



- Actinomyces timonensis
- Actinomyces johnsonii
- Methylobacterium rhodesianum
- Actinomyces sp._oral_taxon_175
- Mogibacterium timidum
- Rothia aeria
- Streptococcus australis
- Neisseria pharyngis
- Olsenella uli
- Streptococcus infantis
- Slackia exigua
- Atopobium parvulum
- Actinomyces sp._oral_taxon_896
- Actinomyces israelii
- Filifactor alocis
- Treponema sp._oral_taxon_237
- Propionibacterium acidifaciens
- Veillonella parvula_group
- Actinomyces sp._oral_taxon_448
- Streptococcus dentisani
- Streptococcus constellatus
- Olsenella sp._oral_taxon_807
- Streptococcus sp._oral_taxon_066
- Streptococcus sp._oral_taxon_431
- Sphingobium japonicum
- Neisseria flavescens|subflava
- Treponema lecithinolyticum
- Rothia mucilaginosus
- Treponema socranskii
- Peptostreptococcaceae_[X1][G-1] [Eubacterium]_infirum
- Actinomyces massiliensis
- Streptococcus sp._oral_taxon_058
- Mogibacterium diversum
- Porphyromonas gingivalis
- Tannerella forsythia
- Actinomyces sp._oral_taxon_170
- Brochothrix thermosphacta
- Actinomyces odontolyticus
- Peptostreptococcaceae_[X1][G-5] [Eubacterium]_saphenum
- Abiotrophia defectiva
- Methylobacterium sp._Oral_Taxon_B84
- Actinomyces gerencseriae
- Atopobium sp._oral_taxon_199
- Streptococcus mitis
- Actinomyces sp._oral_taxon_897
- Noviherbaspirillum suwonense
- Streptococcus gordonii
- Streptococcus sanguinis
- Treponema sp._oral_taxon_258
- Actinomyces sp._oral_taxon_525
- TM7_[G-5] sp._oral_taxon_356
- Planococcus antarcticus
- Clostridiales_[G] sp._Oral_Taxon_G74
- Peptoniphilaceae_[G-1] sp._oral_taxon_113
- Actinomyces sp._oral_taxon_180
- Atopobium rimae
- Rhizobium rhizogenes_Oral_Taxon_D34
- Actinomyces oricola
- Streptococcus sp._oral_taxon_423
- TM7_[G-1] sp._oral_taxon_346
- Actinomyces naeslundii
- Selenomonas sputigena
- Actinomyces cardiffensis
- Streptococcus downei
- Actinomyces dentalis
- Rothia dentocariosa
- Actinomyces sp._oral_taxon_169
- Peptostreptococcus stomatis
- Actinomyces meyeri
- Brevundimonas diminuta
- Propionibacterium propionicum
- Anaerolineae_[G-1] sp._oral_taxon_439
- TM7_[G-1] sp._oral_taxon_349
- Haemophilus parainfluenzae
- Solobacterium moorei
- Actinomyces georgiae
- Peptostreptococcaceae_[X1][G-9] [Eubacterium]_brachy
- Actinomyces sp._oral_taxon_414
- Bacteroidales_[G-2] sp._oral_taxon_274
- Desulfohalobium sp._oral_taxon_041
- Fusobacterium nucleatum_subsp._vincentii
- Fretibacterium fastidiosum
- Streptococcus sp._oral_taxon_064
- Parvimonas micra
- Propionibacterium acnes
- Fusobacterium sp._oral_taxon_203
- Parvimonas sp._oral_taxon_110
- Streptococcus anginosus
- Carnobacterium divergens
- Porphyromonas endodontalis
- Actinobaculum sp._oral_taxon_183
- Granulicatella adiacens
- Actinomyces oris
- Actinomyces timonensis_nov_93.927%
- Pseudomonas stutzeri_Oral_Taxon_A68_nov_83.229%
- TM7_[G-1] sp._oral_taxon_349_nov_97.788%
- Acinetobacter sp._oral_taxon_408_nov_92.480%
- Actinomyces israelii_nov_96.857%
- Reyranella massiliensis_soli
- Mogibacterium multispecies_sppn1_2_nov_91.393%

Species

Samples