



Group005
 Baseline
 Final F -PMA



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

Family

- F28914.S40
- F28914.S43
- F28914.S433
- F28914.S436
- F28914.S438
- F28914.S596
- F28914.S595
- F28914.S118
- F28914.S280
- F28914.S279
- F28914.S052
- F28914.S049
- F28914.S050
- F28914.S115
- F28914.S274
- F28914.S120
- F28914.S278
- F28914.S031
- F28914.S056
- F28914.S065
- F28914.S064
- F28914.S062
- F28914.S035
- F28914.S032
- F28914.S058
- F28914.S059
- F28914.S041
- F28914.S037
- F28914.S042
- F28914.S070
- F28914.S072
- F28914.S071
- F28914.S047
- F28914.S043
- F28914.S048
- F28914.S029
- F28914.S026
- F28914.S027

Samples