

Species

SP1 Porphyromonas gingivalis
 SP10 Prevotella oulorum
 SP100 Enterococcus gallinarum
 SP102 Fusobacterium nucleatum_subsp._vincentii
 SP103 Enterobacter cancerogenus
 SP11 Brochothrix thermosphacta
 SP110 Lactobacillus johnsonii
 SP111 Shigella flexneri
 SP112 Tannerella forsythia
 SP115 Anaerostipes caccae
 SP116 Bacillus licheniformis
 SP117 Peptostreptococcus stomatis
 SP118 Tannerella serpentinae
 SP12 Fusobacterium nucleatum_subsp._animalis
 SP120 Staphylococcus ureilyticus
 SP121 Psychrobacter cibarius
 SP126 Kocuria rhizophila
 SP127 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
 SP128 Streptococcus mitis
 SP129 Pseudomonas fragi
 SP13 Fusobacterium nucleatum
 SP132 Phocaeicola sartorii
 SP133 Corynebacterium pilbarensis
 SP134 Sphingomonas echinoides
 SP136 Finegoldia magna
 SP142 Citrobacter koseri
 SP143 Chryseobacterium gambrii
 SP148 Acinetobacter radioresistens
 SP15 Enterobacter hormaechei
 SP150 Treponema sp._HMT_257
 SP154 Corynebacterium macginleyi
 SP156 Stenotrophomonas maltophilia
 SP159 Akkermansia muciniphila
 SP165 Lactobacillus kisonensis
 SP167 Campylobacter concisus
 SP17 Klebsiella pneumoniae
 SP170 Sphingobium yanoikuyae
 SP173 Enterobacter asburiae
 SP179 Acinetobacter junii
 SP18 Ligilactobacillus murinus
 SP180 Aerococcus urinaequi
 SP182 Streptococcus thermophilus
 SP183 Corynebacterium tuberculoearicum
 SP187 Actinomyces sp._HMT_414
 SP19 Erysipelotrichaceae_[G-1] bacterium_MOT-189
 SP194 Cutibacterium namnetense
 SP195 Bacteroides caccae

SP198 Leptotrichia trevisanii
 SP2 Streptococcus sp._HMT_423
 SP20 Kosakonia sacchari
 SP200 Micrococcus aloeverae
 SP203 Bergeyella sp._HMT_900
 SP207 Streptococcus sanguinis
 SP208 Actinomyces sp._HMT_897
 SP209 Selenomonas sp._HMT_137
 SP211 Corynebacterium xerosis
 SP213 Leptotrichia buccalis
 SP214 Bacteroidaceae_[G-1] bacterium_HMT_272
 SP216 Actinomyces dentalis
 SP220 Bacteroidetes_[G-3] bacterium_HMT_365
 SP225 Corynebacterium mucifaciens
 SP23 Streptococcus australis
 SP231 Treponema amylovorum
 SP232 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
 SP234 Acidovorax defluvi
 SP243 Acinetobacter modestus
 SP244 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
 SP26 Corynebacterium amycolatum
 SP28 Enterobacter cloacae
 SP29 Prevotella saccharolytica
 SP30 Sphingomonas leidy
 SP32 Filifactor alocis
 SP33 Veillonella parvula
 SP36 Pseudomonas marginalis
 SP37 Enterococcus faecalis
 SP39 Faecalibaculum rodentium
 SP4 Bulleidia extructa
 SP41 Cutibacterium acnes
 SP42 Acinetobacter johnsonii
 SP46 Acinetobacter tjernbergiae
 SP47 Lawsonella clevelandensis
 SP51 Campylobacter gracilis
 SP54 Pseudomonas weihenstephanensis
 SP56 Psychrobacter urativorans
 SP58 Corynebacterium stationis
 SP60 Arthrobacter psychrolactophilus
 SP61 Enterobacter kobei
 SP64 Actinidia eriantha
 SP65 Limosilactobacillus reuteri
 SP66 Treponema sp._HMT_237
 SP68 Mammaliococcus lentus
 SP69 Neisseriaceae_[G-1] bacterium_MOT-031
 SP7 Enterobacter mori
 SP70 Streptococcus thoraltensis

SP71 Porphyromonas catoniae
 SP72 Bacteroidetes_[G-5] bacterium_HMT_505
 SP73 Streptococcus danieliae
 SP74 Treponema denticola
 SP76 Arthrobacter psychrochitinophilus
 SP78 Lautropia mirabilis
 SP79 Streptococcus oralis_subsp._tigurinus_clade_070
 SP8 Proteus mirabilis
 SP81 Pelomonas saccharophila
 SP86 Atopobium vaginae
 SP87 Peptoniphilus indolicus
 SP89 Selenomonas noxia
 SP90 Corynebacterium godavarianum
 SP93 Treponema sp._HMT_230
 SP95 Prevotella oris
 SPN10 Duncaniella freteri_nov_89.775%
 SPN108 Faecalibaculum rodentium_nov_96.715%
 SPN120 Duncaniella freteri_nov_89.113%
 SPN126 Sphaerobacter thermophilus_nov_82.265%
 SPN127 Duncaniella freteri_nov_89.718%
 SPN129 Pyrinomonas methylalipatogenes_nov_96.809%
 SPN130 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.387%
 SPN131 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
 SPN132 Duncaniella freteri_nov_90.612%
 SPN133 Duncaniella freteri_nov_88.800%
 SPN14 Peptoniphilus indolicus_nov_97.699%
 SPN15 Acinetobacter lwoffii_nov_96.289%
 SPN2 Enterobacter mori_nov_97.951%
 SPN20 Actinomyces israelii_nov_94.882%
 SPN23 Lawsonibacter asaccharolyticus_nov_91.116%
 SPN24 Parasutterella excrementihominis_nov_94.578%
 SPN28 Veillonellaceae_[G-1] bacterium_HMT_129_nov_96.356%
 SPN37 Duncaniella freteri_nov_91.853%
 SPN38 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.634%
 SPN4 Duncaniella freteri_nov_92.915%
 SPN52 Streptococcus azizii_nov_95.171%
 SPN6 Acinetobacter johnsonii_nov_97.531%
 SPN8 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.045%
 SPP10 Veillonella dispar_purvula
 SPP12 Staphylococcus argenteus_aureus_roterodami
 SPP13 Staphylococcus capitis_epidermidis
 SPP14 Pseudomonas antarctica_meridiana
 SPP15 Staphylococcus saprophyticus_xylosum
 SPP4 Fusobacterium nucleatum_nucleatum_subsp._animalis
 SPP6 Acinetobacter calcoaceticus_pittii
 SPP8 Psychrobacter arcticus_fjordensis
 SPP9 Psychrobacter fjordensis_okhotskensis