

SP133 Prevotella pallens	SP29 Fusobacterium nucleatum	SP63 Saccharibacteria (TM7)_[G-1] bacterium_HMT_346
SP134 Anaeroglobus geminatus	SP290 Selenomonas sp._HMT_892	SP64 Leptotrichia buccalis
SP135 Actinomyces sp._HMT_170	SP291 Streptococcus chosunense	SP65 Prevotella sp._HMT_300
SP136 Capnocytophaga granulosa	SP292 Selenomonas flueggei	SP66 Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum
SP137 Peptococcus sp._HMT_167	SP293 Fretibacterium sp._HMT_361	SP68 Oribacterium sp._HMT_078
SP139 Fusobacterium canifelinum	SP294 Neisseria elongata	SP69 Leptotrichia hongkongensis
SP14 Prevotella sp._HMT_317	SP295 Prevotella sp._HMT_314	SP7 Capnocytophaga sp._HMT_864
SP140 Peptostreptococcus stomatis	SP3 Actinomyces sp._HMT_897	SP70 Leptotrichia sp._HMT_392
SP142 Saccharibacteria (TM7)_[G-5] bacterium_HMT_356	SP30 Eikenella corrodens	SP71 Clostridiales_[F-1][G-1] bacterium_HMT_093
SP143 Kingella denitrificans	SP302 Actinomyces oricola	SP73 Leptotrichia wadei
SP144 Streptococcus mitis	SP303 Selenomonas sp._HMT_481	SP74 Treponema socranskii
SP145 Gemella sanguinis	SP304 Actinomyces gerencseriae	SP75 Lachnospiraceae_[G-8] bacterium_HMT_500
SP146 Fretibacterium sp._HMT_360	SP305 Prevotella sp._HMT_526	SP76 Prevotella loeschii
SP15 Porphyromonas endodontalis	SP306 Prevotella marshii	SP8 Veillonella parvula
SP150 Actinomyces sp._HMT_175	SP31 Prevotella melaninogenica	SP80 Capnocytophaga sp._HMT_336
SP151 Streptococcus sp._HMT_066	SP310 Shuttleworthia satellites	SP81 Megaspheara micronuciformis
SP153 Corynebacterium durum	SP311 Neisseria bacilliformis	SP82 Bacteroidales_[G-2] bacterium_HMT_274
SP154 Selenomonas noxia	SP312 Leptotrichia sp._HMT_225	SP83 Fusobacterium sp._HMT_204
SP156 Rothia dentocariosa	SP314 Leptotrichia shahii	SP84 Capnocytophaga sputigena
SP158 Peptidiphaga gingivicola	SP315 Streptococcus periodonticum	SP85 Selenomonas artemidis
SP159 Capnocytophaga sp._HMT_902	SP316 Lautropia dentalis	SP86 Neisseria subflava
SP16 Veillonella dispar	SP317 Treponema sp._HMT_238	SP89 Leptotrichia sp._HMT_215
SP160 Prevotella intermedia	SP319 Capnocytophaga sp._HMT_338	SP9 Bacteroidetes_[G-3] bacterium_HMT_365
SP161 Prevotella baroniae	SP32 Fretibacterium fastidiosum	SP93 Prevotella multiformis
SP162 Prevotella denticola	SP322 Prevotella sp._HMT_301	SP94 Alloprevotella rava
SP163 Haemophilus haemolyticus	SP324 Streptococcus cristatus	SP95 Ruminococcaceae_[G-2] bacterium_HMT_085
SP164 Veillonellaceae_[G-1] bacterium_HMT_155	SP325 Schaalii sp._HMT_877	SP97 Ralstonia sp._HMT_406
SP166 Bacteroidetes_[G-3] bacterium_HMT_280	SP329 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369	SP98 Veillonellaceae_[G-1] bacterium_HMT_129
SP167 Stomatobaculum sp._HMT_097	SP33 Selenomonas sputigena	SPN100 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
SP168 Streptococcus parasanguinis_clade_411	SP334 Peptococcus sp._HMT_168	SPN103 Sphingomonas hankookensis_nov_97.685%
SP169 Neisseria flavescens	SP336 Rothia mucilaginosa	SPN112 Selenomonas noxia_nov_96.634%
SP17 Capnocytophaga endodontalis	SP338 Prevotella oralis	SPN122 Propionivibrio dicarboxylicus_nov_93.802%
SP170 Selenomonas sp._HMT_137	SP339 Bacteroidetes_[G-5] bacterium_HMT_505	SPN132 Actinomyces israelii_nov_96.825%
SP171 Tannerella serpentiniformis	SP34 Lachnanaerobaculum gingivalis	SPN133 Actinomyces dentalis_nov_97.787%
SP172 Tannerella sp._HMT_286	SP340 Streptococcus sobrinus	SPN140 Corynebacterium matruchotii_nov_97.764%
SP173 Haemophilus sp._HMT_036	SP342 Saccharibacteria(TM7)_[G-1] bacterium_HMT_349	SPN147 Caproiciproducens galactitolivorans_nov_90.042%
SP174 Selenomonas sp._HMT_134	SP345 Cardiobacterium valvarum	SPN154 Porphyromonas sp._HMT_284_nov_97.746%
SP177 Porphyromonas pasteri	SP347 Alloprevotella sp._HMT_473	SPN162 Capnocytophaga gingivalis_nov_97.689%
SP178 Bacteroidetes_[G-5] bacterium_HMT_511	SP349 Neisseria mucosa	SPN17 Treponema sp._HMT_927_nov_91.411%
SP179 Fusobacterium sp._HMT_370	SP35 Lachnanaerobaculum umeaense	SPN171 Lacrimispora xylanolytica_nov_88.613%
SP18 Aggregatibacter sp._HMT_458	SP351 Lancefieldella parvula	SPN178 Campylobacter rectus_nov_97.826%
SP183 Stomatobaculum sp._HMT_373	SP353 Streptococcus australis	SPN186 Treponema sp._HMT_237_nov_97.566%
SP184 Solobacterium moorei	SP355 Veillonellaceae_[G-1] bacterium_HMT_150	SPN196 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.918%
SP186 Campylobacter showae	SP359 Actinomyces timonensis	SPN205 Treponema sp._HMT_242_nov_97.769%
SP187 Prevotella pleuritidis	SP36 Actinomyces oris	SPN211 Leptotrichia hofstadii_nov_96.970%
SP188 Treponema sp._HMT_257	SP360 Streptococcus anginosus	SPN214 Actinomyces israelii_nov_94.882%
SP19 Actinomyces dentalis	SP366 Leptotrichia goodfellowii	SPN215 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.685%
SP190 Peptostreptococcaceae_[XII][G-6] [Eubacterium]_minutum	SP37 Fusobacterium nucleatum_subsp._animalis	SPN225 Selenomonas noxia_nov_97.835%
SP191 Peptostreptococcaceae_[XII][G-2] bacterium_HMT_091	SP371 Treponema medium	SPN235 Parvimonas sp._HMT_110_nov_97.895%
SP195 Fusobacterium sp._HMT_203	SP375 Capnocytophaga sp._HMT_332	SPN245 Capnocytophaga gingivalis_nov_96.008%
SP197 Mogibacterium diversum	SP378 Bacteroidaceae_[G-1] bacterium_HMT_272	SPN25 Sneathia sanguinegens_nov_90.792%
SP198 Johnsonella ignava	SP379 Streptococcus oralis_subsp._tigurinus_clade_070	SPN253 Soehngenella saccharolytica_nov_84.855%
SP199 Streptococcus sinensis	SP38 Bergeyella sp._HMT_322	SPN262 Prevotella sp._HMT_292_nov_97.755%
SP20 Veillonellaceae_[G-1] bacterium_HMT_132	SP381 Treponema sp._HMT_247	SPN269 Actinomyces timonensis_nov_97.769%
SP200 Kingella oralis	SP382 Scardovia wiggisiae	SPN283 Saccharibacteria (TM7)_[G-1] bacterium_HMT_346_nov_97.769%
SP201 Bilophila wadsworthia	SP384 Streptococcus mutans	SPN293 Aggregatibacter actinomycetemcomitans_nov_97.947%
SP202 Treponema sp._HMT_234	SP385 Capnocytophaga sp._HMT_323	SPN301 Corynebacterium matruchotii_nov_97.755%
SP203 Schaalii sp._HMT_178	SP387 Veillonella rogosae	SPN302 Selenomonas sp._HMT_137_nov_97.030%
SP204 Streptococcus cristatus_clade_578	SP388 Gemella bergeri	SPN309 Selenomonas sp._HMT_919_nov_97.426%
SP205 Neisseria perflava	SP389 Neisseria sicca	SPN318 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.086%
SP206 Cryptobacterium curtum	SP39 Streptococcus gordonii	SPN325 Kingella oralis_nov_97.536%
SP207 Prevotella veroralis	SP390 Pseudoramibacter alactolyticus	SPN336 Treponema maltophilum_nov_97.769%
SP208 Tannerella sp._HMT_808	SP393 Saccharibacteria (TM7)_[G-1] bacterium_HMT_869	SPN346 Wolinella succinogenes_nov_97.550%
SP210 Prevotella sp._HMT_292	SP396 Treponema sp._HMT_263	SPN35 Actinomyces sp._HMT_169_nov_97.951%
SP212 Capnocytophaga leadbetteri	SP397 Capnocytophaga haemolytica	SPN355 Prevotella sp._HMT_317_nov_97.536%
SP213 Desulfomicrobium orale	SP4 Fusobacterium nucleatum_subsp._vincentii	SPN364 Veillonellaceae_[G-1] bacterium_HMT_918_nov_96.154%
SP214 Fusobacterium periodonticum	SP40 Streptococcus sanguinis	SPN370 Brachymonas denitrificans_nov_92.418%