



- Streptococcus salivarius
- Leptotrichia wadei
- Leptotrichia hongkongensis
- Aggregatibacter sp.\_HMT\_458
- Capnocytophaga leadbetteri
- Prevotella sp.\_HMT\_317
- Anaeroglobus geminatus
- Peptidiphaga sp.\_HMT\_183
- Selenomonas sp.\_HMT\_137
- Prevotella sp.\_HMT\_314
- Tannerella sp.\_HMT\_286
- Leptotrichia sp.\_HMT\_392
- Actinomyces naeslundii
- Lancefieldella rimae
- Cardiobacterium hominis
- Sphingomonas oligophenolica
- Arachnia propionica
- Tannerella serpentiformis
- Streptococcus chosunense
- Prevotella oulorum
- Streptococcus intermedius
- Arachnia rubra
- Selenomonas flueggei
- Capnocytophaga sp.\_HMT\_864
- Gemella morbillorum
- Campylobacter gracilis
- Rothia dentocariosa
- Scardovia wiggsiae
- Saccharibacteria\_(TM7)\_[G-2] bacterium\_HMT\_350
- Prevotella veroralis
- Prevotella sp.\_HMT\_292
- Veillonella parvula
- Actinomyces sp.\_HMT\_175
- Schaalia sp.\_HMT\_180
- Dialister invisus
- Prevotella salivae
- Peptostreptococcus stomatis
- Leptotrichia sp.\_HMT\_498
- Solobacterium moorei
- Prevotella intermedia
- Stomatobaculum longum
- Neisseria elongata
- Streptococcus sp.\_HMT\_064
- Streptococcus mitis
- Neisseria flavescens
- Selenomonas noxia
- Porphyromonas endodontalis
- Prevotella oris
- Selenomonas sputigena
- Bacteroidales\_[G-2] bacterium\_HMT\_274
- Prevotella sp.\_HMT\_300
- Stenotrophomonas maltophilia
- Streptococcus oralis\_subsp\_tigurinus\_clade\_070
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_348
- Neisseria sp.\_HMT\_020
- Aggregatibacter segnis
- Prevotella sp.\_HMT\_820
- Porphyromonas pasteri
- Saccharibacteria\_(TM7)\_[G-5] bacterium\_HMT\_356
- Selenomonas infelix
- Gracilibacteria\_(GN02)\_[G-2] bacterium\_HMT\_873
- Streptococcus gordonii
- Peptostreptococcaceae\_[XII][G-4] bacterium\_HMT\_369
- Kingella oralis
- Campylobacter concisus
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_869
- Prevotella melaninogenica
- Haemophilus parainfluenzae
- Leptotrichia hofstadii
- Granulicatella adiacens
- Corynebacterium matruchotii
- Campylobacter showae
- Capnocytophaga granulosa
- Actinomyces sp.\_HMT\_171
- Fusobacterium nucleatum
- Treponema socranskii
- Parvimonas micra
- Prevotella maculosa
- Alloprevotella tannerae
- Prevotella denticola
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_957
- Streptococcus oralis
- Actinomyces sp.\_HMT\_169
- Saccharibacteria\_(TM7)\_[G-4] bacterium\_HMT\_355
- Tannerella forsythia
- Peptostreptococcaceae\_[XII][G-1] [Eubacterium]\_infirmum
- Bacteroidetes\_[G-3] bacterium\_HMT\_280
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_346
- Bacteroides zoogloformans\_nov\_96.578%
- Aggregatibacter sp.\_HMT\_458\_nov\_97.826%
- Sphingomonas jeddahensis\_nov\_97.447%
- Saccharibacteria\_(TM7)\_[G-3] bacterium\_HMT\_351\_nov\_92.871%
- Neisseria sp.\_HMT\_020\_nov\_97.619%
- Prevotella melaninogenica\_nov\_97.170%
- Streptococcus sanguinis\_nov\_97.538%
- Selenomonas noxia\_nov\_96.337%
- Selenomonas sp.\_HMT\_137\_nov\_97.901%
- Corynebacterium durum\_nov\_97.384%
- Selenomonas sp.\_HMT\_479\_nov\_96.756%
- Veillonella dispar\_parvula

Samples