Task.

There are 22 samples collected of chicken’s intestinal in these datasets and 771 different **Operational Taxonomic Units (OUT)**, the observed abundance of each feature can be observed in Figure 1. The microbiome is constituted by Bacteria in every sample, also the figure shows that the there is mayor abundance of the Phylum *Firmicutes*.

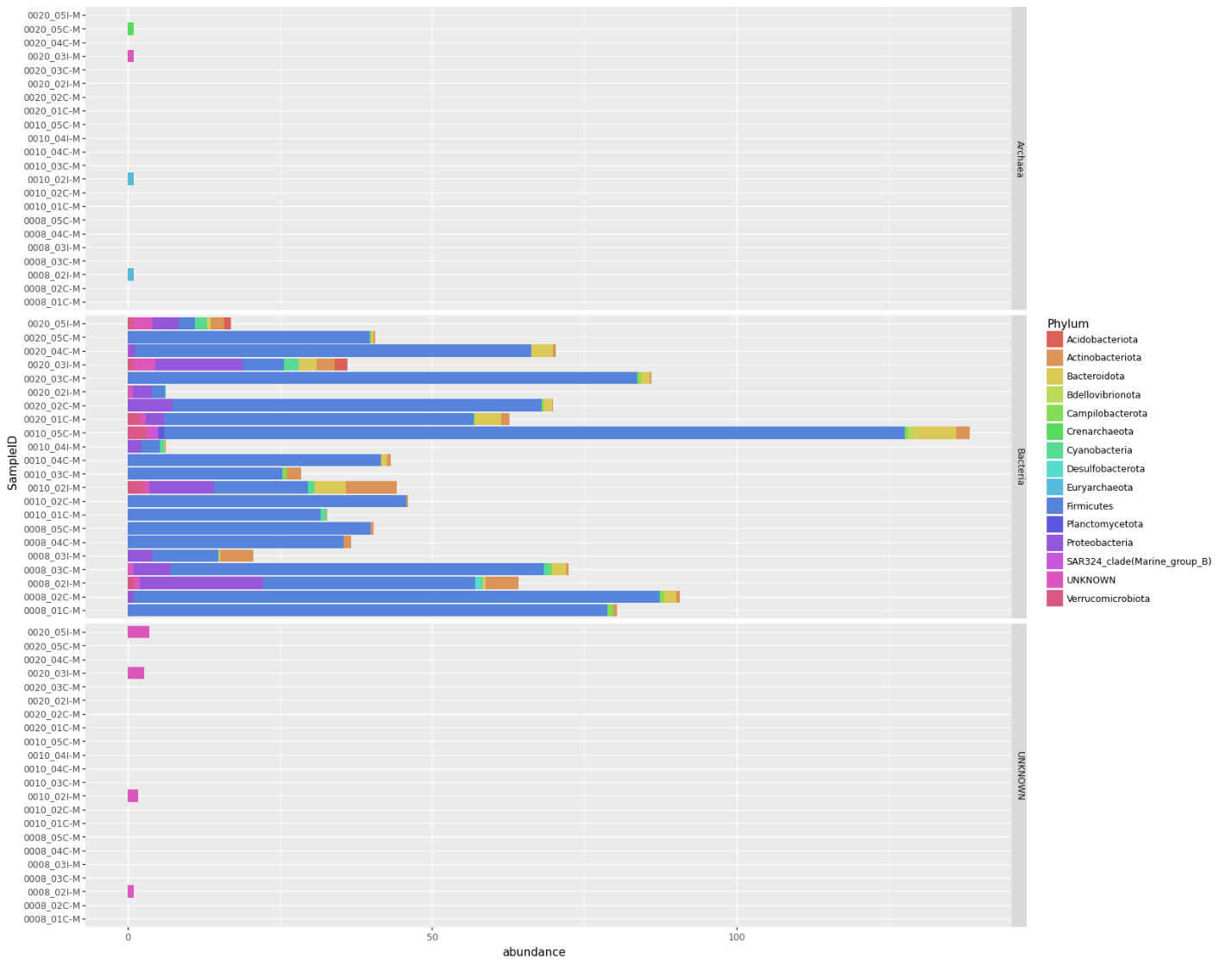


Figure .

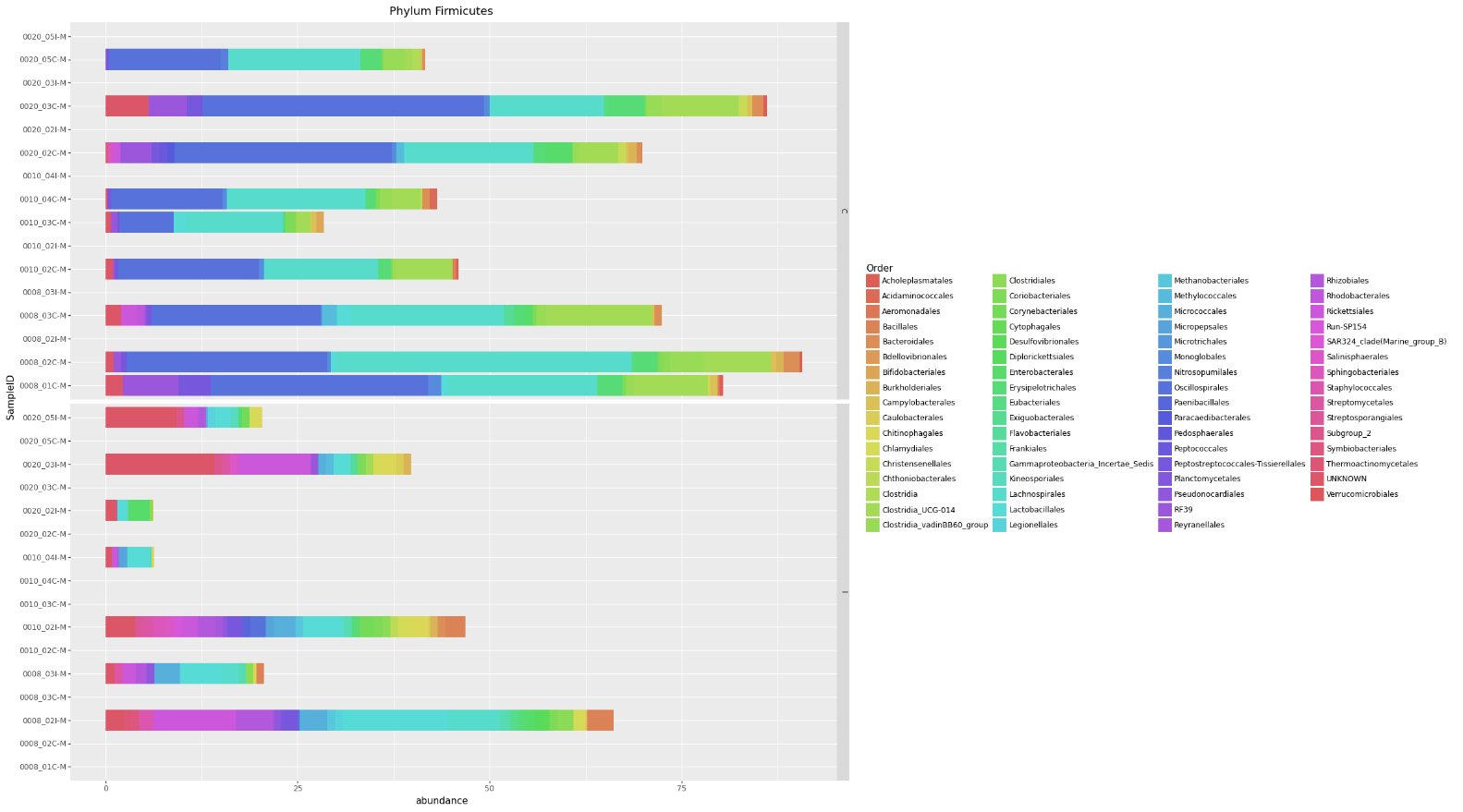


Figure . Phylum Firmicutes.

