Supplementary Table S1. Salient Features of Polymerase Chain Reaction Primer Pairs Used in this Study

Target	*Primer: sequence (5′–3′)*	*Product (bp)*	*Reference*
bla$_{CTX\text{-}M}$ gp1[a]	CTX-M Gp1.F: GGAATCTGACGCTGGGTAAA CTX-M Gp1.R: GGTTGAGGCTGGGTGAAGTA	232	16
bla$_{CTX\text{-}M}$ gp1[a]	CTX-M-F: TTCGTCTCTTCCAGAATAAGG[b] CTX-M-R: CAGCACTTTTGCCGTCTAAG	968	43
bla$_{CTX\text{-}M}$ gp2	MultiCTXMGp2_for: CGTTAACGGCACGATGAC MultiCTXMGp1-2_rev: CGATATCGTTGGTGGTRCCAT	404	13
bla$_{CTX\text{-}M}$ gp9	MultiCTXMGp9_for: TCAAGCCTGCCGATCTGGT MultiCTXMGp9_rev: TGATTCTCGCCGCTGAAG	561	13
bla$_{CTX\text{-}M}$ gp8/25	CTX-Mg8/25_for: AACRCRCAGACGCTCTAC CTX-Mg8/25_rev: TCGAGCCGGAASGTGTYAT	326	13
16S rDNA	Bac331F: TCCTACGGGAGGCAGCAGT Bac797R: GGACTACCAGGGTATCTAATCCTGTT	466	37
Vector MCS	M13F: GTTTTCCCAGTCACGAC M13R: CAGGAAACAGCTATGAC	Variable	Promega

[a]*bla*$_{CTX\text{-}M}$ gp1 primers from Ellem *et al.*[16] were used for quantitative PCR, while *bla*$_{CTX\text{-}M}$ gp1 primers from Pfeifer *et al.*[43] were used for conventional PCR and cloning.

[b]The originally published CTX-M-F primer in Pfeifer *et al.*[43] had an unintended G at the 5′ end (Yvonne Pfeifer, personal communication), which was removed here.

Promega, Corp. 2014. pGEM-T and pGEM-T Easy Vector Systems Technical Manual.

PCR, polymerase chain reaction.