

## 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%