



- Species
- Leptothrix sp._HMT_025
 - Peptidiphaga sp._HMT_183
 - Johnsonella sp._HMT_166
 - Peptostreptococcaceae_[XII][G-6] [Eubacterium]_nodatum
 - Prevotella intermedia
 - Alloprevotella tanneræ
 - Bifidobacterium dentium
 - Bordetella trematum
 - Roseomonas hibiscisoli
 - Bacteroidales_[G-2] sp._Oral_Taxon_274
 - Fusobacterium nucleatum_subsp._vincentii
 - Peptostreptococcus stomatis
 - Fretibacterium sp._HMT_359
 - Actinomyces gerencseriae
 - Lautropia mirabilis
 - Prevotella oralis
 - Treponema socranskii
 - Selenomonas artemidis
 - Kingella oralis
 - Cardiobacterium hominis
 - Prevotella sp._HMT_317
 - Corynebacterium matruchotii
 - Treponema sp._HMT_238
 - Fretibacterium sp._HMT_362
 - Haemophilus parainfluenzae
 - Gemella morbillorum
 - Arachnia propionica
 - Dialister invisus
 - Prevotella nigrescens
 - Schaalia meyeri
 - Selenomonas sp._HMT_146
 - Bacteroidetes_[G-5] bacterium_HMT_511
 - Fusobacterium sp._HMT_370
 - Capnocytophaga leadbetteri
 - Filifactor alocis
 - Solobacterium moorei
 - Enterococcus cecorum
 - Pseudomonas aeruginosa
 - Porphyromonas gingivalis
 - Rothia dentocariosa
 - Veillonellaceae_[G-1] bacterium_HMT_129
 - Porphyromonas endodontalis
 - Streptococcus anginosus
 - Fretibacterium sp._HMT_360
 - Nocardioides dilutus
 - Veillonella parvula
 - Selenomonas diana
 - Acidovorax temperans
 - Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
 - Treponema denticola
 - Acinetobacter sp._Oral_Taxon_A58
 - Treponema sp._HMT_237
 - Sphingobium yanoikuyae
 - Acinetobacter johnsonii
 - Prevotella oris
 - Tannerella forsythia
 - Fretibacterium fastidiosum
 - Sphingomonas oryzae
 - Corynebacterium durum
 - Selenomonas noxia
 - Prevotella melaninogenica
 - Actinomyces sp._HMT_170
 - Selenomonas sputigena
 - Campylobacter gracilis
 - Fusobacterium hwasookii
 - Lachnospiraceae_[G-8] bacterium_HMT_500
 - Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.968%
 - Treponema sp._HMT_927_nov_94.839%
 - Saccharibacteria_(TM7)_[G-1] bacterium_HMT_488_nov_97.065%
 - Porphyromonas sp._Oral_Taxon_B43_nov_97.619%
 - Arachnia propionica_nov_97.768%
 - Curvibacter fontanus_nov_97.645%
 - Litorilinea aerophila_nov_92.777%
 - Muribaculum intestinale_nov_92.857%
 - Sphingosinella cucumeris_nov_96.136%
 - Fusobacterium nucleatum_nucleatum_ss_animalis
 - Capnocytophaga endodontalis_sp._HMT_326
 - Fusobacterium nucleatum_nucleatum_subsp._vincentii
 - Granulicatella adiacens_paradiacens
 - Lautropia dentalis_mirabilis
 - Neisseria flava_macacae_mucosa_sicca_sp._TM101
 - Eubacterium_[XII][G-7]_Peptoanaerobacter_Peptostreptococcaceae
 - Fusobacterium nucleatum_ss_vincentii_nucleatum_subsp._vincentii
 - Streptococcus sanguinis_sp._Oral_Taxon_B66
 - Paraburkholderia caledonica_fungorum_strydomiana
 - Fusobacterium nucleatum_nucleatum_ss_vincentii_nucleatum_subsp._vincentii
 - Aggregatibacter aphrophilus_paraphrophilus
 - Cutibacterium_Propionibacterium_acnes_sp._str._LG
 - Lachnoanaerobaculum gingivalis_umeaense
 - Fusobacterium nucleatum_nucleatum_ss_nucleatum_nucleatum_subsp._vincentii
 - Pseudomonas fildesensis_veronii
 - Selenomonas sp._HMT_138_sp._HMT_892
 - Micrococcus aloeverae_luteus
 - Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431
 - Pseudomonas alcaliphila_chengduensis_mendocina_oleovorans_torreyi
 - Veillonella dispar_parvula
 - Sphingomonas kyeongiense_leidyi
 - Campylobacter rectus_showae
 - multigenus multispecies_sppn2_2_nov_97.210%
 - Nakamurella multispecies_sppn7_2_nov_95.260%

Samples