



Group005
 Baseline
 Final E -PMA



- Leptotrichia sp._HMT_215
- Fusobacterium nucleatum
- Aggregatibacter sp._HMT_513
- Haemophilus pittmaniae
- Veillonella tobetsuensis
- Gemella haemolysans
- Schaalia odontolytica
- Streptococcus australis
- Streptococcus oralis_subsp._tigurinus_clade_070
- Neisseria sicca
- Streptococcus mitis
- Streptococcus oralis_subsp._tigurinus_clade_071
- Klebsiella pneumoniae
- Eikenella corrodens
- Streptococcus parasanguinis_clade_411
- Streptococcus gordonii
- Streptococcus sanguinis
- Aggregatibacter aphrophilus
- Veillonella denticariosi_dispar_parvula
- Veillonella dispar
- Prevotella melaninogenica
- Campylobacter concisus_nov_96.963%
- Aggregatibacter segnis
- Rothia mucilaginosa
- Gemella sanguinis
- Haemophilus sputorum
- Neisseria subflava
- Salmonella enterica
- Streptococcus sp._HMT_064
- Enterobacter cancerogenus
- Rothia dentocariosa
- Neisseria elongata
- Neisseria mucosa
- Veillonella parvula
- Veillonella atypica
- Fusobacterium periodonticum
- Streptococcus parasanguinis_parasanguinis_clade_721
- Granulicatella adiacens
- Streptococcus salivarius
- Neisseria perflava
- Veillonella rogosae
- Porphyromonas pasteri
- Campylobacter concisus
- Streptococcus oralis
- Citrobacter koseri
- Veillonella dispar_parvula
- Neisseria flava
- Neisseria flavescens
- Haemophilus parainfluenzae
- Streptococcus sp._HMT_423

Species

- F28914.S422
- F28914.S437
- F28914.S487
- F28914.S486
- F28914.S482
- F28914.S486
- F28914.S425
- F28914.S428
- F28914.S028
- F28914.S029
- F28914.S030
- F28914.S036
- F28914.S035
- F28914.S057
- F28914.S047
- F28914.S043
- F28914.S046
- F28914.S070
- F28914.S069
- F28914.S068
- F28914.S035
- F28914.S030
- F28914.S041
- F28914.S037
- F28914.S061
- F28914.S063
- F28914.S065
- F28914.S051
- F28914.S052
- F28914.S054
- F28914.S107
- F28914.S110
- F28914.S111
- F28914.S268
- F28914.S269
- F28914.S272

Samples