



Supplemental Figure 4. Four nucleotide substitutions within the CroR-dependent regulatory sequence motif in the promoter of *pbp4(5)* disrupts CroR-dependent regulation in *E. faecalis* CK221. Abundance of *pbp4(5)* transcripts from *E. faecalis* CK221 wild-type, $\Delta croR$ (SB45), $\Delta pbp4(5)$ (JL640), and *pbp4(5)* ATAA (ST8) mutant strains grown to exponential phase with exposure (or not) to 16 μ g/ml vancomycin to activate CroS/R. *pbp4(5)* RNA abundance analyzed by RTqPCR. Error bars represent standard deviation of a minimum of two independent cultures analyzed in triplicate; *indicates a P value of <0.05, as determined by a two-tailed, unpaired parametric *t*-test.