

SP133 Prevotella pallens	SP291 Streptococcus chosunense	SP63 Saccharibacteria (TM7) [G-1] bacterium_HMT_346
SP134 Anaeroglobus geminatus	SP292 Selenomonas flueggei	SP64 Leptotrichia buccalis
SP135 Actinomyces sp. HMT_170	SP293 Fretibacterium sp. HMT_361	SP65 Prevotella sp. HMT_300
SP136 Capnocytophaga granulosa	SP294 Neisseria elongata	SP66 Peptostreptococcaceae [XII][G-5] [Eubacterium]_saphenum
SP137 Peptococcus sp. HMT_167	SP295 Prevotella sp. HMT_314	SP68 Oribacterium sp. HMT_078
SP139 Fusobacterium canifelinum	SP3 Actinomyces sp. HMT_897	SP69 Leptotrichia hongkongensis
SP14 Prevotella sp. HMT_317	SP30 Eikenella corrodens	SP7 Capnocytophaga sp. HMT_864
SP140 Peptostreptococcus stomatis	SP302 Actinomyces oricola	SP70 Leptotrichia sp. HMT_392
SP142 Saccharibacteria (TM7) [G-5] bacterium_HMT_356	SP303 Selenomonas sp. HMT_481	SP71 Clostridiales [F-1][G-1] bacterium_HMT_093
SP143 Kingella denitrificans	SP304 Actinomyces gerencseriae	SP73 Leptotrichia wadei
SP144 Streptococcus mitis	SP305 Prevotella sp. HMT_526	SP74 Treponema socranskii
SP145 Gemella sanguinis	SP306 Prevotella marshii	SP75 Lachnospiraceae [G-8] bacterium_HMT_500
SP146 Fretibacterium sp. HMT_360	SP31 Prevotella melaninogenica	SP76 Prevotella loeschii
SP15 Porphyromonas endodontalis	SP310 Shuttleworthia satellites	SP8 Veillonella parvula
SP150 Actinomyces sp. HMT_175	SP311 Neisseria bacilliformis	SP80 Capnocytophaga sp. HMT_336
SP151 Streptococcus sp. HMT_066	SP312 Leptotrichia sp. HMT_225	SP81 Megaspheara micronuciformis
SP153 Corynebacterium durum	SP314 Leptotrichia shahii	SP82 Bacteroidales [G-2] bacterium_HMT_274
SP154 Selenomonas noxia	SP315 Streptococcus periodonticum	SP83 Fusobacterium sp. HMT_204
SP156 Rothia dentocariosa	SP316 Lautropia dentalis	SP84 Capnocytophaga sputigena
SP158 Peptidiphaga gingivicola	SP317 Treponema sp. HMT_238	SP85 Selenomonas artemidis
SP159 Capnocytophaga sp. HMT_902	SP319 Capnocytophaga sp. HMT_338	SP86 Neisseria subflava
SP16 Veillonella dispar	SP32 Fretibacterium fastidiosum	SP89 Leptotrichia sp. HMT_215
SP160 Prevotella intermedia	SP322 Prevotella sp. HMT_301	SP9 Bacteroidetes [G-3] bacterium_HMT_365
SP161 Prevotella baroniae	SP324 Streptococcus cristatus	SP93 Prevotella multiformis
SP162 Prevotella denticola	SP325 Schaalii sp. HMT_877	SP94 Alloprevotella rava
SP163 Haemophilus haemolyticus	SP329 Peptostreptococcaceae [XI][G-4] bacterium_HMT_369	SP95 Ruminococcaceae [G-2] bacterium_HMT_085
SP164 Veillonellaceae [G-1] bacterium_HMT_155	SP33 Selenomonas sputigena	SP97 Ralstonia sp. HMT_406
SP166 Bacteroidetes [G-3] bacterium_HMT_280	SP334 Peptococcus sp. HMT_168	SP98 Veillonellaceae [G-1] bacterium_HMT_129
SP167 Stomatobaculum sp. HMT_097	SP336 Rothia mucilaginosa	SPN100 Muribaculaceae [G-1] bacterium_MOT-129_nov_88.105%
SP168 Streptococcus parasanguinis_clade_411	SP338 Prevotella oralis	SPN103 Sphingomonas hankookensis_nov_97.685%
SP169 Neisseria flavescens	SP339 Bacteroidetes [G-5] bacterium_HMT_505	SPN112 Selenomonas noxia_nov_96.634%
SP17 Capnocytophaga endodontalis	SP34 Lachnoanaerobaculum gingivalis	SPN122 Propionivibrio dicarboxylicus_nov_93.802%
SP170 Selenomonas sp. HMT_137	SP340 Streptococcus sobrinus	SPN132 Actinomyces israelii_nov_96.825%
SP171 Tannerella serpentiniformis	SP342 Saccharibacteria (TM7) [G-1] bacterium_HMT_349	SPN133 Actinomyces dentalis_nov_97.787%
SP172 Tannerella sp. HMT_286	SP345 Cardiobacterium valvarum	SPN140 Corynebacterium matruchotii_nov_97.764%
SP173 Haemophilus sp. HMT_036	SP347 Alloprevotella sp. HMT_473	SPN147 Caproiciproducens galactitolivorans_nov_90.042%
SP174 Selenomonas sp. HMT_134	SP349 Neisseria mucosa	SPN154 Porphyromonas sp. HMT_284_nov_97.746%
SP177 Porphyromonas pasteri	SP35 Lachnoanaerobaculum umeaense	SPN162 Capnocytophaga gingivalis_nov_97.689%
SP178 Bacteroidetes [G-5] bacterium_HMT_511	SP351 Lancefieldella parvula	SPN17 Treponema sp. HMT_927_nov_91.411%
SP179 Fusobacterium sp. HMT_370	SP353 Streptococcus australis	SPN171 Lacrimispora xylanolytica_nov_88.613%
SP18 Aggregatibacter sp. HMT_458	SP355 Veillonellaceae [G-1] bacterium_HMT_150	SPN178 Campylobacter rectus_nov_97.826%
SP183 Stomatobaculum sp. HMT_373	SP359 Actinomyces timonensis	SPN186 Treponema sp. HMT_237_nov_97.566%
SP184 Solobacterium moorei	SP36 Actinomyces oris	SPN196 Lachnospiraceae [G-14] bacterium_MOT-182_nov_92.918%
SP186 Campylobacter showae	SP360 Streptococcus anginosus	SPN205 Treponema sp. HMT_242_nov_97.769%
SP187 Prevotella pleuritidis	SP366 Leptotrichia goodfellowii	SPN211 Leptotrichia hofstadii_nov_96.970%
SP188 Treponema sp. HMT_257	SP37 Fusobacterium nucleatum_subsp._animalis	SPN214 Actinomyces israelii_nov_94.882%
SP19 Actinomyces dentalis	SP371 Treponema medium	SPN215 Muribaculaceae [G-1] bacterium_MOT-129_nov_85.685%
SP190 Peptostreptococcaceae [XI][G-6] [Eubacterium]_minutum	SP375 Capnocytophaga sp. HMT_332	SPN225 Selenomonas noxia_nov_97.835%
SP191 Peptostreptococcaceae [XI][G-2] bacterium_HMT_091	SP378 Bacteroidaceae [G-1] bacterium_HMT_272	SPN235 Parvimonas sp. HMT_110_nov_97.895%
SP195 Fusobacterium sp. HMT_203	SP379 Streptococcus oralis_subsp._tigurinus_clade_070	SPN245 Capnocytophaga gingivalis_nov_96.008%
SP197 Mogibacterium diversum	SP38 Bergelyella sp. HMT_322	SPN25 Sneathia sanguinegens_nov_90.792%
SP198 Johnsonella ignava	SP381 Treponema sp. HMT_247	SPN253 Soehngenella saccharolytica_nov_84.855%
SP199 Streptococcus sinensis	SP382 Scardovia wiggisiae	SPN262 Prevotella sp. HMT_292_nov_97.755%
SP20 Veillonellaceae [G-1] bacterium_HMT_132	SP384 Streptococcus mutans	SPN269 Actinomyces timonensis_nov_97.769%
SP200 Kingella oralis	SP385 Capnocytophaga sp. HMT_323	SPN283 Saccharibacteria (TM7) [G-1] bacterium_HMT_346_nov_97.76%
SP201 Bilophila wadsworthia	SP387 Veillonella rogosa	SPN293 Aggregatibacter actinomycetemcomitans_nov_97.947%
SP202 Treponema sp. HMT_234	SP388 Gemella bergeri	SPN301 Corynebacterium matruchotii_nov_97.755%
SP203 Schaalii sp. HMT_178	SP389 Neisseria sicca	SPN302 Selenomonas sp. HMT_137_nov_97.030%
SP204 Streptococcus cristatus_clade_578	SP39 Streptococcus gordonii	SPN309 Selenomonas sp. HMT_919_nov_97.426%
SP205 Neisseria perflava	SP390 Pseudoramibacter alactolyticus	SPN318 Clostridiales [F-1][G-1] bacterium_HMT_093_nov_84.086%
SP206 Cryptobacterium curtum	SP393 Saccharibacteria (TM7) [G-1] bacterium_HMT_869	SPN325 Kingella oralis_nov_97.536%
SP207 Prevotella veroralis	SP396 Treponema sp. HMT_263	SPN336 Treponema maltophilum_nov_97.769%
SP208 Tannerella sp. HMT_808	SP397 Capnocytophaga haemolytica	SPN346 Wolinella succinogenes_nov_97.550%
SP210 Prevotella sp. HMT_292	SP4 Fusobacterium nucleatum_subsp._vincentii	SPN35 Actinomyces sp. HMT_169_nov_97.951%
SP212 Capnocytophaga leadbetteri	SP40 Streptococcus sanguinis	SPN355 Prevotella sp. HMT_317_nov_97.536%
SP213 Desulfomicrobium orale		SPN364 Veillonellaceae [G-1] bacterium_HMT_918_nov_96.154%
SP214 Fusobacterium periodonticum		SPN370 Brachymonas denitrificans_nov_92.418%