

Table S1. Primers and probes targeting antimicrobial resistance genes (ARGs) and mobile genetic elements (MGEs).

Target	Primers and probes sequences	Reference
aac(6')-Ib	F-AACTTGCGAGCGATCCGA R-TGGCGTGTTTGAACCATGTAC P-TACCTTGCTTCTCAAACCCCGCTTTCTC	[22]
aac(6')-Ib-cr	F-AACTTGCGAGCGATCCGA R-TGGCGTGTTTGAACCATGTAC P-MGB-TACGGTACCTTGCTCT	[22]
aac(3)-IIa	F-AGGACATCGTGACGTTCCG R-AACCTGAAGGCTCGCAAGAG P-CGGAGTGGTTCGAAATGCTTCTCAAG	[22]
aac(3)-IV	F-ATCGGTCAGCTTCTCAACCT R-TCGTCCAGACCTGACCAC P-TCCACAGCTCCTTCCGTAGCGTCC	[22]
armA	F-GAAAGAGTCGCAACATTAAATGACTT R-GATTGAAGCCACAACCAAAATCT P-TCAAACATGTCTCATCTATT	[23]
ant(2'')-Ia	F-CGGCGAGCTCGAGGC R-ATTCATACGCTTCGTCTGCC P-TCATGGAGGAGTTGGACTATGGATTCTTAGCG	[22]
ant(3'')-Ia	F-ACGTTGTCCCGCATTTGG R-TCCTTCGGCGCGATTTT P-MGB-ACAGCGCAGTAACC	[22]
aph(3')-Ia	F-ACCGAGGCAGTTCATAGGA R-ACCTTATTTTTGACGAGGGGAAA P-ATCCTGGTATCGGTCTGCGATTCCGA	[22]
strA	F-TCAATCCCGACTTCTTACCGG R-CACCATGGCAAACAACCATATC P-MGB-TGGCTCGTGTCGAAC	[22]
strB	F-ATCGCTTTCGAGCTTTGTTTC R-TCCGAGGCATTGCTCATCATT P-CGTCCACGCGGCGATTATAGCC	[22]
blaTEM	F-GGTATTATCCCGTRTTGACGC R-GTCCCTCCGATCGTTGTCAGAA P-ACWCACCAGTCACAGAAAAGCATCTTMCG	This study
blaCTX-M-1group	F-AATCTGACGCTGGGTAAAGCR R-CGTTGGTGGTRCCATAGYCAC P-AGTCCWGCCTGAATRCTCGCTGCA	This study
blaCTX-M-2group	F-CGACGCTACCCCTGCTATT R-CATCACCTTACTGGTACTGCA P-AACGCCAAGCCGACCTCCCGAACTT	This study
blaCTX-M-8group	F-TAAGCGGATGATGCTAATGACAA R-AACGTCGTTGCGCTGCGCA P-CACTCCCCAGCAACAGCGAAATACAG	This study
blaCTX-M-9group	F-CGTGGCTCAAAGGCAATACGA R-ATAGGTCACCAGAACCAGCG P-ACCGCAATATCATTTGGTGGTGCCGTAG	This study
blaKPC2/3	F-AGCGGCAGCAGTTTGTTGAT R-ACGGCCAACACAATAGGT	[22]

blaNDM1/2	P-CAGTCGGAGACAAAACCGGAACCTGC F-CGCAACACAGCCTGACTTT R-CAGCCACCAAAAAGCGATGTC	[22]
blaPER1	P-CCAGCAACCGCGCCCAACTTTGG F-GGCTAAGGTTTTACAGAATACCT R-GTCCACCAACCAGTTCAAATAAC	This study
blaVIM	P-CGTATCAGGGAGACSAGTTTAGTGTTC F1-GAGATTCCCACGCACTCTCTAGA F2-GAGATTCCCACGATTCTCTAGA	[24]
blaSHV	R-AATGCGCAGCACCAGGATAG P-ACGCAGTGCCTTCGGTCCAGT F-TGACRAACAGCTGGAGCGAA	This study
blaCMY-1	R-GCTGTTATCGCTCAYGGTAATG P-TCCACTATMGCCAGCAGSATCTGG F-TGCTCAAGGATGGCAAGGC	This study
blaCMY-2	R-TCCCTTGACCACCGCATAG P-AGGGTCGCAGTCAGGGTCTTGCTCA F-AACCCTCAGGAATGAGTTACG	This study
blaCMY-3	R-CAAGTTGTCCCGGAGAAACG P-CAATGACCAGACGCGTCCTGCAACC F-GCTAACTCCAGYATTGGTCTG	This study
floR	R-CCAGGTATGCGCCAGTTTTAA P-GACGCGTYTGGTCAATGCCTCTTC F-AGATCGGATTCAGCTTTGCC	[22]
qnrA	R-CAAAGGACTTCGCGAAACG P-CGCCACTGTCGCGCTTGTAATGATC F-AGGATTGCAGTTTCATTGAAAGC	[25]
qnrB1	R-TGAACTCTATGCCAAAGCAGTTG P-TATGCCGATCTGCGCGA F-GTAGCGCATATATCACGAATACCAATC	[22]
qnrS	R-ATCTGAACCACTGAACGTCGC P-AAGTCGTGTTGGAAAAGTGTGAGCTGTGG F-CGACGWGCTAACTTGCGTGA	[22]
mcr-1	R-GGCATTGTTGGAACTTGCA P-TCATTGAACAGGGTGATATCGAAGGCTGC F-TGGCGTTCAGCAGTCATTAT	[26]
mcr-2	R-AGCTTACCCACCGAGTAGAT P-AGTTTCTTTCGCGTGCATAAGCCG F-TTGTCGTGCTGTTATCCTATCG	This study
mcr-3	R-CCGTGCCATAAGTATCGGTAAA P-ACTGATTATGGGTGCGGTGACGAG F-CTTGCWGAACCAATCCCATTAC	This study
mcr-4	R-TTACCYCGAGCYGTTTCACC P-TAAGCCCACGTTGATGTTYCTGGT F-GCTGGCTTGGGTATCTAATGAC	This study
mcr-5	R-AAGTCCTAGCAAACGTGCGAAC P--ACAGCGAGCAGAACAGGGCGGCTTT F-GACTCTGTTCAATCAGCCTTGTC	This study
mcr-9	R-CCCAGTGCAAAGGAGCATTAG P-TCACCCTGCTGTGCAATGGCGTGT F-TGTTTATCACTGGCTCACTGGT	This study

	R-GGATGGCAGGAAGGATACC	
	P-ACCGGTAAAGACGAACCATATAATAATTGAGC	
sul1	F-CCGTTGGCCTTCCTGTAAAG	[22]
	R-TTGCCGATCGCGTGAAGT	
	P-CAGCGAGCCTTGCGGCGG	
sul2	F-CGGCTGCGCTTCGATT	[22]
	R-CGCGCGCAGAAAGGATT	
	P-MGB-TGCTTCTGTCTGTTTCG	
sul3	F-TAGGCTGCAAAGATAGGGAAAATG	[22]
	R-CAACGCCCACTTCAGTTGTATC	
	P-TGGAGCAGATGTGATTGATTGTTGGGAGC	
tetA	F-TCTGAGCACTGTCGCGCTC	[22]
	R-ACAGGTGCGCAGGCAAA	
	P-TGGTTCACCTCGAACGACGTCACCG	
tetB	F-TGTGGCAGGAAGAATAGCCAC	[22]
	R-ATCCCTGTAAAGCACCTTGCTG	
	P-TTGGCTGGTGGTGGGATCGCTT	
dfrA1	F-TCGAAGAATGGAGTTATCGGGA	[22]
	R-TGTCAGATGTAAACTTGAACGTGTTA	
	P-TGGGAGCATTACCCAACCGAAAGTATGC	
dfrA12	F-TGCTGCGATGGGAGCC	[22]
	R-GACGGTTCGGTAGAGGCTTG	
	P-TTCGCAGACTCACTGAGGGAAAAGTCGT	
dfrA17	F-GCGTAATCGGTAGTGGTCCCTGA	[22]
	R-TTCCGTTCTTTGACACTACTGCAT	
	P-TTGACTCTATGGGTGTTCTTCCAAATCGCA	
IncA/C	F- GAACCAAAGACAAAGACCTGG	This study
	R-TGCTCACGCTTTTGGTCCCAA	
	P-CTCGACGTWATCAAAGACTCACCGCAAATG	
IncFIA	F-ATGGAYCTGTCATGGCGGAA	This study
	R-AACGAACAGATACAGCAKACG	
	P-CGTAAAAATAGCCCCGGRATYGTCCAGT	
IncFIB-AP001918	F-AACTGTTCCGGCGAGGTGGA	This study
	R-CAGCTCTTCCGTGGCATCA	
	P-CACTGACACCAAACAATAACAACACCGTAC	
IncFIB-K	F-ACTCGGTGTAATCCAGATAACC	This study
	R-AGCGCAAGGAATCAGCCCA	
	P-ATCGCCTTACGCACCGTGTGGTTC	
IncFIB-pLF82pHCM2	F-TGAACTCAAACAGCTGTCTCTG	This study
	R-GTCCAGGCGYTTGCGTAAC	
	P-AGCCGAAGGGTACGARGACATAAGGATC	
IncFIB-pQIL	F-TCGATGGAGACTGACTTCAAG	This study
	R-CAGACCAGATACAACACCATTC	
	P-CTTCCATTACGCGAGTTTGCTACATTMTGC	
IncFIB-S	F-CTCCGGTGAAATCGTAAATCTC	This study
	R-GATCTCCACTTTATCAAACCCC	
	P-CTCTGATGCGGCTTGAGTATTTGTTCC	
IncFII	F-GCACACCATCCTGCACTTAY	This study
	R-CTATGCGGGGAGTATAGTYATG	
	P-CRCAGAAGGAGYGAGCACAGAAAGAAGTC	
	F-TAAGTGTTGGCTCGGTTGCATA	This study

IncHI1A- AF250878	R-TGCTAGCGCCTCCTGAAATC P-TCTACAAACATAAAACAAGCGAACTCGCTCA	
IncHI1B-CIT-27	F-GAAAACCGATCTCTTTAAGCTGG R-TTCGAAGCTCAATACGCTGTAG P-ACAGGCCCTGTAYCTGTATCTGGCTGA	This study
IncFIA(HI1)	F-TTGCCCTCAGGAAGCGTTG R-CGCCTCAGTCAGCTCATTGTA P-AGTCTGTCACACTTCACATGGACCTGTC	This study
IncHI1B-MAR	F-TCTTCCCGTCGACACGATAC R-CACACCCATTTCCAGTTCATC P-TCCGGATTTGGCCTGATAATGACCTCATC	This study
IncHI2-1	F-ACCTTGCATCGGTAGTGCTTG R-CGATACAGTTCCACATAGAAG P-CTCGKAGTGGTCGGTTCACAGTAACG	This study
IncHI2-2	F-CATGATGTGTACCTTACTGGCA R-AACCGGAAGCCGTATTTCG P-CATTGTCCTCCGAGTGTGCGTTCAGC	This study
IncI2	F-GGCTTGAACATCGTTGATCGAT R-CCTGGAAAASACGAGATTACG R-AGGCAAGTTTTCCGCGATTGTAAATGCAG	This study
IncI-I1	F-GGACGGCAGAATGCGCCAT R-TGGGGGTTTTTCCTTTTATGGC P-ATCTTCRGTCTTCTGACTTACTGYCCG	This study
IncK/B/O/Z	F-TGCGCCATAAGGCATTTCAGG R-CACAAGCTGTCGCTTATGGC P-CGGTTTCATATATACTTATCCCGTATTCTGTG	This study
IncLM-pOXA48- AF550415	F-AGCGCCAGGAACAGAAGAGT R-GCGAACCCGCTTTTACATAGTA P-CGGAGTTCCAGAGAGAGTACCGGA	This study
IncL/M-pMU407	F-CTGAAGAGTAAACTGACAGACG R-CACCTGCCGCTCAACTACT P-CACAAGCCCTCCGACTACAAACGACTA	This study
IncN-1	F-TTTGAGTTAAGCCGTCTGCG R-GGATCGTCCGTGGATTTCTGA P-ATCGCGGAGCGAGTAGGTGGTGAAC	This study
IncN-2	F-GTTTCTTCACCTTATGGGGATG R-CAGCTGCAAACGACACGCG P-CACAGGGAACCCGTCATAAAACGCTG	This study
IncQ-1	F-GTGCCTGGAAGAACTCAGAG R-AGGGCCAGCATGGATTACC P-CTAACCACCAACTGCCTGCAAAGGAG	This study
IncR	F-CCAGCCACGGACGTTTAAAC R-TGTGGTTATGCCTCATGCAGG P-CCGTTGATTGCGCCAGAGCATTTACC	This study
IncX4-1	F-GACTTCCCTTTCTCGCTTAAC R-TAGAGTTTATCTGGCACATGGC P-TTTAGCAGCCATAGGCAATGTTGACAGCG	This study
IS1294	F-GTCAACTGTCCGCTCAATAATG R-CTTGCCAGACATCAGTAATGGA P-CCGATAACACACCGGCAACAACAAGA	This study
IS26-1	F-TGTTGCAAATAGTCGGTGGTG	This study

IS26-2	R-CATTCACTCCGCGTTCAGC	This study
	P-ATTCACTCCGCGTTCAGCCAGCATCT	
	F-ATGGAGCTGCACATGAACCC	
IS903	R-CTCACGGTAACTGATGCCG	This study
	P-TTTGCAGTACCAGCGTACGGCCAC	
	F-TTCAGGCAATACGCACGCTTT	
IS903B	R-ACAACAGATTACAACCGTCGC	This study
	P-TTCGTCATTTTGTTCAGCGCTCGTACCA	
	F-GTTGCATCTGGCCGTTGAC	
IS4321-L	R-CCGCAGTTCATCGTGACAGA	This study
	P-TGCAGACCTGTCGCTGAACAATGTGAC	
	F-CCGGCGTCTGAATGGGATT	
IS4321-R	R-AAGCTGGCCAGAATAGCCTG	This study
	P-CGCCTGATGAACCTCCAGAAAAATATACG	
	F-TCTCAGCAGGCAATGCGNG	
IS6100	R-CATCTTCGTCCAAC TTTCGTTTC	This study
	P-TGCATCGTGTCCGTGAATCCYTGGT	
	F-TCAGAGGTAGGCTGTCGCT	
ISAb24	R-AGGCCGATCACGGAAAGCT	This study
	P-CTGCGAAATGGTGGTTGAGCATGCC	
	F-CCTCAATCCTTACCTGATGATG	
ISAb125	R-GACTCGTCTTCAATCAGCTGT	This study
	P-CGGGTAAACCAGCAACTGTCGGAC	
	F-TGGTCTCCTCAGCAAATAGCAA	
ISAp11	R-TCTTGAAC TTTCGGCTGGGC	This study
	P-TCAACGATACGAGCATTACCCAAGGGTG	
	F-GAGTACTTCTACCGACATCT	
ISCR1	R-CTAGAGCGTGTCTGTTTCAGC	This study
	P-CGCGAGAAATGGTGTCTGGAGCTACGT	
	F-AGTCAATCGCCCACTCAAACA	
ISCR3/14/27	R-GATGACAATCCTCGCAGCG	This study
	P-CCTCACTCGAAGCCCAAGGTCAAC	
	F-ATGGATGGGCTGCGGATGA	
ISEc33	R-GCTTGTGGCTTTTCGTGTGCT	This study
	P-ATSACCTACCGCATCGCCACCGG	
	F-ACGACGCCTGATGCAGAAAG	
ISEcp1-1	R-AACGGATGAACGGGCACAACA	This study
	P-CCATGCTRATGCTTCATCGGGGTGAAC	
	F-GTAGAAGGTCATCAAGAAGGTG	
ISEcp1-2	R-GATCATTTCCGCAGCACCG	This study
	P-CTAAGAAACTGGGAAACCGATGCTACAATATC	
	F-TACCGGATTTGTAAGAAGTGGC	
ISSEn4	R-ATCGTCTGATTTGATGTTAGCAAC	This study
	P-AATACTTACACTGCAAACGGTGCTGCGG	
	F-GCATGAACGATATCGCTGAAC	
Tn2	R-CATGGTGTGAGTTGCGCTTA	This study
	P-CCAGTGCTCATTCAAACATTCATCGCGAA	
	F-AGATCAGTTGGGTGCACGAG	
ISKpn14	R-TTCTGAGAATAGTGTATGCGGC	This study
	P-ACAGCGGTAAGATCCTTGAGAGTTTTTCGC	
	F-GTGGCTTCAGTCTCCGTCT	

ISKpn26	R-GAGAAGCGGCATAAGTGAAGG P-CTGTCCCTCCTGTTCCGCTACTGAA F-GAATTTCCGCCACCTGCTC R-CCACTGATTGCCTTTCTTGGT	This study
ISKpn19	P-TGTTCAAGACCATCAATCGCTGGCTGG F-ATACTCAACACCATATCCGGTG R-GAGTCGTCATTTTCATAGGAGATC P-CTGTTTGGCGTTATTCCGGTATCCGG	This study
ISKpn27	F-GGAAATCAGGAACTCGACGCT R-GGCTGGATTGATGAACTCGC P-ACGTCACCCGCCAAACCATCCGAAA	This study
intI1	F-GCCTTGATGTTACCCGAGAG R-GATCGGTCTGAATGCGTGT P-CGACGCCCTTGAGCGGAAGTATC	Modified from [27]
intI2	F-GACGGCTACCCTCTGTTATCTC R-TGCACCATAACAGCAGCGTA P-TGCGAGTATCCATAACCTGCAAAATGCGTT	Modified from [27]
intI3	F-GCCACCACTTGTGTTGAGGA R-GGATGTCTGTGCCTGCTTG P-TGCAGCAAGTGGGTGGCGAATGAGT	Modified from [27]

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