



Group006
 D Final -PMA
 E Final -PMA



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

- F28914.S426
- F28914.S590
- F28914.S587
- F28914.S591
- F28914.S429
- F28914.S586
- F28914.S592
- F28914.S583
- F28914.S427
- F28914.S577
- F28914.S424
- F28914.S582
- F28914.S421
- F28914.S419
- F28914.S580
- F28914.S581
- F28914.S097
- F28914.S265
- F28914.S258
- F28914.S259
- F28914.S257
- F28914.S264
- F28914.S104
- F28914.S102
- F28914.S266
- F28914.S268
- F28914.S272
- F28914.S112
- F28914.S111
- F28914.S107
- F28914.S098
- F28914.S100

Family

Samples