

Species

- SP1 Porphyromonas gingivalis
- SP10 Tannerella sp._HMT_286
- SP100 Gemella morbillorum
- SP101 Stomatobaculum sp._HMT_097
- SP102 Oribacterium sp._HMT_078
- SP103 Capnocytophaga gingivalis
- SP104 Campylobacter concisus
- SP105 Actinomyces sp._HMT_448
- SP106 Lachnoanaerobaculum saburreum
- SP107 Peptidiphaga sp._HMT_183
- SP108 Actinomyces johnsonii
- SP110 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- SP111 Streptococcus sobrinus
- SP112 Cutibacterium acnes
- SP113 Leptotrichia sp._HMT_392
- SP114 Peptostreptococcus stomatis
- SP115 Prevotella sp._HMT_314
- SP116 Escherichia coli
- SP117 Streptococcus oralis_subsp._tigurinus_clade_070
- SP118 Actinomyces sp._HMT_171
- SP119 Alloprevotella sp._HMT_473
- SP12 Streptococcus sanguinis
- SP120 Actinomyces gerenceriae
- SP121 Lachnospiraceae_[G-3] bacterium_HMT_100
- SP122 Leptotrichia sp._HMT_221
- SP123 Enterococcus faecalis
- SP124 Pseudomonas aeruginosa
- SP125 Porphyromonas pasteri
- SP128 Arachnia propionica
- SP129 Prevotella sp._HMT_475
- SP130 Actinomyces sp._HMT_175
- SP131 Porphyromonas cationiae
- SP132 Schaalia sp._HMT_877
- SP133 Bergeyella sp._HMT_322
- SP134 Selenomonas artemidis
- SP135 Leptotrichia hongkongensis
- SP136 Phyllobacterium myrsinacearum
- SP137 Streptococcus oralis_subsp._dentisani_clade_058
- SP138 Actinomyces dentalis
- SP143 Leptotrichia sp._HMT_225
- SP144 Acidovorax temperans
- SP146 Actinomyces oris
- SP147 Neisseria sicca
- SP148 Granulicatella elegans
- SP149 Schaalia sp._HMT_178
- SP150 Prevotella loescheii
- SP151 Erythrobacter aureus
- SP152 Streptococcus constellatus
- SP153 Actinomyces naeslundii
- SP154 Capnocytophaga ochracea
- SP156 Actinomyces sp._HMT_414
- SP16 Capnocytophaga leadbetteri
- SP160 Streptococcus vestibularis
- SP161 Haemophilus haemolyticus
- SP162 Streptococcus anginosus
- SP165 Peptidiphaga gingivicola
- SP166 Selenomonas noxia
- SP169 Lancefieldella parvula
- SP17 Kingella oralis
- SP170 Porphyromonas endodontalis
- SP171 Halomonas alkaliantarctica
- SP174 Stutzerimonas stutzeri
- SP175 Schaalia meyeri
- SP178 Neisseria cinerea
- SP18 Lachnoanaerobaculum gingivalis
- SP184 Prevotella copri
- SP185 Selenomonas sputigena
- SP188 Actinomyces viscosus
- SP19 Granulicatella adiacens
- SP190 Phocaeicola vulgatus
- SP191 Parvimonas micra
- SP193 Capnocytophaga sp._HMT_412
- SP2 Corynebacterium otitidis
- SP20 Streptococcus mutans
- SP201 Lachnoanaerobaculum orale
- SP202 Halomonas venusta
- SP209 Herbaspirillum huttiense
- SP22 Actinomyces sp._HMT_170
- SP23 Prevotella melaninogenica
- SP24 Halomonas titanicae
- SP25 Capnocytophaga granulosa
- SP26 Streptococcus chosunense
- SP27 Capnocytophaga sp._HMT_380
- SP28 Listeria monocytogenes
- SP29 Campylobacter showae
- SP3 Bacillus halotolerans
- SP30 Alloprevotella sp._HMT_308
- SP31 Streptococcus intermedius
- SP32 Streptococcus oralis
- SP33 Solobacterium moorei
- SP34 Capnocytophaga sp._HMT_335
- SP35 Truepera radiovictrix
- SP36 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
- SP37 Salmonella enterica
- SP38 Eikenella corrodens
- SP39 Lautropia mirabilis
- SP4 Fusobacterium nucleatum
- SP40 Neisseria subflava
- SP42 Prevotella intermedia
- SP43 Actinomyces sp._HMT_525
- SP45 Streptococcus sp._HMT_064
- SP46 Gemella haemolysans
- SP47 Leptotrichia buccalis
- SP48 Actinomyces sp._HMT_897
- SP49 Streptococcus gordonii
- SP5 Streptococcus mitis
- SP50 Prevotella nigrescens
- SP52 Staphylococcus warneri
- SP53 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP54 Leptotrichia sp._HMT_212
- SP55 Leptotrichia sp._HMT_215
- SP56 Prevotella sp._HMT_317
- SP57 Prevotella outorum
- SP58 Stomatobaculum longum
- SP59 Ruminococcaceae_[G-1] bacterium_HMT_075
- SP6 Veillonella parvula
- SP60 Parvimonas sp._HMT_110
- SP61 Rothia aera
- SP62 Corynebacterium matruchotii
- SP63 Abiotrophia defectiva
- SP64 Escherichia fergusonii
- SP65 Veillonella sp._HMT_780
- SP66 Ralstonia pickettii
- SP67 Schaalia odontolytica
- SP68 Aggregatibacter sp._HMT_513
- SP69 Ottowia sp._HMT_894
- SP7 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870
- SP70 Porphyromonas sp._HMT_278
- SP71 Streptococcus cristatus
- SP72 Leptotrichia wadei
- SP73 Prevotella oris
- SP74 Capnocytophaga sputigena
- SP75 Haemophilus sp._HMT_036
- SP76 Dialister invisus
- SP77 Peptoanaerobacter [Eubacterium] yurii
- SP78 Cardiobacterium valvarum
- SP79 Neisseria elongata
- SP8 Limosilactobacillus fermentum
- SP80 Prevotella denticola
- SP81 Rothia dentocariosa
- SP82 Campylobacter gracilis
- SP83 Shigella sonnei
- SP84 Schaalia sp._HMT_180
- SP85 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- SP86 Scardovia wiggisiae
- SP87 Aggregatibacter segnis
- SP88 Alloprevotella tannerae
- SP89 Rothia mucilaginosa
- SP9 Allobacillus halotolerans
- SP90 Haemophilus parainfluenzae
- SP91 Streptococcus salivarius
- SP92 Catonella morbi
- SP93 Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- SP94 Actinomyces massiliensis
- SP95 Prevotella salivae
- SP96 Olsenella sp._HMT_807
- SP97 Capnocytophaga endodontalis
- SP99 Actinomyces sp._HMT_169
- SPN1 Abiotrophia defectiva_nov_97.037%
- SPN11 Frondibacter mangrovi_nov_92.771%
- SPN15 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870_nov_97.093%
- SPN2 Actinomyces israelii_nov_96.514%
- SPN22 Faecalibacterium prausnitzii_nov_96.341%
- SPN27 Aeromicrobium panaciterrae_nov_96.327%
- SPN3 Actinomyces sp._HMT_175_nov_97.830%
- SPN33 Faecalibacterium prausnitzii_nov_97.755%
- SPN39 Veillonella sp._HMT_780_nov_97.519%
- SPN44 Bergeyella zoohelcum_nov_92.673%
- SPN50 Nocardiopsis nikkonensis_nov_97.250%
- SPN54 Prevotella veroralis_nov_96.780%
- SPN60 Corynebacterium matruchotii_nov_97.839%
- SPN61 Actinomyces naeslundii_nov_97.417%
- SPN62 Actinomyces viscosus_nov_95.833%
- SPN63 Capnocytophaga gingivalis_nov_97.276%
- SPN64 Selenomonas noxia_nov_96.154%
- SPN65 Porphyromonas cationiae_nov_97.909%
- SPN66 Actinomyces massiliensis_nov_96.344%
- SP2 Staphylococcus argenteus_aureus_rotterdamii
- SPP3 Prevotella sp._HMT_292_sp._HMT_300
- SPP4 Haemophilus seminalis_sp._HMT_908
- SPP8 Veillonella dispar_parvula