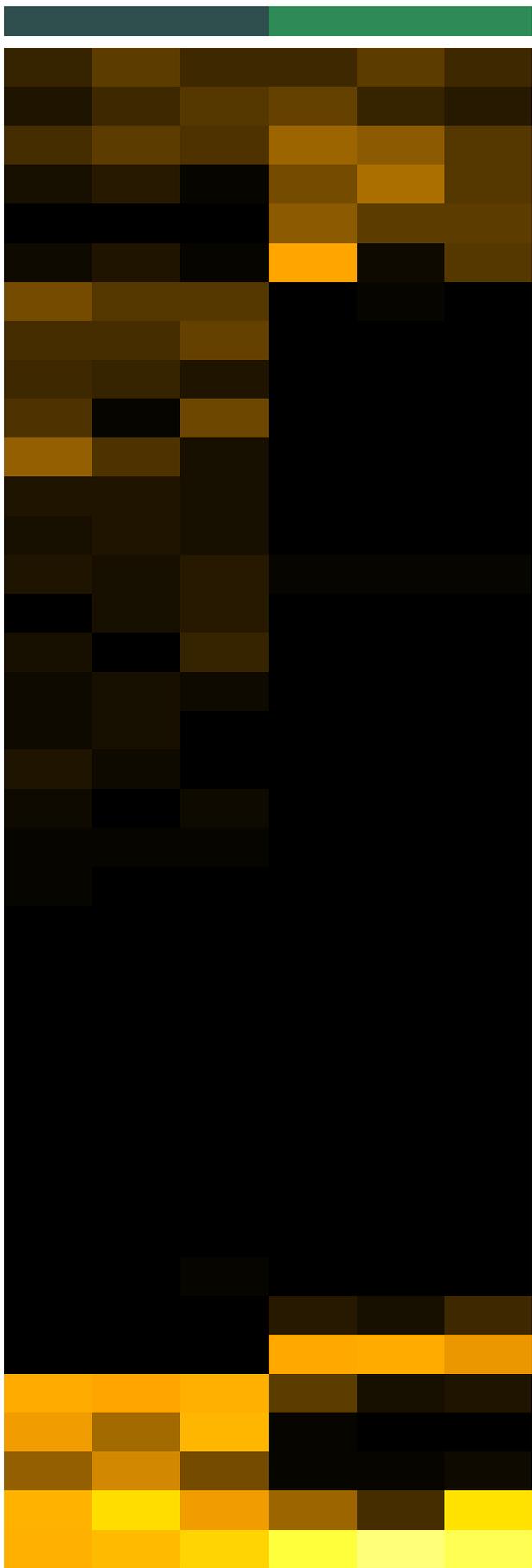
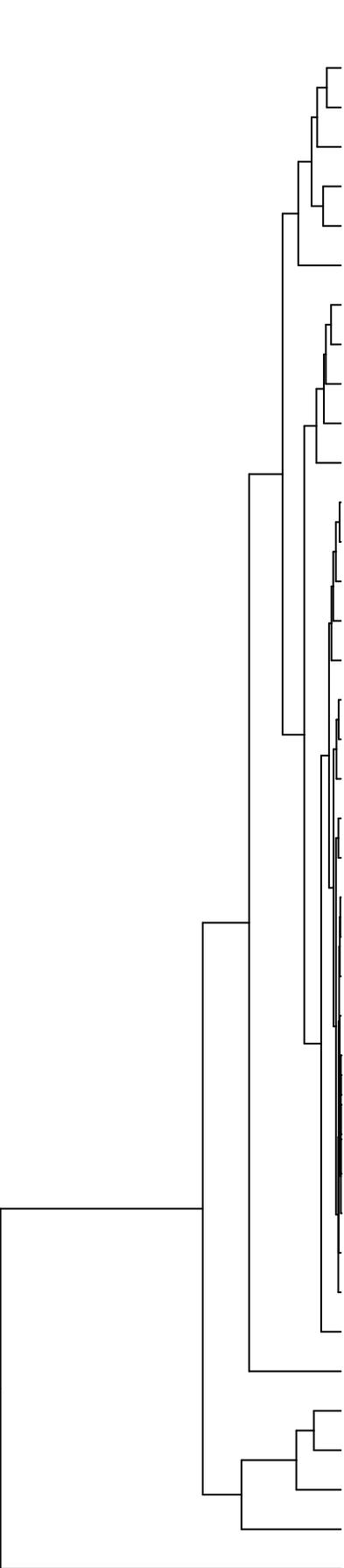




Group  
 Human Microbiome Day 0  
 Static Day 3 LR



- Fusobacteriaceae
- Fusobacteriaceae
- Porphyromonadaceae
- Gemellaceae
- Weeksellaceae
- Neisseriaceae
- Saccharibacteria\_(TM7)\_[F-1]
- Flavobacteriaceae
- Carnobacteriaceae
- Actinomycetaceae
- Selenomonadaceae
- Burkholderiaceae
- Micrococcaceae
- Absconditabacteria\_(SR1)\_[F-1]
- Leptotrichiaceae
- Actinomycetaceae
- Bacteroidales\_[F-2]
- Pasteurellaceae
- Campylobacteraceae
- Propionibacteriaceae
- Cardiobacteriaceae
- Lachnospiraceae\_[XIV]
- Lachnospiraceae\_[XIVa]
- Peptoniphilaceae
- Atopobiaceae\_Coriobacteriaceae
- Enterobacteriaceae
- Ruminococcaceae
- Peptostreptococcaceae
- Lachnospiraceae
- Lachnospiraceae\_Lachnospiraceae\_[XIV]
- Porphyromonadaceae
- Comamonadaceae
- Peptostreptococcaceae\_[XI]
- Erysipelotrichaceae
- Prevotellaceae
- Leptotrichiaceae
- Flavobacteriaceae
- Veillonellaceae
- Streptococcaceae

F7293.S01  
 F7293.S02  
 F7293.S03  
 F7293.S06  
 F7293.S09  
 F7293.S12

Samples

Family