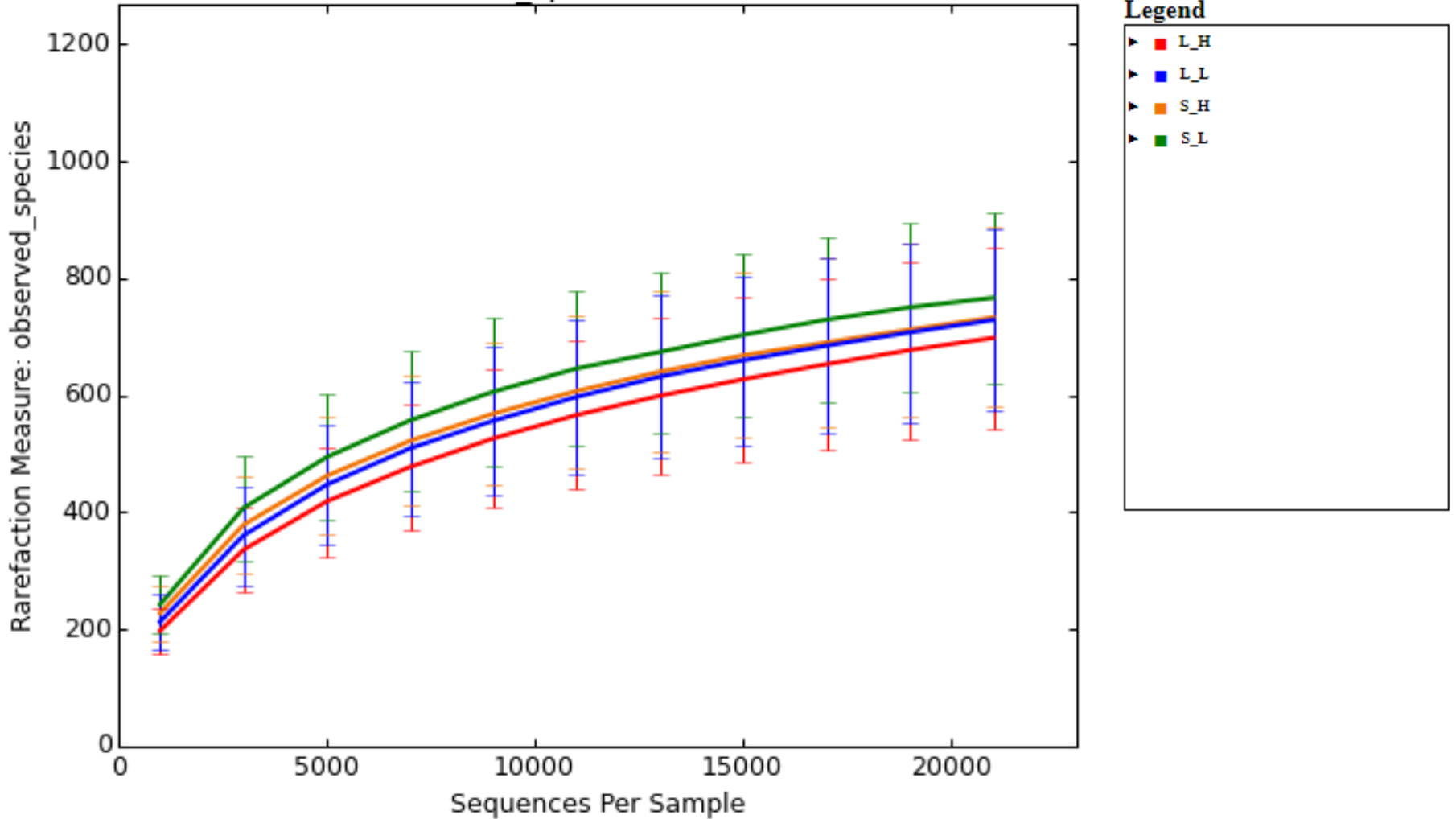


observed species: Treatment



Supplementary Figure S1. Rarefaction curves of observed species (i.e., OTUs) clustered at the 97% identity based on the 16S rRNA gene sequences of rumen solid (S) and liquid (L) digesta of Simmental bulls phenotypically divergent for residual feed intake. High RFI = H, Low RFI = L