Table S3. Oligonucleotide Sequences, primers, and targets for Polymerase Chain Reaction Amplification of Antibiotic resistance genes and integrons in *Staphylococcus aureus* Isolates

Gene	Primer name	Oligonucleotide sequence (5′→3’)	Amplicon size (bp)	Reference
oqxA	oqxA-F	GGGGAAACATTCACCTGAC	407	This study
	oqxA-R	ACGGGAGACGAGGTTGGTA		
oqxB	oqxB-F	TTCTCCCCCGGCGGGAAGTAC	513	This study
	oqxB-R	CTCGGCCATTTTGGCGCGTA		
ermA	ermA-F	GTTCAAGAACAATCAATACAGAG	421	1
	ermA-R	GGATCAGGAAAAGGACATTTTAC		
ermB	ermB-F	GAAAAGGTACTCAACCAAATA	639	1
	ermB-R	AGTAACGGTACTTAAATTGTTTAC		
ermC	ermC-F	GCTAATATTGTTTAAATCGTCAATTCC	572	1
	ermC-R	GGATCAGGAAAAGGACATTTTAC		
blaZ	blaZ-F	TCAAACAGTTCACATGCC	800	This study
	blaZ-R	TTCATTACACTCTGGCG		
vanA	vanA-F	CATGACGTATCGGTAAAATC	885	2
	vanA -R	ACCGGGCAGCGTATTGAC		
fosB	fosB-F	ACCGGTACTTTACAAGAGCGT	660	This study
	fosB-R	AACAGCACCATCACTTCCTT		
rpoB	rpoB-F	CCGTATCGGTTTATCAAGAATG	432	This study
	rpoB-R	TCAACTTTACGATATGGTGTTT		
cfr	cfr-F	TGAAGTATAAAGCAGGTTGGGAGTCA	746	3
	cfr-R	ACCATATAATTGACCACAAGCAGC		
tetK	tetK-F	TAGGGGGAATAATAGCACATT	587	4
	tetK-R	AATCCGCCCATAACAAATA		
tetM	tetM-F	GAACTCGAACAAGAGGAAAGC	717	4
	tetM-R	ATGGAAGCCCAGAAAGGAT		
aac(6’)-Ib-cr	aac(6’)-Ib-cr -F	TTGCGATGCTCTATGAGTGGCTA	482	This study
	aac(6’)-Ib-cr -R	CTCGAATGCCTGGCGTGTTT		
qepA	qepA-F	CTGCAGGTACTGCGTCATG	403	This study
	qepA-R	CGTGTTGCTGGAGTTCTTC		
qnrA	qnrA-F	CAAGAGGATTTCTCACGCCAG	628	This study
	qnrA-R	AATCCGGCAGCACTATTACTCC		
qnrB	qnrB-F	AGCGGCACTGAATTTATCGG	418	This study
	qnrB-R	CGCAATGTGTGAAGTTTGCT		
qnrC	qnrC-F	GGGTT GTACA TTTAT TGAATCG	449	5

	qnrC-R	AATCC ACTTT ACGAG GTTCT		
qnrD	qnrD-F	CACGAGATCAATTTACGGGGAA	572	This study
	qnrD-R	CGCCTGCTCTCCATCCAAC		
qnrS	qnrS-F	CATTGAACAGGGTGATATCGAA	395	This study
	qnrS-R	ATAAATTGGCACCCTGTAGGC		
gyrA	gyrA-F	AATGAACAAGGTATGACACC	203	This study
	gyrA-R	TACGCGCTTCAGTATAACGC		
gyrB	gyrB-F	CAGCGTTAGATGTAGCAAGC	231	This study
	gyrB-R	CCGATTCCTGTACCAAATGC		
grlA	grlA-F	ACTTGAAGATGTTTTAGGTGAT	538	This study
	grlA-R	TTAGGAAATCTTGATGGCAA		
grlB	grlB-F	CGATTAAAGCACAACAAGCAAG	353	This study
	grlB-R	CATCAGTCATAATAATTACTC		
acc(6')-aph(2″)	acc(6')-aph(2″)-F	CCAAGAGCAATAAGGGCATACC	675	6
	acc(6')-aph(2″)-R	ACCCTCAAAAACTGTTGTTGC		
ant(4')-Ia	ant(4')-Ia-F	GGAAGCAGAGTTCAGCCATG	266	6
	ant(4')-Ia-R	TGCCTGCATATTCAAACAGC		
aphA	aphA-F	ACAGCCGGTATAAAGGGACCACC	382	7
	aphA-R	AAAATCATACAGCTCGCGCGGATC		
vga(A)	vga(A)-F	CTTCAATTGGGATCCTCAGGATAGG	631	8
	vga(A)-R	GTTATGGTACCTTCTTGTTAGG		
vga(B)	vga(B)-F	GAATAAGGCGCAAGGAATGA	601	9
	vga(B)-R	TAGCTTGGCAAAAGCAACCT		
vga(C)	vga(C)-F	TAAGTTCATCGGAAGCAA	671	9
	vga(C)-R	GGATTCAAACGCCTCTAT		
vga(D)	vga(D)-F	CAACTGGAGCGAGCTGTTA	201	10
	vga(D)-R	GACAGCCGGATAATCTTTTG		
lnu(A)	lnu(A)-F	GGTGGCTGGGGGGTAGATGTATTAACTGG	-	11
	lnu(A)-R	GCTTCTTTTGAAATACATGGTATTTTTCGATC		
lnu(B)	lnu(B)-F	CCTACCTATTGTTTGTGGAA	-	11
	lnu(B)-R	ATAACGTTACTCTCCTATTC		
lnu(C)	lnu(C)-F	ACTGTCGCAGAGCAGGAAAGCC	-	11
	lnu(C)-R	AGCATCTACACCCCAGCCACCA		
lnu(D)	lnu(D)-F	ACGGAGGGATCACATGGTAAATAAAGC	-	11
	lnu(D)-R	CCTGTCTTTATCGTCCTTCCAAACCGT		
bcrA	bcrA-F	CCGCAATGAAAATGATGTTG	584	12

	bcrA-R	TGCGGCTATCTTACCATCTG		
bcrB	bcrB-F	AAAGAAACCGACTGCTGATA	489	12
	bcrB-R	GCTTACTTGTATAGCAGAGA		
bcrD	bcrD-F	GCGAAGCGTTTAAGGAAATG	482	12
	bcrD-R	TGGCACAGCAAGAAAGAATG		
bcrR	bcrR-F	TAACGCAGGAACAACTTGC	461	12
	bcrR-R	CAAAGCGGTAATGGTGAGG		
class I	class I-F	CCTCCCGCACGATGATC	288	13
	class I-R	TCCACGCATCGTCAGGC		
class II	class II-F	GTAGCAAACGAGTGACGAAATG	788	13
	class II-R	CACGGATATGCGACAAAAAGGT		
class III	class III-F	GCCTCCGGCAGCGACTTTCAG	979	13
	class III-R	ACGGATCTGCCAAACCTGACT		
Tn 552	Tn 552-F	CTAACTAATTTTTCTGAAGCCAAACG	1360	7
	Tn 552-R	TTTAAGTGTTTTCTTCTCTGACTACG		
Tn 5801	Tn 5801-F	CCGATATTGAGCCTATTGATGTG	722	14
	Tn 5801-R	GTCCATACGTTCCTAAAGTCGTC		
Tn 916-like	Tn 916-like-F	GCCATGACCTATCTTATA	1057	14
	Tn 916-like-R	CTAGATTGCGTCCAA		
optrA	optrA-F	GCACCAGACCAATACGATACAA	794	This study
	optrA-R	TCCTTCTTAACCTTCTCCTTCTCA		
fexA	fexA-F	TTGGGAAGAATGGTTCAGGG	977	This study
	fexA-R	ATCGGCTCAGTAGCATCACG		

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