

Species

SP131 Leptotrichia sp._HMT_223
SP132 Prevotella oulorum
SP133 Prevotella pallens
SP134 Anaeroglobus geminatus
SP135 Actinomyces sp._HMT_170
SP136 Capnocytophaga granulosa
SP137 Peptococcus sp._HMT_167
SP139 Fusobacterium canifilinum
SP14 Prevotella sp._HMT_317
SP140 Peptostreptococcus stomatis
SP142 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
SP143 Kingella denitrificans
SP144 Streptococcus mitis
SP145 Gemella sanguinis
SP146 Fretibacterium sp._HMT_360
SP15 Porphyromonas endodontalis
SP150 Actinomyces sp._HMT_175
SP151 Streptococcus sp._HMT_066
SP153 Corynebacterium durum
SP154 Selenomonas noxia
SP156 Rothia dentocariosa
SP158 Peptidiphaga gingivicola
SP159 Capnocytophaga sp._HMT_902
SP16 Veillonella dispar
SP160 Prevotella intermedia
SP161 Prevotella baroniae
SP162 Prevotella denticola
SP163 Haemophilus haemolyticus
SP164 Veillonellaceae_[G-1] bacterium_HMT_155
SP166 Bacteroidetes_[G-3] bacterium_HMT_280
SP167 Stomatobaculum sp._HMT_097
SP168 Streptococcus parasanguinis_clade_411
SP169 Neisseria flavescens
SP17 Capnocytophaga endodontalis
SP170 Selenomonas sp._HMT_137
SP171 Tannerella serpentinae
SP172 Tannerella sp._HMT_286
SP173 Haemophilus sp._HMT_036
SP174 Selenomonas sp._HMT_134
SP177 Porphyromonas pasteri
SP178 Bacteroidetes_[G-5] bacterium_HMT_511
SP179 Fusobacterium sp._HMT_370
SP18 Aggregatibacter sp._HMT_458
SP183 Stomatobaculum sp._HMT_373
SP184 Solobacterium moorei
SP186 Campylobacter showae
SP187 Prevotella pleuritidis
SP188 Treponema sp._HMT_257
SP19 Actinomyces dentalis
SP190 Peptostreptococcaceae_[XII][G-6] [Eubacterium]_minutum
SP191 Peptostreptococcaceae_[XII][G-2] bacterium_HMT_091
SP196 Fusobacterium sp._HMT_203
SP197 Mogibacterium diversum
SP198 Johnsonella ignava
SP199 Streptococcus sinensis
SP20 Veillonellaceae_[G-1] bacterium_HMT_132
SP200 Kingella oralis
SP201 Bilophila wadsworthia
SP202 Treponema sp._HMT_234
SP203 Schaalia sp._HMT_178
SP204 Streptococcus cristatus_clade_578
SP205 Neisseria perflava
SP206 Cryptobacterium curtum
SP207 Prevotella veroralis
SP208 Tannerella sp._HMT_808
SP210 Prevotella sp._HMT_292
SP212 Capnocytophaga leadbetteri
SP213 Desulfomicrobium orale
SP214 Fusobacterium periodonticum
SP215 Actinomyces sp._HMT_169
SP216 Neisseria oralis

SP200 Actinomyces phisconii
SP289 Selenomonas sp._HMT_138
SP29 Fusobacterium nucleatum
SP290 Selenomonas sp._HMT_892
SP291 Streptococcus chosunense
SP292 Selenomonas flueggei
SP293 Fretibacterium sp._HMT_361
SP294 Neisseria elongata
SP295 Prevotella sp._HMT_314
SP3 Actinomyces sp._HMT_897
SP30 Eikenella corrodens
SP302 Actinomyces oricola
SP303 Selenomonas sp._HMT_481
SP304 Actinomyces gerencseriae
SP305 Prevotella sp._HMT_526
SP306 Prevotella marshii
SP31 Prevotella melanigenica
SP310 Shuttleworthia satellites
SP311 Neisseria bacilliformis
SP312 Leptotrichia sp._HMT_225
SP314 Leptotrichia shahii
SP315 Streptococcus periodonticum
SP316 Lautropia dentalis
SP317 Treponema sp._HMT_238
SP319 Capnocytophaga sp._HMT_338
SP32 Fretibacterium fastidiosum
SP322 Prevotella sp._HMT_301
SP324 Streptococcus cristatus
SP325 Schaalia sp._HMT_877
SP329 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
SP33 Selenomonas sputigena
SP334 Peptococcus sp._HMT_168
SP336 Rothia mucilaginosa
SP338 Prevotella oralis
SP339 Bacteroidetes_[G-5] bacterium_HMT_505
SP34 Lachnanaerobaculum gingivialis
SP340 Streptococcus sobrinus
SP342 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
SP345 Cardiobacterium valvarum
SP347 Alloprevotella sp._HMT_473
SP349 Neisseria mucosa
SP35 Lachnanaerobaculum umeaense
SP351 Lancefieldella parvula
SP353 Streptococcus australis
SP355 Veillonellaceae_[G-1] bacterium_HMT_150
SP359 Actinomyces timonensis
SP36 Actinomyces oris
SP360 Streptococcus anginosus
SP366 Leptotrichia goodfellowii
SP37 Fusobacterium nucleatum_subsp._animalis
SP371 Treponema medium
SP375 Capnocytophaga sp._HMT_332
SP378 Bacteroidaceae_[G-1] bacterium_HMT_272
SP379 Streptococcus oralis_subsp._tigurinus_clade_070
SP38 Bergeyella sp._HMT_322
SP381 Treponema sp._HMT_247
SP382 Scardovia wiggsiae
SP384 Streptococcus mutans
SP385 Capnocytophaga sp._HMT_323
SP387 Veillonella rogosa
SP388 Gemella bergeri
SP389 Neisseria sicca
SP39 Streptococcus gordonii
SP390 Pseudoramibacter alactolyticus
SP393 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869
SP396 Treponema sp._HMT_263
SP397 Capnocytophaga haemolytica
SP4 Fusobacterium nucleatum_subsp._vincentii
SP40 Streptococcus sanguinis
SP402 Anaerolineae_[G-1] bacterium_HMT_439
SP404 Peptostreptococcaceae_[XIII][G-1] [Eubacterium]_infirum

SP61 Selenomonas sp._HMT_130
SP62 Treponema denticola
SP63 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
SP64 Leptotrichia buccalis
SP65 Prevotella sp._HMT_300
SP66 Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum
SP68 Orbibacterium sp._HMT_078
SP69 Leptotrichia hongkongensis
SP7 Capnocytophaga sp._HMT_864
SP70 Leptotrichia sp._HMT_392
SP71 Clostridiales_[F-1][G-1] bacterium_HMT_093
SP73 Leptotrichia wadei
SP74 Treponema socranskii
SP75 Lachnospiraceae_[G-8] bacterium_HMT_500
SP76 Prevotella loeschii
SP8 Veillonella parvula
SP80 Capnocytophaga sp._HMT_336
SP81 Megaspheera micronuciformis
SP82 Bacteroidales_[G-2] bacterium_HMT_274
SP83 Fusobacterium sp._HMT_204
SP84 Capnocytophaga sputigena
SP85 Selenomonas artemidis
SP86 Neisseria subflava
SP89 Leptotrichia sp._HMT_215
SP9 Bacteroidetes_[G-3] bacterium_HMT_365
SP93 Prevotella multiformis
SP94 Alloprevotella rava
SP95 Ruminococcaceae_[G-2] bacterium_HMT_085
SP97 Ralstonia sp._HMT_406
SP98 Veillonellaceae_[G-1] bacterium_HMT_129
SPN100 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
SPN103 Sphingomonas hankookensis_nov_97.685%
SPN112 Selenomonas noxia_nov_96.634%
SPN122 Propionivibrio dicarboxylicus_nov_93.802%
SPN132 Actinomyces israelii_nov_96.825%
SPN133 Actinomyces dentalis_nov_97.787%
SPN140 Corynebacterium matruchotii_nov_97.764%
SPN147 Caproiciproducens galactitolivorans_nov_90.042%
SPN154 Porphyromonas sp._HMT_284_nov_97.746%
SPN162 Capnocytophaga gingivialis_nov_97.689%
SPN17 Treponema sp._HMT_927_nov_91.411%
SPN171 Lacrimispora xylanolytica_nov_88.613%
SPN178 Campylobacter rectus_nov_97.826%
SPN186 Treponema sp._HMT_237_nov_97.566%
SPN196 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.918%
SPN205 Treponema sp._HMT_242_nov_97.769%
SPN211 Leptotrichia hofstadii_nov_96.970%
SPN214 Actinomyces israelii_nov_94.882%
SPN215 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.685%
SPN225 Selenomonas noxia_nov_97.835%
SPN235 Parvimonas sp._HMT_110_nov_97.895%
SPN245 Capnocytophaga gingivialis_nov_96.008%
SPN25 Sneathia sanguinigena_nov_90.792%
SPN253 Soehngenia saccharolytica_nov_84.855%
SPN262 Prevotella sp._HMT_292_nov_97.755%
SPN269 Actinomyces timonensis_nov_97.769%
SPN283 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.768%
SPN293 Aggregatibacter actinomycetemcomitans_nov_97.947%
SPN301 Corynebacterium matruchotii_nov_97.755%
SPN302 Selenomonas sp._HMT_137_nov_97.030%
SPN309 Selenomonas sp._HMT_919_nov_97.426%
SPN318 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.086%
SPN325 Kingella oralis_nov_97.536%
SPN336 Treponema maltophilum_nov_97.769%
SPN346 Wolinella succinogenes_nov_97.550%
SPN35 Actinomyces sp._HMT_169_nov_97.951%
SPN355 Prevotella sp._HMT_317_nov_97.536%
SPN364 Veillonellaceae_[G-1] bacterium_HMT_918_nov_96.154%
SPN370 Brachymonas denitrificans_nov_92.418%
SPN378 Actinomyces sp._HMT_169_nov_97.988%
SPN385 Streptococcus sanguinis_nov_97.783%