

Species

- SP136 Actinomyces sp._HMT_169
- SP137 Prevotella vespertina
- SP138 Gemella morbillorum
- SP139 Prevotella veroralis
- SP14 Kingella denitrificans
- SP140 Streptococcus sp._HMT_064
- SP142 Treponema sp._HMT_231
- SP143 Mogibacterium diversum
- SP144 Prevotella denticola
- SP145 Fusobacterium sp._HMT_203
- SP148 Fusobacterium sp._HMT_204
- SP149 Schaalia odontolytica
- SP15 Corynebacterium matruchotii
- SP150 Capnocytophaga sp._HMT_901
- SP151 Shuttleworthia satelles
- SP153 Veillonella rogosae
- SP154 Prevotella sp._HMT_314
- SP155 Prevotella micans
- SP156 Peptostreptococcaceae_[G-9] [Eubacterium]_brachy
- SP157 Pseudoleptotrichia goodfellowii
- SP158 Stomatobaculum sp._HMT_097
- SP159 Treponema sp._HMT_257
- SP16 Capnocytophaga leadbetteri
- SP160 Prevotella sp._HMT_317
- SP161 Peptococcus sp._HMT_168
- SP162 Neisseria elongata
- SP163 Streptococcus sanguinis
- SP165 Butyrivibrio sp._HMT_455
- SP166 Haemophilus parahaemolyticus
- SP167 Peptococcus sp._HMT_167
- SP169 Neisseria cinerea
- SP17 Granulicatella adiacens
- SP173 Leptotrichia sp._HMT_392
- SP175 Neisseria sicca
- SP176 Streptococcus intermedius
- SP179 Capnocytophaga sp._HMT_878
- SP180 Gemella haemolysans
- SP181 Porphyromonas gingivalis
- SP182 Lachnoanaerobaculum saburreum
- SP183 Prevotella jejuni

- SP267 Prevotella pleuritidis
- SP268 Actinomyces sp._HMT_171
- SP273 Fusobacterium nucleatum_subsp._vincentii
- SP275 Neisseria flava
- SP277 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_871
- SP279 Capnocytophaga sp._HMT_332
- SP28 Stomatobaculum longum
- SP282 Peptostreptococcus stomatis
- SP283 Fusobacterium hwasookii
- SP285 Aggregatibacter sp._HMT_512
- SP287 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345
- SP288 Campylobacter showae
- SP29 Streptococcus downii
- SP293 Treponema denticola
- SP296 Veillonella sp._HMT_917
- SP3 Porphyromonas pasteri
- SP30 Leptotrichia buccalis
- SP302 Aggregatibacter sp._HMT_949
- SP305 Haemophilus sputorum
- SP31 Veillonella parvula
- SP311 Ottowia sp._HMT_894
- SP312 Prevotella intermedia
- SP317 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- SP318 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351
- SP32 Veillonella dispar
- SP320 Cloacibacterium sp._HMT_206
- SP33 Streptococcus oralis
- SP332 Rothia mucilaginosa
- SP336 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- SP34 Streptococcus infantis_clade_431
- SP340 Streptococcus sp._HMT_074
- SP343 Campylobacter gracilis
- SP344 Fusobacterium sp._HMT_248
- SP347 Pseudomonas aeruginosa
- SP35 Pseudostreptobacillus hongkongensis
- SP36 Streptococcus gordonii
- SP365 Haemophilus influenzae
- SP37 Streptococcus oralis_subsp._dentisani_clade_058
- SP370 Neisseria sp._HMT_018
- SP38 Veillonella sp._HMT_780

- SP85 Lachnospiraceae_[G-2] bacterium_HMT_096
- SP86 Fusobacterium periodonticum
- SP87 Abiotrophia defectiva
- SP88 Aggregatibacter sp._HMT_458
- SP89 Leptotrichia sp._HMT_218
- SP9 Oribacterium parvum
- SP90 Filifactor alocis
- SP91 Streptococcus sp._HMT_057
- SP92 Streptococcus oralis_subsp._tigurinus_clade_071
- SP93 Streptococcus anginosus
- SP94 Alloprevotella sp._HMT_914
- SP95 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- SP96 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872
- SP97 Alloprevotella rava
- SP98 Prevotella oulorum
- SP99 Selenomonas sp._HMT_136
- SPN100 Corynebacterium matruchotii_nov_97.546%
- SPN101 Pseudoleptotrichia goodfellowii_nov_91.416%
- SPN102 Capnocytophaga gingivalis_nov_97.899%
- SPN103 Prevotella jejuni_nov_97.755%
- SPN104 Porphyromonas sp._HMT_284_nov_97.746%
- SPN106 Streptobacillus notomysis_nov_93.856%
- SPN107 Corynebacterium durum_nov_97.280%
- SPN108 Granulicatella elegans_nov_97.988%
- SPN109 Porphyromonas catoniae_nov_97.746%
- SPN110 Peptoanaerobacter [Eubacterium] yurii_nov_90.795%
- SPN111 Streptococcus sanguinis_nov_97.782%
- SPN112 Fusobacterium hwasookii_nov_97.783%
- SPN146 Schaalia sp._HMT_180_nov_97.930%
- SPN148 Streptococcus parasanguinis_clade_411_nov_97.773%
- SPN15 Prevotella histicola_nov_97.347%
- SPN156 Alloprevotella sp._HMT_308_nov_97.347%
- SPN169 Ruminococcaceae_[G-1] bacterium_HMT_075_nov_91.453%
- SPN184 Alloprevotella sp._HMT_914_nov_97.551%
- SPN188 Veillonella sp._HMT_780_nov_97.628%
- SPN210 Selenomonas sputigena_nov_97.000%
- SPN220 Pseudoleptotrichia goodfellowii_nov_92.060%
- SPN231 Moraxella boevei_nov_92.807%
- SPN240 Lachnospiraceae_[G-2] bacterium_HMT_088_nov_93.096%
- SPN25 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870_nov_96.994%