

Species

SP103 Roseburia intestinalis	SP323 Capnocytophaga leadbetteri
SP104 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346	SP325 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
SP106 Schaalia sp._HMT_180	SP341 Phascolarctobacterium faecium
SP108 Haemophilus parainfluenzae	SP350 Coprococcus catus
SP109 Bacteroidales_[G-2] sp._Oral_Taxon_274	SP355 Granulicatella elegans
SP113 Rothia aera	SP363 Eubacterium ramulus
SP116 Rothia dentocariosa	SP368 Bifidobacterium bifidum
SP118 Abiotrophia defectiva	SP38 Prevotella stercorea
SP119 Bacteroides fragilis	SP384 TM7_[G] sp._Oral_Taxon_A56
SP124 Fusobacterium periodonticum	SP386 Eubacterium ventriosum
SP126 Butyrivibrio crossotus	SP388 Lachnospirillum sp._str._L2_50
SP132 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349	SP392 Corynebacterium argentoratense
SP133 Schaalia sp._HMT_172	SP40 Massiliaprevotella massiliensis
SP136 Neisseria elongata	SP400 Alistipes finegoldii
SP137 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356	SP403 Akkermansia muciniphila
SP155 Peptostreptococcus stomatis	SP407 Eubacterium siraeum
SP158 Tannerella forsythia	SP41 Phocaicola vulgatus
SP160 Holdemanella biformis	SP419 Duodenibacillus massiliensis
SP163 Prevotella salivae	SP426 Veillonella atypica
SP164 Bacteroides caccae	SP43 Porphyromonas pasteri
SP171 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351	SP435 Selenomonas noxia
SP175 Blautia obeum	SP439 Eubacterium hallii
SP179 Bergeyella sp._HMT_322	SP53 Faecalibacterium prausnitzii
SP182 Veillonellaceae_[G-1] bacterium_HMT_155	SP56 Neisseria subflava
SP183 Prevotella histicola	SP59 Streptococcus anginosus
SP185 Fusicatenaibacter saccharivorans	SP60 Ruminococcaceae_[G-1] bacterium_HMT_075
SP186 Peptostreptococcaceae_[XII][G-1] [Eubacterium] sulci	SP66 Peptostreptococcaceae_[XII][G-9] [Eubacterium] brachy
SP188 Mycoplasma faucium	SP68 Parvimonas micra
SP193 Agathobaculum butyriciproducens	SP70 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352
SP206 Oribacterium sinus	SP76 Actinomyces odontolyticus
SP207 Barnesiella intestinhominis	SP77 Dorea longicatena
SP208 Megasphaera elsdenii	SP78 Subdoligranulum variabile
SP222 Prevotella melaninogenica	SP79 Collinsella aerofaciens
SP223 Ruminococcaceae_[G-2] bacterium_HMT_085	SP81 Solobacterium moorei
SP224 Odoribacter splanchnicus	SP86 Blautia faecis
SP225 Methanobrevibacter smithii	SP89 Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873
SP227 Erysipelotrichaceae_[G-1] bacterium_HMT_905	SP93 Bifidobacterium longum
SP23 Gemella sanguinis	SP98 Streptococcus thermophilus
SP234 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345	SPN109 Acetanaerobacterium elongatum_nov_92.081%
SP236 Campylobacter concisus	SPN119 Oscillibacter ruminantium_nov_94.144%
SP238 Parabacteroides distasonis	SPN120 Oscillibacter valericigenes_nov_94.344%
SP24 Ruminococcus callidus	SPN129 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_93.946%
SP240 Parabacteroides merdae	SPN140 Prevotella stercorea_nov_97.403%
SP241 Actinomyces graevenitzi	SPN144 Ruminococcus callidus_nov_94.369%
SP243 Alistipes putredinis	SPN149 Eubacterium coprostanoligenes_nov_97.059%
SP248 Porphyromonas endodontalis	SPN170 Kineothrix alysioides_nov_96.825%
SP25 Rothia muclaginoso	SPN182 Faecalibacterium prausnitzii_nov_97.964%
SP252 Porphyromonas sp._HMT_285	SPN193 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_89.038%
SP254 Mediterraneibacter faecis	SPN194 Oscillibacter ruminantium_nov_93.919%
SP259 Lactobacillus rogosae	SPN2 Alloprevotella rava_nov_92.857%
SP26 Eubacterium rectale	SPN205 Lachnospira pectinoschiza_nov_96.372%
SP263 Romboutsia timonensis	SPN216 Prevotella copri_nov_97.397%
SP266 Coprococcus eutactus	SPN226 Lachnospira eligens_nov_94.785%
SP268 Capnocytophaga gingivalis	SPN23 unclassified_Ruminococcaceae sp._str._D16_nov_93.243%
SP27 Prevotella copri	SPN236 Eubacterium ruminantium_nov_97.279%
SP270 Bilophila wadsworthia	SPN246 Butyrivibrio sp._HMT_090_nov_92.795%
SP273 Dorea formicigenerans	SPN249 Sporobacter termitidis_nov_90.562%
SP275 Gemella haemolysans	SPN255 Paraprevotella clara_nov_97.609%
SP28 Blautia luti	SPN264 Prevotella colorans_nov_93.492%
SP280 Corynebacterium durum	SPN275 Sporobacter termitidis_nov_93.243%
SP283 Stomatobaculum sp._HMT_097	SPN289 Christensenella massiliensis_nov_90.766%
SP288 Alistipes shahii	SPN301 Acutalibacter muris_nov_96.599%
SP29 Lachnospira eligens	SPN322 Prevotellamassilia timonensis_nov_97.397%
SP295 Lachnospirillum sp._SS3/4	SPN334 Eubacterium coprostanoligenes_nov_92.777%
SP297 Coprococcus comes	SPN335 Kineothrix alysioides_nov_96.614%
SP30 Prevotella pallens	SPN346 Asteroleplasma anaerobium_nov_80.300%
SP301 Ruminococcus bromii	SPN35 Succinivibrio dextrinosolvens_nov_97.079%
SP307 Roseburia inulinivorans	SPN358 Eubacterium coprostanoligenes_nov_95.711%
SP309 Catenibacterium mitsuokai	SPN366 Prevotella oralis_nov_92.641%

SPN377 Dialister succinatiphilus_nov_96.567%
SPN39 Duncaniella freteri_nov_87.069%
SPN390 Prevotella copri_nov_96.537%
SPN399 Oscillibacter ruminantium_nov_93.049%
SPN409 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.422%
SPN418 Ruminococcus bromii_nov_97.528%
SPN427 Bergeyella sp._HMT_907_nov_97.397%
SPN436 Dialister succinatiphilus_nov_96.360%
SPN444 Oscillibacter ruminantium_nov_95.701%
SPN45 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.968%
SPN455 Prevotella copri_nov_96.963%
SPN462 Fusicatenaibacter saccharivorans_nov_97.279%
SPN467 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_97.072%
SPN536 Lactrimispora xylanolytica_nov_97.285%
SPN55 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_91.011%
SPN567 Monoglobus pectinilyticus_nov_90.112%
SPN610 Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
SPN621 Lachnospirillum pacaense_nov_96.145%
SPN635 Sporobacter termitidis_nov_92.584%
SPN64 Ruminoclostridium methylpentosum_nov_89.462%
SPN75 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_92.809%
SPN87 Agathobaculum desmolans_nov_97.738%
SPN96 Oscillibacter ruminantium_nov_92.601%
SPN98 Holdemanella biformis_nov_97.854%
SPP1 Bacteroides sp._str._4136_uniformis
SPP104 Anaerostipes Lachnospirillum unclassified_Lachnospiraceae hadrus_sp._SSC/2_sp._str._cont1.81
SPP105 Lactobacillus_Ligilactobacillus ruminis
SPP109 Neisseria flavescens_flavescens[subflava]
SPP11 Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_...(20 species)
SPP111 Neisseria flavescens[subflava_perflava]
SPP113 Streptococcus sanguinis_sp._HMT_074
SPP117 Blautia_Mediterraneibacter [Ruminococcus] lactaris_lactaris
SPP118 Atopobium_Lancefieldella parvula_parvulum
SPP12 Roseburia faecalis_faecis
SPP134 Granulicatella adiacens_padiacens
SPP135 Actinomyces_Schaalia odontolytica_odontolyticus
SPP137 Blautia_Mediterraneibacter [Ruminococcus] gnnavus_gnavus
SPP138 Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torqueus
SPP14 Lachnospirillum sp._SM4/1_sp._str._M62/1
SPP15 Bacteroides_Phocaicola dorei_sp._Oral_Taxon_D25_sp._str._219_sp._str._313_...(4 species)
SPP17 Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibulari_...(4 species)
SPP18 Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
SPP19 Bifidobacterium adolescentis_faecale
SPP26 Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
SPP27 Gemella morbillorum_morbillum
SPP29 Blautia obeum_wexlerae
SPP39 Neisseria flava_macacae_mucosa_sicca_sp._TM101
SPP5 Actinomyces_Schaalia odontolyticus_sp._HMT_180
SPP54 Parasutterella_Sutterella excrementihominis_sp._str._cont1.66
SPP65 Selenomonas sp._HMT_136_sp._Oral_Taxon_149
SPP60 Veillonella rogosae_sp._str._NVG100cf
SPP67 Bifidobacterium catenulatum_pseudocatenulatum
SPP68 Bacteroides koreensis_kribbi
SPP70 Veillonella dispar_parvula
SPP71 Veillonella parvula_tobetsuensis
SPP74 Bacteroides_Phocaicola plebeius
SPP79 Oribacterium parvum_sp._str._ACB1
SPP8 Mogibacterium diversum_neglectum_vescum
SPP87 Actinomyces sp._HMT_169_sp._str._ChDCB197
SPP95 Streptococcus parasanguinis_parasanguinis_L_parasanguinis_clade_...(4 species)
SPPN1 Ruminococcus multispecies_sppn1_2_nov_95.023%
SPPN106 Roseburia multispecies_sppn106_3_nov_96.833%
SPPN2 Christensenella multispecies_sppn2_2_nov_91.216%
SPPN27 Clostridium multispecies_sppn27_2_nov_97.959%
SPPN34 Mobiluncus multispecies_sppn34_2_nov_97.817%
SPPN46 multigenus multispecies_sppn46_2_nov_92.568%
SPPN52 Roseburia multispecies_sppn52_2_nov_97.072%
SPPN68 Lachnospirillum multispecies_sppn68_2_nov_97.279%