



- Aggregatibacter actinomycetemcomitans
- Arachnia propionica
- Leptotrichia sp._HMT_392
- Fusobacterium periodonticum
- Lachnospiraceae_[G-2] bacterium_HMT_088
- Catonella morbi
- Fusobacterium hwasookii
- Corynebacterium matruchotii
- Prevotella falsenii
- Campylobacter showae
- Fusobacterium nucleatum
- Tannerella forsythia
- Porphyromonas pasteri
- Streptococcus cristatus_clade_578
- Fusobacterium nucleatum_subsp._vincentii
- Fusobacterium simiae
- Fusobacterium sp._HMT_204
- Parvimonas sp._HMT_110
- 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%
- 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 endodontalis_nov_95.560%
- Capnocytophaga gingivalis_nov_97.689%
- Pseudoleptotrichia goodfellowii_nov_92.241%
- Capnocytophaga sputigena_nov_96.822%
- Streptococcus gordonii_nov_94.036%
- Eikenella sp._HMT_011_nov_96.728%
- Aggregatibacter sp._HMT_898_nov_94.490%
- Capnocytophaga sp._HMT_332_nov_95.763%
- Capnocytophaga sp._HMT_412_nov_95.597%
- Neisseria flava_nov_97.947%
- Campylobacter rectus_nov_96.957%
- Propionivibrio dicarboxylicus_nov_93.802%
- Porphyromonas sp._HMT_278_nov_96.107%
- Porphyromonas sp._HMT_275_nov_97.125%
- Prevotella sp._HMT_317_nov_97.947%
- Peptococcus sp._HMT_167_nov_94.336%
- Fusobacterium hwasookii_sp._HMT_204
- Porphyromonas multispecies_sppn2_2_nov_96.115%
- Capnocytophaga multispecies_sppn22_2_nov_96.855%
- multigenus multispecies_sppn9_2_nov_96.800%

Species

- F25123.S35
- F25123.S34
- F25123.S33
- F25123.S32
- F25123.S31
- F25123.S40
- F25123.S39
- F25123.S38
- F25123.S37
- F25123.S36

Samples