



- Species
- Porphyromonas gingivalis
  - Gemella morbillorum
  - Capnocytophaga gingivalis
  - Campylobacter concisus
  - Actinomyces sp.\_HMT\_448
  - Lachnoanaerobaculum saburrium
  - Peptidiphaga sp.\_HMT\_183
  - Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_957
  - Streptococcus sobrinus
  - Leptotrichia sp.\_HMT\_392
  - Escherichia coli
  - Streptococcus oralis\_subsp.\_tigurinus\_clade\_070
  - Alloprevotella sp.\_HMT\_473
  - Streptococcus sanguinis
  - Actinomyces gerencseriae
  - Leptotrichia sp.\_HMT\_221
  - Enterococcus faecalis
  - Pseudomonas aeruginosa
  - Porphyromonas pasteri
  - Actinomyces sp.\_HMT\_175
  - Bergeyella sp.\_HMT\_322
  - Selenomonas artemidis
  - Leptotrichia hongkongensis
  - Actinomyces dentalis
  - Actinomyces oris
  - Neisseria sicca
  - Granulicatella elegans
  - Schaalia sp.\_HMT\_178
  - Prevotella loescheii
  - Actinomyces naeslundii
  - Actinomyces sp.\_HMT\_414
  - Capnocytophaga leadbetteri
  - Haemophilus haemolyticus
  - Kingella oralis
  - Schaalia meyeri
  - Neisseria cinerea
  - Actinomyces viscosus
  - Granulicatella adiacens
  - Streptococcus mutans
  - Actinomyces sp.\_HMT\_170
  - Prevotella melaninogenica
  - Capnocytophaga granulosa
  - Streptococcus chosunense
  - Listeria monocytogenes
  - Campylobacter showae
  - Bacillus halotolerans
  - Streptococcus intermedius
  - Streptococcus oralis
  - Truepera radiovictrix
  - Salmonella enterica
  - Lautropia mirabilis
  - Fusobacterium nucleatum
  - Neisseria subflava
  - Prevotella intermedia
  - Streptococcus sp.\_HMT\_064
  - Gemella haemolysans
  - Leptotrichia buccalis
  - Actinomyces sp.\_HMT\_897
  - Streptococcus gordonii
  - Streptococcus mitis
  - Prevotella nigrescens
  - Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_348
  - Leptotrichia sp.\_HMT\_212
  - Leptotrichia sp.\_HMT\_215
  - Prevotella oulorum
  - Ruminococcaceae\_[G-1] bacterium\_HMT\_075
  - Veillonella parvula
  - Corynebacterium matruchotii
  - Abiotrophia defectiva
  - Escherichia fergusonii
  - Veillonella sp.\_HMT\_780
  - Schaalia odontolytica
  - Ottowia sp.\_HMT\_894
  - Porphyromonas sp.\_HMT\_278
  - Streptococcus cristatus
  - Leptotrichia wadei
  - Prevotella oris
  - Capnocytophaga sputigena
  - Peptoanaerobacter [Eubacterium] yurii
  - Neisseria elongata
  - Limosilactobacillus fermentum
  - Rothia dentocariosa
  - Campylobacter gracilis
  - Shigella sonnei
  - Schaalia sp.\_HMT\_180
  - Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_346
  - Aggregatibacter segnis
  - Alloprevotella tanneriae
  - Allobacillus halotolerans
  - Haemophilus parainfluenzae
  - Streptococcus salivarius
  - Actinomyces massiliensis
  - Capnocytophaga endodontalis
  - Actinomyces sp.\_HMT\_169
  - Frondebacter mangrovi\_nov\_92.771%
  - Saccharibacteria\_(TM7)\_[G-6] bacterium\_HMT\_870\_nov\_97.093%
  - Actinomyces sp.\_HMT\_175\_nov\_97.830%
  - Veillonella sp.\_HMT\_780\_nov\_97.519%
  - Staphylococcus argenteus\_aureus\_roterodami
  - Veillonella dispar\_parvula

Samples