

Species

SP139 Leptotrichia sp._HMT_225	SP286 Bacteroidetes_[G-5] bacterium_HMT_511	SP79 Prevotella sp._HMT_472
SP14 Alloprevotella sp._HMT_912	SP287 Streptococcus cristatus	SP8 Weeksellaceae_[G-1] sp._HMT_900
SP140 Prevotella loescheii	SP29 Streptococcus oralis	SP80 Leptotrichia sp._HMT_417
SP142 Streptococcus infantis_clade_431	SP290 Fretibacterium sp._HMT_360	SP81 Capnocytophaga sp._HMT_380
SP144 Gemella sanguinis	SP291 Actinomyces sp._HMT_414	SP82 Streptococcus sanguinis
SP145 Fusobacterium hwasookii	SP296 Neisseria perflava	SP83 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
SP146 Campylobacter showae	SP297 Neisseria subflava	SP84 Streptococcus parasanguinis_clade_411
SP148 Actinomyces sp._HMT_525	SP298 Neisseria flavescens	SP85 Pseudoleptotrichia sp._HMT_221
SP149 Capnocytophaga sp._HMT_412	SP299 Capnocytophaga sp._HMT_878	SP86 Peptostreptococcaceae_[G-7] [Eubacterium]_yurii_subsp._yurii_&_margaretiae
SP150 Solobacterium moorei	SP3 Haemophilus parainfluenzae	SP89 Johnsonella sp._HMT_166
SP152 Prevotella melaninogenica	SP300 Leptotrichia sp._HMT_498	SP90 Gemella haemolysans
SP153 Rothia aeria	SP302 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874	SP93 Veillonella atypica
SP154 Schaalia meyeri	SP304 Kingella oralis	SP94 Porphyromonas sp._HMT_278
SP156 Selenomonas artemidis	SP306 Actinomyces sp._HMT_896	SP96 Corynebacterium durum
SP157 Capnocytophaga endodontalis	SP308 Aggregatibacter segnis	SP97 Fusobacterium nucleatum_subsp._vincentii
SP158 Neisseria cinerea	SP309 Neisseria mucosa	SP98 Fusobacterium nucleatum_subsp._animalis
SP159 Capnocytophaga sp._HMT_864	SP31 Actinomyces dentalis	SP99 Leptotrichia sp._HMT_392
SP16 Lachnospiraceae_[G-3] bacterium_HMT_100	SP313 Peptococcus sp._HMT_168	SPN105 Corynebacterium matruchotii_nov_97.546%
SP160 Leptotrichia sp._HMT_463	SP318 Actinomyces israelii	SPN11 Actinomyces israelii_nov_94.882%
SP161 Streptococcus sp._HMT_066	SP319 Tannerella sp._HMT_286	SPN123 Corynebacterium matruchotii_nov_97.160%
SP162 Alloprevotella sp._HMT_913	SP32 Cardiobacterium hominis	SPN166 Neisseria sp._HMT_018_nov_97.942%
SP163 Tannerella serpentinaformis	SP320 Streptococcus sp._HMT_064	SPN176 Actinomyces sp._HMT_175_nov_97.951%
SP164 Streptococcus gordonii	SP322 Aggregatibacter sp._HMT_513	SPN184 Prevotella veroralis_nov_97.342%
SP165 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346	SP326 Haemophilus sputorum	SPN194 Haemophilus haemolyticus_nov_97.947%
SP166 Streptococcus downii	SP327 Actinomyces johnsonii	SPN202 Porphyromonas catoniae_nov_97.947%
SP167 Leptotrichia trevisanii	SP330 Aggregatibacter sp._HMT_949	SPN209 Corynebacterium durum_nov_97.484%
SP168 Prevotella micans	SP332 Peptoanaerobacter [Eubacterium] yurii	SPN210 Lactimyspora xylanolytica_nov_88.613%
SP169 Leptotrichia hofstadii	SP333 Schaalia sp._HMT_877	SPN213 Capnocytophaga gingivalis_nov_96.218%
SP17 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870	SP34 Granulicatella elegans	SPN221 Porphyromonas catoniae_nov_97.746%
SP170 Peptidiphaga sp._HMT_183	SP341 Selenomonas flueggei	SPN229 Actinomyces sp._HMT_169_nov_97.551%
SP172 Capnocytophaga sp._HMT_332	SP342 Fretibacterium sp._HMT_359	SPN237 Corynebacterium matruchotii_nov_97.967%
SP174 Actinomyces timonensis	SP344 Leptotrichia sp._HMT_223	SPN244 Actinomyces israelii_nov_96.838%
SP175 Streptococcus oralis_subsp._tigurinus_clade_070	SP345 Peptidiphaga gingivicola	SPN249 Actinomyces oris_nov_97.796%
SP176 Cardiobacterium valvarum	SP346 Tannerella sp._HMT_808	SPN258 Selenomonas noxia_nov_96.634%
SP178 Veillonellaceae_[G-1] bacterium_HMT_150	SP347 Lancefieldella parvula	SPN267 Alloprevotella sp._HMT_473_nov_97.546%
SP179 Ottowia sp._HMT_894	SP35 Leptotrichia buccalis	SPN275 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348_nov_95.089%
SP180 Prevotella pallens	SP352 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345	SPN283 Selenomonas sp._HMT_137_nov_97.030%
SP182 Prevotella multiformis	SP359 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957	SPN286 Aggregatibacter aphrophilus_nov_97.347%
SP183 Capnocytophaga sp._HMT_335	SP36 Porphyromonas sp._HMT_930	SPN287 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347_nov_95.759%
SP184 Capnocytophaga sp._HMT_880	SP360 Porphyromonas sp._HMT_931	SPN288 Porphyromonas sp._HMT_932