

Species

- SP106 Saccharibacteria_(TM7)[G-1] bacterium_HMT_347
- SP107 Actinomyces oris
- SP109 Porphyromonas catoniae
- SP114 Actinomyces sp._HMT_897
- SP116 Schaalia odontolytica
- SP118 Actinomyces sp._HMT_175
- SP119 Campylobacter concisus
- SP120 Leptotrichia sp._HMT_392
- SP121 Capnocytophaga gingivalis
- SP122 Abiotrophia defectiva
- SP123 Shigella sonnei
- SP124 Selenomonas infelix
- SP125 Veillonella dispar
- SP127 Absconditabacteria_(SR1)[G-1] bacterium_HMT_874
- SP128 Prevotella loescheii
- SP13 Haemophilus haemolyticus
- SP130 Prevotella salivae
- SP131 Truepera radiovictrix
- SP132 Actinomyces massiliensis
- SP133 Streptococcus constellatus
- SP136 Prevotella sp._HMT_317
- SP14 Bergeyella sp._HMT_322
- SP140 Porphyromonas gingivalis
- SP141 Prevotella sp._HMT_300
- SP142 Saccharibacteria_(TM7)[G-3] bacterium_HMT_351
- SP143 Leptotrichia buccalis
- SP145 Prevotella copri
- SP148 Selenomonas sp._HMT_920
- SP15 Bacteroidales_[G-2] bacterium_HMT_274
- SP150 Campylobacter gracilis
- SP151 Salmonella enterica
- SP153 Saccharibacteria_(TM7)[G-1] bacterium_HMT_348
- SP154 Prevotella oulorum
- SP156 Listeria monocytogenes
- SP158 Peptidiphaga sp._HMT_183
- SP159 Veillonella sp._HMT_780
- SP161 Haemophilus sp._HMT_036
- SP162 Streptococcus oralis_subsp._tigurinus_clade_071
- SP164 Leptotrichia hofstadii
- SP165 Halomonas alkaliantarctica
- SP166 Neisseria elongata
- SP168 Streptococcus chosunense
- SP169 Megasphaera micronuciformis
- SP17 Veillonella parvula
- SP170 Capnocytophaga sp._HMT_864
- SP172 Streptococcus oralis_subsp._dentisani_clade_058
- SP173 Haemophilus parainfluenzae
- SP176 Prevotella sp._HMT_314
- SP177 Streptococcus oralis
- SP178 Oribacterium sp._HMT_078
- SP18 Eikenella corrodens
- SP180 Capnocytophaga granulosa
- SP181 Alloprevotella tannerae
- SP184 Fusobacterium nucleatum_subsp._vincentii
- SP187 Aggregatibacter sp._HMT_513
- SP188 Cutibacterium acnes
- SP189 Ruminococcaceae_[G-1] bacterium_HMT_075
- SP190 Schaalia sp._HMT_180
- SP191 Enterococcus faecalis
- SP192 Neisseria flavescens
- SP193 Streptococcus oralis_subsp._tigurinus_clade_070
- SP194 Bergeyella sp._HMT_900
- SP195 Actinomyces sp._HMT_171
- SP197 Saccharibacteria_(TM7)[G-1] bacterium_HMT_349
- SP198 Dialister invisus
- SP2 Fusobacterium nucleatum_subsp._animalis
- SP200 Leptotrichia sp._HMT_212
- SP201 Prevotella bisticola
- SP218 Selenomonas noxia
- SP219 Prevotella intermedia
- SP22 Schaalia sp._HMT_178
- SP224 Rothia aeria
- SP227 Acidovorax temperans
- SP228 Phocaeicola vulgatus
- SP23 Fusobacterium sp._HMT_204
- SP231 Lachnoanaerobaculum gingivalis
- SP232 Parvimonas sp._HMT_110
- SP24 Catonella morbi
- SP241 Lancefieldella parvula
- SP243 Schaalia georgiae
- SP246 Capnocytophaga sputigena
- SP249 Allobacillus halotolerans
- SP250 Actinomyces sp._HMT_170
- SP257 Alloprevotella sp._HMT_914
- SP259 Streptococcus mitis
- SP261 Streptococcus anginosus
- SP262 Capnocytophaga sp._HMT_335
- SP263 Actinomyces johnsonii
- SP265 Peptostreptococcus stomatis
- SP266 Scardovia wiggisiae
- SP267 Citrobacter koseri
- SP268 Streptococcus sp._HMT_066
- SP269 Aggregatibacter segnis
- SP27 Prevotella oris
- SP270 Schaalia sp._HMT_877
- SP273 Rothia dentocariosa
- SP277 Solobacterium moorei
- SP278 Leptotrichia trevisanii
- SP279 Porphyromonas sp._HMT_278
- SP28 Stomatobaculum sp._HMT_097
- SP280 Streptococcus sp._HMT_064
- SP284 Anaeroglobus geminatus
- SP288 Tannerella serpentiniformis
- SP290 Saccharibacteria_(TM7)[G-1] bacterium_HMT_957
- SP292 Prevotella melaninogenica
- SP293 Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- SP294 Staphylococcus warneri
- SP296 Streptococcus cristatus
- SP297 Cardiobacterium hominis
- SP3 Capnocytophaga sp._HMT_412
- SP304 Ottowia sp._HMT_894
- SP306 Saccharibacteria_(TM7)[G-6] bacterium_HMT_870
- SP31 Actinomyces dentalis
- SP310 Streptococcus vestibularis
- SP312 Pseudomonas aeruginosa
- SP313 Rothia mucilaginosa
- SP315 Streptococcus cristatus_clade_578
- SP317 Capnocytophaga sp._HMT_863
- SP320 Streptococcus sp._HMT_423
- SP321 Veillonella atypica
- SP322 Actinomyces sp._HMT_414
- SP323 Capnocytophaga sp._HMT_324
- SP328 Olsenella sp._HMT_807
- SP33 Saccharibacteria_(TM7)[G-1] bacterium_HMT_346
- SP333 Actinomyces gerencseriae
- SP336 Capnocytophaga sp._HMT_902
- SP34 Prevotella denticola
- SP343 Herbaspirillum huttiense
- SP344 Streptococcus mutans
- SP348 Qipengyuania seohaensis
- SP35 Prevotella maculosa
- SP358 Neisseria cinerea
- SP36 Selenomonas artemidis
- SP365 Porphyromonas endodontalis
- SP367 Alloprevotella sp._HMT_473
- SP369 Ralstonia pickettii
- SP40 Lautropia mirabilis
- SP408 Schaalia meyeri
- SP410 Shigella boydii
- SP45 Streptococcus salivarius
- SP46 Treponema sp._HMT_237
- SP49 Leptotrichia hongkongensis
- SP5 Fusobacterium nucleatum
- SP50 Granulicatella elegans
- SP51 Leptotrichia sp._HMT_221
- SP52 Neisseria subflava
- SP53 Corynebacterium matruchotii
- SP56 Actinomyces naeslundii
- SP58 Leptotrichia sp._HMT_417
- SP59 Leptotrichia wadei
- SP6 Gemella haemolysans
- SP60 Peptidiphaga gingivicola
- SP62 Streptococcus sobrinus
- SP63 Peptostreptococcaceae_[XIII][G-7] [Eubacterium]_yurii_subsp._yurii_&_margaretiae
- SP64 Porphyromonas pasteri
- SP66 Lachnoanaerobaculum umeaense
- SP67 Cardiobacterium valvarum
- SP69 Campylobacter showae
- SP70 Prevotella sp._HMT_472
- SP71 Prevotella sp._HMT_942
- SP79 Fusobacterium hwasookii
- SP8 Treponema socranskii
- SP80 Prevotella sp._HMT_475
- SP81 Actinomyces sp._HMT_169
- SP84 Alloprevotella sp._HMT_308
- SP85 Gemella morbillorum
- SP86 Capnocytophaga sp._HMT_338
- SP88 Capnocytophaga leadbetteri
- SP89 Bacillus halotolerans
- SP9 Lachnoanaerobaculum saburreum
- SP91 Granulicatella adiacens
- SP92 Lachnospiraceae_[G-3] bacterium_HMT_100
- SP93 Leptotrichia sp._HMT_225
- SP95 Actinomyces sp._HMT_525
- SP97 Selenomonas sputigena
- SPN1 Bergeyella zoohelcum_nov_92.593%
- SPN11 Frondibacter mangrovi_nov_92.484%
- SPN15 Saccharibacteria_(TM7)[G-6] bacterium_HMT_870_nov_96.994%
- SPN2 Porphyromonas gingivalis_nov_96.319%
- SPN25 Porphyromonas gingivalis_nov_94.672%
- SPN3 Actinomyces sp._HMT_175_nov_97.746%
- SPN31 Porphyromonas gingivalis_nov_97.131%
- SPN36 Aeromicrobium panaciterrae_nov_96.104%
- SPN4 Actinomyces israelii_nov_96.647%
- SPN41 Actinomyces israelii_nov_94.882%
- SPN48 Veillonella sp._HMT_780_nov_97.228%
- SPN5 Actinomyces naeslundii_nov_97.760%
- SPN52 Saccharibacteria_(TM7)[G-1] bacterium_HMT_957_nov_97.550%
- SPN6 Schaalia odontolytica_nov_90.083%
- SPN60 Nocardopsis nikkonensis_nov_97.257%
- SPN61 Actinomyces sp._HMT_171_nov_96.813%
- SPN62 Actinomyces sp._HMT_169_nov_97.992%
- SPN63 Prevotella veroralis_nov_97.342%
- SPN64 Faecalibacterium prausnitzii_nov_96.976%
- SPN65 Lactimipora xylanolytica_nov_88.613%
- SPN66 Porphyromonas sp._HMT_284_nov_97.746%
- SPN7 Campylobacter concisus_nov_97.397%
- SPN8 Leptotrichia sp._HMT_225_nov_96.739%
- SPP15 Fusobacterium nucleatum_subsp._animalis
- SPP19 Halomonas alkaliphila_titanicae
- SPP25 Streptococcus infantis_infantis_clade_638
- SPP3 Veillonella dispar_parvula
- SPP5 Staphylococcus argenteus_aureus_roterodami
- SPP6 Prevotella sp._HMT_292_sp._HMT_300