Supplementary Figure 5: Alpha diversity of rabbit eukaryotic microbial communities comparing different locations of the GI tract. The figure compares the number of observed species (Observed), species richness (Chao1), diversity (Shannon), and evenness (Simpson) between different locations in the foregut and hindgut. Overall, the alpha diversity of the eukaryotic microbiota was far lower than the alpha diversity of the bacterial microbiota. Please see supplementary Tables 7 and 8 for the genus-level taxonomic composition and statistical comparisons.