

Species

SP142 Mogibacterium timidum
SP143 Prevotella sp._HMT_475
SP144 Veillonella atypica
SP145 Selenomonas sp._HMT_137
SP146 Veillonella denticariosi
SP147 Prevotella denticola
SP148 Capnocytophaga sp._HMT_864
SP149 Treponema sp._HMT_268
SP15 Schaalii cardiffensis
SP150 Solobacterium moorei
SP153 Aggregatibacter sp._HMT_949
SP155 Treponema sp._HMT_231
SP156 Aggregatibacter sp._HMT_512
SP157 Prevotella veroralis
SP158 Fretibacterium sp._HMT_360
SP159 Porphyromonas catoniae
SP16 Dialister invisus
SP160 Stomatobaculum longum
SP162 Lachnoanaerobaculum umeaense
SP163 Lachnoanaerobaculum gingivalis
SP164 Prevotella pallens
SP165 Haemophilus haemolyticus
SP166 Fusobacterium sp._HMT_370
SP167 Kingella denitrificans
SP168 Prevotella intermedia
SP169 Streptococcus downii
SP17 Lachnospiraceae_[G-8] bacterium_HMT_500
SP170 Fusobacterium periodonticum
SP174 Schaalii sp._HMT_180
SP175 Veillonellaceae_[G-1] bacterium_HMT_132
SP176 Leptotrichia hongkongensis
SP177 Treponema sp._HMT_234
SP178 Selenomonas infelix
SP179 Treponema sp._HMT_257
SP18 Actinomyces oricola
SP180 Megaspheara sp._HMT_123
SP181 Prevotella salivae
SP183 Neisseria oralis
SP184 Stomatobaculum sp._HMT_373
SP185 Selenomonas sp._HMT_138
SP186 Selenomonas sp._HMT_920
SP187 Selenomonas flueggei
SP188 Bergelyella sp._HMT_900
SP19 Treponema sp._HMT_230
SP191 Alloprevotella tannerae
SP192 Aggregatibacter sp._HMT_513
SP193 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870
SP196 Treponema sp._HMT_238
SP197 Actinomyces dentalis
SP198 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345
SP2 Prevotella oris
SP20 Parvimonas sp._HMT_110
SP200 Lachnoanaerobaculum saburreum
SP201 Peptostreptococcus stomatis
SP202 Selenomonas noxia
SP203 Veillonellaceae_[G-1] bacterium_HMT_145
SP204 Neisseria flavescens
SP205 Streptococcus anginosus
SP206 Actinomyces israelii
SP207 Prevotella sp._HMT_315
SP208 Campylobacter concisus
SP209 Treponema sp._HMT_237
SP21 Prevotella sp._HMT_314
SP211 Neisseria subflava
SP214 Treponema sp._HMT_951
SP216 Leptotrichia hofstadii
SP219 Streptococcus sp._HMT_423
SP22 Streptococcus sp._HMT_066
SP220 Gemella haemolysans
SP222 Streptococcus mutans

SP313 Mogibacterium diversum
SP315 Haemophilus sp._HMT_036
SP316 Aggregatibacter sp._HMT_898
SP319 Leptotrichia sp._HMT_223
SP32 Lautropia mirabilis
SP321 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874
SP322 Hellaella seregens
SP323 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869
SP325 Streptococcus constellatus
SP327 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
SP328 Cardiobacterium valvarum
SP329 Peptostreptococcaceae_[XII][G-1] bacterium_HMT_383
SP33 Selenomonas sp._HMT_919
SP330 Selenomonas sp._HMT_149
SP332 Actinomyces massiliensis
SP333 Ruminococcaceae_[G-1] bacterium_HMT_075
SP335 Leptotrichia shahii
SP338 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872
SP339 Capnocytophaga sp._HMT_863
SP34 Streptococcus australis
SP340 Schaalii sp._HMT_172
SP342 Schaalii sp._HMT_877
SP345 Capnocytophaga sp._HMT_902
SP346 Peptoniphilaceae_[G-1] bacterium_HMT_113
SP347 Gemella sanguinis
SP351 Johnsonella ignava
SP352 Bacteroidetes_[G-5] bacterium_HMT_507
SP354 Cryptobacterium curtum
SP355 Capnocytophaga sp._HMT_336
SP357 Neisseria flava
SP358 Lancefieldella parvula
SP36 Fretibacterium fastidiosum
SP360 Actinomyces johnsonii
SP361 Catonella sp._HMT_451
SP363 Streptococcus vestibularis
SP367 Actinomyces timonensis
SP369 Porphyromonas sp._HMT_930
SP37 Leptotrichia sp._HMT_498
SP373 Treponema sp._HMT_490
SP375 Streptococcus sp._HMT_061
SP376 Prevotella oralis
SP377 Mycoplasma faucium
SP379 Oribacterium sp._HMT_078
SP38 Alloprevotella sp._HMT_473
SP380 Capnocytophaga haemolytica
SP381 Peptostreptococcaceae_[XII][G-2] bacterium_HMT_091
SP382 Rothia aera
SP385 Streptococcus sp._HMT_074
SP386 Bacteroides zoogloeiformans
SP387 Phocaeicola abscessus
SP388 Leptotrichia sp._HMT_909
SP390 Peptostreptococcaceae_[XII][G-7] bacterium_HMT_081
SP398 Peptostreptococcaceae_[XII][G-1] [Eubacterium]_infirmum
SP4 Capnocytophaga leadbetteri
SP40 Actinomyces oris
SP400 Bergelyella sp._HMT_206
SP401 Lancefieldella rimae
SP403 Porphyromonas gingivalis
SP404 Olsenella uli
SP405 Bacteroidaceae_[G-1] bacterium_HMT_272
SP407 Aggregatibacter paraprohilus
SP409 Tannerella sp._HMT_808
SP41 Ruminococcaceae_[G-2] bacterium_HMT_085
SP410 Mollicutes_[G-2] bacterium_HMT_906
SP411 Treponema sp._HMT_270
SP412 Neisseria sp._HMT_018
SP415 Erysipelotrichaceae_[G-1] bacterium_HMT_905
SP416 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
SP420 Treponema sp._HMT_254
SP421 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955

SP82 Lachnospiraceae_[G-3] bacterium_HMT_100
SP84 Prevotella pleuritidis
SP85 Veillonellaceae_[G-1] bacterium_HMT_155
SP86 Fusobacterium nucleatum_subsp._vincentii
SP87 Treponema maltophilum
SP88 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351
SP89 Bacteroidetes_[G-5] bacterium_HMT_505
SP9 Alloprevotella sp._HMT_913
SP90 Fusobacterium sp._HMT_203
SP91 Porphyromonas endodontalis
SP92 Prevotella marshii
SP93 Prevotella melaninogenica
SP94 Aggregatibacter sp._HMT_458
SP96 Prevotella baroniae
SP97 Veillonella rogosa
SP98 Streptococcus sp._HMT_064
SP99 Prevotella maculosa
SPN103 Selenomonas sp._HMT_137_nov_97.426%
SPN108 Sneathia sanguinegens_nov_90.792%
SPN113 Leptotrichia hofstadii_nov_96.970%
SPN114 Treponema sp._HMT_490_nov_96.545%
SPN125 Porphyromonas sp._HMT_284_nov_97.336%
SPN135 Treponema sp._HMT_258_nov_97.556%
SPN146 Corynebacterium durum_nov_97.899%
SPN15 Treponema maltophilum_nov_97.769%
SPN150 Lachnospiraceae_[G-8] bacterium_HMT_500_nov_97.895%
SPN155 Brooklawia cerclae_nov_97.291%
SPN165 Selenomonas sp._HMT_442_nov_97.595%
SPN176 Anaerocolumna cellulolytica_nov_88.333%
SPN186 Bacteroidetes_[G-5] bacterium_HMT_505_nov_96.250%
SPN198 Eikenella corrodens_nov_97.137%
SPN210 Wolinella succinogenes_nov_97.550%
SPN220 Lacrimispora xylanolytica_nov_88.430%
SPN221 Porphyromonas catoniae_nov_97.546%
SPN241 Treponema sp._HMT_508_nov_97.751%
SPN251 Anaerotaenia torta_nov_89.121%
SPN259 Aggregatibacter aphrophilus_nov_97.347%
SPN262 Haploplasma axanthum_nov_84.917%
SPN274 Treponema sp._HMT_252_nov_94.919%
SPN286 Treponema sp._HMT_239_nov_97.972%
SPN295 Prevotella sp._HMT_317_nov_97.746%
SPN304 Actinomyces sp._HMT_171_nov_96.813%
SPN305 Neisseria sp._HMT_018_nov_97.526%
SPN306 Treponema sp._HMT_927_nov_91.411%
SPN307 Porphyromonas catoniae_nov_97.951%
SPN308 Treponema sp._HMT_518_nov_97.546%
SPN309 Mollicutes_[G-1] bacterium_HMT_504_nov_97.917%
SPN310 Treponema denticola_nov_95.927%
SPN311 Bacteroidetes_[G-5] bacterium_HMT_505_nov_97.708%
SPN312 Schaalii sp._HMT_178_nov_97.737%
SPN313 Capnocytophaga gingivalis_nov_97.899%
SPN315 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349_nov_97.773%
SPN316 Selenomonas sp._HMT_138_nov_97.233%
SPN317 Veillonellaceae_[G-1] bacterium_HMT_145_nov_97.967%
SPN318 Actinomyces israelii_nov_97.605%
SPN319 Propionivibrio dicarboxylicus_nov_93.802%
SPN320 Selenomonas sp._HMT_149_nov_97.813%
SPN321 Selenomonas noxia_nov_96.634%
SPN322 Ottowia sp._HMT_894_nov_96.868%
SPN323 Selenomonas infelix_nov_97.822%
SPN326 Synergistes jonesii_nov_96.484%
SPN327 Slackia heliotrinireducens_nov_91.775%
SPN328 Prevotella sp._HMT_443_nov_97.955%
SPN329 Corynebacterium matruchotii_nov_97.325%
SPN33 Actinomyces israelii_nov_94.882%
SPN330 Selenomonas artemidis_nov_97.628%
SPN331 Hoppeia youngheungensis_nov_80.249%
SPN332 Corynebacterium matruchotii_nov_97.967%
SPN333 Streptococcus cristatus_nov_97.576%
SPN335 Actinomyces sp._HMT_169_nov_97.992%