

## Species

SP1 Streptococcus salivarius	SP17 Dialister invisus	SP67 Streptococcus cristatus
SP10 Leptotrichia wadei	SP171 Streptococcus oralis_subsp._dentisani_clade_398	SP68 Capnocytophaga gingivalis
SP100 Prevotella buccae	SP172 Capnocytophaga sp._HMT_336	SP69 Haemophilus parainfluenzae
SP101 Leptotrichia hongkongensis	SP180 Prevotella salivae	SP7 Leptotrichia hofstadii
SP103 Aggregatibacter sp._HMT_458	SP182 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_103	SP70 Granulicatella adiacens
SP104 Capnocytophaga leadbetteri	SP184 Shuttleworthia satelles	SP71 Corynebacterium matruchotii
SP105 Eikenella corrodens	SP19 Peptostreptococcus stomatis	SP72 Campylobacter showae
SP107 Lachnoanaerobaculum gingivalis	SP192 Schaalia sp._HMT_172	SP73 Filifactor alocis
SP108 Prevotella sp._HMT_317	SP194 Leptotrichia sp._HMT_498	SP75 Capnocytophaga granulosa
SP109 Anaeroglobus geminatus	SP2 Solobacterium moorei	SP76 Leptotrichia sp._HMT_212
SP11 Peptidiphaga sp._HMT_183	SP20 Prevotella intermedia	SP77 Actinomyces sp._HMT_171
SP110 Selenomonas sp._HMT_137	SP21 Veillonella atypica	SP78 Klebsiella michiganensis
SP111 Prevotella sp._HMT_314	SP22 Leptotrichia sp._HMT_223	SP79 Peptoanaerobacter [Eubacterium] yurii
SP112 Tannerella sp._HMT_286	SP23 Porphyromonas sp._HMT_275	SP8 Fusobacterium nucleatum
SP113 Leptotrichia sp._HMT_392	SP24 Prevotella saccharolytica	SP80 Treponema socranskii
SP114 Alloprevotella sp._HMT_473	SP25 Veillonella dispar	SP81 Parvimonas micra
SP115 Actinomyces naeslundii	SP26 Stomatobaculum longum	SP82 Prevotella maculosa
SP116 Lancefieldella rimae	SP27 Neisseria elongata	SP83 Alloprevotella tannerae
SP117 Cardiobacterium hominis	SP28 Fusobacterium canifelinum	SP84 Streptococcus anginosus
SP119 Bergeyella sp._HMT_900	SP29 Neisseria sicca	SP85 Prevotella denticola
SP12 Sphingomonas oligophenolica	SP3 Streptococcus sp._HMT_064	SP86 Phocaeicola abscessus
SP120 Arachnia propionica	SP30 Streptococcus mitis	SP87 Prevotella nanceiensis
SP122 Lachnospiraceae_[G-3] bacterium_HMT_100	SP31 Neisseria flavescens	SP88 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
SP123 Tannerella serpentiniformis	SP32 Selenomonas noxia	SP89 Cardiobacterium valvarum
SP124 Streptococcus chosunense	SP33 Porphyromonas endodontalis	SP9 Streptococcus oralis
SP125 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345	SP34 Porphyromonas catoniae	SP91 Actinomyces sp._HMT_169
SP127 Prevotella oulorum	SP35 Ottowia sp._HMT_894	SP92 Saccharibacteria_(TM7)_[G-4] bacterium_HMT_355
SP128 Capnocytophaga sp._HMT_380	SP36 Neisseria subflava	SP93 Tannerella forsythia
SP129 Streptococcus oralis_subsp._dentisani_clade_058	SP37 Prevotella oris	SP94 Peptostreptococcaceae_[XII][G-1] [Eubacterium]_infirmum
SP13 Streptococcus intermedius	SP38 Leptotrichia sp._HMT_215	SP95 Ruminococcaceae_[G-2] bacterium_HMT_085
SP130 Lautropia mirabilis	SP39 Selenomonas sputigena	SP96 Capnocytophaga sputigena
SP131 Treponema maltophilum	SP4 Bacteroidales_[G-2] bacterium_HMT_274	SP97 Bacteroidetes_[G-3] bacterium_HMT_280
SP132 Arachnia rubra	SP40 Prevotella sp._HMT_300	SP98 Lachnoanaerobaculum orale
SP133 Selenomonas flueggei	SP41 Streptococcus sanguinis	SP99 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
SP134 Fusobacterium nucleatum_subsp._vincentii	SP42 Fusobacterium periodonticum	SPN1 Porphyromonas sp._HMT_284_nov_97.436%
SP136 Streptococcus sp._HMT_056	SP43 Streptococcus vestibularis	SPN11 Bergeyella zoohelcum_nov_92.673%
SP137 Capnocytophaga sp._HMT_864	SP44 Capnocytophaga endodontalis	SPN15 Bacteroides zoogloformans_nov_96.578%
SP138 Catonella morbi	SP45 Capnocytophaga ochracea	SPN2 Aggregatibacter sp._HMT_458_nov_97.826%
SP139 Rothia aeria	SP46 Stenotrophomonas maltophilia	SPN22 Actinomyces sp._HMT_448_nov_96.718%
SP14 Gemella morbillorum	SP47 Streptococcus oralis_subsp._tigurinus_clade_070	SPN27 Prevotella sp._HMT_305_nov_93.688%
SP140 Lachnospiraceae_[G-8] bacterium_HMT_500	SP48 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348	SPN3 Sphingomonas jeddahensis_nov_97.447%
SP141 Campylobacter gracilis	SP49 Leptotrichia sp._HMT_225	SPN33 Selenomonas sp._HMT_137_nov_96.947%
SP143 Rothia dentocariosa	SP50 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874	SPN39 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_92.871%
SP144 Campylobacter sp._HMT_044	SP51 Neisseria sp._HMT_020	SPN4 Neisseria elongata_nov_96.374%
SP146 Scardovia wiggsiae	SP52 Aggregatibacter segnis	SPN44 Selenomonas sp._HMT_137_nov_97.126%
SP147 Prevotella marshii	SP53 Prevotella sp._HMT_820	SPN5 Neisseria sp._HMT_020_nov_97.619%
SP15 Bergeyella sp._HMT_322	SP54 Campylobacter rectus	SPN51 Leptotrichia sp._HMT_215_nov_97.292%
SP150 Saccharibacteria_(TM7)_[G-2] bacterium_HMT_350	SP55 Porphyromonas pasteri	SPN55 Actinomyces naeslundii_nov_97.593%
SP153 Porphyromonas gingivalis	SP56 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356	SPN6 Prevotella melaninogenica_nov_97.170%
SP154 Prevotella loeschei	SP57 Selenomonas infelix	SPN63 Streptococcus sanguinis_nov_97.538%
SP155 Fretibacterium sp._HMT_360	SP58 Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873	SPN66 Selenomonas noxia_nov_96.337%
SP157 Prevotella veroralis	SP59 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351	SPN71 Selenomonas sp._HMT_137_nov_97.901%
SP159 Prevotella sp._HMT_292	SP6 Streptococcus gordonii	SPN72 Corynebacterium durum_nov_97.384%
SP16 Veillonella parvula	SP60 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369	SPN73 Abiotrophia defectiva_nov_96.481%
SP160 Streptococcus cristatus_clade_578	SP61 Kingella oralis	SPN74 Selenomonas sputigena_nov_97.026%
SP161 Actinomyces sp._HMT_175	SP62 Campylobacter concisus	SPN75 Selenomonas sp._HMT_479_nov_96.756%
SP163 Veillonella rogosae	SP63 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869	SPP2 Lachnoanaerobaculum gingivalis_umeaense
SP164 Schaalia sp._HMT_180	SP64 Leptotrichia buccalis	SPP3 Selenomonas flueggei_noxia
SP165 Selenomonas sp._HMT_138	SP65 Aggregatibacter sp._HMT_512	SPP6 Veillonella dispar_parvula
SP168 Lachnoanaerobaculum saburreum	SP66 Prevotella melaninogenica	