Species

SP147 Neisseria subflava
SP148 Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
SP149 Leptotrichia wadei
SP15 Slackia exigua
SP150 Prevotella sp._HMT_315
SP151 Streptococcus australis
SP152 Leptotrichia sp._HMT_392
SP153 Selenomonas dianae
SP154 Prevotella nigrescens
SP155 Selenomonas noxia
SP156 Leptotrichia sp._HMT_463
SP157 Streptococcus mitis
SP159 Prevotella veroralis
SP16 Veillonellaceae_[G−1] bacterium_HMT_155
SP160 Actinomyces sp._HMT_897
SP161 Tannerella sp._HMT_286
SP162 Peptococcus sp._HMT_167
SP163 Selenomonas sp._HMT_920
SP165 Stomatobaculum sp._HMT_097
SP166 Prevotella salivae
SP167 Capnocytophaga gingivalis
SP168 Bacteroidales_[G−2] bacterium_HMT_274
SP169 Selenomonas sp._HMT_892
SP17 Prevotella jejuni
SP170 Eggerthia catenaformis
SP172 Riemerella sp._HMT_322
SP173 Rothia dentocariosa
SP174 Selenomonas sp._HMT_137
SP177 Solobacterium moorei
SP18 Schaalia meyeri
SP182 Abiotrophia defectiva
SP183 Fusobacterium sp._HMT_370
SP185 Prevotella melaninogenica
SP186 Lachnoanaerobaculum sp._HMT_496
SP187 Lachnoanaerobaculum sp._HMT_083
SP188 Treponema sp._HMT_237
SP189 Peptostreptococcus stomatis
SP19 Johnsonella sp._HMT_166
SP190 Selenomonas sp._HMT_136
SP191 Erysipelotrichaceae_[G−1] bacterium_HMT_905
SP311 Streptococcus oralis_subsp._tigurinus_clade_071
SP312 Streptococcus vestibularis
SP313 Aggregatibacter sp._HMT_898
SP315 Lancefieldella rimae
SP316 Actinomyces dentalis
SP318 Lachnoanaerobaculum gingivalis
SP32 Schaalia odontolytica
SP321 Peptoniphilaceae_[G−1] bacterium_HMT_113
SP322 Actinomyces timonensis
SP323 Schaalia cardiffensis
SP324 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_869
SP325 Treponema sp._HMT_234
SP326 Saccharibacteria_(TM7)_[G−4] bacterium_HMT_355
SP327 Tannerella sp._HMT_916
SP328 Ottowia sp._HMT_894
SP329 Pseudoramibacter alactolyticus
SP331 Peptoanaerobacter [Eubacterium] yurii
SP334 Prevotella marshii
SP336 Centipeda periodontii
SP338 Stomatobaculum sp._HMT_910
SP339 Leptotrichia sp._HMT_212
SP34 Haemophilus haemolyticus
SP342 Rothia aeria
SP343 Veillonellaceae_[G−1] bacterium_HMT_135
SP346 Tannerella serpentiformis
SP347 Actinomyces oricola
SP349 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
SP350 Neisseria flava
SP351 Peptoanaerobacter stomatis
SP353 Prevotella sp._HMT_475
SP355 Treponema sp._HMT_238
SP356 Treponema sp._HMT_253
SP358 Treponema sp._HMT_268
SP359 Capnocytophaga sp._HMT_323
SP36 Veillonellaceae_[G−1] bacterium_HMT_132
SP360 Capnocytophaga sp._HMT_412
SP361 Streptococcus chosunense
SP362 Peptostreptococcaceae_[G−6] [Eubacterium]_minutum
SP363 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
SP366 Granulicatella elegans
SP83 Lancefieldella parvula
SP84 Schaalia sp._HMT_180
SP85 Prevotella sp._HMT_300
SP86 Eikenella corrodens
SP88 Prevotella sp._HMT_317
SP89 Alloprevotella sp._HMT_473
SP9 Fusobacterium nucleatum_subsp._animalis
SP90 Gemella sanguinis
SP91 Actinomyces sp._HMT_448
SP92 Selenomonas sp._HMT_919
SP93 Lachnoanaerobaculum umeaense
SP94 Neisseria oralis
SP96 Treponema sp._HMT_230
SP97 Treponema denticola
SP98 Bacteroidetes_[G−3] bacterium_HMT_280
SP99 Campylobacter showae
SPN103 Actinomyces israelii_nov_96.838%
SPN109 Cardiobacterium hominis_nov_97.510%
SPN111 Prevotella veroralis_nov_97.959%
SPN121 Porphyromonas catoniae_nov_97.951%
SPN128 Actinomyces israelii_nov_94.882%
SPN136 Corynebacterium matruchotii_nov_97.769%
SPN145 Selenomonas sp._HMT_481_nov_97.822%
SPN152 Hoppeia youngheungensis_nov_80.457%
SPN161 Actinomyces sp._HMT_171_nov_96.813%
SPN162 Prevotella sp._HMT_300_nov_97.955%
SPN172 Actinomyces sp._HMT_897_nov_97.769%
SPN178 Desulfovibrio legallii_nov_94.343%
SPN184 Tannerella forsythia_nov_97.308%
SPN192 Sneathia sanguinegens_nov_90.578%
SPN199 Lacrimispora xylanolytica_nov_88.613%
SPN207 Corynebacterium durum_nov_97.484%
SPN210 Actinomyces sp._HMT_175_nov_97.751%
SPN213 Corynebacterium matruchotii_nov_97.955%
SPN215 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_348_nov_95.089%
SPN223 Kingella oralis_nov_97.536%
SPN232 Tepidisphaera mucosa_nov_92.461%
SPN233 Selenomonas infelix_nov_97.822%
SPN242 Actinomyces naeslundii_nov_97.963%
SPN251 Prevotella oris_nov_97.131%