

Species

SP152 Bacteroidetes_[G-3] bacterium_HMT_365	SP322 Prevotella micans	SP89 Alloprevotella sp._HMT_914
SP153 Leptotrichia hofstadii	SP323 Capnocytophaga sp._HMT_332	SP90 Actinomyces sp._HMT_175
SP154 Solobacterium moorei	SP326 Anaerolineae_[G-1] bacterium_HMT_439	SP91 Lachnoanaerobaculum umeaense
SP155 Veillonella atypica	SP327 Selenomonas artemidis	SP92 Veillonella dispar
SP156 Abiotrophia defectiva	SP33 Mycoplasma salivarium	SP93 Veillonella parvula
SP157 Parvimonas sp._HMT_110	SP330 Treponema sp._HMT_262	SP94 Peptostreptococcaceae_[G-7] bacterium_HMT_081
SP158 Streptococcus oralis_subsp._dentisani_clade_058	SP332 Capnocytophaga sp._HMT_380	SP95 Mitsukella sp._HMT_521
SP16 Treponema denticola	SP333 Porphyromonas sp._HMT_284	SP96 Lachnospiraceae_[G-8] bacterium_HMT_500
SP160 Streptococcus vestibularis	SP337 Peptostreptococcaceae_[G-4] bacterium_HMT_103	SP97 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
SP161 Streptococcus salivarius	SP339 Brevundimonas vesicularis	SP98 Porphyromonas endodontalis
SP162 Streptococcus sp._HMT_064	SP34 Streptococcus parasanguinis_clade_411	SP99 Porphyromonas sp._HMT_275
SP164 Capnocytophaga sp._HMT_336	SP340 Microbacterium hydrothermale	SPN1037 Porphyromonas cationiae_nov_97.959%
SP165 Streptococcus cristatus_clade_578	SP341 Sphingobium yanoikuyae	SPN154 Moraxella nonliquefaciens_nov_92.149%
SP167 Campylobacter gracilis	SP347 Haemophilus parahaemolyticus	SPN191 Micrococcus luteus_nov_97.210%
SP168 Brevundimonas subvibrioides	SP348 Megaspheera micronuciformis	SPN202 Capnocytophaga gingivalis_nov_97.699%
SP169 Alloprevotella rava	SP349 Peptostreptococcaceae_[G-7] [Eubacterium]_yurii_subsp._yurii_&_margaretiae	SPN213 Lachnospiraceae_[G-2] bacterium_HMT_096_nov_87.603%
SP17 Prevotella saccharolytica	SP35 Leptotrichia sp._HMT_392	SPN224 Actinomyces sp._HMT_175_nov_97.760%
SP170 Veillonella rogosa	SP353 Bacteroidetes_[G-5] bacterium_HMT_507	SPN247 Actinomyces sp._HMT_175_nov_97.963%
SP171 Fusobacterium sp._HMT_204	SP354 Prevotella buccae	SPN260 Leptotrichia sp._HMT_215_nov_97.840%
SP173 Eikenella corrodens	SP355 Treponema sp._HMT_247	SPN272 Streptococcus sanguinis_nov_97.791%
SP174 Neisseria perflava	SP356 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957	SPN273 Prevotella oris_nov_97.149%
SP175 Fretibacterium sp._HMT_362	SP358 Tannerella sp._HMT_916	SPN285 Streptococcus oralis_subsp._oralis_nov_97.988%
SP176 Selenomonas noxia	SP359 Phocaeicola abscessus	SPN293 Corynebacterium durum_nov_97.495%
SP177 Arachnia propionica	SP36 Bacteroidetes_[G-5] bacterium_HMT_505	SPN298 Prevotella intermedia_nov_97.172%
SP178 Odoribacter bacterium_HMT_516	SP362 Lachnoanaerobaculum orale	SPN310 Porphyromonas sp._HMT_278_nov_97.764%
SP179 Alloprevotella tannerae	SP363 Rothia dentocariosa	SPN322 Capnocytophaga gingivalis_nov_96.234%
SP18 Prevotella denticola	SP364 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349	SPN334 Prevotella sp._HMT_300_nov_97.764%
SP180 Granulicatella adiacens	SP37 Bifidobacterium dentium	SPN346 Bacteroidetes_[G-5] bacterium_HMT_511_nov_96.894%
SP181 Schaalia odontolytica	SP373 Asticcacaulis excentricus	SPN357 Prevotella melaninogenica_nov_96.559%
SP182 Selenomonas sp._HMT_134	SP38 Treponema sp._HMT_230	SPN370 Capnocytophaga gingivalis_nov_90.586%
SP183 Gemella haemolysans	SP382 Colibacter massiliensis	SPN382 Corynebacterium matruchotii_nov_97.967%
SP184 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356	SP383 Streptococcus sp._HMT_074	SPN393 Porphyromonas sp._HMT_284_nov_97.760%
SP185 Streptococcus sp._HMT_057	SP385 Cardiobacterium valvarum	SPN394 Streptococcus intermedius_nov_97.633%
SP186 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348	SP387 Aerococcus viridans	SPN406 Porphyromonas uenonis_nov_94.479%
SP187 Peptostreptococcaceae_[G-1] [Eubacterium]_infirmum	SP388 Kingella denitrificans	SPN418 Parvimonas sp._HMT_110_nov_97.699%
SP188 Prevotella marshii	SP39 Treponema socranskii	SPN427 Rhodobacter capsulatus_nov_96.305%
SP19 Tannerella forsythia	SP392 Acinetobacter junii	SPN437 Prevotella pleuritidis_nov_97.358%
SP190 Granulicatella elegans	SP393 Johnsonella sp._HMT_166	SPN449 Alloprevotella tannerae_nov_94.737%
SP191 Haemophilus sp._HMT_036	SP394 Streptococcus intermedius	SPN461 Capnocytophaga granulosa_nov_84.948%
SP192 Streptococcus	SP395 Capnocytophaga	SPN462 Capnocytophaga