

Species

- SP114 Veillonella atypica
- SP115 Treponema sp._HMT_270
- SP116 Tannerella serpentiniformis
- SP117 Capnocytophaga sp._HMT_902
- SP118 Fusobacterium sp._HMT_370
- SP119 Olsenella sp._HMT_807
- SP12 Parvimonas micra
- SP120 Ruminococcaceae_[G-2] bacterium_HMT_085
- SP121 Streptococcus oralis_subsp._tigurinus_clade_071
- SP123 Fusobacterium sp._HMT_203
- SP124 Gemella haemolysans
- SP125 Selenomonas artemidis
- SP126 Mogibacterium timidum
- SP127 Aggregatibacter sp._HMT_512
- SP128 Peptostreptococcaceae_[XII][G-7] bacterium_HMT_081
- SP129 Streptococcus oralis_subsp._dantisani_clade_058
- SP13 Capnocytophaga leadbetteri
- SP130 Leptotrichia hongkongensis
- SP131 Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873
- SP132 Capnocytophaga haemolytica
- SP134 Fretibacterium fastidiosum
- SP135 Prevotella marshii
- SP136 Shuttlesworthia satelles
- SP137 Schaalia sp._HMT_180
- SP138 Oribacterium sp._HMT_078
- SP139 Capnocytophaga sp._HMT_336
- SP14 Capnocytophaga gingivalis
- SP140 Capnocytophaga endodontalis
- SP141 Leptotrichia sp._HMT_417
- SP142 Klebsiella michiganensis
- SP143 Selenomonas sp._HMT_919
- SP144 Peptostreptococcaceae_[XII][G-1] [Eubacterium]_infirmum
- SP146 Peptoanaerobacter [Eubacterium] yurii
- SP147 Leptotrichia sp._HMT_498
- SP148 Prevotella sp._HMT_317
- SP15 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345
- SP150 Aggregatibacter aphrophilus
- SP151 Prevotella oulorum
- SP152 Leptotrichia sp._HMT_221
- SP153 Fusobacterium nucleatum_subsp._vincentii
- SP154 Prevotella sp._HMT_292
- SP155 Bacteroidetes_[G-3] bacterium_HMT_280
- SP156 Leptotrichia trevisanii
- SP157 Treponema denticola
- SP159 Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum
- SP16 Enterobacter cloacae
- SP160 Prevotella nigrescens
- SP161 Bacteroidetes_[G-7] bacterium_HMT_911
- SP162 Veillonella rogosa
- SP163 Prevotella buccae
- SP164 Neisseria flavescens
- SP165 Capnocytophaga ochracea
- SP166 Lachnoanaerobaculum saburreum
- SP167 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955
- SP168 Veillonellaceae_[G-1] bacterium_HMT_155
- SP170 Prevotella salivae
- SP171 Capnocytophaga sp._HMT_332
- SP173 Peptostreptococcaceae_[XII][G-2] bacterium_HMT_091
- SP175 Saccharibacteria_(TM7)_[G-2] bacterium_HMT_350
- SP177 Actinomyces massiliensis
- SP178 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351
- SP179 Campylobacter gracilis
- SP18 Prevotella denticola
- SP181 Stomatobaculum longum
- SP182 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- SP184 Tannerella forsythia
- SP185 Lachnospiraceae_[G-7] bacterium_HMT_163
- SP187 Dialister invisus
- SP188 Leptotrichia wadei
- SP189 Selenomonas sp._HMT_481

- SP229 Porphyromonas sp._HMT_284
- SP23 Prevotella intermedia
- SP231 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_103
- SP232 Selenomonas sp._HMT_137
- SP234 Prevotella sp._HMT_472
- SP236 Actinomyces timonensis
- SP237 Kingella oralis
- SP239 Selenomonas sp._HMT_146
- SP24 Haemophilus paraaerolyticus
- SP240 Leptotrichia sp._HMT_219
- SP243 Prevotella oralis
- SP245 Neisseria oralis
- SP248 Streptococcus mutans
- SP249 Campylobacter concisus
- SP25 Prevotella sp._HMT_300
- SP250 Slackia exigua
- SP252 Aggregatibacter segnis
- SP253 Leptotrichia shahii
- SP254 Actinomyces sp._HMT_448
- SP256 Fretibacterium sp._HMT_359
- SP257 Capnocytophaga sp._HMT_338
- SP258 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP259 Lachnoanaerobaculum sp._HMT_083
- SP26 Catonella morbi
- SP260 Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- SP261 Oribacterium sinus
- SP262 Lachnoanaerobaculum gingivalis
- SP263 Lachnoanaerobaculum umeaense
- SP264 Selenomonas fueggei
- SP266 Capnocytophaga granulosa
- SP267 Megasphaera micronuciformis
- SP268 Megasphaera sp._HMT_123
- SP27 Solobacterium moorei
- SP271 Schaalia sp._HMT_877
- SP272 Treponema sp._HMT_231
- SP273 Bacteroidetes_[G-5] bacterium_HMT_511
- SP274 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869
- SP275 Lancefieldella parvula
- SP276 Treponema vincentii
- SP278 Haemophilus sputorum
- SP28 Peptococcus sp._HMT_167
- SP280 Bacteroides heparinolyticus
- SP282 Neisseria mucosa
- SP283 Bergeyella sp._HMT_907
- SP286 Ruminococcaceae_[G-1] bacterium_HMT_075
- SP29 Capnocytophaga sp._HMT_864
- SP290 Pseudoleptotrichia goodfellowii
- SP293 Campylobacter rectus
- SP294 Bulleidia extructa
- SP295 Anaeroglobus geminatus
- SP296 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
- SP298 Streptococcus sp._HMT_423
- SP3 Streptococcus gordonii
- SP30 Selenomonas infelix
- SP301 Peptoniphilus lacrimalis
- SP302 Parvimonas sp._HMT_393
- SP304 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874
- SP305 Treponema lecithinolyticum
- SP307 Olsenella uli
- SP308 Mycoplasma salivarium
- SP31 Streptococcus sanguinis
- SP310 Alloprevotella rava
- SP32 Porphyromonas endodontalis
- SP328 Treponema sp._HMT_247
- SP33 Streptococcus oralis_subsp._tigurinus_clade_070
- SP330 Prevotella micans
- SP331 Desulfobulbus sp._HMT_041
- SP333 Schaalia odontolytica
- SP337 Arachnia rubra
- SP34 Actinomyces sp._HMT_175

- SP46 Porphyromonas pasteri
- SP47 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- SP48 Granulicatella adiacens
- SP5 Fretibacterium sp._HMT_360
- SP50 Prevotella melaninogenica
- SP52 Porphyromonas sp._HMT_275
- SP53 Haemophilus parainfluenzae
- SP54 Streptococcus intermedius
- SP55 Campylobacter showae
- SP56 Bergeyella sp._HMT_900
- SP57 Selenomonas sputigena
- SP58 Treponema maltophilum
- SP59 Alloprevotella tannerae
- SP6 Propionibacterium acidifaciens
- SP60 Veillonella dispar
- SP61 Actinomyces naeslundii
- SP62 Fusobacterium nucleatum
- SP63 Selenomonas noxia
- SP64 Bergeyella sp._HMT_322
- SP65 Centipeda periodontii
- SP66 Mogibacterium diversum
- SP67 Leptotrichia hofstadii
- SP68 Selenomonas sp._HMT_892
- SP69 Filifactor alocis
- SP7 Capnocytophaga sputigena
- SP70 Klebsiella aerogenes
- SP71 Leptotrichia sp._HMT_212
- SP72 Aggregatibacter sp._HMT_458
- SP73 Actinomyces sp._HMT_169
- SP74 Erysipelotrichaceae_[G-1] bacterium_HMT_905
- SP75 Lautropia mirabilis
- SP76 Arachnia propionica
- SP77 Lachnospiraceae_[G-3] bacterium_HMT_100
- SP78 Rothia dentocariosa
- SP8 Ottowia sp._HMT_894
- SP80 Neisseria flava
- SP81 Streptococcus chosunense
- SP83 Tannerella sp._HMT_808
- SP84 Parvimonas sp._HMT_110
- SP87 Prevotella maculosa
- SP88 Lancefieldella rimae
- SP89 Streptococcus cristatus
- SP9 Eikenella corrodens
- SP90 Prevotella loescheii
- SP91 Streptococcus sp._HMT_064
- SP93 Actinomyces sp._HMT_171
- SP94 Prevotella saccharolytica
- SP95 Prevotella sp._HMT_315
- SP96 Prevotella histiicola
- SP97 Streptococcus mitis
- SP98 Peptidiphaga sp._HMT_183
- SP99 Prevotella pleuritidis
- SPN105 Bergeyella zoohelcum_nov_92.593%
- SPN120 Ottowia sp._HMT_894_nov_96.988%
- SPN146 Streptococcus sanguinis_nov_97.782%
- SPN18 Corynebacterium durum_nov_97.787%
- SPN213 Peptidiphaga gingivicola_nov_96.920%
- SPN235 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345_nov_97.934%
- SPN267 Selenomonas sp._HMT_137_nov_97.137%
- SPN28 Selenomonas sp._HMT_892_nov_97.041%
- SPN339 Capnocytophaga gingivalis_nov_90.756%
- SPN347 Bacteroides zoogloeiformans_nov_96.578%
- SPN359 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_92.871%
- SPN371 Prevotella sp._HMT_305_nov_93.661%
- SPN374 Cardiobacterium hominis_nov_88.719%
- SPN383 Centipeda periodontii_nov_97.436%
- SPN39 Prevotella nanceiensis_nov_91.353%
- SPN40 Desulfovibrio legalii_nov_94.343%
- SPN404 Corynebacterium matruchotii_nov_97.951%
- SPN416 Actinomyces sp._HMT_171_nov_96.929%