



Group002
 Saliva P1 -PMA
 Baseline P1 -PMA



- Leptotrichia sp._HMT_215
- Streptococcus australis
- Alloprevotella sp._HMT_473
- Alloprevotella sp._HMT_473_nov_97.347%
- Oribacterium sinus
- Prevotella jejuni
- Alloprevotella sp._HMT_914
- Absconditabacteria_(SR1)_[G-1]_bacterium_HMT_875
- Solobacterium moorei
- Streptococcus parasanguinis_clade_411
- Veillonella dispar
- Gemella haemolysans
- Veillonella denticariosi_dispar_parvula
- Veillonella tobetsuensis
- Gemella sanguinis
- Aggregatibacter segnis
- Campylobacter concisus_nov_96.963%
- Eikenella corrodens
- Streptococcus sanguinis
- Gemella morbillorum
- Prevotella nanceiensis
- Alloprevotella sp._HMT_308
- Prevotella pallens
- Leptotrichia sp._HMT_417
- Streptococcus infantis_infantis_clade_638
- Streptococcus mitis
- Granulicatella adiacens
- Citrobacter koseri
- Veillonella atypica
- Veillonella parvula
- Neisseria subflava
- Aggregatibacter aphrophilus
- Streptococcus parasanguinis_parasanguinis_clade_721
- Neisseria elongata
- Haemophilus sputorum
- Neisseria mucosa
- Rothia mucilaginos
- Schaalia odontolytica
- Streptococcus salivarius
- Neisseria flava
- Campylobacter concisus
- Veillonella dispar_parvula
- Streptococcus sp._HMT_423
- Neisseria perflava
- Veillonella rogosae
- Fusobacterium periodonticum
- Porphyromonas pasteri
- Prevotella melaninogenica
- Neisseria flavescens
- Haemophilus parainfluenzae

Species

- F28914.S028
- F28914.S027
- F28914.S029
- F28914.S025
- F28914.S030
- F28914.S026
- F28914.S004
- F28914.S002
- F28914.S006
- F28914.S001
- F28914.S003
- F28914.S005
- F28914.S042
- F28914.S039
- F28914.S040
- F28914.S041
- F28914.S038
- F28914.S037

Samples