

Table S1: PCR primer sequences.

Gene	Primer sequence (5' to 3')	Gene target	Product size (bp)	Reference
<i>ehlyA-F</i>	(F) GCATCATCAAGCGTACGTTCC (R) AATGAGCCAAGCTGGTTAACGT	<i>hlyA</i>	534	
<i>eaeA</i>	(F) GACCCGGCACAAGCATAAGC (R) CCACCTGCAGCAACAAGAGG	<i>eaeA</i>	384	(Paton and Paton, 1998)
<i>stx1</i>	(F) ATAAATGCCATTGCTTACTAC (R) AGAACGCCACTGAGATCATC	<i>stx1</i>	180	
<i>stx2</i>	(F) GGCACTGTCTGAAACTGCTCC (R) TCGCCAGTTATCTGACATTCTG	<i>stx2</i>	255	
<i>stx2a</i>	(F) AGATATCGACCCCTTGAAG (R) GTCAACCTTCACTGTAAATG	<i>Stx2a</i>	969	(He et al., 2012)
<i>uidA</i>	(F) CTGGTATCAGCGCGAAGTCT (R) AGCGGGTAGATATCACACTC	<i>uidA</i>	600	(Anbazhagan et al., 2010)
<i>rfbO157</i>	(F) CGGACATCCATGTGATATGG (R) TTGCCTATGTACAGCTAATCC	<i>rfbO157</i>	259	(Morin et al., 2004)
Streptomycin	(F) TATCCAGCTAACGCGGAAC (R) ATTGCGGACTACCTTGGTC	<i>aadA1</i>	447	(Momtaz et al., 2012)
Chloramphenicol	(F) AGTTGCTCAATGTACCTATAACC (R) TTGTAATTCAATTAGCATTCTGCC	<i>cataI</i>	547	(Van et al., 2008)
Tetracycline A	(F) GGTCACTCGAACGACGTCA (R) CTGTCGACAAGTTGCATGA	<i>tetA</i>	577	
Tetracycline B	(F) CCTCAGCTCTAACGCGTG (R) GCACCTTGCTGATGACTCTT	<i>tetB</i>	634	
Beta-lactams	(F) ATGAGTATTCAACATTCCTGTGTC (R) TTACCAATGCTTAATCAGTGAGG	<i>blaTEM-1</i>	861	(Haidinger et al., 2003)
Biofilm-related	(F) ACTCTGACTTGACTATTACC (R) AGATGCAGTCTGGTCAAC	<i>csgA</i>	200	
Biofilm-related	(F) TTTCGATTGTCTGGCTGTATG (R) CTTCAGATTCAAGCGCTGTC	<i>crl</i>	250	
Biofilm-related	(F) TTATCGCCTGAGGTTATCGTTGC (R) TCTTCAGGCTCTATTATTCTGGATAT	<i>csgD</i>	501	(Ogasawara et al., 2010)
Biofilm-related	(F) TATCGCTGGATTACTGGCAAC (R) TAGGACGCTGACGTGTCTTATC	<i>rpos</i>	1900	(Olsen et al., 1993)
Biofilm-related	(F) TGCAGAACGGATAAGCCGTGG (R) GCAGTCACCTGCCCTCCGGA	<i>fimH</i>	506	(Johnson and Stell, 2000)
Biofilm-related	(F) CCGGCAGGCAATGGGTACA (R) CAGCTCTCACAAATCTGGCGAC	<i>flu</i>	385	(Restieri et al., 2007)
Biofilm-related	(F) ATATCGGCTAAAGTGGAACAGGTCC (R) TGCTTCCCTGGCATTAACTTCACC	<i>ehaA^a</i>	627	
Biofilm-related	(F) AGTGCATGACTACTGATTGTGCTG (R) CACATTAAACAAACCGCTCTGG	<i>ehaA^b</i>	887	(Friedrich et al., 2002)

Table S2: Reference values used to classify isolates as susceptible, intermediate resistant and resistant.

Antimicrobial agent	Disk content	S	I	R
Ampicillin	10 µg	≥ 17	14-16	≤ 13
Amoxicillin-clavulanate	20/10 µg	≥ 18	14-17	≤ 13
Ampicillin-sulbactam	10/10 µg	≥ 15	12-14	≤ 11
Cefepime	30 µg	≥ 25	19-24	≤ 18
Cefotaxime	30 µg	≥ 26	23-25	≤ 22
Cefoxitin	30 µg	≥ 18	15-17	≤ 14
Ceftazidime	30 µg	≥ 21	18-20	≤ 17
Aztreonam	30 µg	≥ 21	18-20	≤ 17
Imipenem	10 µg	≥ 23	20-22	≤ 19
Meropenem	10 µg	≥ 23	20-22	≤ 19
Gentamicin	10 µg	≥ 15	13-14	≤ 12
Streptomycin	10 µg	≥ 15	12-14	≤ 11
Tetracycline	30 µg	≥ 15	12-14	≤ 11
Nalidixic acid	30 µg	≥ 19	14-18	≤ 16
Norfloxacin	10 µg	≥ 17	13-16	≤ 12
Trimethoprim-sulfamethoxazole	1.25/23.75µg	≥ 16	11-15	≤ 10
Chloramphenicol	30 µg	≥ 18	13-17	≤ 12
Colistin	10 µg	≥ 17	13-17	≤ 12

S= Susceptible, I= Intermediate and R= Resistance.

Table S3: Non-O157 *E. coli* isolates sequenced in this study.

Isolate ID	Serogroup	Number of contigs	Total length (bp)	GC (%)	N50
3	O99:H9	116	4686813	50.79	222406
11	O156:H25	299	5181977	50.43	94802
12	O108:H2	103	4881051	50.72	186185
14	O136:H30	128	4564407	50.92	105160
15	O99:H9	103	4686475	50.78	260468
22	wzx-Onovel24:H20	94	4964550	50.53	478048
25	O140:H21	247	5507075	50.47	148987
30	O102:H4	376	5244869	50.65	87095
32	O129:H23	203	5501418	50.53	240397
37	O17:H18	139	5169910	50.45	186852
38	O76:H34	219	4812816	50.88	114298
42	O26:H11	494	5526211	50.4	101076
50	O129:H23	204	5502410	50.52	211833
56	O154:H10	144	4706567	50.66	96083
60	O116:H21	195	5081941	50.7	177288
64	wzx-Onovel5:H19	137	5056230	50.74	161129
65	wzx-Onovel5:H19	135	5058237	50.74	236278
67	O87:H7	127	4764651	50.63	195836
68	O129:H21	171	5105006	50.67	156897
69	O26:H11	478	5562801	50.4	112464
72	O26:H11	107	4610054	50.96	161578
76	O163:H19	303	5171039	50.57	117460
77	O40:H19	156	4930994	50.7	151142
80	O22:H21	87	4795733	50.72	185546

Table S4: Annotation of virulence factors for non-O157 *E. coli* isolates.

Isolate ID	Serogroup	Virulence factor related genes																			
		<i>ehaA</i>	<i>ehaB</i>	<i>epeA</i>	<i>agn43</i>	<i>espP</i>	<i>hlyA</i>	<i>subA</i>	<i>east1</i>	<i>cdtA</i>	<i>espA</i>	<i>chuA</i>	<i>fimH</i>	<i>aslA</i>	<i>ibeB</i>	<i>ompA</i>	<i>toxB</i>	<i>paa</i>	<i>ecpA</i>	<i>csgA</i>	<i>eaeA</i>
3	O99:H9	-	+	-	-	-	-	-	-	-	-	-	+	+	+	-	-	-	+	+	-
11	O156:H25	+	+	-	+	+	+	-	+	-	+	-	+	-	+	-	-	+	+	+	+
12	O108:H2	+	+	-	+	-	-	-	-	-	-	-	+	-	+	-	-	+	+	+	-
14	O136:H30	-	+	-	-	-	-	-	-	-	-	-	+	-	+	-	-	-	-	+	-
15	O99:H9	-	+	-	-	-	-	-	-	-	-	-	+	+	+	-	-	-	+	+	-
22	wzx- Onovel24:H20	-	+	-	-	-	-	-	-	-	-	-	+	+	+	-	-	-	+	+	-
25	O140:H21	+	-	-	-	-	-	-	-	-	-	-	-	+	-	+	-	-	-	+	-
30	O102:H4	-	+	-	-	-	-	-	-	-	-	-	+	-	+	-	-	-	.	+	-
32	O129:H23	-	+	-	-	-	-	-	-	-	-	-	+	+	+	-	-	-	+	+	-
37	O17:H18	-	+	-	+	+	+	+	-	-	-	-	+	+	+	+	-	-	+	+	-
38	O76:H34	-	+	-	-	-	-	-	-	-	-	-	-	+	+	+	-	-	+	+	-
42	O26:H11	+	+	-	+	+	+	-	+	-	+	-	+	-	+	-	+	+	+	+	+
50	O129:H23	-	+	-	-	-	-	-	-	-	-	-	+	+	+	-	-	-	+	+	-
56	O154:H10	-	+	-	-	-	-	-	-	-	-	-	-	+	-	+	-	-	+	+	-
60	O116:H21	+	+	+	-	+	+	+	-	-	-	-	-	+	-	+	-	-	+	+	-
64	wzx- Onovel5:H19	-	+	+	+	+	+	+	-	-	-	-	-	+	-	+	-	-	+	+	-
65	wzx- Onovel5:H19	-	+	+	+	+	+	+	-	-	-	-	-	+	-	+	-	-	-	+	-
67	O87:H7	+	+	-	-	-	-	-	-	-	-	-	-	+	-	+	-	-	-	+	-
68	O129:H21	+	+	+	-	+	+	+	-	-	-	-	-	+	-	+	-	-	+	+	-
69	O26:H11	+	+	-	+	+	+	-	+	-	+	-	+	-	+	-	+	+	+	+	+
72	O26:H11	-	+	-	-	-	-	-	-	-	-	-	+	-	+	-	-	-	+	+	-
76	O163:H19	+	+	+	-	-	-	-	-	-	+	-	-	+	-	+	-	-	+	+	-
77	O40:H19	+	+	-	-	+	+	+	-	-	-	-	-	+	-	+	-	-	+	+	-
80	O22:H21	+	+	-	-	+	-	-	-	-	-	-	-	-	+	-	+	-	-	+	-

+ = gene present, - = gene absent

Autotransporter protein genes= (*ehaA*, *ehaB*, *epeA* and *agn43*), serine protease gene= (*espP*), toxins (*hlyA*, *subA*, *east1* and *cdtA*), type III translocated gene (*espA*), iron uptake gene (*chuA*), invasins genes (*fimH*, *aslA*, *ibeB* and *ompA*) and adhesins genes= (*toxB*, *eaeA*, *ecpA*, *paa* and *csgA*).

Table S5: Annotation results of antimicrobial resistance for non-O157 *E. coli*.

Isolate ID	Serogroup	Antimicrobial related genes															
		TEM-105	TEM-150	AadA1	aadA2	aadA5	sul2	sul3	tet(A)	tet(C)	tet(D)	cmlA6	dfrA12	linG	floR	ugd	vgaC
3	O99:H9	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
11	O156:H25	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
12	O108:H2	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
14	O136:H30	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
15	O99:H9	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
22	wzx-Onovel24:H20	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
25	O140:H21	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
30	O102:H4	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+
32	O129:H23	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
37	O17:H18	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
38	O76:H34	-	+	-	-	-	+	-	-	+	-	-	-	-	-	+	+
42	O26:H11	-	-	-	-	-	+	-	+	-	+	-	-	-	-	+	+
50	O129:H23	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
56	O154:H10	+	-	+	+	-	-	+	+	-	+	+	+	+	-	+	-
60	O116:H21	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
64	wzx-Onovel5:H19	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
65	wzx-Onovel5:H19	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
67	O87:H7	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
68	O129:H21	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
69	O26:H11	-	-	-	-	-	+	-	+	-	+	-	-	-	-	+	+
72	O26:H11	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
76	O163:H19	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
77	O40:H19	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-
80	O22:H21	-	-	-	-	-	+	+	-	-	+	-	-	-	-	+	-

+ = gene present, - = gene absent

Beta-lactams (TEM-105 and TEM-150), streptomycin (aadA1, aadA2 aadA5), sulfonamide (sul2 and sul3), tetracycline (tetACD), phenicol (cmlA6 and floR), diaminopyrimidine (dfrA12), lincosamides (linG), polymyxin (ugd) and streptogramin (vgaC).

Table S6: Annotated results of plasmids search for non-O157 *E. coli*.

Isolate ID	Serogroup	Plasmids types															
3	O99:H9	-	-	Col156_1	-	-	ColRNAI_1	-	-	ColpVC_1	-	IncB/O/K/Z_2	-	IncB/O/K/Z_3	-	IncB/O/K/Z_4	-
11	O156:H25	+	-	'	-	-	'	-	-	'	-	'	-	'	-	IncFIB(AP001918)_1	-
12	O108:H2	-	-	-	-	-	-	-	-	-	-	-	-	-	-	IncFIB(K)_1_Kpn3	-
14	O136:H30	-	+	-	-	-	-	-	-	-	-	-	-	-	-	IncFIC(FII)_1	-
15	O99:H9	-	-	+	-	-	-	-	-	-	-	-	-	-	-	IncFII(pHN7A8)_1_pH	-
22	wzx-Onovel24:H20	-	.	-	-	-	-	-	+	+	-	-	+	-	-	N7A8	-
25	O140:H21	-	-	-	-	-	-	-	+	+	-	-	+	-	-	IncH2_1	-
30	O102:H4	-	-	-	-	-	-	-	-	+	-	-	+	-	-	Incl1_1_Alpha	-
32	O129:H23	-	+	-	-	-	-	-	-	+	-	-	-	-	-	Incl2_1_Delta	-
37	O17:H18	-	-	-	-	-	-	-	-	+	-	-	+	-	-	IncR_1	-
38	O76:H34	-	+	-	-	-	-	-	-	-	+	-	-	-	-	IncY_1	-
42	O26:H11	-	+	-	-	+	-	-	-	+	-	-	-	-	-	-	-
50	O129:H23	-	+	-	-	-	-	-	-	+	-	-	-	-	-	-	-
56	O154:H10	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
60	O116:H21	-	+	-	+	-	-	-	-	+	-	-	+	-	-	-	-
64	wzx-Onovel5:H19	-	+	-	+	-	-	-	-	+	-	-	-	-	-	-	-
65	wzx-Onovel5:H19	-	+	-	+	-	-	-	-	+	-	-	-	-	-	-	-
67	O87:H7	-	-	-	-	-	-	-	+	+	-	-	+	-	-	-	-
68	O129:H21	-	+	-	+	-	-	-	-	+	-	-	+	-	-	-	-
69	O26:H11	-	+	-	-	+	-	-	-	+	-	-	-	-	-	-	-
72	O26:H11	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
76	O163:H19	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-
77	O40:H19	-	+	-	-	-	-	-	+	+	-	-	+	-	-	-	-
80	O22:H21	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+

+ = gene present, - = gene absent.

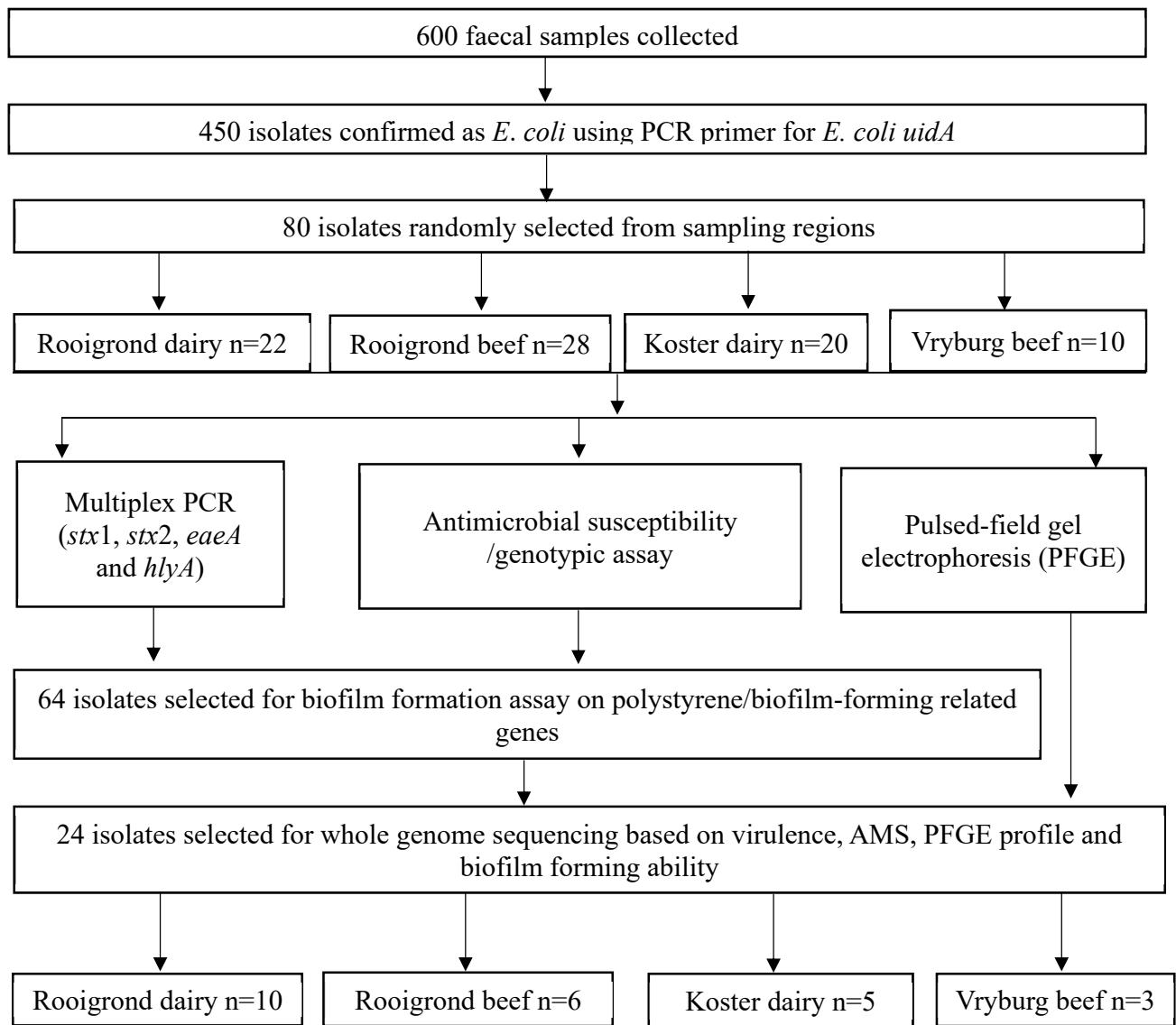


Figure S1: A flow chart illustrating bacterial characterization.

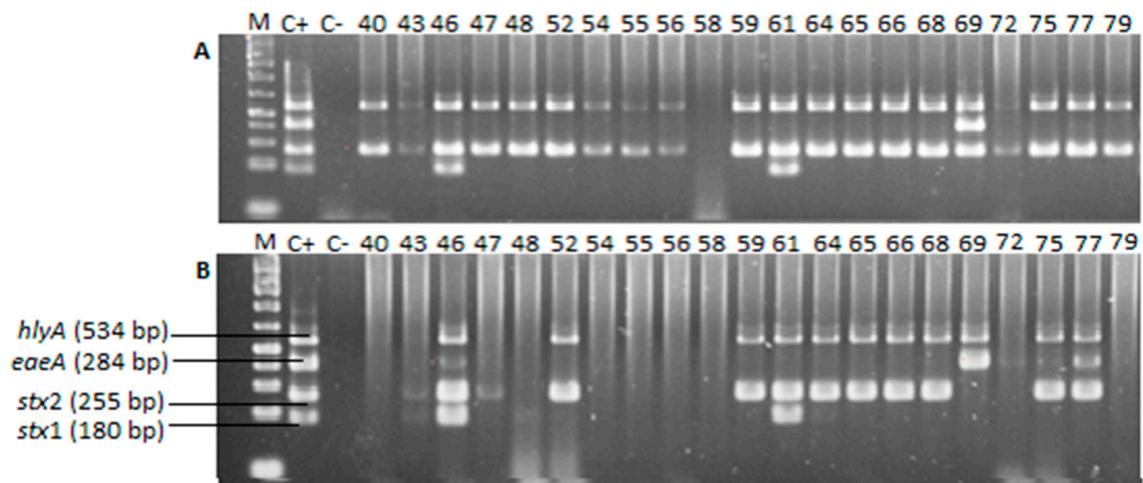


Figure S2: Multiplex gel image of amplicons indicating no detection of *stx2* (gel B). M indicates molecular marker 1kb plus, C+ the positive control, C- the negative control and isolate number (40-79).