

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 Schaaliala 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 serpentiformis	● 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 Lacrimispora xylanolytica_nov_88.613%
● SP169 Leptotrichia hofstadii	● SP333 Schaaliala 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%