

Species

- SP117 Corynebacterium durum
- SP118 Neisseria bacilliformis
- SP119 Capnocytophaga gingivalis
- SP12 Actinomyces sp._HMT_896
- SP120 Prevotella oulorum
- SP121 Haemophilus parainfluenzae
- SP123 Centipeda periodontii
- SP124 Selenomonas sp._HMT_126
- SP125 Eikenella corrodens
- SP129 Campylobacter concisus
- SP13 Limosilactobacillus mucosae
- SP130 Pseudoleptotrichia goodfellowii
- SP131 Actinomyces gerencseriae
- SP132 Actinomyces sp._HMT_448
- SP133 Prevotella sp._HMT_292
- SP137 Veillonellaceae_[G-1] bacterium_HMT_132
- SP138 Treponema denticola
- SP139 Treponema lecithinolyticum
- SP14 Lactobacillus delbrueckii
- SP140 Cryptobacterium curtum
- SP141 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
- SP142 Catonella morbi
- SP143 Gemella sanguinis
- SP144 Selenomonas diana
- SP145 Bilophila wadsworthia
- SP146 Streptococcus constellatus
- SP147 Actinomyces israelii
- SP149 Streptococcus chosunense
- SP15 Capnocytophaga sp._HMT_864
- SP151 Treponema sp._HMT_231
- SP152 Actinomyces sp._HMT_897
- SP153 Porphyromonas endodontalis
- SP154 Leptotrichia hongkongensis
- SP155 Scardovia inopinata
- SP159 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- SP16 Propionibacterium acidifaciens
- SP160 Treponema sp._HMT_237
- SP161 Veillonellaceae_[G-1] bacterium_HMT_150
- SP162 Lactocaseibacillus casei
- SP163 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347

- SP215 Streptococcus oralis_subsp._tigurinus_clade_070
- SP216 Tractidigestivibacter scatoligenes
- SP218 Streptococcus downei
- SP219 Lachnospiraceae_[G-3] bacterium_HMT_100
- SP22 Treponema vincentii
- SP220 Prevotella multiformis
- SP221 Treponema sp._HMT_247
- SP223 Fretibacterium sp._HMT_359
- SP226 Liquorilactobacillus nagelii
- SP228 Aggregatibacter sp._HMT_458
- SP229 Prevotella saccharolytica
- SP23 Shuttleworthia satelles
- SP231 Leptotrichia sp._HMT_223
- SP24 Lactocaseibacillus rhamnosus
- SP246 Leptotrichia sp._HMT_212
- SP25 Parascardovia denticolens
- SP257 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP258 Lactobacillus crispatus
- SP26 Actinomyces sp._HMT_414
- SP265 Gemella morbillorum
- SP266 Prevotella melaninogenica
- SP27 Peptidiphaga gingivicola
- SP272 Capnocytophaga ochracea
- SP274 Blautia luti
- SP275 Selenomonas sp._HMT_136
- SP279 Dialister pneumosintes
- SP28 Veillonellaceae_[G-1] bacterium_HMT_155
- SP280 Prevotella baroniae
- SP281 Johnsonella ignava
- SP284 Peptoanaerobacter stomatis
- SP285 Colibacter massiliensis
- SP29 Actinomyces sp._HMT_169
- SP3 Limosilactobacillus fermentum
- SP30 Fusobacterium nucleatum
- SP305 Schaalia odontolytica
- SP307 Faecalibacterium prausnitzii
- SP308 Schaalia georgiae
- SP31 Actinomyces dentalis
- SP32 Streptococcus sanguinis

- SP66 Prevotella denticola
- SP67 Selenomonas noxia
- SP68 Treponema sp._HMT_270
- SP7 Stomatobaculum longum
- SP70 Prevotella maculosa
- SP72 Bacteroidetes_[G-5] bacterium_HMT_511
- SP73 Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- SP75 Atopobium sp._HMT_416
- SP76 Lachnoanaerobaculum orale
- SP77 Alloprevotella tannerae
- SP79 Neisseria mucosa
- SP8 Dialister invisus
- SP80 Streptococcus cristatus
- SP81 Actinomyces sp._HMT_170
- SP83 Phocaeicola abscessus
- SP84 Prevotella nigrescens
- SP85 Lactobacillus gasseri
- SP86 Lachnospiraceae_[G-7] bacterium_HMT_163
- SP87 Lancefieldella parvula
- SP88 Peptoanaerobacter [Eubacterium] yurii
- SP9 Granulicatella adiacens
- SP92 Selenomonas sp._HMT_149
- SP93 Lachnoanaerobaculum saburreum
- SP94 Mitsukella sp._HMT_131
- SP95 Arachnia propionica
- SP96 Treponema maltophilum
- SP98 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- SP99 Olsenella sp._HMT_807
- SPN103 Actinomyces israelii_nov_96.501%
- SPN110 Actinomyces sp._HMT_171_nov_96.737%
- SPN116 Actinomyces naeslundii_nov_97.238%
- SPN128 Prevotella oulorum_nov_90.207%
- SPN13 Veillonellaceae_[G-1] bacterium_HMT_132_nov_97.076%
- SPN14 Prevotella denticola_nov_97.727%
- SPN140 Corynebacterium durum_nov_97.576%
- SPN153 Actinomyces viscosus_nov_96.395%
- SPN159 Fusobacterium nucleatum_nov_97.217%
- SPN165 Prevotella sp._HMT_300_nov_97.053%
- SPN176 Paraprevotella xylaniphila_nov_86.038%
- SPN188 Selenomonas sputigena_nov_90.538%