

Species

- SP10 Porphyromonas catoniae
- SP100 Streptococcus mutans
- SP101 Tannerella forsythia
- SP104 Prevotella oralis
- SP105 Stomatobaculum longum
- SP107 Granulicatella adiacens
- SP108 Granulicatella paradiacens
- SP109 Actinomyces sp._Oral_Taxon_171
- SP110 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- SP111 Prevotella salivae
- SP116 Veillonella atypica
- SP118 Catonella morbi
- SP12 Prevotella oris
- SP120 Capnocytophaga gingivalis
- SP121 Bifidobacterium dentium
- SP122 Campylobacter showae
- SP124 Fretibacterium fastidiosum
- SP126 Prevotella sp._HMT_300
- SP128 Streptococcus sp._str._2136FAA
- SP129 Prevotella veroralis
- SP130 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- SP135 Propionibacterium acidifaciens
- SP139 Lachnoanaerobaculum orale
- SP140 Haemophilus parainfluenzae
- SP142 Oribacterium sp._Oral_Taxon_78
- SP144 Streptococcus sp._Oral_Taxon_71
- SP146 Leptotrichia buccalis
- SP147 Actinomyces viscosus
- SP15 Streptococcus cristatus
- SP152 Porphyromonas gingivalis
- SP153 Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum
- SP154 Streptococcus sp._str._C150
- SP155 Porphyromonas endodontalis
- SP158 Bergeyella sp._HMT_322
- SP160 Actinomyces sp._HMT_169
- SP165 Veillonellaceae_[G-1] bacterium_HMT_155
- SP166 Scardovia wiggisiae
- SP17 Streptococcus sanguinis
- SP170 Actinomyces odontolyticus
- SP171 Actinomyces sp._HMT_171
- SP172 Actinomyces naeslundii
- SP175 Streptococcus anginosus
- SP176 Kingella oralis
- SP18 Leptotrichia sp._AF189244.1
- SP181 Actinomyces sp._Oral_Taxon_178
- SP183 Gemella haemolysans
- SP184 Porphyromonas pasteri
- SP189 Olsenella sp._Oral_Taxon_807
- SP193 Rothia dentocariosa
- SP198 Streptococcus vestibularis
- SP200 Eikenella corrodens
- SP201 Actinomyces sp._str._ChDCB197
- SP202 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
- SP204 Prevotella multiformis
- SP205 Campylobacter gracilis
- SP21 Selenomonas sputigena
- SP212 Streptococcus oralis_subsp._tigurinus_clade_070
- SP215 Neisseria sicca
- SP217 Lancefieldella rimae
- SP218 Tannerella sp._HMT_286
- SP220 Leptotrichia sp._HMT_417
- SP224 Fusobacteria_[G-1] sp._Oral_Taxon_A71
- SP233 Prevotella pleuritidis
- SP234 Filifactor alocis
- SP24 Corynebacterium matruchotii
- SP242 Cardiobacterium hominis
- SP245 Lautropia mirabilis
- SP249 Actinomyces oris
- SP256 Abiotrophia defectiva
- SP259 Leptotrichia sp._HMT_215
- SP26 Actinomyces johnsonii
- SP260 Arachnia propionica
- SP262 Actinomyces sp._Oral_Taxon_180
- SP268 Capnocytophaga endodontalis
- SP27 Alloprevotella tannerae
- SP271 Leptotrichia sp._HMT_498
- SP272 Actinomyces sp._HMT_414
- SP278 Shuttleworthia satelles
- SP279 Solobacterium moorei
- SP28 Dialister invisus
- SP286 Actinomyces dentalis
- SP287 Anaeroglobus geminatus
- SP29 Prevotella sp._HMT_317
- SP293 Actinomyces israelii
- SP294 Fretibacterium sp._HMT_360
- SP296 Peptostreptococcus stomatis
- SP303 Streptococcus sp._HMT_064
- SP31 Parvimonas micra
- SP310 Actinomyces massiliensis
- SP311 Treponema denticola
- SP318 Rothia mucilaginosa
- SP32 Aggregatibacter sp._HMT_458
- SP320 Capnocytophaga granulosa
- SP324 Mogibacterium timidum
- SP33 Capnocytophaga sputigena
- SP332 Gemella morbillorum
- SP34 Capnocytophaga leadbetteri
- SP35 Bacteroidales_[G-2] sp._Oral_Taxon_274
- SP36 Prevotella loescheii
- SP361 Actinomyces sp._HMT_448
- SP369 Neisseria elongata
- SP374 Streptococcus salivarius
- SP40 Treponema sp._HMT_231
- SP41 Prevotella intermedia
- SP417 Olsenella uli
- SP42 Alloprevotella rava
- SP422 Peptidiphaga sp._HMT_183
- SP44 Neisseria mucosa
- SP442 Streptococcus sp._Oral_Taxon_B66
- SP45 Gemella morbillum
- SP465 Rothia aerea
- SP49 Leptotrichia wadei
- SP5 Streptococcus parasanguinis_II
- SP51 Fusobacterium nucleatum
- SP52 Streptococcus mitis
- SP53 Streptococcus constellatus
- SP55 Prevotella maculosa
- SP56 Streptococcus sp._str._C300
- SP58 Streptococcus oralis
- SP6 Veillonellaceae_[G-1] bacterium_HMT_150
- SP60 Prevotella denticola
- SP62 Actinomyces sp._HMT_175
- SP63 Treponema socranskii
- SP64 Streptococcus gordonii
- SP65 Prevotella sp._HMT_314
- SP67 Campylobacter concisus
- SP68 Corynebacterium durum
- SP69 Actinomyces sp._HMT_170
- SP70 Treponema sp._HMT_237
- SP72 Parvimonas sp._Oral_Taxon_110
- SP73 Actinomyces sp._HMT_525
- SP76 Actinomyces sp._Oral_Taxon_848
- SP79 Prevotella melaninogenica
- SP8 Selenomonas noxia
- SP82 Fusobacterium nucleatum_ss_polymorphum
- SP84 Fusobacterium nucleatum_ss_animalis
- SP86 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- SP88 Veillonella parvula
- SP9 Prevotella oulorum
- SP92 Megasphaera micronuciformis
- SP94 Streptococcus intermedius
- SP95 Actinomyces sp._HMT_897
- SP97 Lachnospiraceae_[G-3] bacterium_HMT_100
- SP98 Capnocytophaga ochracea
- SP99 Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- SPN109 Actinomyces sp._Oral_Taxon_171_nov_97.974%
- SPN129 Actinomyces viscosus_nov_95.849%
- SPN13 Actinomyces sp._HMT_175_nov_97.633%
- SPP1 Fusobacterium nucleatum_ss_vincentii_nucleatum_subsp._vincentii
- SPP17 Streptococcus salivarius_vestibularis
- SPP26 Atopobium_Lancefieldella_parvula_parvulum
- SPP3 Granulicatella adiacens_paradiacens
- SPP6 Eubacterium_[XIVa][G-1]_Lachnoanaerobaculum_saburreum
- SPP60 Fusobacterium nucleatum_nucleatum_ss_animalis