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Table S1. Additional antibiotic and genomic traits of the clinical isolates of ESBL-producing *Escherichia coli* strains

Sample	Strain	MIC/(mg/mL)									Beta-lactamase content	DNA profile	
		CAZ	CTX	CRO	FEP	IMI	MEM	GM	CIP	TZP		Plasmid type	PFGE cluster
1	3459	32 (R)	16 (R)	128 (R)	2 (I)	0.06 (S)	0.06 (S)	4 (S)	128 (R)	4 (S)	CTX-M-15, TEM-1	X, FIA	E XXIV
2	7377	8 (I)	8 (R)	>128 (R)	64 (R)	0.06 (S)	0.06 (S)	64 (R)	>128 (R)	4 (S)	CTX-M-15, TEM-1	FIA	E XVIIIb
3	16416	4 (S)	16 (R)	128 (R)	4 (R)	0.06 (S)	0.06 (S)	64 (R)	128 (R)	2 (S)	CTX-M-15	X, FIA	E XXXII
4	1358	64 (R)	>128 (R)	>128 (R)	64 (R)	0.06 (S)	0.06 (S)	128 (R)	>128 (R)	32 (S)	CTX-M-15, TEM-1	X	E IX
5	23407	64 (R)	>128 (R)	>128 (R)	64 (R)	0.06 (S)	0.06 (S)	128 (R)	128 (R)	2 (S)	CTX-M-15, TEM-1	L/M	E XXVII
6	3263	16 (R)	128 (R)	64 (R)	64 (R)	0.06 (S)	0.06 (S)	64 (R)	>128 (R)	1 (S)	CTX-M-15	FIA	E II
8	21646	16 (R)	64 (R)	16 (R)	4 (R)	0.06 (S)	0.06 (S)	1 (S)	>128 (R)	0.5 (S)	CTX-M-15	X	E XXVII
11	8537	8 (I)	128 (R)	>128 (R)	64 (R)	0.06 (S)	0.06 (S)	128 (R)	>128 (R)	4 (S)	CTX-M-15	NEG	E XIX
12	4145	128 (R)	>128 (R)	>128 (R)	16 (R)	0.25 (S)	0.25 (S)	2 (S)	32 (R)	2 (S)	CTX-M-15, TEM-1	N	E XIV

ESBL=extended-spectrum beta-lactamase, MIC=minimum inhibitory concentration, CAZ=ceftazidime, CTX=cefotaxime, CRO=ceftriaxone, FEP=cefepime, IMI=imipenem, MEM=meropenem, GM=gentamicin, CIP=ciprofloxacin, TZP=piperacillin-tazobactam. R=resistant, I=intermediately sensitive, S=susceptible. The isolation and characterisation of the *E. coli* strains was reported earlier (29)