

Species

- SP1 Lachnospiraceae [G-9] bacterium_HMT_924
- SP10 Lactobacillus fermentum
- SP100 Actinomyces sp._HMT_169
- SP103 Haemophilus parainfluenzae
- SP104 Bacteroidales [G-2] sp._Oral_Taxon_274
- SP105 Selenomonas sp._HMT_134
- SP106 Peptostreptococcaceae [XI][G-9] [Eubacterium]_brachy
- SP107 Actinomyces sp._HMT_175
- SP109 Capnocytophaga granulosa
- SP11 Streptococcus sp._str._C300
- SP110 Capnocytophaga gingivalis
- SP111 Streptococcus parasanguinis_I
- SP112 Bacteroidales [G-2] bacterium_HMT_274
- SP113 Haemophilus haemolyticus
- SP114 Selenomonas sp._Oral_Taxon_E83
- SP115 Lautropia mirabilis
- SP12 Saccharibacteria (TM7) [G-1] bacterium_HMT_348
- SP13 Lactobacillus jensenii
- SP14 Porphyromonas endodontalis
- SP15 Saccharibacteria (TM7) [G-1] bacterium_HMT_346
- SP16 Gemella haemolysans
- SP17 Streptococcus mitis
- SP18 Corynebacterium mucifaciens
- SP19 Corynebacterium sp._Oral_Taxon_B71
- SP2 Gardnerella vaginalis
- SP20 Streptococcus chosunense
- SP21 Saccharibacteria (TM7) [G-1] bacterium_HMT_349
- SP22 Gardnerella swidsinskii
- SP25 Bergeyella sp._HMT_900
- SP26 Streptococcus sp._HMT_064
- SP27 Capnocytophaga leadbetteri
- SP28 Lactobacillus iners
- SP29 Arthrobacter humicola
- SP3 Gardnerella leopoldii
- SP30 Gemella moribillum
- SP31 Enterococcus faecalis
- SP32 Prevotella bivia
- SP33 Aerococcus christensenii_Oral_Taxon_D44
- SP34 Prevotella micans
- SP35 Streptococcus salivarius
- SP36 Cardiobacterium hominis
- SP37 Lactobacillus salivarius
- SP38 Limosilactobacillus coleohominis
- SP39 Parvimonas sp._HMT_110
- SP4 Streptococcus sanguinis
- SP40 Lactobacillus gasseri
- SP41 Capnocytophaga endodontalis
- SP42 Veillonella parvula
- SP43 Granulicatella elegans
- SP44 Granulicatella adiacens
- SP45 Gemella sp._Oral_Taxon_B93
- SP46 Haemophilus sp._HMT_036
- SP47 Saccharibacteria (TM7) [G-6] bacterium_HMT_870
- SP48 Parvimonas sp._Oral_Taxon_110
- SP49 Campylobacter showae
- SP5 Staphylococcus epidermidis
- SP51 Parvimonas micra
- SP52 Streptococcus oralis_subsp._tigurinus_clade_070
- SP53 Capnocytophaga sputigena
- SP54 Enterococcus sp._str._F95
- SP55 Streptococcus sp._Oral_Taxon_71
- SP56 Streptococcus cristatus
- SP57 Ruminococcaceae [G-1] bacterium_HMT_075
- SP59 Selenomonas sputigena
- SP6 Corynebacterium appendicis
- SP60 Streptococcus oralis
- SP61 Capnocytophaga sp._HMT_864
- SP62 Saccharibacteria (TM7) [G-5] bacterium_HMT_356
- SP65 Saccharibacteria (TM7) [G-8] bacterium_HMT_955
- SP66 Veillonella sp._HMT_780
- SP67 Selenomonas sp._Oral_Taxon_137
- SP68 Lachnospiraceae [G] sp._Oral_Taxon_B32
- SP69 Selenomonas sp._HMT_137
- SP7 Lactobacillus crispatus
- SP70 Tannerella sp._HMT_286
- SP71 Gemella morbillorum
- SP72 Sneathia amnii [Not_Validly_Published]
- SP73 Actinomyces naeslundii
- SP74 Saccharibacteria (TM7) [G-1] bacterium_HMT_957
- SP75 Actinomyces sp._HMT_171
- SP76 Streptococcus oralis_subsp._dentisani_clade_058
- SP77 Treponema denticola
- SP78 Fusobacterium nucleatum
- SP79 Streptococcus agalactiae
- SP8 Porphyromonas pasteri
- SP80 Actinomyces johnsonii
- SP81 Megaspheera genomosp.
- SP82 Saccharibacteria (TM7) [G-3] bacterium_HMT_351
- SP84 Streptococcus oralis_subsp._tigurinus_clade_071
- SP85 Bergeyella sp._HMT_322
- SP86 Absconditabacteria (SR1) [G-1] bacterium_HMT_345
- SP87 Sneathia sanguinegens
- SP88 Granulicatella paradiacens
- SP89 Mogibacterium vescum
- SP9 Corynebacterium striatum
- SP90 Fretibacterium sp._HMT_359
- SP91 Mycoplasma hominis
- SP92 Selenomonas sp._HMT_133
- SP93 Capnocytophaga ochracea
- SP94 Actinomyces oris
- SP96 Campylobacter gracilis
- SP97 Prevotella melaninogenica
- SPN1 Lactobacillus jensenii_nov_96.230%
- SPN10 Selenomonas sp._HMT_133_nov_96.947%
- SPN11 Anaerococcus lactolyticus_nov_97.271%
- SPN12 Gordonibacter pamelaeeae_nov_88.867%
- SPN13 Prevotella colorans_nov_96.212%
- SPN14 Streptococcus cristatus_nov_97.363%
- SPN15 Selenomonas sp._Oral_Taxon_137_nov_97.786%
- SPN16 Granulicatella elegans_nov_97.180%
- SPN17 Streptobacillus notomytis_nov_93.555%
- SPN18 Capnocytophaga gingivalis_nov_97.471%
- SPN19 Lactobacillus gasseri_nov_96.948%
- SPN2 Lactobacillus vaginalis_nov_93.889%
- SPN20 Bacteroidetes [G-3] bacterium_HMT_281_nov_88.583%
- SPN21 Parvimonas sp._Oral_Taxon_110_nov_97.466%
- SPN22 Saccharibacteria (TM7) [G-5] bacterium_HMT_356_nov_93.435%
- SPN23 Actinomyces sp._HMT_448_nov_96.346%
- SPN24 Saccharibacteria (TM7) [G-1] bacterium_HMT_957_nov_96.774%
- SPN3 Parvimonas micra_nov_91.456%
- SPN4 Limosilactobacillus coleohominis_nov_93.381%
- SPP2 Streptococcus pneumoniae_sp._str._C300_sp._str._M334
- SPP3 Corynebacterium pseudogenitalium_tuberculostearicum
- SPP4 Streptococcus pneumoniae_sp._str._M334
- SPP5 Staphylococcus argenteus_aureus
- SPPN1 Corynebacterium multispecies_sppn1_2_nov_96.607%