



- Group
- Saliva Healthy Control
 - Saliva Periodontitis
- Species
- Mogibacterium diversum_neglectum_vescum
 - Veillonellaceae_[G-1] bacterium_HMT_155
 - Erysipelotrichaceae_[G-1] bacterium_HMT_905
 - Selenomonas noxia
 - Escherichia_Shigella_coli_dysenteriae_fergusonii_flexneri_sonnei
 - Christensenella multispecies_sppn2_2_nov_91.216%
 - Catenibacterium mitsuokai
 - Bacteroides fragilis
 - Roseburia intestinalis
 - Butyrivibrio crossotus
 - Holdemanella biformis
 - Bacteroides caccae
 - Blautia obeum
 - Fusicatenibacter saccharivorans
 - Agathobaculum butyriciproducens
 - Barnesiella intestinihominis
 - Megasphaera elsdenii
 - Odoribacter splanchnicus
 - Methanobrevibacter smithii
 - Parabacteroides distasonis
 - Ruminococcus callidus
 - Parabacteroides merdae
 - Alistipes putredinis
 - Mediterraneibacter faecis
 - Lactobacillus rogosae
 - Eubacterium rectale
 - Neisseria flava_macacae_mucosa_sicca_sp_TM101
 - Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_97.072%
 - Prevotella histicola
 - Ruminococcaceae_[G-2] bacterium_HMT_085
 - TM7_[G] sp_Oral_Taxon_A56
 - Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.968%
 - Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873
 - Neisseria elongata
 - Corynebacterium argentoratense
 - Mycoplasma faucium
 - Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
 - Parvimonas micra
 - Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
 - Tannerella forsythia
 - Stomatobaculum sp_HMT_097
 - Corynebacterium durum
 - Capnocytophaga leadbetteri
 - Rothia aeria
 - Prevotella salivae
 - Veillonella atypica
 - Campylobacter concisus
 - Actinomyces odontolyticus
 - Streptococcus parasanguinis parasanguinis_l parasanguinis_clade_431
 - Fusobacterium periodonticum
 - Oribacterium sinus
 - Veillonella rogosae_sp_str_NVG100cf
 - Prevotella pallens
 - Prevotella melaninogenica
 - Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345
 - Porphyromonas sp_HMT_285
 - Mobiluncus multispecies_sppn34_2_nov_97.817%
 - Bacteroidales_[G-2] sp_Oral_Taxon_274
 - Oribacterium parvum_sp_str_ACB1
 - Neisseria subflava
 - Abiotrophia defectiva
 - Actinomyces_Schaalia odontolytica_odontolyticus
 - Schaalia sp_HMT_180
 - Neisseria flavescens[subflava_perflava
 - Veillonella parvula_tobetsuensis
 - Schaalia sp_HMT_172
 - Actinomyces sp_HMT_169_sp_str_ChDCB197
 - Gemella morbillorum_morbillum
 - Bergeyella sp_HMT_907_nov_97.397%
 - Granulicatella elegans
 - Porphyromonas endodontalis
 - Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
 - Parvimonas sp_HMT_110_sp_HMT_393_sp_Oral_Taxon_110
 - Ruminococcaceae_[G-1] bacterium_HMT_075
 - Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
 - Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
 - Peptostreptococcaceae_[XII][G-1] [Eubacterium]_sulci
 - Streptococcus sanguinis_sp_HMT_074
 - Streptococcus anginosus
 - Selenomonas sp_HMT_136_sp_Oral_Taxon_149
 - Gemella sanguinis
 - Haemophilus parainfluenzae
 - Streptococcus salivarius_sp_str_ACS2_sp_str_C150_vestibulari
 - Bergeyella sp_HMT_322
 - Capnocytophaga gingivalis
 - Neisseria flavescens_flavescens[subflava
 - Actinomyces graevenitzi
 - Veillonella dispar_parvula
 - Gemella haemolysans
 - Peptostreptococcus stomatis
 - Porphyromonas pasteri
 - Rothia dentocariosa
 - Solobacterium moorei
 - Atopobium_Lancefieldella parvula_parvulum
 - Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351
 - Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352
 - Actinomyces_Schaalia odontolyticus_sp_HMT_180
 - Rothia mucilaginosa
 - Granulicatella adiacens_padiaciens
 - Streptococcus cristatus_gwangujuense_infantis_infantis_clade_431

Samples