

Determining outliers.

The data was explored first to detect outlier samples. Alpha diversity was used to flag samples that were extreme or aberrant compared to the majority. Descriptive statistics on measures of alpha diversity (Observed ASVs, Shannon Diversity, InvSimpson and Chao1) on the full dataset before and after rarefying were compared. After noting a bimodal distribution in alpha diversity measures, the dataset was split into Kadaknath and Commercial broiler prior to descriptive statistics.

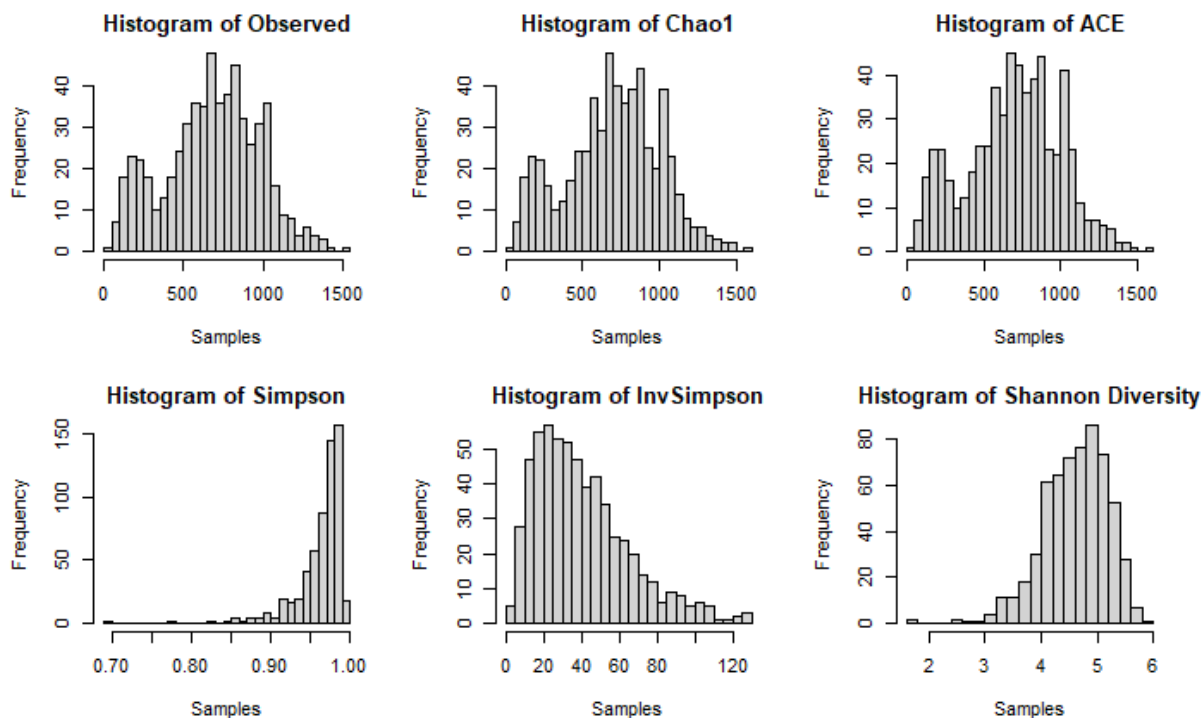


Figure 1 Histogram of alpha diversity measures on unrarefied dataset, with both breeds combined shows that there is a bimodal distribution.

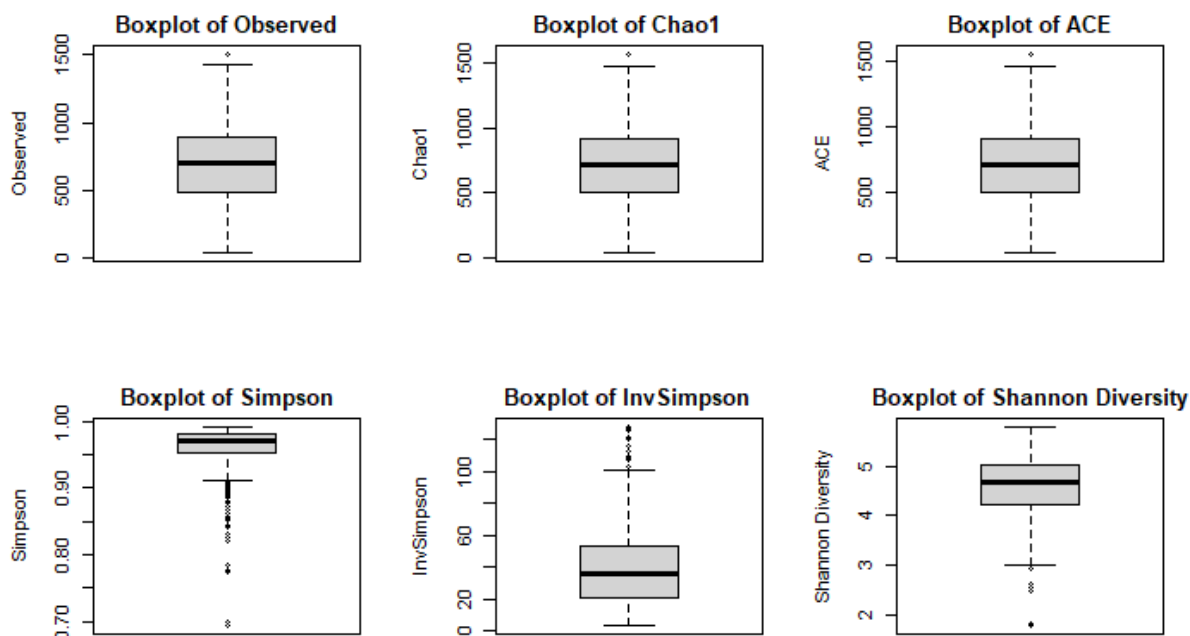
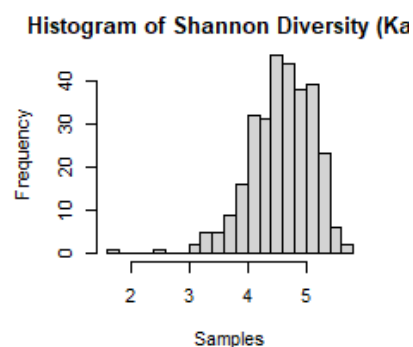
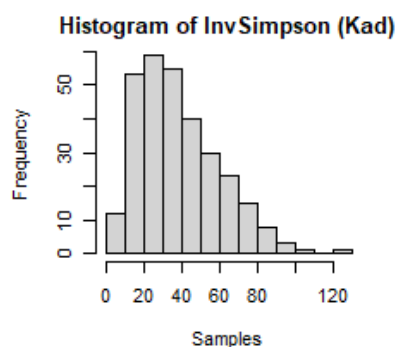
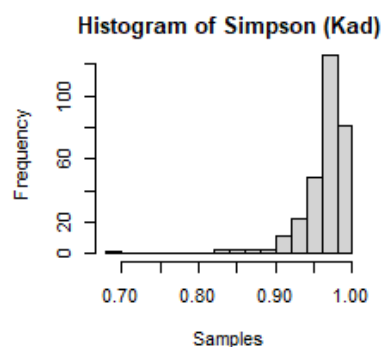
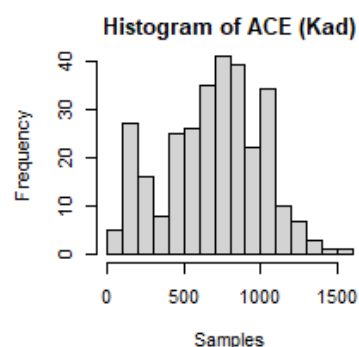
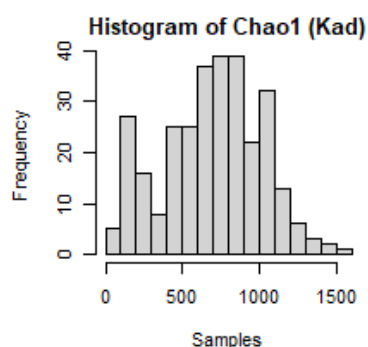
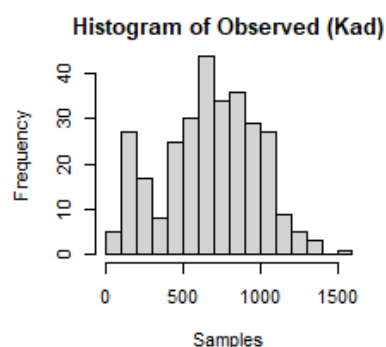
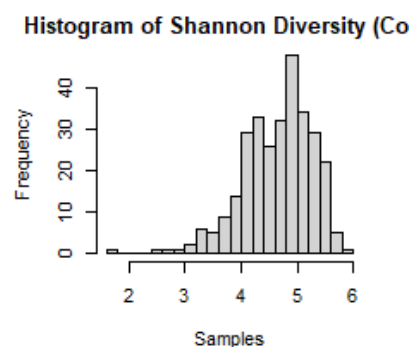
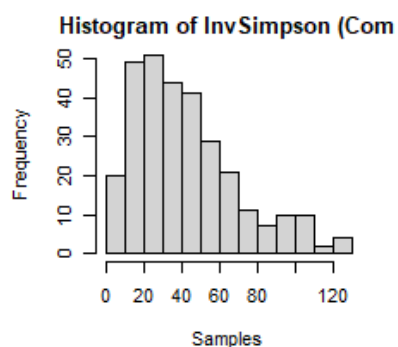
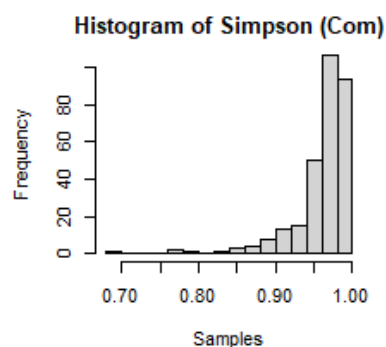
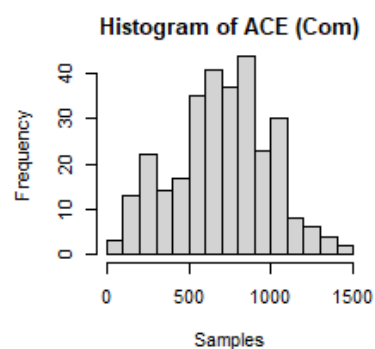
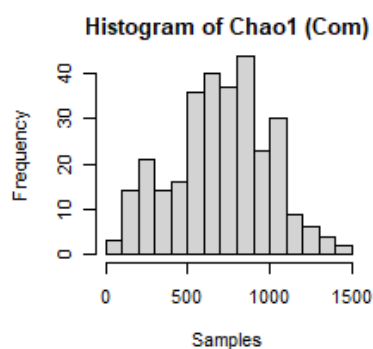
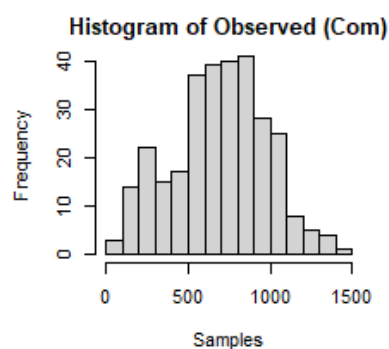


Figure 2 Boxplot of alpha diversity on unrarefied dataset, with breeds combined.



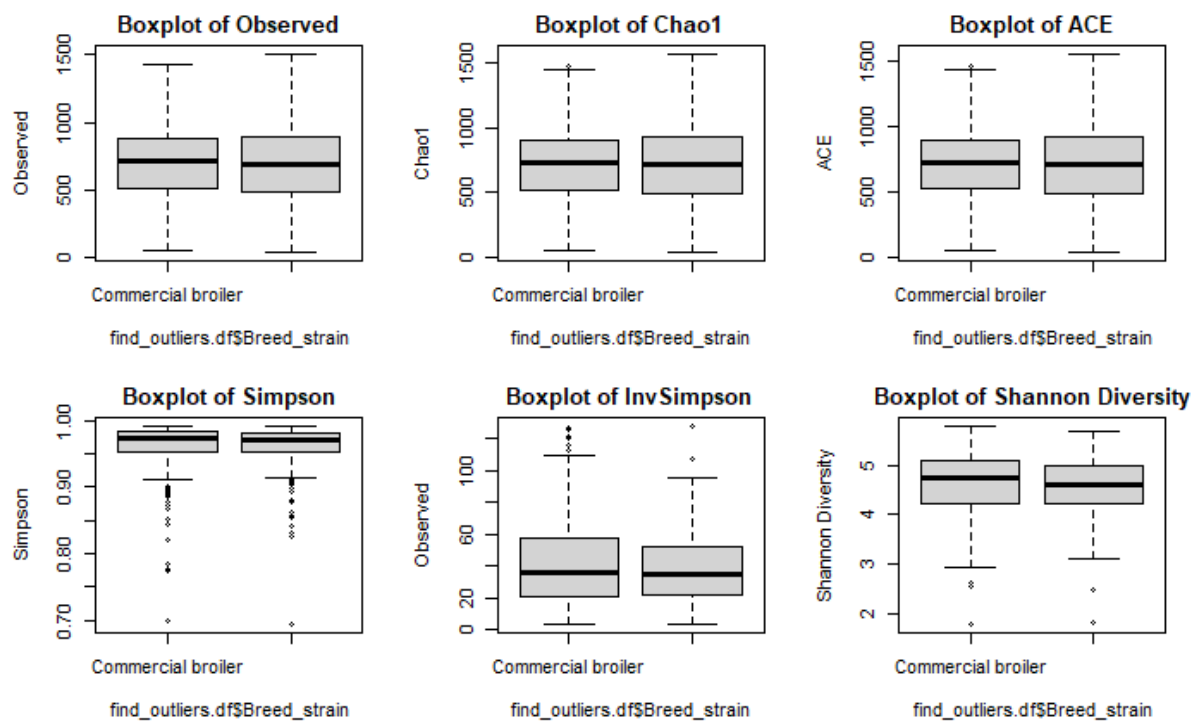


Figure 3 Boxplot of alpha diversity measures on unrarefied dataset, with samples split by breed.

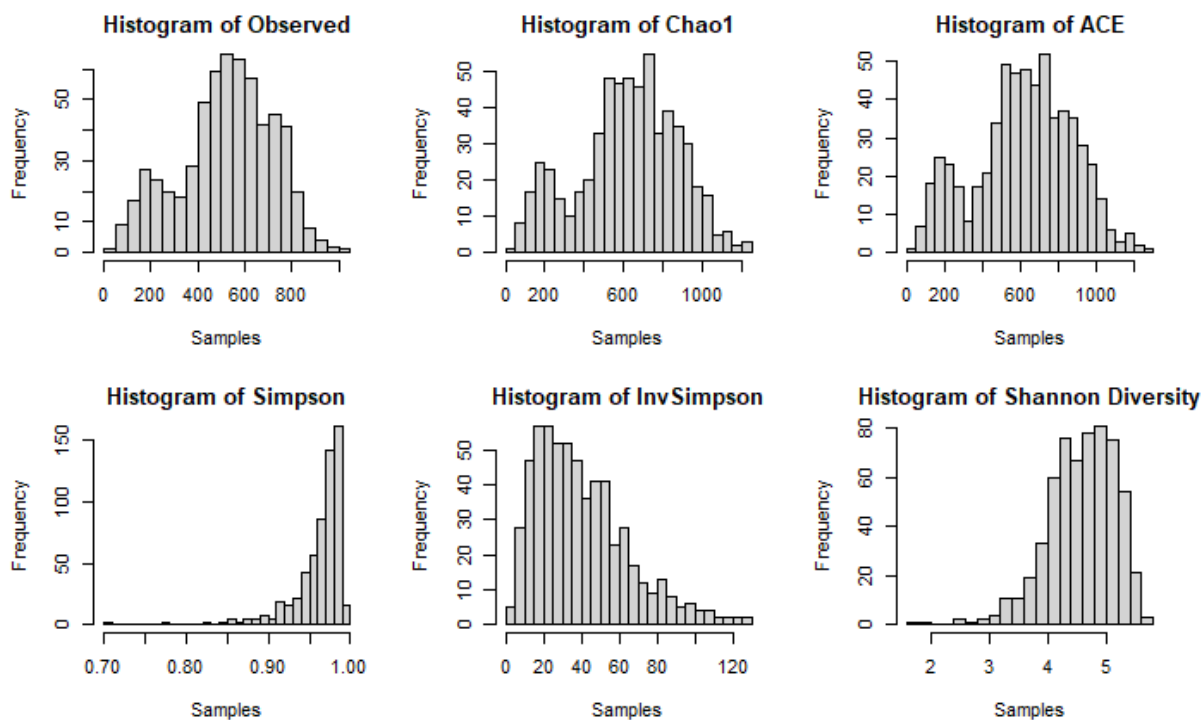


Figure 4 Histogram of alpha diversity measures on rarefied, combined dataset.

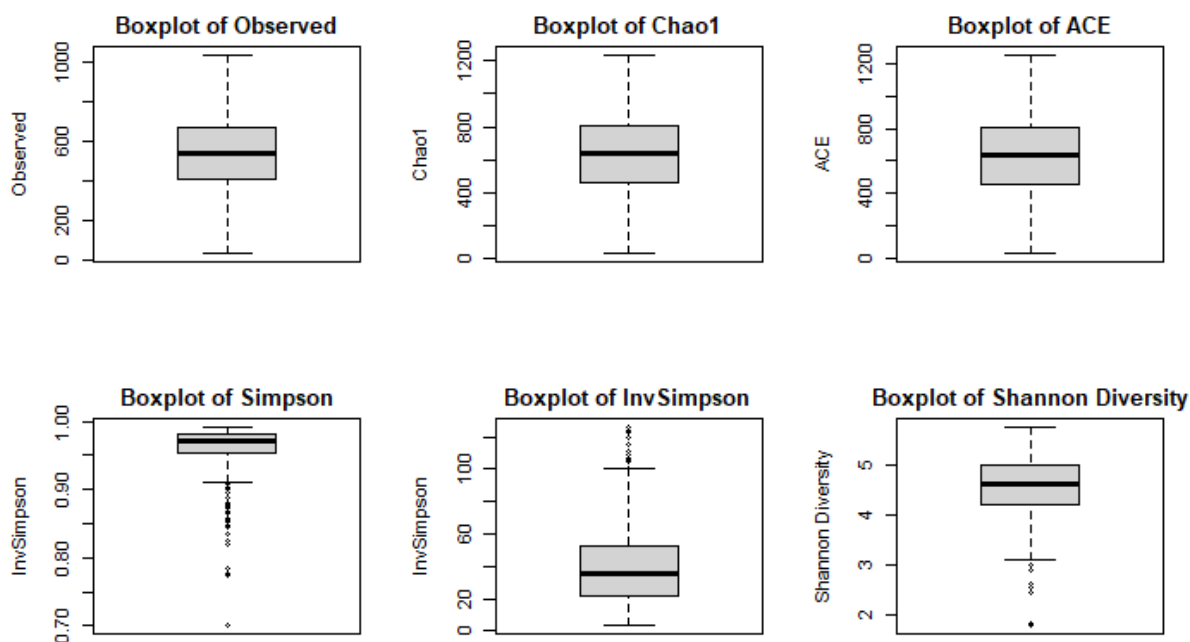


Figure 5 Boxplot of alpha diversity measures on rarefied, combined dataset.