

- SP171 Treponema socransii
- SP174 Treponema sp._oral_taxon_231
- SP175 Moraxella osloensis
- SP176 Streptococcus sp._oral_taxon_074
- SP177 Catonella morbi
- SP178 Streptococcus sp._str_2136FAA
- SP179 Agrobacterium tumefaciens
- SP18 Peptostreptococcaceae_[X1][G-1] [Eubacterium]_infirmum
- SP180 Selenomonas diana
- SP181 Gardnerella vaginalis
- SP182 Actinomyces massiliensis
- SP183 Staphylococcus auricularis
- SP184 Johnsonella ignava
- SP185 Fusobacteria_[G-1] sp._Oral_Taxon_A71
- SP186 TM7_[G-1] sp._oral_taxon_348
- SP187 Streptococcus sp._oral_taxon_058
- SP188 Fusobacterium periodonticum
- SP189 Cryptobacterium curtum
- SP19 Mogibacterium diversum
- SP190 Peptostreptococcaceae_[X1][G-7] [Eubacterium]_yurii_subsp._yurii_&_margaretiae
- SP191 Staphylococcus epidermidis
- SP192 Granulicatella sp._Oral_Taxon_D03
- SP193 Alloprevotella sp._oral_taxon_914
- SP194 Streptococcus peroris
- SP195 Aggregatibacter sp._oral_taxon_458
- SP196 Porphyromonas gingivalis
- SP197 Bacteroides heparinolyticus
- SP198 Scardovia wiggisiae
- SP199 Fusobacterium naviforme
- SP2 Tannerella forsythia
- SP20 Veillonellaceae_[G-1] sp._oral_taxon_132
- SP200 Leptotrichia sp._oral_taxon_219
- SP201 Streptococcus sp._oral_taxon_070
- SP202 Corynebacterium durum
- SP203 Klebsiella pneumoniae
- SP204 Veillonellaceae_[G-1] sp._oral_taxon_135
- SP207 TM7_[G-4] sp._oral_taxon_355
- SP208 Capnocytophaga sp._oral_taxon_326
- SP209 Porphyromonas sp._oral_taxon_279
- SP21 Actinomyces sp._oral_taxon_170
- SP210 Prevotella intermedia
- SP211 Bacteroidetes_[G-3] sp._oral_taxon_365
- SP212 Cardiobacterium valvarum
- SP213 Peptostreptococcaceae_[X1][G-2] sp._oral_taxon_091
- SP214 Brochothrix thermosphacta
- SP215 Bulleidia extructa
- SP216 Leptotrichia sp._oral_taxon_392
- SP217 SR1_[G-1] sp._oral_taxon_345
- SP218 Stomatobaculum sp._oral_taxon_373
- SP219 TM7_[G-1] sp._oral_taxon_869
- SP22 Actinomyces odontolyticus
- SP220 Gemella sanguinis
- SP221 Kocuria sp._Oral_Taxon_B56
- SP222 Actinomycetales_[G] sp._Oral_Taxon_C05
- SP223 Finegoldia magna
- SP224 Streptococcus sp._oral_taxon_057
- SP225 Peptostreptococcaceae_[X1][G-5] [Eubacterium]_saphenum
- SP226 Chryseobacterium treverense
- SP227 Selenomonas artemidis
- SP228 Streptococcus parasanguinis_I
- SP229 Leptotrichia sp._oral_taxon_498
- SP23 Abiotrophia defectiva
- SP230 Methylobacterium sp._Oral_Taxon_B84
- SP231 Planococcus kocurii
- SP232 Clostridiales_[F-1][G-1] sp._oral_taxon_093
- SP233 Prevotella maculosa
- SP234 TM7_[G-2] sp._oral_taxon_350
- SP383 Delftia sp._Oral_Taxon_A59
- SP384 Paracoccus chinensis
- SP385 Williamsia muralis
- SP386 Lachnoanaerobaculum sp._oral_taxon_083
- SP387 Peptostreptococcaceae_[G] sp._Oral_Taxon_B61
- SP388 Corynebacterium matruchotii
- SP389 Shuttleworthia satelles
- SP39 Actinomyces sp._oral_taxon_180
- SP390 Prevotella pleuritidis
- SP391 Corynebacterium aurimucosum
- SP392 Alloprevotella sp._oral_taxon_473
- SP393 Propionibacterium avidum
- SP394 Brevundimonas subvibrioides
- SP395 Enterococcus casseliflavus
- SP396 Leptotrichia hofstadii
- SP397 SR1_[G-1] sp._oral_taxon_875
- SP398 Ochrobactrum anthropi
- SP399 Aggregatibacter segnis
- SP4 Atopobium rimae
- SP40 Rhizobium rhizogenes_Oral_Taxon_D34
- SP400 Stenotrophomonas rhizophila
- SP401 Capnocytophaga gingivalis
- SP402 Bergeyella sp._oral_taxon_900
- SP403 Brachy bacterium rhamnosum
- SP404 Treponema sp._oral_taxon_257
- SP405 Bacillus sp._Oral_Taxon_C03
- SP406 Pyramidobacter piscicolens
- SP407 Moraxella sp._Oral_Taxon_B07
- SP408 Microbacterium foliorum
- SP409 Prevotella buccae
- SP41 Actinomyces oricola
- SP410 Pedobacter agri
- SP411 Treponema sp._oral_taxon_262
- SP412 Bacteroidetes_[G-3] sp._oral_taxon_280
- SP413 Afipia broomeae
- SP414 Prevotella sp._oral_taxon_314
- SP415 Prevotella micans
- SP416 Prevotella sp._oral_taxon_526
- SP417 Selenomonas sp._oral_taxon_501
- SP418 Brevundimonas staleyii
- SP419 Capnocytophaga sputigena
- SP42 Propionibacterium sp._oral_taxon_194
- SP420 Selenomonas sp._Oral_Taxon_G78
- SP421 Corynebacterium singulare
- SP422 Prevotella dentalis
- SP423 Veillonellaceae_[G-1] sp._Oral_Taxon_155
- SP424 Aeribacillus pallidus
- SP425 Prevotella sp._oral_taxon_300
- SP426 Aggregatibacter sp._oral_taxon_513
- SP427 Fusobacterium canifelinum
- SP428 Selenomonas sp._oral_taxon_149
- SP429 Neisseria sp._oral_taxon_020
- SP429 TM7_[G-1] sp._oral_taxon_347
- SP430 Duganella zoogloeoides
- SP431 Kribbella albertsoniae
- SP432 Dialister pneumosintes
- SP433 Deinococcus geothermalis
- SP434 Erysipelotrichaceae_[G-1] sp._oral_taxon_904
- SP435 Prevotella sp._oral_taxon_315
- SP436 Modestobacter roseus
- SP437 Corynebacterium sp._Oral_Taxon_D28
- SP438 Prevotella sp._oral_taxon_313
- SP439 Anoxybacillus flavithermus
- SP44 TM7_[G-1] sp._oral_taxon_952
- SP440 Prevotella baroniae
- SP441 Lachnospiraceae_[G] sp._Oral_Taxon_B18
- SP442 TM7_[G-6] sp._oral_taxon_870
- SPN137 Actinomyces sp._oral_taxon_180_nov_97.531%
- SPN138 Slackia exigua_nov_85.043%
- SPN139 Actinomyces sp._oral_taxon_172_nov_97.342%
- SPN14 Veillonellaceae_[G-1] sp._oral_taxon_129_nov_97.984%
- SPN140 TM7_[G-1] sp._oral_taxon_952_nov_96.909%
- SPN141 Actinomyces sp._oral_taxon_180_nov_97.938%
- SPN142 Alphaproteobacteria_[G] sp._Oral_Taxon_B70_nov_78.523%
- SPN143 Peptoniphilaceae_[G-1] sp._oral_taxon_113_nov_97.292%
- SPN144 Variovorax paradoxus_nov_94.227%
- SPN145 Actinomyces oricola_nov_97.421%
- SPN146 Clostridiales_[F-1][G-2] sp._oral_taxon_402_nov_80.973%
- SPN147 Streptococcus tigurinus_nov_94.800%
- SPN148 Pseudoramibacter alactolyticus_nov_97.107%
- SPN149 TM7_[G-1] sp._oral_taxon_348_nov_95.133%
- SPN15 Veillonellaceae_[G-1] sp._oral_taxon_918_nov_96.386%
- SPN150 Granulicatella adiacens_nov_97.206%
- SPN151 Ottowia sp._oral_taxon_894_nov_80.573%
- SPN152 Firmicutes_[G] sp._Oral_Taxon_C68_nov_85.921%
- SPN153 Comamonadaceae_[G] sp._Oral_Taxon_B87_nov_78.279%
- SPN154 Actinomyces sp._oral_taxon_175_nov_97.967%
- SPN155 Actinomyces sp._oral_taxon_171_nov_96.450%
- SPN156 Slackia exigua_nov_89.339%
- SPN157 Olsenella uli_nov_97.872%
- SPN158 Olsenella sp._oral_taxon_807_nov_97.872%
- SPN159 Mogibacterium timidum_nov_95.851%
- SPN16 Flexibacter flexilis_nov_82.255%
- SPN160 Hymenobacter roseosalivarius_nov_91.176%
- SPN161 Actinomyces israelii_nov_94.922%
- SPN162 Neisseria sp._oral_taxon_018_nov_97.959%
- SPN163 Afipia broomeae_nov_92.466%
- SPN164 Actinomyces israelii_nov_97.804%
- SPN165 Actinomyces sp._oral_taxon_178_nov_89.400%
- SPN166 Actinomyces sp._oral_taxon_170_nov_97.843%
- SPN167 Peptostreptococcaceae_[X1][G-3] sp._oral_taxon_950_nov_96.47%
- SPN168 Agrobacterium tumefaciens_nov_94.470%
- SPN169 Dietzia sp._Oral_Taxon_D12_nov_90.717%
- SPN17 Abiotrophia defectiva_nov_97.030%
- SPN170 Actinomyces sp._oral_taxon_525_nov_97.200%
- SPN18 Actinobaculum sp._oral_taxon_183_nov_93.673%
- SPN19 Rhizobium loti_nov_90.930%
- SPN2 Dietzia cinnamea_nov_91.006%
- SPN20 Mogibacterium timidum_nov_96.888%
- SPN21 Burkholderia cepacia_nov_95.277%
- SPN22 Cryptobacterium curtum_nov_88.223%
- SPN23 Actinomyces sp._oral_taxon_178_nov_96.334%
- SPN24 Propionibacterium sp._oral_taxon_915_nov_92.077%
- SPN25 panacarvi panacarvi_nov_78.842%
- SPN26 Actinomyces dentalis_nov_96.627%
- SPN27 Porphyromonas cationiae_nov_97.760%
- SPN28 Selenomonas sp._oral_taxon_146_nov_97.456%
- SPN29 Brevundimonas diminuta_nov_92.449%
- SPN3 Sphingomonas sp._oral_taxon_004_nov_80.320%
- SPN30 Actinomyces timonensis_nov_93.927%
- SPN31 Selenomonas sp._oral_taxon_481_nov_97.446%
- SPN32 Prevotella sp._oral_taxon_300_nov_97.976%
- SPN33 Actinomyces oricola_nov_97.065%
- SPN34 Anaerococcus lactolyticus_nov_91.023%
- SPN35 Acinetobacter sp._oral_taxon_408_nov_92.308%
- SPN36 Lachnospiraceae_[G-2] sp._oral_taxon_096_nov_88.187%
- SPN37 Gemella sanguinis_nov_81.854%
- SPN38 Anaerococcus prevotii_nov_96.033%
- SPN39 Defluviobacter lusatiensis_nov_89.862%
- SPN4 Actinomyces timonensis_nov_92.944%
- SPN40 Actinomyces sp._oral_taxon_448_nov_96.450%
- SPN41 Streptococcus infantis_nov_88.016%
- SPN42 Tannerella forsythia_nov_97.737%
- SPN43 Mogibacterium timidum_nov_95.851%