



Group005
 Baseline
 Final I -PMA

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

Family

F28914.S007
 F28914.S464
 F28914.S459
 F28914.S462
 F28914.S458
 F28914.S608
 F28914.S304
 F28914.S143
 F28914.S300
 F28914.S297
 F28914.S139
 F28914.S140
 F28914.S041
 F28914.S037
 F28914.S042
 F28914.S070
 F28914.S072
 F28914.S071
 F28914.S047
 F28914.S043
 F28914.S048
 F28914.S029
 F28914.S026
 F28914.S027
 F28914.S066
 F28914.S061
 F28914.S063
 F28914.S053
 F28914.S051
 F28914.S054
 F28914.S056
 F28914.S031
 F28914.S035
 F28914.S032
 F28914.S058
 F28914.S059

Samples