**Table S1.** Protein level comparisons between groups. Statistics were performed on the output of Fragpipe using the MSStatsTMT 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 r statix 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.