

SP104 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346	SP325 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369	SPN39 Duncaniella freteri_nov_87.069%
SP106 Schaalia sp._HMT_180	SP341 Phascolarctobacterium faecium	SPN390 Prevotella copri_nov_96.537%
SP108 Haemophilus parainfluenzae	SP350 Coprococcus catus	SPN399 Oscillibacter ruminantium_nov_93.049%
SP109 Bacteroidales_[G-2] sp._Oral_Taxon_274	SP355 Granulicatella elegans	SPN409 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.422%
SP113 Rothia aeria	SP363 Eubacterium ramulus	SPN418 Ruminococcus bromii_nov_97.528%
SP116 Rothia dentocariosa	SP368 Bifidobacterium bifidum	SPN427 Bergeyella sp._HMT_907_nov_97.397%
SP118 Abiotrophia defectiva	SP38 Prevotella stercorea	SPN436 Dialister succinatiphilus_nov_96.360%
SP119 Bacteroides fragilis	SP384 TM7_[G] sp._Oral_Taxon_A56	SPN444 Oscillibacter ruminantium_nov_95.701%
SP124 Fusobacterium periodonticum	SP386 Eubacterium ventriosum	SPN45 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.968%
SP126 Butyrivibrio crossotus	SP388 Lactoclostridium sp._str._L2_50	SPN455 Prevotella copri_nov_96.963%
SP132 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349	SP392 Corynebacterium argentoratense	SPN462 Fusicatenibacter saccharivorans_nov_97.279%
SP133 Schaalia sp._HMT_172	SP40 Massiliiprevotella massiliensis	SPN467 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_97.072%
SP136 Neisseria elongata	SP400 Alistipes finegoldii	SPN436 Lacrimispora xylanolytica_nov_97.285%
SP137 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356	SP403 Akkermansia muciniphila	SPN55 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_91.011%
SP155 Peptostreptococcus stomatis	SP407 Eubacterium siraeum	SPN567 Monoglobus pectinilyticus_nov_90.112%
SP158 Tannerella forsythia	SP41 Phocaecicola vulgatus	SPN610 Erysipelatoclostridium_[Clostridium] spiroforme_nov_93.333%
SP160 Holdemanella biformis	SP419 Duodenibacillus massiliensis	SPN621 Lactoclostridium pacaense_nov_96.145%
SP163 Prevotella salivae	SP426 Veillonella atypica	SPN635 Sporobacter termitidis_nov_92.584%
SP164 Bacteroides caccae	SP43 Porphyromonas pasteri	SPN64 Ruminoclostridium methylpentosum_nov_89.462%
SP171 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351	SP435 Selenomonas noxia	SPN75 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_92.809%
SP175 Blautia obeum	SP439 Eubacterium hallii	SPN87 Agathobaculum desmolans_nov_97.738%
SP179 Bergeyella sp._HMT_322	SP53 Faecalibacterium prausnitzii	SPN96 Oscillibacter ruminantium_nov_92.601%
SP182 Veillonellaceae_[G-1] bacterium_HMT_155	SP56 Neisseria subflava	SPN98 Holdemanella biformis_nov_97.854%
SP183 Prevotella histicola	SP59 Streptococcus anginosus	SPP1 Bacteroides sp._str._4136_uniformis
SP185 Fusicatenibacter saccharivorans	SP60 Ruminococcaceae_[G-1] bacterium_HMT_075	SPP104 Anaerostipes_Lactoclostridium_unclassified_Lachnospiraceae_hadrus_sp._SSC/2_sp._
SP186 Peptostreptococcaceae_[XII][G-1] [Eubacterium]_sulci	SP66 Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy	SPP105 Lactobacillus_Ligilactobacillus_ruminis
SP188 Mycoplasma faucium	SP68 Parvimonas micra	SPP109 Neisseria flavescens_flavescens subflava
SP193 Agathobaculum butyriciproducens	SP70 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352	SPP11 Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_...(20 species)
SP206 Oribacterium sinus	SP76 Actinomyces odontolyticus	SPP111 Neisseria flavescens subflava_perflava
SP207 Barnesiella intestinihominis	SP77 Dorea longicatena	SPP113 Streptococcus sanguinis_sp._HMT_074
SP208 Megasphaera elsdenii	SP78 Subdoligranulum variabile	SPP117 Blautia_Mediterraneibacter [Ruminococcus] lactaris_lactaris
SP222 Prevotella melaninogenica	SP79 Collinsella aerofaciens	SPP118 Atopobium_Lancefieldella parvula_parvulum
SP223 Ruminococcaceae_[G-2] bacterium_HMT_085	SP81 Solobacterium moorei	SPP12 Roseburia faecalis_faecis
SP224 Odoribacter splanchnicus	SP86 Blautia faecis	SPP134 Granulicatella adiacens_paradiacens
SP225 Methanobrevibacter smithii	SP89 Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873	SPP135 Actinomyces_Schaalia odontolytica_odontolyticus
SP227 Erysipelotrichaceae_[G-1] bacterium_HMT_905	SP93 Bifidobacterium longum	SPP137 Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
SP23 Gemella sanguinis	SP98 Streptococcus thermophilus	SPP138 Blautia_unclassified_Lachnospiraceae_sp._str._cont1.79_torques
SP234 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345	SPN109 Acetanaerobacterium elongatum_nov_92.081%	SPP14 Lactoclostridium sp._SM4/1_sp._str._M62/1
SP236 Campylobacter concisus	SPN119 Oscillibacter ruminantium_nov_94.144%	SPP15 Bacteroides_Phocaecicola dorei_sp._Oral_Taxon_D25_sp._str._219_sp._str._313_...(4 spe
SP238 Parabacteroides distasonis	SPN120 Oscillibacter valericigenes_nov_94.344%	SPP17 Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibulari_...(4 species)
SP24 Ruminococcus callidus	SPN129 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_93.946%	SPP18 Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
SP240 Parabacteroides merdae	SPN140 Prevotella stercorea_nov_97.403%	SPP19 Bifidobacterium adolescentis_faecale
SP241 Actinomyces graevenitzi	SPN144 Ruminococcus callidus_nov_94.369%	SPP26 Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
SP243 Alistipes putredinis	SPN149 Eubacterium coprostanoligenes_nov_97.059%	SPP27 Gemella morbillorum_morbillum
SP248 Porphyromonas endodontalis	SPN170 Kineothrix alysoides_nov_96.825%	SPP29 Blautia obeum_wexlerae
SP25 Rothia mucilaginosa	SPN182 Faecalibacterium prausnitzii_nov_97.964%	SPP39 Neisseria flava_macacae_mucosa_sicca_sp._TM101
SP252 Porphyromonas sp._HMT_285	SPN193 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_89.038%	SPP5 Actinomyces_Schaalia odontolyticus_sp._HMT_180
SP254 Mediterraneibacter faecis	SPN194 Oscillibacter ruminantium_nov_93.919%	SPP54 Parasutterella_Sutterella excrementihominis_sp._str._cont1.66
SP259 Lactobacillus rogosae	SPN2 Alloprevotella rava_nov_92.857%	SPP56 Selenomonas sp._HMT_136_sp._Oral_Taxon_149
SP26 Eubacterium rectale	SPN205 Lachnospira pectinoschiza_nov_96.372%	SPP60 Veillonella rogosae_sp._str._NVG100cf
SP263 Romboutsia timonensis	SPN216 Prevotella copri_nov_97.397%	SPP67 Bifidobacterium catenulatum_pseudocatenulatum
SP266 Coprococcus eutactus	SPN226 Lachnospira eligens_nov_94.785%	SPP68 Bacteroides koreensis_kribbi
SP268 Capnocytophaga gingivalis	SPN23 unclassified_Ruminococcaceae_sp._str._D16_nov_93.243%	SPP70 Veillonella dispar_parvula
SP27 Prevotella copri	SPN236 Eubacterium ruminantium_nov_97.279%	SPP71 Veillonella parvula_tobetsuensis
SP270 Bilophila wadsworthia	SPN246 Butyrivibrio sp._HMT_090_nov_92.795%	SPP74 Bacteroides_Phocaecicola plebeius
SP273 Dorea formicigenerans	SPN249 Sporobacter termitidis_nov_90.562%	SPP79 Oribacterium parvum_sp._str._ACB1
SP275 Gemella haemolysans	SPN255 Paraprevotella clara_nov_97.609%	SPP8 Mogibacterium diversum_neglectum_vescum
SP28 Blautia luti	SPN264 Prevotella colorans_nov_93.492%	SPP87 Actinomyces sp._HMT_169_sp._str._ChDCB197
SP280 Corynebacterium durum	SPN275 Sporobacter termitidis_nov_93.243%	SPP95 Streptococcus parasanguinis parasanguinis_I parasanguinis_clade_...(4 species)
SP283 Stomatobaculum sp._HMT_097	SPN289 Christensenella massiliensis_nov_90.766%	SPPN1 Ruminococcus multispecies_sppn1_2_nov_95.023%
SP288 Alistipes shahii	SPN301 Acutalibacter muris_nov_96.599%	SPPN106 Roseburia multispecies_sppn106_3_nov_96.833%
SP29 Lachnospira eligens	SPN322 Prevotellamassilia timonensis_nov_97.397%	SPPN2 Christensenella multispecies_sppn2_2_nov_91.216%
SP295 Lactoclostridium sp._SS3/4	SPN334 Eubacterium coprostanoligenes_nov_92.777%	SPPN27 Clostridium multispecies_sppn27_2_nov_97.959%
SP297 Coprococcus comes	SPN335 Kineothrix alysoides_nov_96.614%	SPPN34 Mobiluncus multispecies_sppn34_2_nov_97.817%
SP30 Prevotella pallens	SPN346 Asteroleplasma anaerobium_nov_80.300%	SPPN46 multigenus multispecies_sppn46_2_nov_92.568%
SP301 Ruminococcus bromii	SPN35 Succinivibrio dextrinosolvens_nov_97.079%	SPPN52 Roseburia multispecies_sppn52_2_nov_97.072%
SP307 Roseburia inulinivorans	SPN358 Eubacterium coprostanoligenes_nov_95.711%	SPPN68 Lactoclostridium multispecies_sppn68_2_nov_97.279%