

Species

SP130 Actinomyces sp._HMT_169	SP207 Prevotella piculidis	SP65 Lachnospiraceae_[G-2] bacterium_HMT_056
SP137 Prevotella vespertina	SP268 Actinomyces sp._HMT_171	SP86 Fusobacterium periodonticum
SP138 Gemella morbillorum	SP273 Fusobacterium nucleatum_subsp._vincentii	SP87 Abiotrophia defectiva
SP139 Prevotella veroralis	SP275 Neisseria flava	SP88 Aggregatibacter sp._HMT_458
SP14 Kingella denitrificans	SP277 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_871	SP89 Leptotrichia sp._HMT_218
SP140 Streptococcus sp._HMT_064	SP279 Capnocytophaga sp._HMT_332	SP9 Oribacterium parvum
SP142 Treponema sp._HMT_231	SP28 Stomatobaculum longum	SP90 Filifactor alocis
SP143 Mogibacterium diversum	SP282 Peptostreptococcus stomatis	SP91 Streptococcus sp._HMT_057
SP144 Prevotella denticola	SP283 Fusobacterium hwasookii	SP92 Streptococcus oralis_subsp._tigurinus_clade_071
SP145 Fusobacterium sp._HMT_203	SP285 Aggregatibacter sp._HMT_512	SP93 Streptococcus anginosus
SP148 Fusobacterium sp._HMT_204	SP287 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345	SP94 Alloprevotella sp._HMT_914
SP149 Schaalia odontolytica	SP288 Campylobacter showae	SP95 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
SP15 Corynebacterium matruchotii	SP29 Streptococcus downii	SP96 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872
SP150 Capnocytophaga sp._HMT_901	SP293 Treponema denticola	SP97 Alloprevotella rava
SP151 Shuttleworthia satelles	SP296 Veillonella sp._HMT_917	SP98 Prevotella oulorum
SP153 Veillonella rogosae	SP3 Porphyromonas pasteri	SP99 Selenomonas sp._HMT_136
SP154 Prevotella sp._HMT_314	SP30 Leptotrichia buccalis	SPN100 Corynebacterium matruchotii_nov_97.546%
SP155 Prevotella micans	SP302 Aggregatibacter sp._HMT_949	SPN101 Pseudoleptotrichia goodfellowii_nov_91.416%
SP156 Peptostreptococcaceae_[G-9] [Eubacterium]_brachy	SP305 Haemophilus sputorum	SPN102 Capnocytophaga gingivalis_nov_97.899%
SP157 Pseudoleptotrichia goodfellowii	SP31 Veillonella parvula	SPN103 Prevotella jejuni_nov_97.755%
SP158 Stomatobaculum sp._HMT_097	SP311 Ottowia sp._HMT_894	SPN104 Porphyromonas sp._HMT_284_nov_97.746%
SP159 Treponema sp._HMT_257	SP312 Prevotella intermedia	SPN106 Streptobacillus notomitis_nov_93.856%
SP16 Capnocytophaga leadbetteri	SP317 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346	SPN107 Corynebacterium durum_nov_97.280%
SP160 Prevotella sp._HMT_317	SP318 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351	SPN108 Granulicatella elegans_nov_97.988%
SP161 Peptococcus sp._HMT_168	SP32 Veillonella dispar	SPN109 Porphyromonas catoniae_nov_97.746%
SP162 Neisseria elongata	SP320 Cloacibacterium sp._HMT_206	SPN110 Peptoanaerobacter [Eubacterium] yurii_nov_90.795%
SP163 Streptococcus sanguinis	SP33 Streptococcus oralis	SPN111 Streptococcus sanguinis_nov_97.782%
SP165 Butyrivibrio sp._HMT_455	SP332 Rothia mucilaginosa	SPN112 Fusobacterium hwasookii_nov_97.783%
SP166 Haemophilus parahaemolyticus	SP336 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356	SPN146 Schaalia sp._HMT_180_nov_97.930%
SP167 Peptococcus sp._HMT_167	SP34 Streptococcus infantis_clade_431	SPN148 Streptococcus parasanguinis_clade_411_nov_97.773%
SP169 Neisseria cinerea	SP340 Streptococcus sp._HMT_074	SPN15 Prevotella histicola_nov_97.347%
SP17 Granulicatella adiacens	SP343 Campylobacter gracilis	SPN156 Alloprevotella sp._HMT_308_nov_97.347%
SP173 Leptotrichia sp._HMT_392	SP344 Fusobacterium sp._HMT_248	SPN169 Ruminococcaceae_[G-1] bacterium_HMT_075_nov_91.453%
SP175 Neisseria sicca	SP347 Pseudomonas aeruginosa	SPN184 Alloprevotella sp._HMT_914_nov_97.551%
SP176 Streptococcus intermedius	SP35 Pseudostreptobacillus hongkongensis	SPN188 Veillonella sp._HMT_780_nov_97.628%
SP179 Capnocytophaga sp._HMT_878	SP36 Streptococcus gordonii	SPN210 Selenomonas sputigena_nov_97.000%
SP180 Gemella haemolysans	SP365 Haemophilus influenzae	SPN220 Pseudoleptotrichia goodfellowii_nov_92.060%
SP181 Porphyromonas gingivalis	SP37 Streptococcus oralis_subsp._dentisani_clade_058	SPN231 Moraxella boeveyi_nov_92.807%
SP182 Lachnoanaerobaculum saburreum	SP370 Neisseria sp._HMT_018	SPN240 Lachnospiraceae_[G-2] bacterium_HMT_088_nov_93.096%
SP183 Prevotella intermedia	SP38 Veillonella sp._HMT_780	SPN25 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_872_nov_99.994%