



Group005
 Baseline
 Final I -PMA



- Veillonella denticariosi_dispar_parvula
- Veillonella dispar
- Prevotella melaninogenica
- Campylobacter concisus_nov_96.963%
- Veillonella tobetsuensis
- Aggregatibacter sp._HMT_513
- Leptotrichia sp._HMT_215
- Fusobacterium nucleatum
- Gemella sanguinis
- Neisseria sicca
- Haemophilus sputorum
- Streptococcus oralis
- Streptococcus oralis_subsp_tigurinus_clade_070
- Rothia mucilaginosa
- Streptococcus gordonii
- Streptococcus sanguinis
- Schaalia odontolytica
- Streptococcus australis
- Streptococcus parasanguinis_clade_411
- Enterobacter asburiae
- Enterobacter mori_nov_97.951%
- Aggregatibacter aphrophilus
- Eikenella corrodens
- Aggregatibacter segnis
- Neisseria subflava
- Neisseria elongata
- Neisseria mucosa
- Veillonella parvula
- Enterobacter cancerogenus
- Klebsiella aerogenes
- Raoultella planticola
- Raoultella ornithinolytica_planticola
- Salmonella enterica
- Veillonella rogosae
- Neisseria perflava
- Porphyromonas pasteri
- Campylobacter concisus
- Streptococcus salivarius
- Streptococcus parasanguinis_parasanguinis_clade_721
- Granulicatella adiacens
- Fusobacterium periodonticum
- Veillonella atypica
- Streptococcus sp._HMT_423
- Enterobacter hormaechei
- Klebsiella pneumoniae
- Citrobacter koseri
- Veillonella dispar_parvula
- Neisseria flava
- Haemophilus parainfluenzae
- Neisseria flavescens

Species

- F28914.S459
- F28914.S457
- F28914.S464
- F28914.S463
- F28914.S608
- F28914.S606
- F28914.S036
- F28914.S035
- F28914.S057
- F28914.S058
- F28914.S047
- F28914.S043
- F28914.S046
- F28914.S070
- F28914.S069
- F28914.S068
- F28914.S028
- F28914.S029
- F28914.S030
- F28914.S297
- F28914.S302
- F28914.S303
- F28914.S139
- F28914.S137
- F28914.S138
- F28914.S144
- F28914.S033
- F28914.S053
- F28914.S051
- F28914.S054
- F28914.S064
- F28914.S066
- F28914.S062
- F28914.S042
- F28914.S040
- F28914.S038

Samples