

Prevalence and Molecular Characterization of Vancomycin Variable *Enterococcus faecium* Isolated From Clinical Specimens

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Supplemental Data Table S1. Nucleotide sequences of PCR primers used in this study

Primer	Sequence (5'→3')	Covered region (gene)	Amplicon size (bp)
42F 349R	ATT TTC CTG ACG AAT CCC TCG TCG GAA AAC AAG GTG AGC TTA GA	Transposase	308
164F 921R	AAC CTA AGG GCG ACA TAT GGT G AAA AGG AGC CAC CAT CTA CCG	Transposase	758
949F 1913R	GCATGTAGTGATGAAACACCTAGCTGC CGTCCTGCCGACTATGATTATTT	Transposase	965
1871F 2976R	ACCGTTTTTGCAGTAAGTCTAAAT TGAAGATGAATGGATACTGGGGACC	Transposase	1,106
2524F 3726R	CTTGCTTCCCACACCATT AGCCCTAGATACATTAGTAATT	Transposase– resolvase	1,203
3514F 3978R	ACT GTA ATG GCT GGT GTT AAC CAT AGT TAT CAC CCC TTT CAC TAT	Resolvase– <i>vanR</i>	465
3907F 4794R	ATG CTT ATA AAT TCG GCC C ATC CAA TCC CCA AGT TTC CC	<i>vanR</i> – <i>vanS</i>	888
3992F 4511R	TTA TTG TGG ATG ATG AAC ATG TCG GAG CTA ACC ACA TTC	<i>vanR</i>	520
4676F 5769R	AAC GAC TAT TCC AAA CTA GAA C GCT GGA AGC TCT ACC CTA AA	<i>vanS</i>	1,094
5235F 7035R	ATA TCA CGT TGG ACA AAG C TTA CGT CAT GCT CCT CTG AG	<i>vanS</i> – <i>vanA</i>	1,801
8082F 8505R	ACT TGG GAT AAT TTC ACC GG TGC GAT TTT GCG CTT CAT TG	<i>vanX</i>	424
8448F 9138R	GAT GAA CGC TCT CAT CAT GC TTC CTG AGA AAA CAG TGC TTC A	<i>vanX</i> – <i>vanY</i>	691
8544F 9580R	GCA TAT AGC CTC GAA TGG TCG TCA AGC TTG ATC CTA C	<i>vanX</i> – <i>vanY</i>	1,037