



- Actinomyces sp._HMT_175_nov_97.633%
- Streptococcus sp._HMT_064
- Actinomyces sp._HMT_448
- Leptotrichia wadei
- Streptococcus cristatus
- Streptococcus chosunense
- Leptotrichia sp._HMT_221
- Schaalia sp._HMT_180
- Capnocytophaga gingivalis
- Prevotella melaninogenica
- Granulicatella adiacens
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Kingella oralis
- Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870_nov_97.093%
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- Prevotella intermedia
- Neisseria cinerea
- Streptococcus oralis_subsp_tigurinus_clade_070
- Campylobacter gracilis
- Schaalia meyeri
- Staphylococcus warneri
- Erythrobacter aureus
- Aeromicrobium panaciterrae_nov_96.122%
- Streptococcus mutans
- Parvimonas micra
- Escherichia coli
- Prevotella nigrescens
- Leptotrichia hongkongensis
- Ottowia sp._HMT_894
- Actinomyces viscosus
- Streptococcus intermedius
- Leptotrichia sp._HMT_392
- Actinomyces oris
- Actinomyces sp._HMT_169
- Actinomyces sp._HMT_170
- Streptococcus sobrinus
- Streptococcus salivarius
- Abiotrophia defectiva
- Leptotrichia sp._HMT_212
- Neisseria sicca
- Lautropia mirabilis
- Corynebacterium matruchotii
- Alloprevotella sp._HMT_473
- Rothia dentocariosa
- Streptococcus oralis
- Fusobacterium nucleatum
- Streptococcus gordonii
- Streptococcus sanguinis
- Schaalia odontolytica
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- Capnocytophaga leadbetteri
- Escherichia fergusonii
- Shigella sonnei
- Pseudomonas aeruginosa
- Prevotella oris
- Neisseria subflava
- Listeria monocytogenes
- Limosilactobacillus fermentum
- Staphylococcus argenteus_aureus_roterodami
- Enterococcus faecalis
- Veillonella dispar_parvula
- Veillonella parvula
- Streptococcus mitis
- Bacillus halotolerans
- Salmonella enterica
- Frondebacter mangrovi_nov_92.771%
- Allobacillus halotolerans
- Porphyromonas gingivalis
- Truepera radiovictrix

F-12829_S01
 F-12829_S02
 F-12829_S03
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 F-12829_S24
 F-12829_S34
 F-12829_S35
 F-12829_S36

Species

Samples