

Figure S1: Schematic overview of the allocation

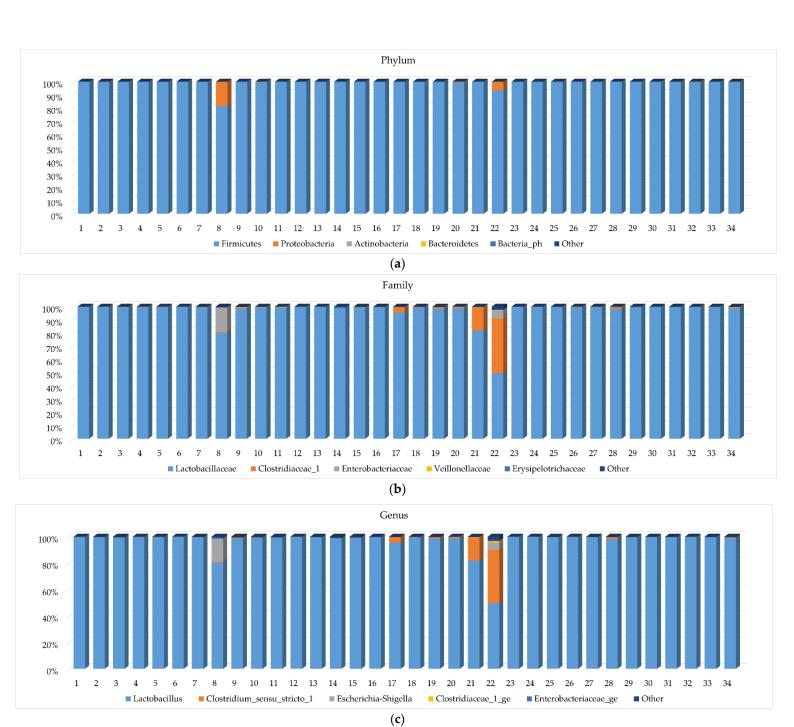


Figure S2: Bacterial community compositions present in digesta of the last 25% of small intestinal length for each individual pig. The cumulated histograms show the relative abundance of the identified taxa at phylum (a), family (b) and genus (c) level.