

Species

- SP157 Streptococcus gordonii
- SP158 Porphyromonas catoniae
- SP159 Actinomyces sp._HMT_896
- SP16 Prevotella sp._HMT_317
- SP160 Selenomonas sp._HMT_149
- SP161 Johnsonella ignava
- SP162 Ruminococcaceae_[G-1] bacterium_HMT_075
- SP165 Leptotrichia shahii
- SP166 Fusobacterium sp._HMT_203
- SP167 Streptococcus oralis_subsp._oralis
- SP169 Rothia mucilaginosa
- SP171 Fretibacterium sp._HMT_362
- SP172 Prevotella sp._HMT_306
- SP173 Parvimonas sp._HMT_393
- SP175 Riemerella sp._HMT_322
- SP176 Tannerella forsythia
- SP178 Desulfovibrio sp._HMT_040
- SP18 Prevotella denticola
- SP180 Neisseria flavescens
- SP181 Bacteroidetes_[G-3] bacterium_HMT_365
- SP182 Actinomyces israelii
- SP183 Peptostreptococcaceae_[G-5] [Eubacterium]_saphenum
- SP184 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP186 Streptococcus constellatus
- SP187 Leptotrichia buccalis
- SP189 Tannerella sp._HMT_808
- SP19 Dialister invisus
- SP190 Peptostreptococcus stomatis
- SP191 Arachnia propionica
- SP193 Actinomyces oris
- SP194 Bacteroidetes_[G-5] bacterium_HMT_505
- SP195 Haemophilus haemolyticus
- SP197 Schaalia meyeri
- SP198 Peptostreptococcaceae_[G-9] [Eubacterium]_brachy
- SP2 Selenomonas sputigena
- SP20 Bacteroidetes_[G-3] bacterium_HMT_280
- SP200 Streptococcus sanguinis
- SP201 Veillonella rogosae
- SP202 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351
- SP203 Prevotella maculosa

- SP330 Prevotella pallens
- SP331 Neisseria elongata
- SP334 Treponema vincentii
- SP335 Treponema sp._HMT_230
- SP336 Peptostreptococcaceae_[G-6] [Eubacterium]_minutum
- SP337 Rothia dentocariosa
- SP338 Limosilactobacillus fermentum
- SP34 Veillonella dispar
- SP342 Schaalia sp._HMT_877
- SP345 Lactobacillus gasseri
- SP346 Mycoplasma salivarium
- SP347 Prevotella sp._HMT_304
- SP348 Capnocytophaga sp._HMT_380
- SP35 Fusobacterium nucleatum_subsp._animalis
- SP352 Leptotrichia sp._HMT_392
- SP354 Streptococcus sinensis
- SP356 Treponema sp._HMT_268
- SP357 Actinomyces timonensis
- SP358 Peptostreptococcaceae_[G-1] [Eubacterium]_infirmum
- SP359 Neisseria subflava
- SP36 Tannerella sp._HMT_286
- SP361 Streptococcus mutans
- SP37 Selenomonas sp._HMT_892
- SP370 Colibacter massiliensis
- SP372 Treponema sp._HMT_262
- SP374 Schaalia georgiae
- SP375 Treponema lecithinolyticum
- SP38 Porphyromonas endodontalis
- SP380 Bacteroidetes_[G-5] bacterium_HMT_511
- SP382 Aggregatibacter sp._HMT_512
- SP387 Leptotrichia sp._HMT_215
- SP388 Treponema sp._HMT_236
- SP389 Actinomyces sp._HMT_414
- SP39 Prevotella oris
- SP390 Aggregatibacter sp._HMT_513
- SP392 Prevotella buccae
- SP395 Streptococcus sp._HMT_056
- SP40 Schaalia sp._HMT_180
- SP400 Aggregatibacter aphrophilus
- SP403 Treponema denticola

- SP75 Slackia exigua
- SP76 Streptococcus oralis_subsp._dentisani_clade_058
- SP77 Prevotella histicola
- SP772 Butyrivibrio sp._HMT_080
- SP79 Leptotrichia trevisanii
- SP799 Alkalibacterium pelagium
- SP8 Fusobacterium periodonticum
- SP80 Leptotrichia hongkongensis
- SP81 Lachnospiraceae_[G-3] bacterium_HMT_100
- SP82 Prevotella veroralis
- SP84 Haemophilus sp._HMT_036
- SP85 Anaeroglobus geminatus
- SP86 Gemella haemolysans
- SP87 Streptococcus oralis_subsp._tigurinus_clade_071
- SP89 Prevotella oulorum
- SP892 Bacteroidetes_[G-4] bacterium_HMT_509
- SP9 Neisseria perflava
- SP90 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
- SP91 Fretibacterium sp._HMT_358
- SP92 Actinomyces sp._HMT_170
- SP923 Enterococcus italicus
- SP93 Abiotrophia defectiva
- SP94 Capnocytophaga sp._HMT_864
- SP95 Fretibacterium fastidiosum
- SP98 Selenomonas sp._HMT_137
- SP99 Selenomonas sp._HMT_138
- SPN1034 Lachnospiraceae_[G-2] bacterium_HMT_096_nov_87.629%
- SPN1074 Actinomyces israelii_nov_97.426%
- SPN1084 Actinomyces israelii_nov_94.912%
- SPN1129 Corynebacterium durum_nov_96.894%
- SPN1217 Parvimonas sp._HMT_110_nov_97.912%
- SPN1311 Actinomyces sp._HMT_897_nov_97.787%
- SPN1414 Actinomyces sp._HMT_175_nov_97.967%
- SPN160 Pedobacter sp._HMT_933_nov_79.637%
- SPN1607 Lachnoanaerobaculum sp._HMT_496_nov_88.125%
- SPN1706 Campylobacter rectus_nov_97.840%
- SPN1795 Treponema maltophilum_nov_97.787%
- SPN1886 Streptococcus parasanguinis_clade_411_nov_97.992%
- SPN1887 Tannerella forsythia_nov_97.325%
- SPN1989 Alkalihalobacillus clausii_nov_88.340%