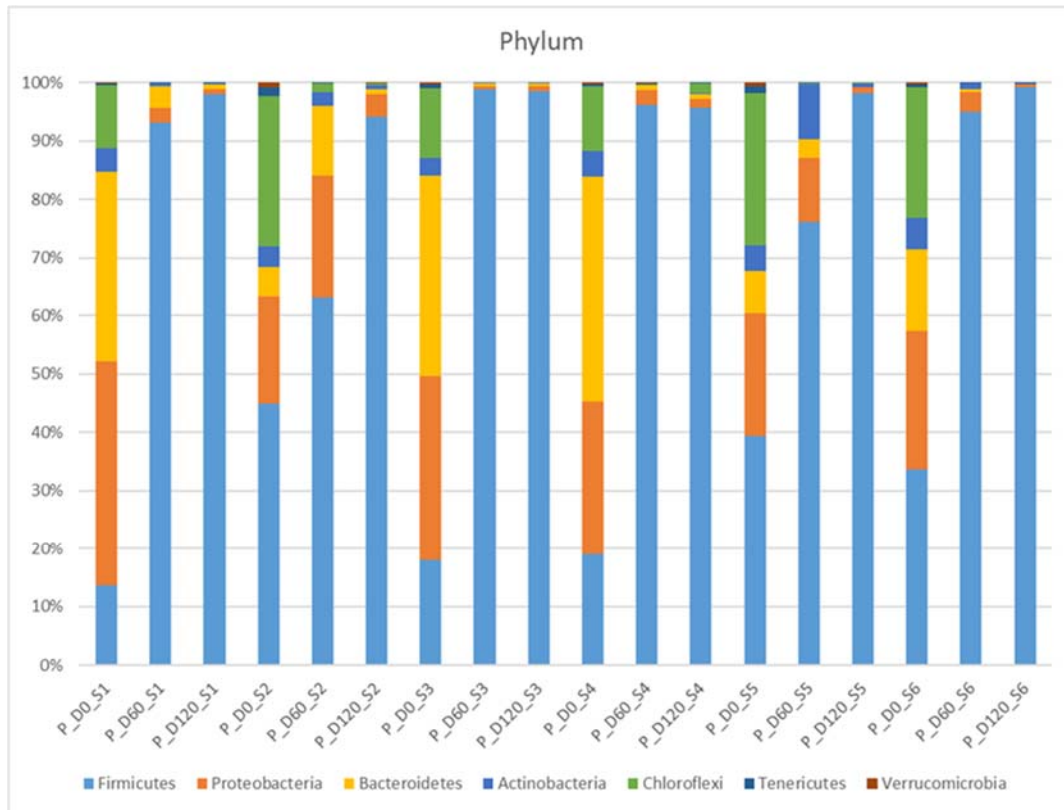


## Supplementary Material



**Supplementary Figure 1.** Relative abundance of bacterial phyla in the different treatments assayed (S1-S6), obtained through metagenetic analysis of 16S rRNA gene at the initial (D0), middle (D60) and the end (D120) of fermentation process.