**Supplement**

***tuf* gene PCR and Sequencing**

Isolates suspected to be viridans group *Streptococcus* underwent PCR to amplify a 761 base pair segment of the *tuf* gene, followed by Sanger sequencing.5

Purified genomic DNA was extracted from colonies using the PrepMan Ultra Extraction kit. PCR was performed using the AmpliTaq Gold Fast PCR kit with 0.1 µg genomic DNA and 0.25 uM of each primer.5 The PCR mixtures were initially denatured at 95°C for 10 minutes and then underwent 35 cycles of amplification (3 sec of denaturing at 96°C, 3 sec of annealing at 50°C, and 15 sec of elongation at 68°C) with a final extension cycle of 72°C for 10 seconds. The PCR products were purified using the ExoSAP-IT PCR Product Cleanup Reagent. Cycle Sequencing was performed with the ABI BigDye Terminator v1.1 Cycle Sequencing Kit using 20 ng of PCR product and 3.2 pmol of the *tuf* gene primer. The cycle sequencing mixtures were initially denatured at 96°C for 1 minute and then underwent 25 cycles of amplification (10 sec of denaturing at 96°C, 5 sec of annealing at 50°C, and 4 min of elongation at 60°C). The extension products were purified using the CentiSep Spin Column. Sequences were obtained using the ABI 3500 Genetic Analyzer and analyzed using Lasergene SeqMan Pro software (DNASTAR, Madison, WI). We looked for homology with NCBI sequences using the BLASTn program.