

es

- SP126 Selenomonas sp._HMT_134
- SP127 Peptococcus sp._HMT_167
- SP129 Leptotrichia sp._HMT_392
- SP13 Stomatobaculum sp._HMT_097
- SP131 Prevotella sp._HMT_472
- SP133 Aggregatibacter sp._HMT_512
- SP135 Rothia dentocariosa
- SP136 Prevotella sp._HMT_315
- SP138 Streptococcus koreensis
- SP139 Granulicatella elegans
- SP14 Fusobacterium nucleatum
- SP140 Gemella morbillorum
- SP141 Campylobacter concisus
- SP142 Saccharibacteria (TM7)_[G-3] bacterium_HMT_351
- SP143 Actinidia eriantha
- SP144 Gemella sanguinis
- SP145 Prevotella pallens
- SP146 Lancefieldella parvula
- SP147 Bacteroidetes_[G-5] bacterium_HMT_505
- SP148 Streptococcus sp._HMT_056
- SP149 Streptococcus gordonii
- SP151 Schaalia odontolyticus
- SP152 Streptococcus downii
- SP153 Leptotrichia sp._HMT_215
- SP154 Corynebacterium matruchotii
- SP157 Lachnoanaerobaculum orale
- SP158 Prevotella nigrescens
- SP159 Leptotrichia sp._HMT_225
- SP16 Porphyromonas endodontalis
- SP160 Leptotrichia hongkongensis
- SP161 Streptococcus oralis
- SP162 Streptococcus oralis_subsp._tigurinus_clade_071
- SP165 Mitsuoella sp._HMT_521
- SP166 Shuttleworthia satelles
- SP167 Haemophilus parahaemolyticus
- SP168 Capnocytophaga endodontalis
- SP169 Anaeroglobus geminatus
- SP17 Kingella oralis
- SP170 Abiotrophia defectiva
- SP171 Leptotrichia sp._HMT_218
- SP174 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872
- SP175 Treponema denticola
- SP176 Selenomonas flueggei
- SP177 Schaalia odontolytica
- SP178 Leptotrichia wadei
- SP179 Lautropia mirabilis
- SP18 Streptococcus australis
- SP180 Prevotella denticola
- SP183 Arachnia propionica
- SP184 Fusobacterium sp._HMT_204
- SP185 Schaalia lingnae_[Not_Validly_Published]
- SP186 Treponema sp._HMT_951
- SP187 Streptococcus oralis_subsp._dentisani_clade_058
- SP188 Eikenella corrodens
- SP189 Lachnoanaerobaculum saburream
- SP19 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP190 Leptotrichia sp._HMT_498
- SP191 Lachnospiraceae_[G-8] bacterium_HMT_500
- SP192 Prevotella histicola
- SP193 Bacteroidales_[G-2] bacterium_HMT_274
- SP194 Alloprevotella tannerae
- SP195 Streptococcus periodonticum
- SP196 Prevotella sp._HMT_314
- SP20 Fretibacterium sp._HMT_359
- SP200 Prevotella saccharolytica
- SP201 Filifactor alocis
- SP202 Parvimonas micra
- SP203 Alloprevotella sp._HMT_914

- SP267 Bergeyella sp._HMT_900
- SP268 Streptococcus parasanguinis_clade_411
- SP27 Selenomonas sp._HMT_136
- SP271 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
- SP272 Streptococcus anginosus
- SP274 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- SP276 Treponema socranskii
- SP278 Oribacterium sp._HMT_078
- SP28 Prevotella sp._HMT_475
- SP280 Mycoplasma faucium
- SP282 Prevotella micans
- SP283 Schaalia meyeri
- SP285 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- SP286 Alloprevotella sp._HMT_913
- SP29 Porphyromonas sp._HMT_275
- SP292 Actinomyces sp._HMT_175
- SP294 Bulleidia extructa
- SP295 Prevotella intermedia
- SP296 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352
- SP298 Prevotella sp._HMT_526
- SP299 Cardiobacterium valvarum
- SP3 Capnocytophaga leadbetteri
- SP30 Streptococcus infantis_clade_431
- SP300 Clostridiales_[F-1][G-1] bacterium_HMT_093
- SP303 Actinomyces sp._HMT_170
- SP304 Treponema sp._HMT_230
- SP305 Lancefieldella rimae
- SP306 Prevotella sp._HMT_304
- SP307 Streptococcus sp._HMT_061
- SP309 Selenomonas sp._HMT_892
- SP31 Prevotella maculosa
- SP312 Prevotella vespertina
- SP313 Capnocytophaga sp._HMT_336
- SP315 Neisseria sicca
- SP32 Megasphaera micronuciformis
- SP320 Scardovia wiggisiae
- SP324 Butyrivibrio sp._HMT_455
- SP327 Aggregatibacter sp._HMT_949
- SP328 Tannerella serpentiniformis
- SP329 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_infirmum
- SP33 Porphyromonas gingivalis
- SP330 Campylobacter gracilis
- SP332 Leptotrichia hofstadii
- SP333 Campylobacter sp._HMT_044
- SP334 Prevotella sp._HMT_305
- SP340 Bifidobacterium dentium
- SP341 Campylobacter showae
- SP343 Gardnerella vaginalis
- SP345 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- SP347 Peptostreptococcaceae_[XI][G-6] [Eubacterium]_nodatum
- SP349 Tannerella forsythia
- SP35 Tannerella sp._HMT_808
- SP353 Alloscardovia omnicolens
- SP354 Kingella denitrificans
- SP355 Peptostreptococcaceae_[XI][G-7] [Eubacterium]_yurii_subsp._yurii_&_margaretiae
- SP36 Capnocytophaga ochracea
- SP366 Peptostreptococcaceae_[XI][G-5] [Eubacterium]_saphenum
- SP37 Porphyromonas pasteri
- SP375 Mogibacterium timidum
- SP379 Saccharibacteria_(TM7)_[G-4] bacterium_HMT_355
- SP38 Actinomyces graevenitzi
- SP380 Prevotella aurantiaca
- SP383 Prevotella sp._HMT_306
- SP385 Fusobacterium sp._HMT_370
- SP388 Streptococcus oralis_subsp._tigurinus_clade_070
- SP39 Oribacterium sinus
- SP390 Ottowia sp._HMT_894
- SP392 Neisseria cinerea

- SP62 Selenomonas artemidis
- SP64 Lachnoanaerobaculum gingivalis
- SP66 Solobacterium moorei
- SP67 Veillonella sp._HMT_780
- SP68 Streptococcus sp._HMT_423
- SP69 Actinomyces sp._HMT_171
- SP7 Prevotella sp._HMT_317
- SP70 Streptococcus intermedius
- SP71 Selenomonas sp._HMT_138
- SP72 Peptostreptococcus stomatis
- SP73 Fusobacterium sp._HMT_203
- SP74 Actinomyces oris
- SP75 Haemophilus haemolyticus
- SP76 Parvimonas sp._HMT_110
- SP77 Stomatobaculum longum
- SP78 Oribacterium parvum
- SP79 Oribacterium asaccharolyticum
- SP8 Prevotella oris
- SP80 Neisseria perflava
- SP81 Treponema sp._HMT_257
- SP82 Neisseria flavescens
- SP83 Veillonella dispar
- SP86 Prevotella sp._HMT_300
- SP87 Porphyromonas sp._HMT_284
- SP88 Porphyromonas sp._HMT_930
- SP89 Fusobacterium nucleatum_subsp._animalis
- SP9 Selenomonas sp._HMT_149
- SP90 Actinomyces naeslundii
- SP91 Prevotella sp._HMT_309
- SP92 Leptotrichia sp._HMT_221
- SP93 Lachnoanaerobaculum umeaense
- SP94 Dialister invisus
- SP95 Leptotrichia buccalis
- SP97 Prevotella veroralis
- SP98 Gemella haemolysans
- SP99 Treponema sp._HMT_231
- SPN105 Leptotrichia sp._HMT_215_nov_97.826%
- SPN112 Prevotella sp._HMT_305_nov_94.070%
- SPN124 Capnocytophaga gingivalis_nov_90.546%
- SPN135 Peptostreptococcaceae_[XI][G-5] bacterium_HMT_493_nov_97.776%
- SPN145 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957_nov_97.776%
- SPN15 Bacteroidetes_[G-5] bacterium_HMT_505_nov_96.250%
- SPN154 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_93.776%
- SPN165 Treponema sp._HMT_246_nov_97.342%
- SPN177 Prevotella denticola_nov_97.959%
- SPN189 Capnocytophaga gingivalis_nov_96.429%
- SPN209 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870_nov_96.429%
- SPN23 Alloprevotella sp._HMT_308_nov_97.347%
- SPN252 Selenomonas sputigena_nov_96.800%
- SPN30 Schaalia sp._HMT_180_nov_97.925%
- SPN324 Alloprevotella sp._HMT_473_nov_97.336%
- SPN35 Capnocytophaga gingivalis_nov_97.899%
- SPN359 Campylobacter concisus_nov_97.180%
- SPN430 Porphyromonas sp._HMT_284_nov_97.541%
- SPN458 Prevotella histicola_nov_97.959%
- SPN46 Peptostreptococcus [Eubacterium]_yurii_nov_90.795%
- SPN470 Sneathia sanguinegens_nov_90.792%
- SPN480 Moraxella oblonga_nov_93.776%
- SPN499 Kingella denitrificans_nov_94.251%
- SPN5 Fusobacterium huasookii_nov_97.783%
- SPN511 Simonsiella muelleri_nov_96.728%
- SPN521 Mobiluncus curtisii_nov_96.694%
- SPN532 Prevotella veroralis_nov_97.546%
- SPN57 Prevotella jejunii_nov_97.755%
- SPN68 Leptotrichia hofstadii_nov_96.970%
- SPN77 Treponema sp._HMT_234_nov_86.922%
- SPN89 Streptococcus sanguinis_nov_97.782%
- SPN99 Porphyromonas catenula_nov_97.951%