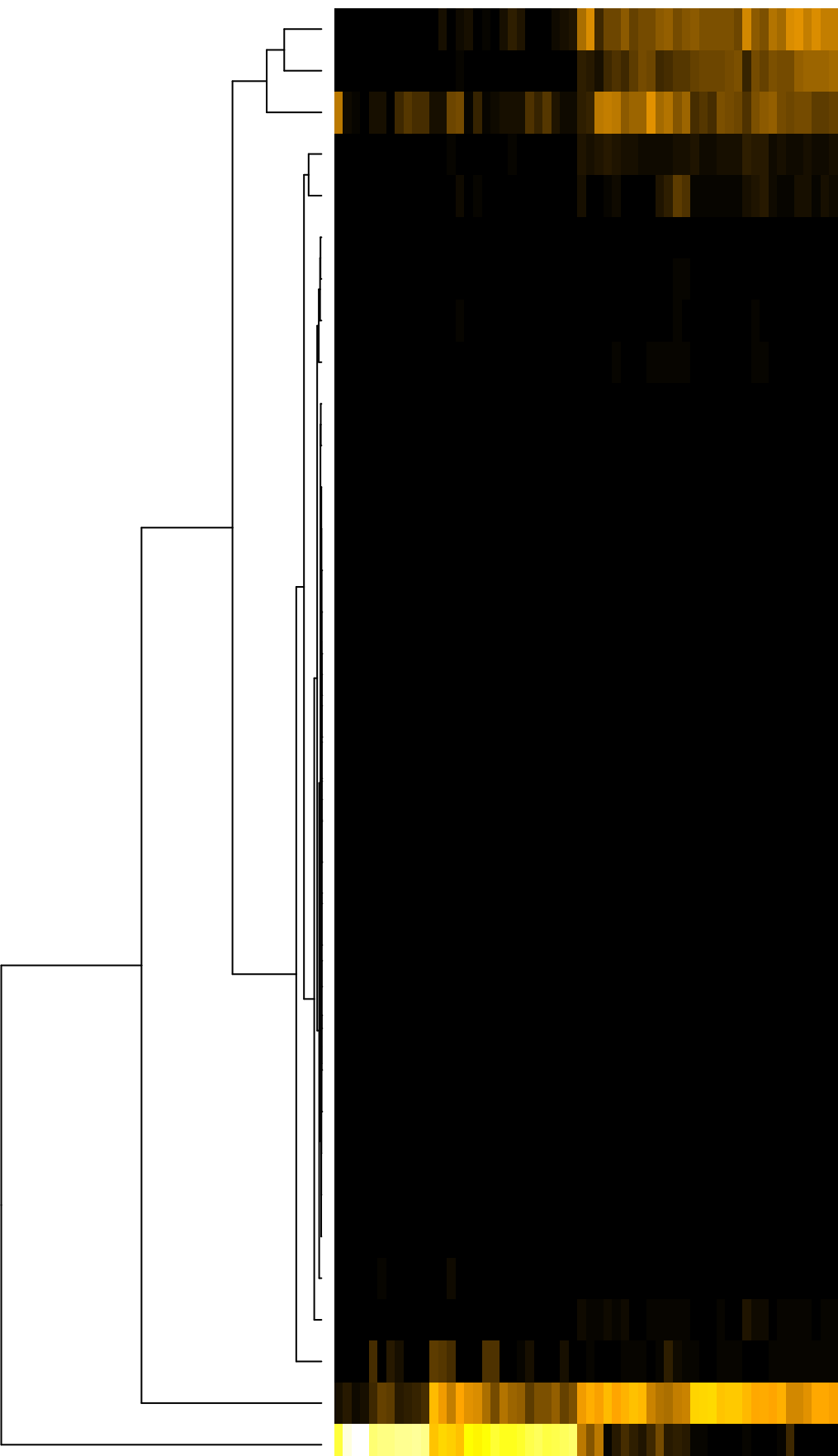


Group006
 I Final -PMA
 C Final -PMA



- Veillonellaceae
- Pasteurellaceae
- Neisseriaceae
- Carnobacteriaceae
- Fusobacteriaceae
- Porphyromonadaceae
- Leptotrichiaceae
- Campylobacteraceae
- Gemellaceae
- Prevotellaceae
- Lachnospiraceae
- Saccharibacteria_(TM7)_[F-1]
- Atopobiaceae
- Flavobacteriaceae
- Moraxellaceae
- Microbacteriaceae
- Sphingomonadaceae
- Rhizobiaceae
- Weeksellaceae
- Hymenobacteraceae
- Saccharibacteria_(TM7)_[F-2]
- Mycobacteriaceae
- Phyllobacteriaceae
- Yersiniaceae
- Absconditabacteria_(SR1)_[F-1]
- Pseudomonadaceae
- Propionibacteriaceae
- Burkholderiaceae
- Erysipelotrichaceae
- Comamonadaceae
- Staphylococcaceae
- Actinomycetaceae
- Micrococcaceae
- Streptococcaceae
- Enterobacteriaceae

- F28914.S461
- F28914.S460
- F28914.S571
- F28914.S573
- F28914.S462
- F28914.S458
- F28914.S416
- F28914.S606
- F28914.S607
- F28914.S572
- F28914.S414
- F28914.S570
- F28914.S605
- F28914.S413
- F28914.S302
- F28914.S297
- F28914.S141
- F28914.S091
- F28914.S142
- F28914.S137
- F28914.S138
- F28914.S096
- F28914.S092
- F28914.S090
- F28914.S144
- F28914.S256
- F28914.S249
- F28914.S250
- F28914.S255

Samples

Family