

- 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 cinnamomea\_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 catoniae\_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%