

Supplementary Materials

Table S1. Oligonucleotide primers used in this work.

Genes	Sequence (5'–3')	Annealing	Amplicon size (bp)	Reference
<i>ddl E. faecalis</i>	F: ATCAAGTACAGTTAGTCTT R: ACGATTCAAAGCTAACTG	50	941	
<i>ddl E. faecium</i>	F: GCAAGGCTTCTTAGAGA R: CATCGTGTAAGCTAACTTC	50	550	[16]
16S	F: AGAGTTTGATCCTGGCTNAG R: GACTACCNNGGTATCTAATCC	50	800	
<i>tet(L)</i>	F: ATTACACTTCCGATTTTCGG R: CATTGGTCTTATTGGTACG	54	1550	[17]
<i>VanA</i>	F: AACAACTTACGCGGCACT R: AAAGTGCGAAAAACCTTGC	55	1550	[21]
<i>ermB</i>	F: GGAACATCTGTGGTATGGCG R: CATTTAACGACGAAACTGGC	52	426	[19]
<i>VanB</i>	F: GAT ATT CAA AGC TCC GCA GC R: GGT ATC TTC CGC ATC CAT CA	55	1000	[22]
<i>tet(M)</i>	F: GTTAAATAGTGTTCTTGGAG R: CTAAGATATGGCTCTAACAA	45	1500	[17]
<i>tet(W)</i>	F: TGG TCC CCT AAT ACA TCG TT R: GCC ATC TTG GTG ATC TCC	55	1550	[18]
<i>vgaA</i>	F: AGT GGT GGT GAA GTA ACA CG R: CTT GTC TCC TCC GCG AAT AC	55	200	[20]
<i>vatD</i>	F: GCT CAA TAG GAC CAG GTG TA R: TCC AGC TAA CAT GTA TGG CG	55	330	
<i>gelE</i>	F: TATGACAATGCTTTTTGGGAT R: AGATGCACCCGAAATAATATA	56	213	
<i>asa1</i>	F: GCACGCTATTACGAACTATGA R: TAAGAAAGAACATCACCACGA	56	375	
<i>esp</i>	F: AGATTTTCATCTTTGATTCTTGG R: AATTGATTCTTTAGCATCTGG	56	510	[23]
<i>cylA</i>	F: ACTCGGGGATTGATAGGC R: GCTGCTAAAGCTGCGCTT	56	688	