

Primers for *E. cloacae* MLST scheme

	Name	Sequence (5'->3')	Position in the target gene
Amplification primers	dnaA-f2	AYAACCCGCTGTTCTBTATGGCGGCAC	500-527*
	dnaA-r	KGCCAGCGCCATCGCCATCTGACGCGG	1222-1248*
	fusA-f2	TCGCGTTCGTTAACAAAATGGACCGTAT	413-440*
	fusA-r2	TCGCCAGACGGCCCAGAGCCAGACCCAT	1291-1318
	gyrB-f	TCGACGAAGCGCTCGCGGGTCACTGTAA	143-170
	gyrB-r	GCAGAACCGCCCCGCGGAGTCCCCTTCCA	1268-1295
	leuS-f2	GATCARCTSCCGGTKATCCTGCCGGAAG	1342-1369*
	leuS-r	ATAGCCGCAATTGCGGTATTGAAGGTCT	2159-2186*
	pyrG-f	AYCCBGAYGTBATTGCRCAYMAGGCGAT	56-83*
	pyrG-r	GCRCGRATYTCVCCCTSHTCGTCCCAGC	563-590*
	rplB-f	GTAAACCGACATCTCCGGGTCGTCGCCA	17-44*
	rplB-r	ACCTTTGGTCTGAACGCCCCACGGAGTT	735-762*
	rpoB-f	AAAAACGTATTCGTAAGGATTTTGGTAA	252-280*
	rpoB-r2	CCAGCAGATCCAGGCTCAGCTCCATGTT	973-1000*
Sequencing primers*	gyrB-r3-seq	GCAGAACCGCCCCGCGGAGTCCCCTTCC	1269-1295*
	gyrB-f3-seq	AAAACCGGTACYATGGTGC GTTTCTGG	484-510*
	fusA-r2-seq	ATCTCTTACGYTTGTTAGCGTGCATCT	1094-1121*

* These primers were used for sequencing of respective amplicons.