

Supplemental material

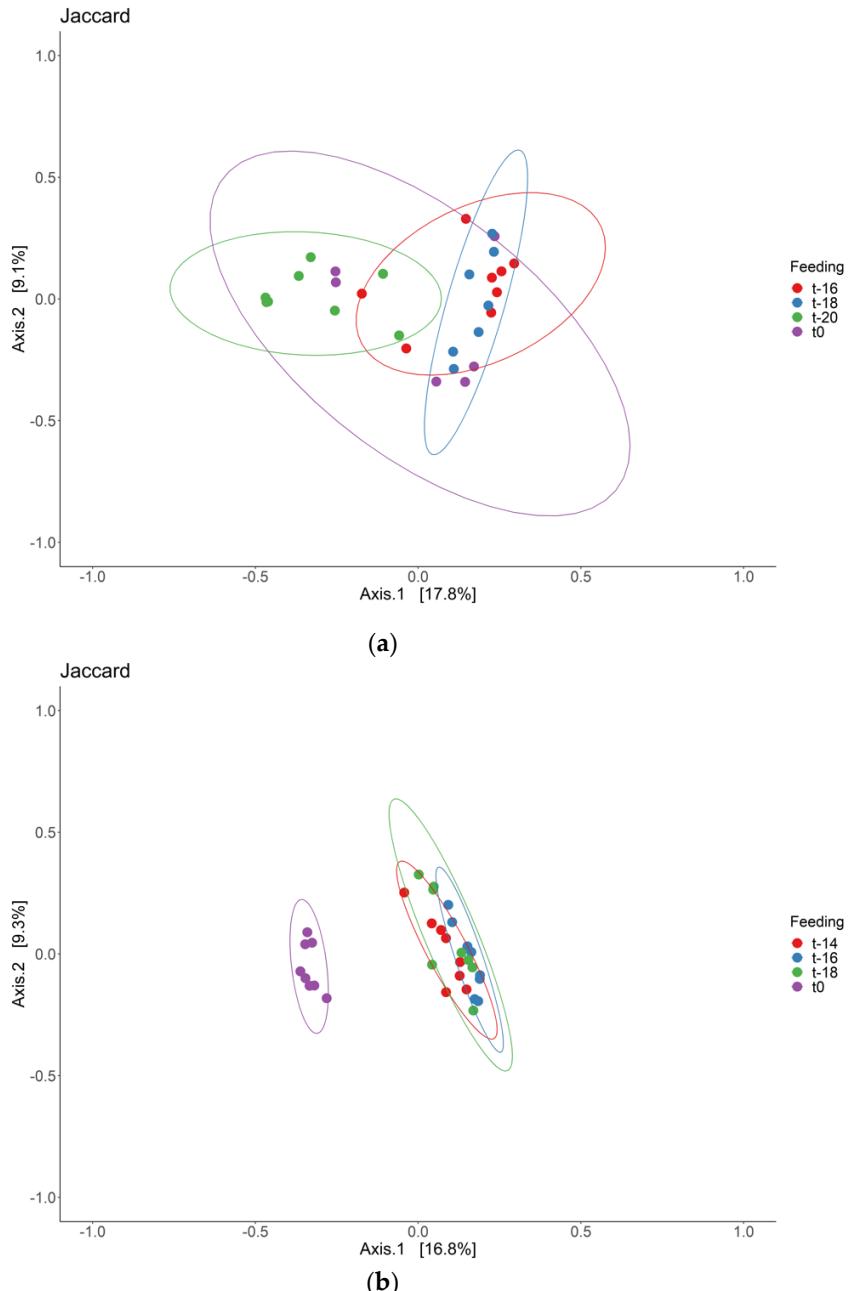
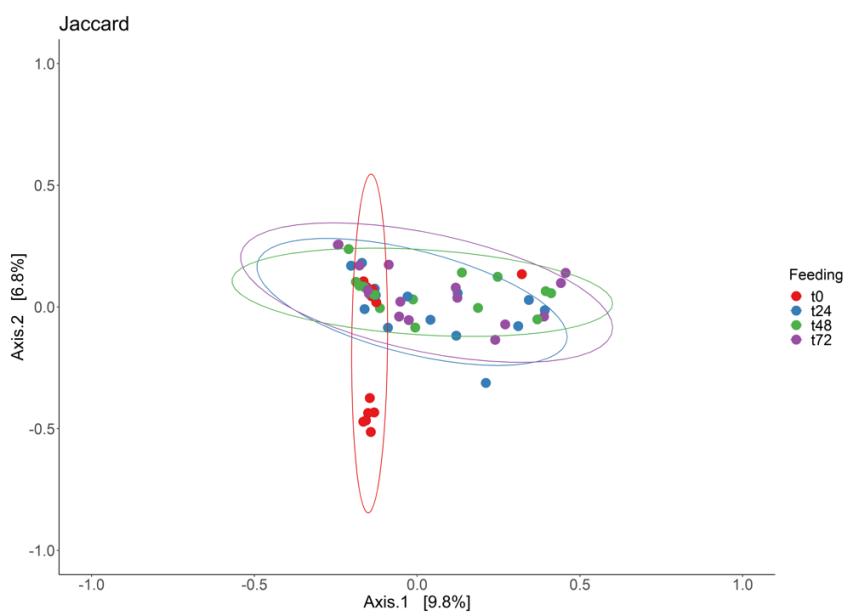
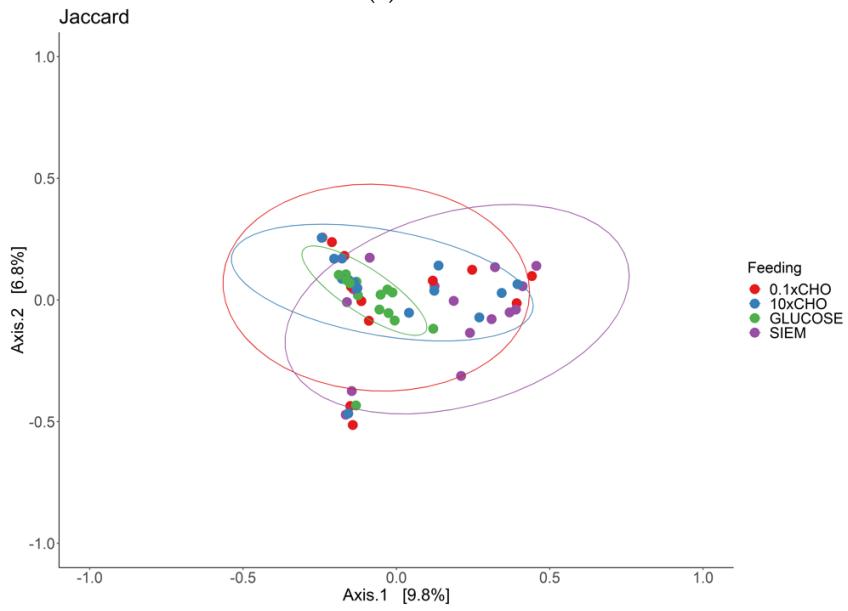


Figure S1: Beta-diversity during the adaption period visualized using PCoA plot with Jaccard.similarities; (a) Fungi; (b) Bacteria. N=8



(a)



(b)

Figure S2: Beta-diversity of the fungal community during the intervention period visualized using PCoA plot with Jaccard similarities; (a) Between time points for all feeding interventions; (b) Between feeding interventions for the time points t24, t48 and t72.

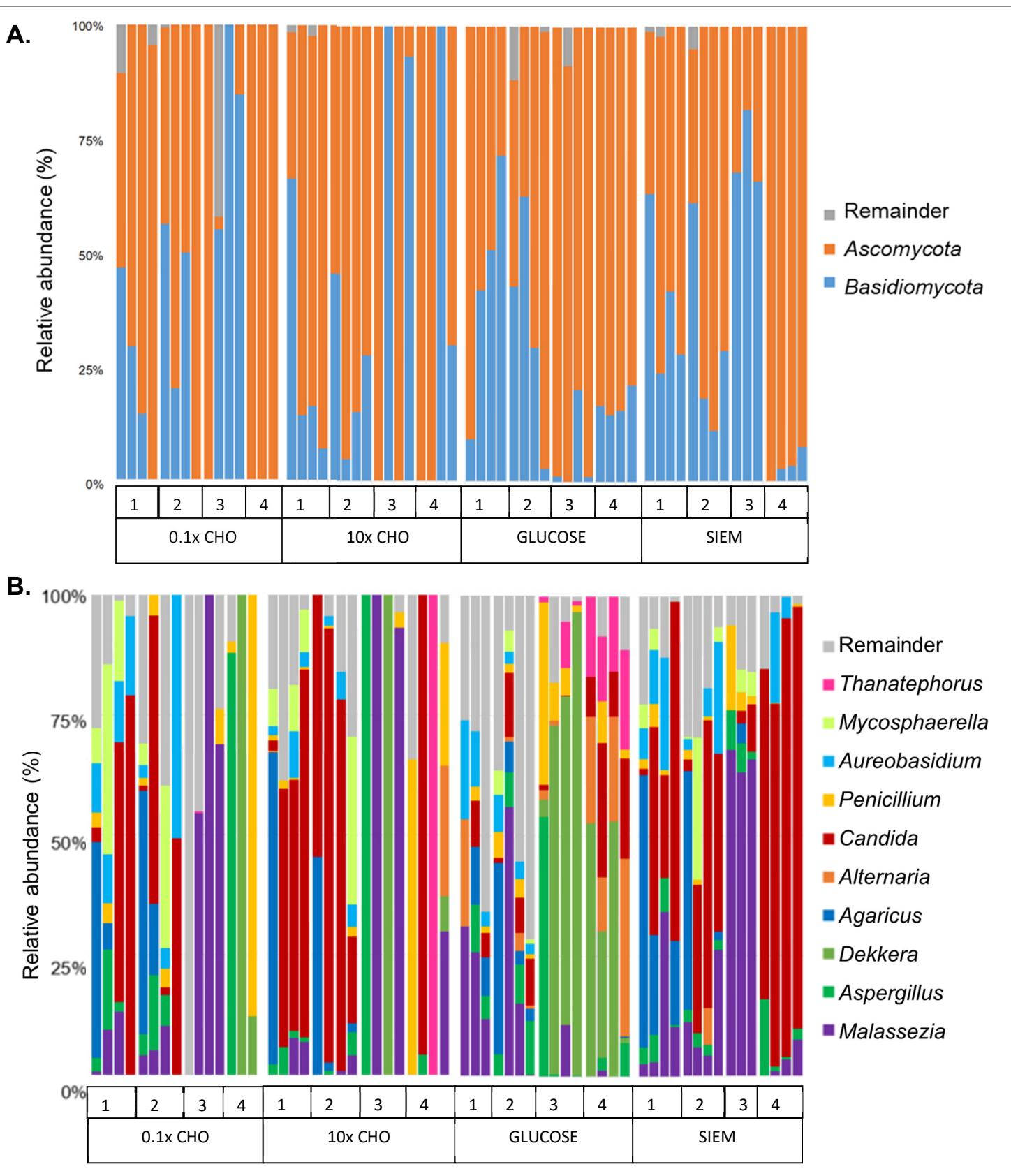


Figure S3: Relative abundance (%) of major groups of fungi during the test period after inoculation in TIM-2; (a) Phyla; (b) Genera; 1,2,3,4 stand for different replicates.

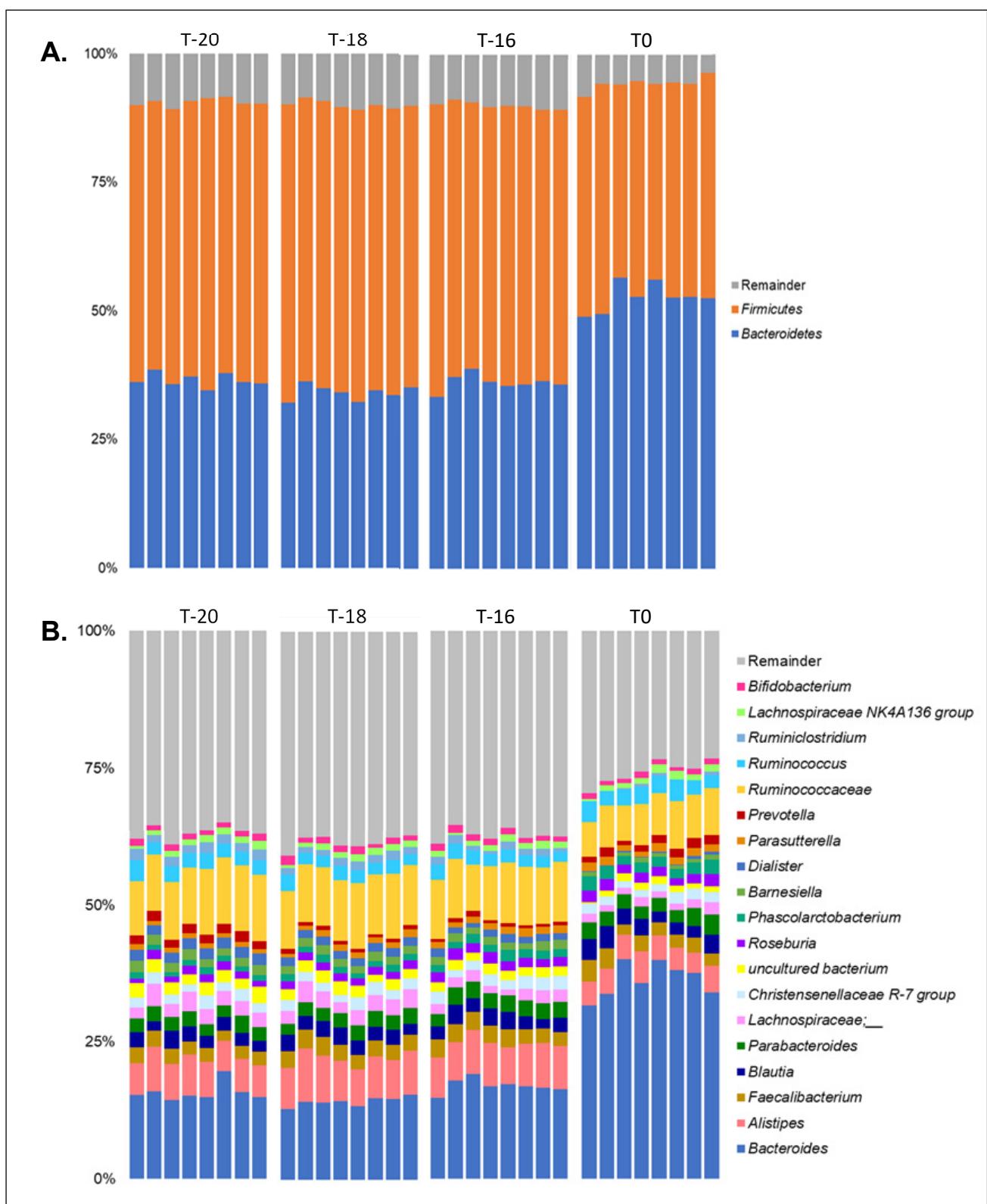


Figure S4: Relative abundance (%) of major groups of bacteria during the adaptation period after inoculation in TIM-2; (a) Phyla; (b) Genera.

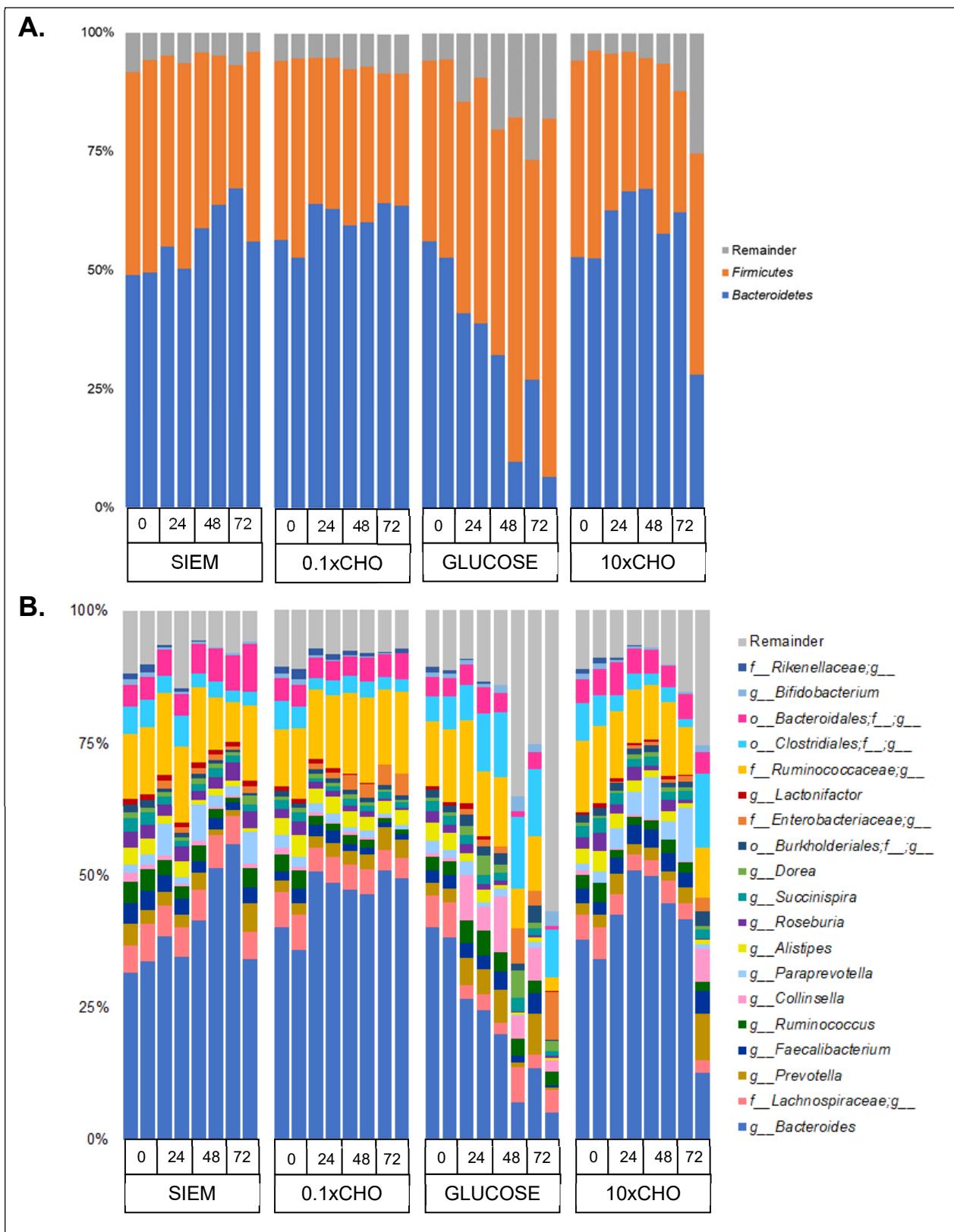


Figure S5: Relative abundance (%) of major groups of bacteria during the test period after inoculation in TIM-2; (a) Phyla; (b) Genera

