

Species

- SP130 Pseudoeptotrichia sp._HMT_221
- SP131 Schaalia georgiae
- SP132 Eikenella corrodens
- SP133 Selenomonas sp._HMT_137
- SP134 Neisseria perflava
- SP135 Neisseria elongata
- SP136 Neisseria sp._HMT_499
- SP137 Streptococcus cristatus_clade_578
- SP14 Corynebacterium durum
- SP140 Peptostreptococcaceae_[G-7] [Eubacterium]_yurii_subsp._yurii_&_margaretiae
- SP142 Selenomonas sp._HMT_920
- SP143 Haemophilus sputorum
- SP144 Kingella denitrificans
- SP145 Porphyromonas catoniae
- SP146 Prevotella saccharolytica
- SP148 Leptotrichia sp._HMT_417
- SP149 Ruminococcaceae_[G-1] bacterium_HMT_075
- SP15 Prevotella veroralis
- SP151 Aggregatibacter segnis
- SP152 Actinomyces sp._HMT_170
- SP153 Tannerella serpentinaformis
- SP154 Lachnoanaerobaculum orale
- SP156 Streptococcus sobrinus
- SP157 Actinomyces israelii
- SP159 Capnocytophaga sp._HMT_863
- SP16 Alloprevotella sp._HMT_913
- SP160 Streptococcus cristatus
- SP162 Prevotella sp._HMT_942
- SP163 Streptococcus oralis
- SP164 Bacteroidales_[G-2] bacterium_HMT_274
- SP165 Prevotella pleuritidis
- SP166 Fusobacterium sp._HMT_204
- SP167 Lachnoanaerobaculum saburreum
- SP168 Prevotella sp._HMT_475
- SP169 Actinomyces sp._HMT_525
- SP17 Actinomyces oris
- SP170 Treponema sp._HMT_231
- SP171 Leptotrichia sp._HMT_212
- SP174 Capnocytophaga sp._HMT_336
- SP175 Catonella morbi

- SP271 Streptococcus periodonticum
- SP272 Pseudoeptotrichia goodfellowii
- SP279 Haemophilus sp._HMT_908
- SP280 Selenomonas sp._HMT_138
- SP281 Neisseria flavescens
- SP283 Absconditibacteria_(SR1)_[G-1] bacterium_HMT_875
- SP284 Rothia aeria
- SP288 Solobacterium moorei
- SP291 Schaalia sp._HMT_172
- SP294 Parvimonas micra
- SP299 Oribacterium sp._HMT_078
- SP3 Cardiobacterium valvarum
- SP30 Weeksellaceae_[G-1] sp._HMT_900
- SP303 Streptococcus peroris
- SP305 Neisseria sicca
- SP306 Leptotrichia sp._HMT_223
- SP31 Pseudoeptotrichia sp._HMT_219
- SP311 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352
- SP314 Leptotrichia sp._HMT_218
- SP32 Fusobacterium sp._HMT_370
- SP320 Streptococcus sp._HMT_074
- SP325 Porphyromonas sp._HMT_278
- SP326 Neisseria bacilliformis
- SP327 Peptostreptococcus stomatis
- SP328 Schaalia sp._HMT_877
- SP329 Actinomyces dentalis
- SP33 Porphyromonas endodontalis
- SP337 Campylobacter gracilis
- SP34 Porphyromonas pasteri
- SP344 Aggregatibacter sp._HMT_513
- SP345 Kingella oralis
- SP35 Actinomyces sp._HMT_171
- SP350 Actinomyces sp._HMT_448
- SP355 Neisseria mucosa
- SP357 Mogibacterium diversum
- SP36 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
- SP360 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- SP362 Cloacibacterium sp._HMT_206
- SP363 Ottowia sp._HMT_894
- SP364 Fusobacterium sp._HMT_248

- SP9 Capnocytophaga gingivalis
- SP90 Prevotella fusca
- SP91 Selenomonas noxia
- SP92 Actinomyces massiliensis
- SP93 Johnsonella ignava
- SP94 Rothia mucilaginosa
- SP95 Tannerella sp._HMT_286
- SP96 Peptococcus sp._HMT_167
- SP97 Alloprevotella sp._HMT_912
- SP99 Leptotrichia sp._HMT_225
- SPN10 Streptococcus gordonii_nov_97.018%
- SPN107 Pseudoeptotrichia goodfellowii_nov_91.416%
- SPN11 Pseudoeptotrichia goodfellowii_nov_97.204%
- SPN167 Corynebacterium durum_nov_97.699%
- SPN192 Actinomyces sp._HMT_175_nov_97.951%
- SPN2 Parvimonas micra_nov_97.053%
- SPN215 Lachnospiraceae_[G-2] bacterium_HMT_088_nov_93.096%
- SPN22 Fusobacterium sp._HMT_370_nov_97.845%
- SPN235 Veillonella sp._HMT_780_nov_97.628%
- SPN27 Selenomonas noxia_nov_97.030%
- SPN276 Leptotrichia sp._HMT_392_nov_97.624%
- SPN284 Streptococcus sobrinus_nov_94.291%
- SPN287 Moraxella oblonga_nov_93.776%
- SPN293 Peptidiphaga gingivicola_nov_96.920%
- SPN299 Streptobacillus notomitis_nov_93.856%
- SPN30 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347_nov_95.759%
- SPN300 Cardiobacterium hominis_nov_97.718%
- SPN306 Actinomyces sp._HMT_171_nov_96.421%
- SPN31 Ruminococcaceae_[G-1] bacterium_HMT_075_nov_91.453%
- SPN311 Alloprevotella sp._HMT_473_nov_97.336%
- SPN317 Cardiobacterium hominis_nov_97.510%
- SPN319 Kingella oralis_nov_97.741%
- SPN323 Porphyromonas sp._HMT_284_nov_97.546%
- SPN326 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348_nov_95.089%
- SPN333 Porphyromonas catoniae_nov_97.746%
- SPN334 Actinomyces sp._HMT_175_nov_97.751%
- SPN338 Streptococcus oralis_subsp._tigrinus_clade_070_nov_97.980%
- SPN340 Capnocytophaga gingivalis_nov_90.546%
- SPN343 Streptococcus sanguinis_nov_97.782%
- SPN345 Capnocytophaga gingivalis_nov_96.218%