

Species

● SP152 Streptococcus australis	● SP322 Saccharibacteria (TM7) [G-1] bacterium_HMT_957	● SPN128 Selenomonas sp._HMT_149_nov_96.947%
● SP153 Leptotrichia trevisanii	● SP320 Fusobacterium sp._HMT_204	● SPN129 Corynebacterium sp._Oral_Taxon_A46_nov_95.898%
● SP154 Leptotrichia shahii	● SP321 Prevotella pleuritidis	● SPN13 Selenomonas artemidis_nov_97.606%
● SP155 Peptoanaerobacter yurii	● SP324 Treponema sp._HMT_951	● SPN130 Actinomyces viscosus_nov_96.234%
● SP156 Lactobacillus salivarius	● SP325 Treponema sp._HMT_270	● SPN131 Corynebacterium durum_nov_96.032%
● SP157 Treponema sp._HMT_268	● SP326 Desulfobulbus sp._HMT_041	● SPN14 Actinomyces sp._Oral_Taxon_178_nov_97.514%
● SP158 Treponema lecithinolyticum	● SP327 Lachnospiraceae [G-7] bacterium_HMT_086	● SPN15 Saccharibacteria (TM7) [G-1] bacterium_HMT_957_nov_97.639%
● SP159 Streptococcus sinensis	● SP328 Veillonellaceae [G-1] bacterium_HMT_155	● SPN16 Streptococcus gordonii_nov_97.782%
● SP16 Actinomyces viscosus	● SP329 Prevotella sp._HMT_942	● SPN17 Treponema denticola_nov_97.170%
● SP160 Peptidiphaga sp._HMT_183	● SP33 Rothia aeria	● SPN18 Corynebacterium sp._Oral_Taxon_A46_nov_97.065%
● SP161 Porphyromonas gingivalis	● SP330 Citrobacter koseri	● SPN19 Selenomonas sp._Oral_Taxon_137_nov_97.059%
● SP162 Actinomyces sp._HMT_170	● SP331 Peptococcus sp._HMT_168	● SPN2 Actinomyces sp._HMT_897_nov_97.461%
● SP163 Gemella moribillum	● SP332 Bulleidia extructa	● SPN20 Streptococcus mitis_nov_97.538%
● SP164 Gemella haemolysans	● SP333 Mogibacterium neglectum	● SPN21 Selenomonas artemidis_nov_97.710%
● SP165 Actinomyces israelii	● SP334 Veillonella sp._HMT_780	● SPN22 Actinomyces viscosus_nov_97.393%
● SP166 Saccharibacteria (TM7) [G-2] bacterium_HMT_350	● SP335 Selenomonas sp._HMT_479	● SPN23 Bacteroidetes [G-5] bacterium_HMT_511_nov_96.393%
● SP167 Actinomyces sp._HMT_896	● SP336 Treponema sp._ovinefootrot	● SPN24 Prevotella enoea_nov_96.680%
● SP168 Lachnoanaerobaculum umeaense	● SP337 Treponema amylovorum	● SPN25 Peptostreptococcaceae [XII][G-2] bacterium_HMT_091_nov_97.194%
● SP169 Actinomyces cardiffensis	● SP338 Saccharibacteria (TM7) [G-1] bacterium_HMT_352	● SPN26 Kingella oralis_nov_97.529%
● SP17 Streptococcus anginosus	● SP339 Prevotella sp._Oral_Taxon_B62	● SPN27 Capnocytophaga sp._Oral_Taxon_A48_nov_94.556%
● SP170 Streptococcus oralis_subsp._tigurinus_clade_071	● SP34 Alloprevotella sp._HMT_473	● SPN28 Peptococcus sp._Oral_Taxon_167_nov_97.798%
● SP171 Actinomyces odontolyticus	● SP340 Streptococcus parasanguinis_I	● SPN29 Streptococcus gordonii_nov_92.818%
● SP172 Prevotella sp._HMT_475	● SP341 NA nodatum	● SPN3 Actinomyces sp._HMT_448_nov_95.969%
● SP173 Actinomyces sp._Oral_Taxon_848	● SP342 Selenomonas sp._Oral_Taxon_G67	● SPN30 Actinomyces massiliensis_nov_97.790%
● SP174 Streptococcus thermophilus	● SP343 Selenomonas sp._HMT_138	● SPN31 Neisseria sp._Oral_Taxon_14_nov_97.154%
● SP175 Porphyromonas sp._HMT_277	● SP344 Shuttleworthia satelles	● SPN32 Capnocytophaga ochracea_nov_97.093%
● SP176 Johnsonella ignava	● SP345 Treponema sp._HMT_257	● SPN33 Neisseria elongata_nov_97.533%
● SP177 Selenomonas flueggei	● SP346 Stomatobaculum sp._HMT_373	● SPN34 Veillonella sp._HMT_780_nov_96.958%
● SP178 Peptostreptococcaceae [XI][G-9] [Eubacterium]_brachy	● SP347 Aggregatibacter sp._HMT_949	● SPN35 Actinomyces massiliensis_nov_89.496%
● SP179 Ottowia sp._HMT_894	● SP348 Streptococcus cristatus_clade_578	● SPN36 Actinomyces israelii_nov_97.393%
● SP18 Fusobacterium periodonticum	● SP349 unclassified_Peptostreptococcaceae oral_sp._str._CM2	● SPN37 Granulicatella adiacens_nov_97.575%
● SP180 Selenomonas artemidis	● SP35 Treponema sp._HMT_237	● SPN38 Anaeroglobus geminatus_nov_95.104%
● SP181 Centipeda periodontii	● SP350 Selenomonas sp._HMT_937	● SPN39 Treponema sp._HMT_257_nov_97.826%
● SP182 Actinomyces timonensis	● SP351 Pseudomonas fluorescens	● SPN4 Leptotrichia sp._HMT_215_nov_97.083%
● SP183 Saccharibacteria (TM7) [G-1] bacterium_HMT_347	● SP352 Tannerella forsythia	● SPN40 Renibacterium salmoninarum_nov_90.979%
● SP184 Gracilibacteria (GN02) [G-2] bacterium_HMT_873	● SP353 Anaerolineae [G-1] bacterium_HMT_439	● SPN41 Prevotella pleuritidis_nov_97.087%
● SP185 Parvimonas sp._HMT_110	● SP354 Mycoplasma faucium	● SPN42 Actinomyces oris_nov_92.435%
● SP186 Veillonella atypica	● SP355 Prevotella jejuni	● SPN43 Actinomyces massiliensis_nov_93.566%
● SP187 Saccharibacteria (TM7) [G-8] bacterium_HMT_955	● SP356 Treponema sp._Oral_Taxon_G85	● SPN44 Veillonellaceae [G-1] sp._Oral_Taxon_155_nov_96.241%
● SP188 Kingella sp._HMT_012	● SP357 Leptotrichia sp._HMT_463	● SPN45 Actinomyces sp._HMT_175_nov_97.830%
● SP189 Capnocytophaga sp._HMT_326	● SP358 Neisseria sp._HMT_523	● SPN46 Actinomyces sp._Oral_Taxon_171_nov_91.728%
● SP19 Streptococcus infantis	● SP359 Fretibacterium sp._HMT_359	● SPN47 Actinomyces israelii_nov_96.685%
● SP190 Megaspheera micronuciformis	● SP36 Capnocytophaga gingivalis	● SPN48 Brachymonas chironomi_nov_91.939%
● SP191 Atopobium rimae	● SP360 Aggregatibacter sp._HMT_512	● SPN49 Pleomorphochaeta multiformis_nov_85.882%
● SP193 Veillonellaceae [G-1] bacterium_HMT_135	● SP361 Selenomonas sp._HMT_136	● SPN5 Selenomonas flueggei_nov_96.750%
● SP194 Alloprevotella sp._HMT_912	● SP362 Capnocytophaga sp._HMT_335	● SPN50 Aggregatibacter sp._HMT_898_nov_88.521%
● SP195 Capnocytophaga granulosa	● SP363 Leptotrichia goodfellowii	● SPN51 Treponema sp._HMT_258_nov_97.843%
● SP196 Streptococcus sobrinus	● SP364 Bifidobacterium dentium	● SPN52 Streptococcus anginosus_nov_97.974%
● SP197 Prevotella sp._HMT_300	● SP365 Selenomonas sp._HMT_919	● SPN53 Leptotrichia sp._HMT_223_nov_93.347%
● SP198 Leptotrichia sp._HMT_221	● SP366 Colibacter massiliensis	● SPN54 Neisseria sp._Oral_Taxon_14_nov_96.584%
● SP199 Veillonellaceae [G-1] sp._Oral_Taxon_155	● SP367 Selenomonas sp._Oral_Taxon_E20	● SPN55 Streptococcus oralis_nov_95.113%
● SP2 Lautropia mirabilis	● SP368 unclassified_Peptostreptococcaceae Oral_sp._str._CM5	● SPN56 Capnocytophaga gingivalis_nov_95.720%
● SP20 Haemophilus parainfluenzae	● SP369 Selenomonas sp._Oral_Taxon_G55	● SPN57 Capnocytophaga gingivalis_nov_97.471%
● SP200 Lachnospiraceae [G-8] bacterium_HMT_500	● SP37 Parvimonas sp._Oral_Taxon_110	● SPN58 Prevotella sp._HMT_443_nov_97.047%
● SP201 Treponema socranskii	● SP370 Alloprevotella sp._HMT_308	● SPN59 Vagococcus penaei_nov_79.167%
● SP202 Filifactor alovis	● SP371 Capnocytophaga sp._HMT_878	● SPN6 Porphyromonas catoniae_nov_97.328%
● SP203 Stomatobaculum longum	● SP372 Veillonellaceae [G-1] bacterium_HMT_132	● SPN60 Herbiconiux ginsengi_nov_83.270%
● SP204 Prevotella baroniae	● SP373 Schaalia sp._HMT_178	● SPN61 Neisseria sp._Oral_Taxon_14_nov_96.970%
● SP205 Fretibacterium fastidiosum	● SP374 Neisseria sp._HMT_499	● SPN62 Actinomyces johnsonii_nov_95.194%
● SP206 Campylobacter gracilis	● SP375 Streptococcus sp._Oral_Taxon_B59	● SPN63 Megaspheera micronuciformis_nov_94.918%
● SP207 Porphyromonas sp._Oral_Taxon_B43	● SP376 Campylobacter sp._HMT_044	● SPN64 Fusobacterium nucleatum_ss_animalis_nov_97.605%
● SP208 Fusobacterium sp._HMT_203	● SP377 Phocaeicola abscessus	● SPN65 Saccharibacillus kuerlensis_nov_78.378%
● SP209 Porphyromonas endodontalis	● SP378 Bacteroidetes [G-3] bacterium_HMT_503	● SPN66 Streptococcus intermedius_nov_92.421%
● SP21 Selenomonas sp._HMT_137	● SP379 Haemophilus sp._HMT_036	● SPN67 Veillonellaceae [G-1] bacterium_HMT_150_nov_97.456%
● SP210 Gemella sanguinis	● SP38 Aggregatibacter segnis	● SPN68 Actinomyces massiliensis_nov_90.842%
● SP211 Bergeyella sp._HMT_900	● SP380 Saccharibacteria (TM7) [G-4] bacterium_HMT_355	● SPN69 Firmicutes [G] sp._Oral_Taxon_C68_nov_86.481%
● SP212 Bacteroidales [G-2] bacterium_HMT_274	● SP387 Schaalia cardiffensis	● SPN7 Actinomyces dentalis_nov_97.885%