



Group005
 Baseline
 Final B -PMA



- Veillonella denticariosi_dispar_parvula
- Veillonella dispar
- Prevotella melaninogenica
- Campylobacter concisus_nov_96.963%
- Aggregatibacter segnis
- Veillonella tobetsuensis
- Gemella haemolysans
- Leptotrichia sp._HMT_215
- Aggregatibacter sp._HMT_513
- Gemella morbillorum
- Streptococcus oralis
- Streptococcus australis
- Gemella sanguinis
- Rothia mucilaginosa
- Haemophilus sputorum
- Neisseria subflava
- Schaalia odontolytica
- Streptococcus parasanguinis_clade_411
- Streptococcus gordonii
- Streptococcus sanguinis
- Eikenella corrodens
- Aggregatibacter aphrophilus
- Enterobacter mori_nov_97.951%
- Enterobacter mori
- Klebsiella aerogenes
- Raoultella planticola
- Streptococcus parasanguinis_parasanguinis_clade_721
- Granulicatella adiacens
- Veillonella parvula
- Campylobacter concisus
- Neisseria elongata
- Neisseria mucosa
- Porphyromonas pasteri
- Veillonella atypica
- Fusobacterium periodonticum
- Veillonella rogosae
- Neisseria perflava
- Streptococcus salivarius
- Streptococcus sp._HMT_423
- Salmonella enterica
- Citrobacter braakii
- Citrobacter murliniaie
- Enterobacter cancerogenus
- Enterobacter asburiae
- Klebsiella pneumoniae
- Neisseria flavescens
- Veillonella dispar_parvula
- Neisseria flava
- Haemophilus parainfluenzae
- Citrobacter koseri

Species

- F28914.S081
- F28914.S083
- F28914.S403
- F28914.S406
- F28914.S409
- F28914.S407
- F28914.S565
- F28914.S562
- F28914.S564
- F28914.S027
- F28914.S025
- F28914.S026
- F28914.S031
- F28914.S085
- F28914.S087
- F28914.S084
- F28914.S034
- F28914.S032
- F28914.S055
- F28914.S057
- F28914.S058
- F28914.S044
- F28914.S046
- F28914.S067
- F28914.S068
- F28914.S069
- F28914.S056
- F28914.S039
- F28914.S041
- F28914.S037
- F28914.S061
- F28914.S063
- F28914.S065
- F28914.S051
- F28914.S050
- F28914.S248
- F28914.S241
- F28914.S244

Samples