

Oligonucleotide sequences used as PCR primers.

Targets	Primers	Nucleotide Sequence (5'-3')	Fragm ent Size	References
CTX-M-1	<i>bla</i> _{CTX-M} -group1-F	GTTACAATGTGTGAGAACAG	1041bp	(1, 2)
	<i>bla</i> _{CTX-M} -group1-R	CCGTTCCGCTATTACAAAC-3		
CTX-M-2	<i>bla</i> _{CTX-M} -group2-F	TGATGAGACATCGCGTTAAG	875bp	(1, 2)
	<i>bla</i> _{CTX-M} -group2-R	TAACCGTCGGTGACGATT		
CTX-M-8	<i>bla</i> _{CTX-M} -group8-F	ATGATGAGACATCGCGTTAAG	865bp	(3)
	<i>bla</i> _{CTX-M} -group8-R	CGGTGACGATTTCGCGGCAG		
CTX-M-9	<i>bla</i> _{CTX-M} -group9-F	ATGGTGACAAAGAGAGTGCA	913bp	(3)
	<i>bla</i> _{CTX-M} -group9-R	TTACAGCCCTTCGGCGATGA		
TEM	<i>bla</i> _{TEM} -F	ATTCTGAAGACGAAAGGGC	1150bp	(1, 2)
	<i>bla</i> _{TEM} -R	ACGCTCAGTGGAACGAAAAC		
SHV	<i>bla</i> _{SHV} -F	CACTCAAGGATGTATTGTG	885bp	(1, 2)
	<i>bla</i> _{SHV} -R	TTAGCGTTGCCAGTGCTCG		
OXA-1	OXA-1-F	ACACAATACATATCAACTTCGC	813bp	(4)
	OXA-1-R	AGTGTGTGTTAGAATGGTGATC		
OXA-2	OXA-2-F	TTCAAGCCAAGGCACGATAG	702bp	(5)
	OXA-2-R	TCCGAGTTGACTGCCGGGTTG		
MOX-1, MOX-2, CMY-1, CMY-8 to CMY-11	MOXMF	GCTGCTCAAGGAGCACAGGAT	520bp	(6)
	MOXMR	CACATTGACATAGGTGTGGTGC		
LAT-1 to LAT-4, CMY-2 to CMY-7, BIL-1	CITMF	TGGCCAGAACTGACAGGCAA	462bp	(6)
	CITMR	TTTCTCCTGAACGTGGCTGGC		
DHA-1, DHA-2	DHAMF	AACTTCACAGGTGTGCTGGGT	405bp	(6)
	DHAMR	CCGTACGCATACTGGCTTG		
FOX-1 to FOX-5b	FOXMF	AACATGGGGTATCAGGGAGATG	190bp	(6)
	FOXMR	CAAAGCGCGTAACCGGATTGG		
CMY-2	CMY-2F	AACACACTGATTGCGTCTGAC	1226bp	(6)
	CMY-2R	CTGGGCCTCATCGTCAGTTA		
sul1	sul1-F	TGGTGACGGTGTTCGGCATTC	789bp	(2, 4, 7)
	sul1-R	GCGAGGGTTCCGAGAAGGTG		
sul2	sul2-F	CGGCATCGTCAACATAACC	722bp	(4, 7)
	sul2-R	GTGTGCGGATGAAGTCAG		
sul3	sul3-F	GAGCAAGATTTTGGAAATCG	990bp	(4, 7)
	sul3-R	CATCTGCAGCTAACCTAGGGCTTGGA		
aadA1 or aadA2	aadA-F	GCAGCGCAATGACATTCTG	282bp	(4, 7)
	aadA-R	ATCCTTCGGCGCGATTTG		
tet(A)	tet(A)-F	GTAATTCTGAGCACTGTCGC	937bp	(4, 7)
	tet(A)-R	CTGTCCTGGACAACATTGCTT		
tet(B)	tet(B)-F	CTCAGTATTCCAAGCCTTG	416bp	(4, 7)

	tet(B)-R	CTAACACTGTCTCCTGTT		
<i>tet(C)</i>	tet(C)-F	TCTAACAAATGCGCTCATCGT	570bp	(4, 7)
	tet(C)-R	GGTTGAAGGCTCTCAAGGGC		
<i>aph(3')-Ia</i>	aph(3')-Ia-F	ATGGGCTCGCGATAATGTC	600bp	(4, 7)
	aph(3')-Ia-R	CTCACCGAGGCAGTCCAT		
<i>aac(3')-IId</i>	AacC2-F	ACTGTGATGGGATACCGGTC	237bp	(4)
	AacC2-R	CTCCGTCAGCGTTTCAGCTA		
<i>aac(3)-IVa</i>	AacC4-F	CTTCAGGATGGCAAGTTGGT	286bp	(4)
	AacC4-R	TCATCTCGTTCTCCGCTCAT		
<i>qepA</i>	qepA-F	GGACATCTACGGCTTCTCG	199bp	(2)
	qepA-R	CAACTGCTTGAGCCCCTAG		
<i>qnrA</i>	qnrA-F	AGAGGATTCTCACGCCAGG	580bp	(2, 8, 9)
	qnrA-R	TGCCAGGCACAGATCTTGAC		
<i>qnrB</i>	qnrB-F	GGMATHGAAATTGCCACTG	264bp	(2, 8, 9)
	qnrB-R	TTTGCYGYYCGCCAGTCGAA		
<i>qnrS</i>	qnrS-F	GCAAGTTCATTGAACAGGGT	428bp	(2, 8, 9)
	qnrS-R	TCTAAACCCTCGAGTCGGCG		
<i>aac(6')-Ib-cr</i>	<i>aac (6')-Ib-cr-F</i>	TTGCGATGCTCTATGAGTGGCTA	482bp	(10)
	<i>aac (6')-Ib-cr-R</i>	CTCGAATGCCTGGCGTGT		
<i>strA</i>	strA-F	CCTGGTGATAACGGCAATT	546bp	(11)
	strA-R	CCAATCGCAGATAGAACGGC		
<i>strB</i>	strB-F	ATCGTCAAGGGATTGAAACC	509bp	(11)
	strB-R	GGATCGTAGAACATATTGGC		
<i>dfrA1, dfrA5,</i>	Dfrla-F	GTGAAACTATCACTAATGG	474bp	(4)
<i>dfrA15, dfrA15b,</i>	Dfrla-R	TTAACCCCTTTGCCAGATT		
<i>dfrA16, dfrA16b</i>				
<i>dfrA14, dfrA6</i>	Dfrlb-F	GAGCAGCTICTITIAAGC	393bp	(4)
	Dfrlb-R	TTAGCCCTTTIICCAATT		
<i>dfrA7, dfrA17</i>	DfrVII-F	TTGAAAATTCATTGATT	474bp	(4)
	DfrVII-R	TTAGCCTTTTCCAAATCT		
<i>dfrA12, dfrA13</i>	DfrXII-F	GGTGCGCAGAAGATTTCGC	319bp	(4)
	DfrXII-R	TGGGAAGAAGGCACCCCTC		
<i>cmlA</i>	CmlA-F	TGTCATTACGGCATACTCG	455bp	(4)
	CmlA-R	ATCAGGCATCCCATTCCAT		
<i>floR</i>	FloR1	CACGTTGAGCCTCTATAT	868bp	(4)
	FloR2	ATGCAGAAGTAGAACGCG		
<i>intI1</i>	<i>intI1-F</i>	CACTCCGGCACCGCCAAC	545bp	(12)
	<i>intI1-R</i>	TTTC		
		GAACGGGCATGCGGATCAGTGAG		
<i>mcr-1</i>	mcr1_320bp_fw	AGTCGTTGTTCTGTGGC	320bp	(13)
	mcr1_320bp_rev	AGATCCTGGTCTCGGCTTG		
<i>mcr-2</i>	mcr2_700bp_fw	CAAGTGTGTTGGTCGCAGTT	715bp	(13)
	mcr2_700bp_rev	TCTAGCCCGACAAGCATACC		
<i>mcr-3</i>	mcr3_900bp_fw	AAATAAAAATTGTTCCGCTTATG	929bp	(13)
	mcr3_900bp_rev	AATGGAGATCCCCGTTTT		

<i>mcr-4</i>	mcr4_1100bp_fw	TCACCTTCATCACTGCGTTG	1116bp	(13)
	mcr4_1100bp_rev	TTGGTCCATGACTACCAATG		
<i>mcr-5</i>	MCR5_FW	ATGCGGTTGTCTGCATTTATC	1644bp	(14)
	MCR5_RV	TCATTGTGGTTGTCCTTTCTG		

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