

Table S1. Genes and primers' sequences utilized for all PCRs in this study.

Gene	Primers' oligonucleotide (5'⇒3')	Amplicon size	Reference
Antimicrobial resistance (AMR) genes			
<i>blaZ</i>	F: CAGTCACATGCCAAAGAG	772 bp	[1]
	R: TACACTCTGGCGGTTTC		
<i>mecA</i>	F: GGGATCATAGCGTCATTATTC	527 bp	[2]
	R: AACGATTGTGACACGATAGCC		
<i>ermA</i>	F: TCTAAAAAGCATGTAAAAGAA	645 bp	[3]
	R: CTTCGATAGTTATTAAATATTAG		
<i>ermB</i>	F: GAAAAGTACTCAACCAAATA	639 bp	[3]
	R: AGTAACGGTACTTAAATTGTTTA		
<i>ermC</i>	F: TCAAAACATAATATAGATAAA	642 bp	[3]
	R: GCTAATATTGTTAAATCGTCAAT		
<i>ermT</i>	F: CCGCCATTGAAATAGATCCT	200 bp	[4]
	R: TTCTGTAGCTGTGCTTCAAAAAA		
<i>mphC</i>	F: ATGACTCGACATAATGAAAT	900 bp	[1]
	R: CTACTCTTCATACCTAACTC		
<i>msrA</i>	F: GCAAATGGTAGGTAAGACAACT	399 bp	[5]
	R: ATCATGTGATGTAAACAAAT		
<i>lnuA</i>	F: GGTGGCTGGGGGTAGATGTATTAACTGG	323 bp	[6]
	R: GCTTCTTTGAAATACATGGTATTTTCGATC		
<i>lnuB</i>	F: CCTACCTATTGTTGTGGAA	499 bp	[7]
	R: ATAACGTTACTCTCCTATTTC		
<i>vgaA</i>	F: AGTGGTGGTGAAGTAACACG	1264 pb	[8]
	R: GGTTCAATACTCAATCGACTGAG		
<i>aac6'-aph2''</i>	F: CCAAGAGCAATAAGGGCATA	220 bp	[9]
	R: CACTATCATAACCACTACCG		
<i>ant4'</i>	F: GCAAGGACCGACAACATTTC	165 bp	[9]
	R: TGGCACAGATGGTCATAACC		

<i>tet</i> (L)	F: CATTGGTCTTATTGGATCG R: ATTACACTCCGATTCCGG	456 bp	[10]
<i>tet</i> (M)	F: GTAAATAGTGTCTTGGAG R: CTAAGATATGGCTCTAACAA	576 bp	[10]
<i>tet</i> (K)	F: TTAGGTGAAGGGTTAGGTCC R: GCAAACTCATTCCAGAAGCA	697 bp	[10]
<i>dfrA</i>	F: CCTTGGCACTTACCAAATG R: CTGAAGATTCGACTTCCC	374 bp	[1]
<i>dfrD</i>	F: TTCTTAATTGTTGCGATGG R: TTAACGAATTCTCTCATATATATG	582 bp	[1]
<i>dfrG</i>	F: TCGGAAGAGCCTTACCTGACAGAA R: CCCTTTGGGCAAATACCTCATTCCA	323 bp	[4]
<i>dfrK</i>	F: GAGAATCCCAGAGGGATTGGG R: CAAGAACGTTTCGCTCATAAA	423 bp	[4]
<i>cat</i> _{pC221}	F: ATTTATGCAATTATGGAAGTTG R: TGAAGCATGGTAACCATCAC	434 bp	[1]
<i>cat</i> _{pC223}	F: GAATCAAATGCTAGTTAACTC R: ACATGGTAACCACATCAC	283 bp	[1]
<i>cat</i> _{pC194}	F: CGACTTTAGTATAACCACAGA R: GCCAGTCATTAGGCCTAT	570 bp	[1]
<i>catA</i>	F: GGATATGAAATTATCCCTC R: CAATCATCTACCCTATGAAT	505 bp	[10]
<i>fexA</i>	F: GTACTTGTAGGTGCAATTACGGCTGA R: CGCATCTGAGTAGGACATAGCGTC	1272 bp	[11]
<i>fexB</i>	F: TTCCCACATTGGTGAAAGGAT R: GCAATTCCCTTTATGGACGTT	816 bp	[12]
<i>cfr</i>	F: TGAAGTATAAAGCAGGTTGGGAGTCA R: ACCATATAATTGACCACAAGCAGC	746 bp	[13]
<i>cfrB</i>	F: TGAGCATATACGAGTAACCTCAAGA R: CGCAAGCAGCGTCTATATCA	293 bp	[14]
<i>cfrD</i>	F: AGAAGTCGCAACAAGTGAGGA R: GCAACTGCATGAGTCAAAGAA	595 bp	[15]
<i>optrA</i>	F: AGGTGGTCAGCGAACTAA	1395 bp	[16]

	R: ATCAACTGTTCCCATTCA		
<i>poxtA</i>	F: TCAATGCAGAGCAGGAAGCA	791 bp	[15]
	R: GGTGGATTACCGACACCGT		
<i>mupA</i>	F: CCCATGGCTTACCGAGTTGA	419 pb	[17]
	R: CCATGGAGCACTATCCGAA		
<i>S. epidermidis</i> housekeeping alleles for MLST			
<i>arcC</i>	F: TGTGATGAGCACGCTACCGTTAG	465 pb	[18]
	R: TCCAAGTAAACCCATCGGTCTG		
<i>aroE</i>	F: CATTGGATTACCTCTTGTTCAGC	430 pb	[18]
	R: CAAGCGAAATCTGTTGGGG		
<i>gti</i>	F: CAGCCAATTCTTTATGACTTT	438 pb	[18]
	R: GTGATTAAAGGTATTGATTGAAT		
<i>mutS</i>	F: GATATAAGAATAAGGGTTGTGAA	412 bp	[18]
	R: GTAATCGTCTCAGTTATCATGTT		
<i>pyrR</i>	F: GTTACTAATACTTTGCTGTGTT	428 bp	[18]
	R: GTAGAACATGAAAGAGACTAAAATGAA		
<i>tpi</i>	F: ATCCAATTAGACGCTTAGTAAC	424 bp	[18]
	R: TTAATGATGCGGCCACCTACA		
<i>yqiL</i>	F: CACGCATAGTATTAGCTGAAG	416 bp	[18]
	R: CTAATGCCTTCATCTTGAGAAATAA		

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