**Table S1.** Protein level comparisons between groups. Statistics were performed on the output of Fragpipe using the MSStatsTMT R package. Three comparisons are reported, healthy volunteers vs EcB infected, E. faecalis vs E. faecium, and mortality vs survival.

**Table S2.** Metabolite comparisons between groups. Statistics were performed on normalized metabolomics data for each feature using t tests as implemented in the rstatix R package. FDR correction was then applied to account for multiple comparisons. Three comparisons are reported, healthy volunteers vs EcB infected, E. faecalis vs E. faecium, and mortality vs survival.