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Supplementary Table 1. Oligonucleotide primers and annealing temperature used in this study.

Function	Genes		Nucleotide sequence	Amplicon size (bp)	Annealing temp. (°C)	Reference
ESBL types	*bla*$_{CTX-M-1}$ group	F	GTTACAATGTGTGAGAAGCAG	1,041	60	(Jouini et al., 2007)
		R	CCGTTTCCGCTATTACAAAC			
	bla$_{CTX-M-2}$ group	F	CGACGCTACCCCTGCTATT	832	60	(Jouini et al., 2007)
		R	CAGAAACCGTGGGTTACGAT			
	bla$_{CTX-M-8}$ group	F	GGCGCTGGAGAAAAGCAG	862	60	(Jouini et al., 2007)
		R	GGTTTTATCCCCGACAACC			
	bla$_{CTX-M-9}$ group	F	GTGACAAAGAGAGTGCAACGG	857	60	(Jouini et al., 2007)
		R	ATGATTCTCGCCGCTGAAGCC			
	bla$_{CTX-M-25}$ group	F	GCACGATGACATTCGGG	327	60	(Jouini et al., 2007)
		R	AACCCACGATGTGGGTAGC			
	bla$_{CMY}$	F	AACACACTGATTGCGTCTGAC	1,226	60	(Jouini et al., 2007)
		R	CTGGGCCTCATCGTCAGTTA			
	bla$_{SHV}$	F	TCGCCTGTGTATTATCTCCC	768	54	(Jouini et al., 2007)
		R	CGCAGATAAATCACCACAATG			
	bla$_{TEM}$	F	TCCGCTCATGAGACAATAACC	1,057	58	(Jouini et al., 2007)
		R	ACGCTCAGTGGAACGAAAAC			
	bla$_{OXA}$	F	ACACAATACATATCAACTTCGC	813	60	(Jouini et al., 2007)
		R	AGTGTGTTTAGAATGGTGATC			
Phylogenetic group	*yjaA*	F	CAAACGTGAAGTGTCAGGAG	288	55	(Clermont et al., 2013)
		R	AATGCGTTCCTCAACCTGTG			
	chuA	F	ATGGTACCGGACGAACCAAC	211	55	(Clermont et al., 2013)
		R	TGCCGCCAGTACCAAAGACA			
	tspE4.C2	F	CACTATTCGTAAGGTCATCC	152	55	(Clermont et al., 2013)
		R	AGTTTATCGCTGCGGGTCGC			
	AceK.f	F	AACGCTATTCGCCAGCTTGC	400	55	(Clermont et al., 2013)
	ArpA1.r	R	TCTCCCCATACCGTACGCTA			
	ArpAgpE.f	F	GATTCCATCTTGTCAAAATATGCC	301	55	(Clermont et al., 2013)
	ArpAgpE.r	R	GAAAAGAAAAAGAATTCCCAAGAG			
	trpAgpC.1	F	AGTTTTATGCCCAGTGCGAG	219	55	(Clermont et al., 2013)
	trpAgpC.2	R	TCTGCGCCGGTCACGCCC			
	trpBA.f	F	CGGCGATAAAGACATCTTCAC	489	55	(Clermont et al., 2013)
	trpBA.r	R	GCAACGCGGCCTGGCGGAAG			
Virulence factors	*fimH*	F	TGCAGAACGGATAAGCCGTGG	508	63	(Gonzalez Moreno et al., 2020)
		R	GCAGTCACCTGCCCTCCGGTA			
	iha	F	CTGGCGGAGGCTCTGAGATCA	827	55	(Aazam et al., 2012)
		R	TCCTTAAGCTCCCGCGGCTGA			
	papC	F	GTGGCAGTATGAGTAATGACCGTTA	200	63	(Zhi et al., 2016)
		R	ATATCCTTTCTGCAGGGATGCAATA			
	csgA	F	ACTCTGACTTGACTATTACC	200	55	(Gonzalez Moreno et al., 2020)
		R	AGATGCAGTCTGGTCAAC			
	astA	F	TGCCATCAACACAGTATATCCG	102	65	(Muller et al., 2007)
		R	ACGGCTTTGTAGTCCTTCCAT			
	hlyA	F	AACAAGGATAAGCACTGTTCTGGCT	1,176	63	(Aazam et al., 2012)
		R	ACCATATAAGCGGTCATTCCCGTCA			
	aat	F	CTGGCGAAAGACTGTATCAT	629	53	(Moyo et al., 2007)
		R	CAATGTATAGAAATCCGCTGTT			
	tsh	F	ACTATTCTCTGCAGGAAGTC	824	55	(Aazam et al., 2012)
		R	CTTCCGATGTTCTGAACGT			
	pic	F	AGCCGTTTCCGCAGAAGCC	1,111	63	(Muller et al., 2007)
		R	AAATGTCAGTGAACCGACGATTGG			
	traT	F	GGTGTGGTGCGATGAGCACAG	290	60	(Gonzalez Moreno et al., 2020)
		R	CACGGTTCAGCCATCCCTGAG			
	ompT	F	ATCTAGCCGAAGAAGGAGGC	559	64	(Aazam et al., 2012)

		R	CCCGGGTCATAGTGTTCATC			
	fyuA	F	TGATTAACCCCGCGACGGGAA	880	63	(Gonzalez Moreno et al., 2020)
		R	CGCAGTAGGCACGATGTTGTA			
	iroNe.coli	F	AAGTCAAAGCAGGGGTTGCCCG	665	63	(Aazam et al., 2012)
		R	GACGCCGACATTAAGACGCAG			
Antimicrobial resistance	*catA*	F	AGTTGCTCAATGTACCTATAACC	547	57	(Saenz et al., 2004)
		R	TTGTAATTCATTAAGCATTCTGCC			
	cmlA	F	CCGCCACGGTGTTGTTGTTATC	698	57	(Saenz et al., 2004)
		R	CACCTTGCCTGCCCATCATTAG			
	floR	F	TATCTCCCTGTCGTTCCAG	399	52	(Saenz et al., 2004)
		R	AGAACTCGCCGATCAATG			
	tetA	F	GCTACATCCTGCTTGCCTTC	210	58	(Saenz et al., 2004)
		R	CATAGATCGCCGTGAAGAG			
	tetB	F	TTGGTTAGGGGCAAGTTTTG	659	56	(Saenz et al., 2004)
		R	GTAATGGGCCAATAACACCG			
	tetD	F	AAACCATTACGGCATTCTGC	787	60	(Saenz et al., 2004)
		R	GACCGGATACACCATCCATC			
	qnrA	F	ATTTCTCA CGCCAGGATTTG	516	53	(Liao et al., 2015)
		R	GATCGGCAAAGGTTAGGTCA			
	qnrB	F	GATCGTGAAAGCCAGAAAGG	469	53	(Liao et al., 2015)
		R	ACGATGCCTGGTAGTTGTCC			
	qnrC	F	GGGTTGTACATTTATTGAATC	447	50	(Liao et al., 2015)
		R	TCCACTTTACGAGGTTCT			
	qnrS	F	ACGACATTCGTCAACTGCAA	417	53	(Liao et al., 2015)
		R	TAAATTGGCACCCTGTAGGC			
	aac(6)-Ib-cr	F	TTGCGATGCTCTATGAGTGGCTA	482	50	(Liao et al., 2015)
		R	CTCGAATGCCTGGCGTGTTT			
	aac(3)-I	F	ACCTACTCCCAACATCAGCC	169	60	(Saenz et al., 2004)
		R	ATATAGATCTCACTACGCGC			
	aac(3)-II	F	ACTGTGATGGGATACGCGTC	237	60	(Saenz et al., 2004)
		R	CTCCGTCAGCGTTTCAGCTA			
	aac(3)-IV	F	CTTCAGGATGGCAAGTTGGT	286	60	(Saenz et al., 2004)
		R	TCATCTCGTTCTCCGCTCAT			
	dfrIa	F	GTGAAACTATCACTAATGG	474	55	(Saenz et al., 2004)
		R	TTAACCCTTTTGCCAGATTT			
	dfrIb	F	GAGCAGCTICTITTIAAAGC	393	60	(Saenz et al., 2004)
		R	TTAGCCCTTTIICCAATTTT			
	dfrII	F	GATCACGTGCGCAAGAAATC	141	50	(Saenz et al., 2004)
		R	AAGCGCAGCCACAGGATAAAT			
	dfrVII	F	TTGAAAATTTCATTGATT	474	55	(Saenz et al., 2004)
		R	TTAGCCTTTTTTCCAAATCT			
	dfrXII	F	GGTGSGCAGAAGATTTTTCGC	319	60	(Saenz et al., 2004)
		R	TGGGAAGAAGGCGTCACCCTC			
Replicon types	IncHI1	F	GGAGCGATGGATTACTTCAGTAC	471	60	(Carattoli et al., 2005)
		R	TGCCGTTTCACCTCGTGAGTA			
	IncHI2	F	TTTCTCCTGAGTCACCTGTTAACAC	644	60	(Carattoli et al., 2005)
		R	GGCTCACTACCGTTGTCATCCT			
	IncI1-Iγ	F	CGAAAGCCGGACGGCAGAA	139	60	(Carattoli et al., 2005)
		R	TCGTCGTTCCGCCAAGTTCGT			
	IncI2	F	CTGTCGGCATGTCTGTCTC	553	55	(Lv et al., 2013)
		R	CTGGCTACCAGTTGCTCTAA			
	IncX1	F	GCTTAGACTTTGTTTTATCGTT	461	62	(Johnson et al., 2012)
		R	TAATGATCCTCAGCATGTGAT			
	IncX2	F	GCGAAGAAATCAAAGAAGCTA	678	63	(Johnson et al., 2012)
		R	TGTTGAATGCCGTTCTTGTCCAG			
	IncX3	F	GTTTTCTCCACGCCCTTGTTCA	351	63	(Johnson et al., 2012)
		R	CTTTGTGCTTGGCTATCATAA			
	IncX4	F	AGCAAACAGGGAAAGGAGAAGACT	569	62	(Johnson et al., 2012)
		R	TACCCCAAATCGTAACCTG			
	IncL/M	F	GGATGAAAACTATCAGCATCTGAAG	785	60	(Carattoli et al., 2005)

	R	CTGCAGGGGCGATTCTTTAGG			
IncFIA	F	CCATGCTGGTTCTAGAGAAGGTG	462	60	(Carattoli et al., 2005)
	R	GTATATCCTTACTGGCTTCCGCAG			
IncFIB	F	GGAGTTCTGACACACGATTTTCTG	702	63	(Carattoli et al., 2005)
	R	CTCCCGTCGCTTCAGGGCATT			
IncFIC	F	GTGAACTGGCAGATGAGGAAGG	262	60	(Carattoli et al., 2005)
	R	TTCTCCTCGTCGCCAAACTAGAT			
IncFIIs	F	CTGTCGTAAGCTGATGGC	270	60	(Carattoli et al., 2005)
	R	CTCTGCCACAAACTTCAGC			
IncA/C	F	GAGAACCAAAGACAAAGACCTGGA	465	60	(Carattoli et al., 2005)
	R	ACGACAAACCTGAATTGCCTCCTT			
IncP	F	CTATGGCCCTGCAAACGCGCCAGAAA	534	60	(Carattoli et al., 2005)
	R	TCACGCGCCAGGGCGCAGCC			
IncK	F	GCGGTCCGGAAAGCCAGAAAAC	160	60	(Carattoli et al., 2005)
	R	TCTTTCACGAGCCCGCCAAA			
IncB/O	F	GCGGTCCGGAAAGCCAGAAAAC	159	60	(Carattoli et al., 2005)
	R	TCTGCGTTCCGCCAAGTTCGA			
IncN	F	GTCTAACGAGCTTACCGAAG	559	55	(Carattoli et al., 2005)
	R	GTTTCAACTCTGCCAAGTTC			

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