



- Aggregatibacter actinomycetemcomitans
- Arachnia propionica
- Fusobacterium periodonticum
- Lachnospiraceae_[G-2] bacterium_HMT_088
- Fusobacterium hwasookii
- Alloprevotella sp._HMT_913
- Prevotella falsenii
- Peptostreptococcus stomatis
- Streptococcus sinensis
- Fusobacterium nucleatum
- Porphyromonas pasteri
- Fusobacterium nucleatum_subsp._vincentii
- Fusobacterium simiae
- Tannerella serpentiformis
- Bacteroidetes_[G-5] bacterium_HMT_511
- Cardiobacterium hominis
- Fusobacterium sp._HMT_204
- Aggregatibacter sp._HMT_513_nov_97.331%
- Leptotrichia sp._HMT_212_nov_93.043%
- Catonella morbi_nov_97.908%
- Abiotrophia defectiva_nov_95.609%
- Bacteroidales_[G-2] bacterium_HMT_274_nov_97.930%
- Leptotrichia sp._HMT_909_nov_96.739%
- Prevotella nanceiensis_nov_90.361%
- Haemophilus parainfluenzae_nov_94.082%
- Aggregatibacter aphrophilus_nov_95.697%
- Streptococcus oralis_subsp._tigurinus_clade_071_nov_96.162%
- Aggregatibacter aphrophilus_nov_96.524%
- Moraxella oblonga_nov_94.387%
- Aggregatibacter actinomycetemcomitans_nov_97.541%
- Capnocytophaga gingivalis_nov_97.689%
- Streptococcus sinensis_nov_91.683%
- Veillonella parvula_nov_95.874%
- Streptococcus gordonii_nov_93.638%
- Streptococcus gordonii_nov_94.036%
- Capnocytophaga sp._HMT_878_nov_97.028%
- Capnocytophaga sp._HMT_412_nov_95.597%
- Lachnospiraceae_[G-3] bacterium_HMT_100_nov_96.868%
- Aggregatibacter sp._HMT_898_nov_97.331%
- Neisseria cinerea_nov_97.951%
- Streptococcus troglodytidis_nov_92.079%
- Riemerella sp._HMT_322_nov_97.942%
- Porphyromonas pasteri_nov_96.939%
- Johnsonella sp._HMT_166_nov_94.154%
- Porphyromonas sp._HMT_275_nov_97.125%
- Alloprevotella sp._HMT_473_nov_95.918%
- Alloprevotella sp._HMT_473_nov_96.311%
- Fusobacterium hwasookii_sp._HMT_204
- Porphyromonas multispecies_sppn2_2_nov_96.115%
- multigenus multispecies_sppn9_2_nov_96.800%

Species

F25123.S45

F25123.S44

F25123.S43

F25123.S42

F25123.S41

F25123.S50

F25123.S49

F25123.S48

F25123.S47

F25123.S46

Samples