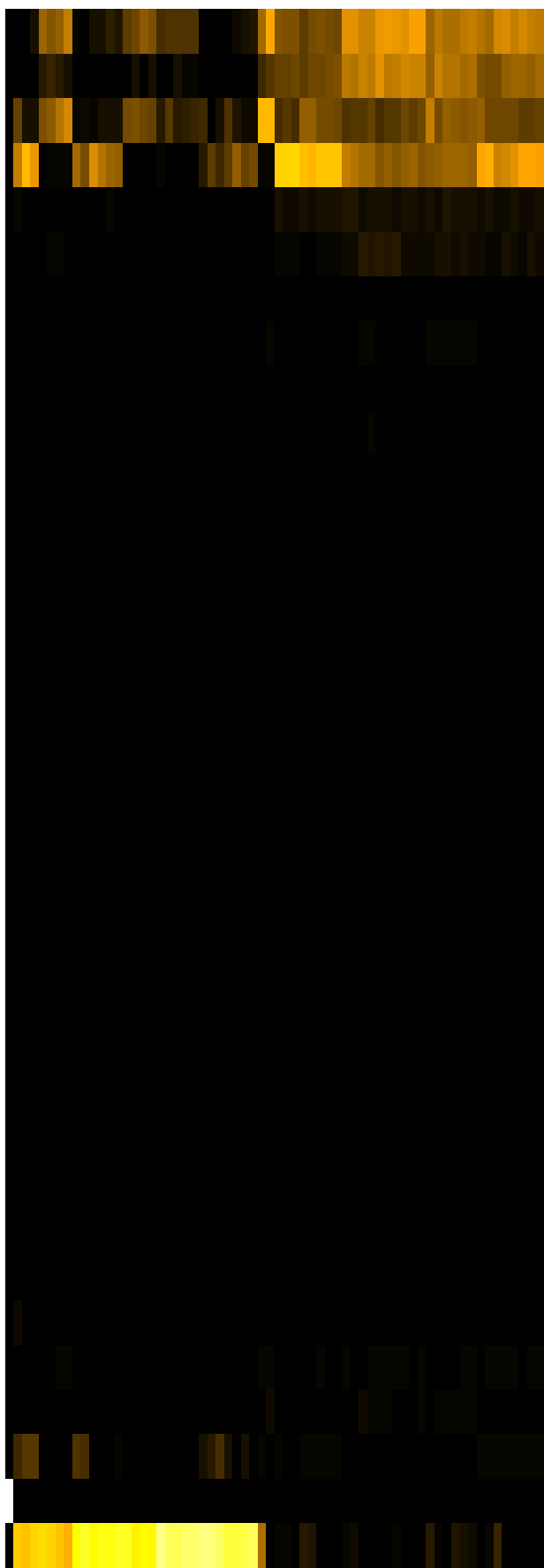


Group006
 D Final -PMA
 C Final -PMA



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

- F28914.S583
- F28914.S575
- F28914.S578
- F28914.S584
- F28914.S574
- F28914.S415
- F28914.S414
- F28914.S420
- F28914.S422
- F28914.S418
- F28914.S579
- F28914.S417
- F28914.S573
- F28914.S413
- F28914.S581
- F28914.S095
- F28914.S094
- F28914.S089
- F28914.S093
- F28914.S264
- F28914.S104
- F28914.S258
- F28914.S259
- F28914.S261
- F28914.S097
- F28914.S098
- F28914.S100
- F28914.S256
- F28914.S249
- F28914.S250
- F28914.S255

Family

Samples