Figure S1. 16s rRNA sequencing identifies specific bacterial differences in mice after stem cell transplant. (**A**) Taxonomic composition of bacteria taxa over time from select fecal specimens of mice over time, pre and post stem cell transplant, and following treatment. Each vertical bar represents 1 individual mouse. (**B**) Linear discriminant analysis effect size measurements (LEfSe) of bacterial taxa effect enriched in groups that received the standard probiotic blend versus the optimized probiotic blend.