



Figure S1. Principal component analysis. Principal component analysis demonstrated a general clustering of the biological replicates that had the same genotype and were exposed to the same bacterium, indicating consistency. When considering just the bacterial exposure, *E. faecalis* vs. *E. coli*, a large separation between the two conditions was evident (PC1). When considering just genotype, the *skpo-1* mutant vs. N2, a more modest separation was observed (PC2).