

Species

- SP130 Prevotella rosea
- SP131 Kingella oralis
- SP132 Fusobacterium naviforme
- SP134 Streptococcus oralis_subsp._dentisani_clade_058
- SP135 Neisseria perflava
- SP138 Alloprevotella tannerae
- SP139 Weeksellaceae_[G-1] sp._HMT_900
- SP14 Fretibacterium sp._HMT_362
- SP140 Centipeda periodontii
- SP141 Pseudoleptotrichia sp._HMT_221
- SP143 Capnocytophaga sp._HMT_323
- SP146 Serratia marcescens
- SP147 Fusobacterium nucleatum_subsp._animalis
- SP149 Fusobacterium sp._HMT_204
- SP15 Prevotella baroniae
- SP151 Ruminococcaceae_[G-2] bacterium_HMT_085
- SP152 Alloprevotella sp._HMT_912
- SP153 Streptococcus infantis_clade_638
- SP154 Streptococcus oralis_subsp._oralis
- SP156 Selenomonas infelix
- SP157 Selenomonas artemidis
- SP16 Veillonella rogosae
- SP160 Veillonellaceae_[G-1] bacterium_HMT_155
- SP161 Schaalia sp._HMT_877
- SP162 Prevotella oulorum
- SP163 Capnocytophaga sp._HMT_338
- SP165 Actinomyces dentalis
- SP166 Dialister invisus
- SP167 Burkholderia cepacia
- SP169 Selenomonas sp._HMT_920
- SP17 Prevotella denticola
- SP170 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- SP172 Pseudoleptotrichia goodfellowii
- SP173 Aggregatibacter segnis
- SP174 Actinomyces johnsonii
- SP178 Schaalia sp._HMT_178
- SP179 Streptococcus salivarius
- SP180 Streptococcus oralis_subsp._tigurinus_clade_071
- SP181 Porphyromonas sp._HMT_275
- SP182 Leptotrichia hofstadii

- SP202 Campylobacter gracilis
- SP263 Selenomonas sp._HMT_137
- SP266 Porphyromonas gingivalis
- SP267 Capnocytophaga sp._HMT_335
- SP268 Selenomonas sp._HMT_479
- SP269 Johnsonella ignava
- SP27 Streptococcus sp._HMT_423
- SP270 Bulleidia extructa
- SP272 Schaalia sp._HMT_180
- SP275 Mycoplasma salivarium
- SP276 Leptotrichia sp._HMT_498
- SP277 Actinomyces massiliensis
- SP278 Aggregatibacter sp._HMT_898
- SP279 Aggregatibacter sp._HMT_513
- SP28 Fusobacterium nucleatum_subsp._vincentii
- SP280 Leptotrichia sp._HMT_223
- SP282 Streptococcus sp._HMT_056
- SP284 Leptotrichia shahii
- SP285 Arachnia rubra
- SP288 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- SP289 Filifactor alocis
- SP29 Fusobacterium nucleatum_subsp._polymorphum
- SP293 Ruminococcaceae_[G-1] bacterium_HMT_075
- SP294 Bacteroidetes_[G-3] bacterium_HMT_280
- SP295 Lancefieldella parvula
- SP296 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870
- SP3 Leptotrichia hongkongensis
- SP30 Prevotella saccharolytica
- SP302 Peptostreptococcus stomatis
- SP304 Lachnospiraceae_[G-7] bacterium_HMT_163
- SP306 Lancefieldella rimae
- SP308 Oribacterium sp._HMT_078
- SP309 Prevotella dentalis
- SP31 Capnocytophaga sputigena
- SP310 Neisseria bacilliformis
- SP311 Streptococcus australis
- SP312 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP314 Lachnoanaerobaculum sp._HMT_083
- SP32 Prevotella intermedia
- SP321 Porphyromonas sp._HMT_277

- SP53 Prevotella melaninogenica
- SP67 Leptotrichia sp._HMT_463
- SP68 Ottowia sp._HMT_894
- SP69 Prevotella salivae
- SP7 Prevotella sp._HMT_317
- SP70 Haemophilus parahaemolyticus
- SP72 Abiotrophia defectiva
- SP73 Kingella denitrificans
- SP74 Porphyromonas catoniae
- SP75 Aggregatibacter sp._HMT_458
- SP76 Tannerella sp._HMT_808
- SP77 Peptococcus sp._HMT_168
- SP78 Prevotella veroralis
- SP79 Gemella morbillorum
- SP8 Gemella haemolysans
- SP80 Leptotrichia hofstadii
- SP83 Prevotella pleuritidis
- SP84 Streptococcus oralis_subsp._tigurinus_clade_070
- SP85 Cardiobacterium valvarum
- SP86 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874
- SP87 Prevotella sp._HMT_472
- SP88 Capnocytophaga sp._HMT_332
- SP89 Catonella morbi
- SP9 Fretibacterium fastidiosum
- SP91 Campylobacter showae
- SP92 Streptococcus sp._HMT_064
- SP94 Actinomyces sp._HMT_171
- SP95 Alloprevotella sp._HMT_473
- SP99 Parvimonas micra
- SPN106 Weeksellaceae_[G-1] sp._HMT_900_nov_90.760%
- SPN11 Porphyromonas sp._HMT_275_nov_97.947%
- SPN125 Prevotella sp._HMT_300_nov_97.347%
- SPN13 Porphyromonas catoniae_nov_97.331%
- SPN149 Cardiobacterium hominis_nov_97.510%
- SPN159 Porphyromonas sp._HMT_284_nov_97.746%
- SPN169 Lachnospiraceae_[G-2] bacterium_HMT_096_nov_87.526%
- SPN179 Corynebacterium matruchotii_nov_97.942%
- SPN188 Streptococcus sanguinis_nov_97.778%
- SPN198 Peptidiphaga gingivicola_nov_96.502%
- SPN199 Capnocytophaga gingivalis_nov_97.925%