST40	CG40	C5	1998	K	O111:H21	0.73	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	-
37 K1027	ST40	CG40	C5	1999	K	O111:H21	0.94	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-	-
38 K1028	ST40	CG40	C5	1999	K	O111:H21	0.99	AA	+	+	+	-	-	-	-	-	+	-	-	-	+	+	-	-	-		

Abbreviations: K, Kagoshima; O, Osaka; Biofilm, biofilm index; EAEC, enteroaggregative Escherichia coli; UPEC, uropathogenic E. coli; ESBL, extended-spectrum  $\beta$ -lactamase; AA, aggregative adherent pattern; aAA, atypical AA, DA, diffuseadherent pattern; NA, no-adherent; NT, not tested.

\*A Clonal group (CG) was defined as a group of closely related sequence type and named in accordance with the dominant ST. †The number of CTX-M column shows the subtype of CTX-M. §ESBL-producing confirmed by antimicrobial susceptibility test.