

SP323 Capnocytophaga leadbetteri	SPN377 Dialister succinatiphilus_nov_96.567%
SP325 Peptostreptococcaceae_[X][G-4] bacterium_HMT_369	SPN39 Duncaniella freteri_nov_87.069%
SP341 Phascolarctobacterium faecium	SPN390 Prevotella copri_nov_96.537%
SP350 Coprococcus catus	SPN399 Oscillibacter ruminantium_nov_93.049%
SP355 Granulicatella elegans	SPN409 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.422%
SP363 Eubacterium ramulus	SPN418 Ruminococcus bromii_nov_97.528%
SP368 Bifidobacterium bifidum	SPN427 Bergeyella sp._HMT_907_nov_97.397%
SP38 Prevotella stercorea	SPN436 Dialister succinatiphilus_nov_96.360%
SP384 TM7_[G] sp._Oral_Taxon_A56	SPN444 Oscillibacter ruminantium_nov_95.701%
SP386 Eubacterium ventriosum	SPN45 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.968%
SP388 Lachnospirillum sp._str._L2_50	SPN455 Prevotella copri_nov_96.963%
SP392 Corynebacterium argentoratense	SPN462 Fusicatibacter saccharivorans_nov_97.279%
SP40 Massilibrevotella massiliensis	SPN467 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_97.072%
SP400 Alistipes finegoldii	SPN536 Lactrimispora xylanolytica_nov_97.285%
SP403 Akkermansia muciniphila	SPN55 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_91.011%
SP407 Eubacterium siraeum	SPN567 Monoglobus pectinilyticus_nov_90.112%
SP41 Phocaeicola vulgatus	SPN610 Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
SP419 Duodenibacillus massiliensis	SPN621 Lachnospirillum pacaense_nov_96.145%
SP426 Veillonella atypica	SPN635 Sporobacter termitidis_nov_92.584%
SP43 Porphyromonas pasteri	SPN64 Ruminoclostridium methylpentosum_nov_89.462%
SP435 Selenomonas noxia	SPN75 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_92.809%
SP439 Eubacterium hallii	SPN87 Agathobaculum desmolans_nov_97.738%
SP53 Faecalibacterium prausnitzii	SPN96 Oscillibacter ruminantium_nov_92.601%
SP56 Neisseria subflava	SPN98 Holdemanella biformis_nov_97.854%
SP59 Streptococcus anginosus	SPP1 Bacteroides sp._str._4136_uniformis
SP60 Ruminococcaceae_[G-1] bacterium_HMT_075	SPP104 Anaerostipes_Lachnospiraceae_unclassified_Lachnospiraceae_hadrus_sp._SSC2/sp._str._cont1.81
SP66 Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy	SPP105 Lactobacillus_Ligilactobacillus_ruminis
SP68 Parvimonas micra	SPP109 Neisseria flavescens_flavescens[subflava
SP70 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352	SPP11 Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_infantis_clade_638_mitis_oralis_oralis_subsp._dentisani_clade_058_oralis_subsp._dentisani_clade_398_oralis_subsp._oralis_oralis_subsp._tigurinus_clade_070_oralis_subsp._tigurinus_clade_071_periodonticum_pneumoniae_sp._
SP76 Actinomyces odontolyticus	SPP111 Neisseria flavescens[subflava_perflava
SP77 Dorea longicatena	SPP113 Streptococcus sanguinis_sp._HMT_074
SP78 Subdoligranulum variabile	SPP117 Blautia_Mediterraneibacter [Ruminococcus] lactaris_lactaris
SP79 Collinsella aerofaciens	SPP118 Atopobium_Lancefieldella parvula_parvulum
SP81 Solobacterium moorei	SPP12 Roseburia faecalis_faecis
SP86 Blautia faecis	SPP134 Granulicatella adiacens_paradiacens
SP89 Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873	SPP135 Actinomyces_Schaalia odontolytica_odontolyticus
SP93 Bifidobacterium longum	SPP137 Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
SP98 Streptococcus thermophilus	SPP138 Blautia_unclassified_Lachnospiraceae_sp._str._cont1.79_torques
SPN109 Acetanaerobacterium elongatum_nov_92.081%	SPP14 Lachnospirillum sp._SM4/1_sp._str._M62/1
SPN119 Oscillibacter ruminantium_nov_94.144%	SPP15 Bacteroides_Phocaeicola dorei_sp._Oral_Taxon_D25_sp._str._219_sp._str._3133FAA
SPN120 Oscillibacter valericigenes_nov_94.344%	SPP17 Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibularis
SPN129 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_93.946%	SPP18 Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
SPN140 Prevotella stercorea_nov_97.403%	SPP19 Bifidobacterium adolescentis_faecale
SPN144 Ruminococcus callidus_nov_94.369%	SPP26 Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
SPN149 Eubacterium coprostanoligenes_nov_97.059%	SPP27 Gemella morbillorum_morbillum
SPN170 Kineothrix alysoides_nov_96.825%	SPP29 Blautia obeum_wexlerae
SPN182 Faecalibacterium prausnitzii_nov_97.964%	SPP39 Neisseria flava_macacae_mucosa_sicca_sp._TM101
SPN193 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_89.038%	SPP5 Actinomyces_Schaalia odontolyticus_sp._HMT_180
SPN194 Oscillibacter ruminantium_nov_93.919%	SPP54 Parasutterella_Sutterella excrementihominis_sp._str._cont1.66
SPN2 Alloprevotella rava_nov_92.857%	SPP56 Selenomonas sp._HMT_136_sp._Oral_Taxon_149
SPN205 Lachnospira pectinoschiza_nov_96.372%	SPP60 Veillonella rogosa_sp._str._NVG100cf
SPN216 Prevotella copri_nov_97.397%	SPP67 Bifidobacterium catenulatum_spudocatenulatum
SPN226 Lachnospira eligens_nov_94.785%	SPP68 Bacteroides koreensis_kribbi
SPN23 unclassified_Ruminococcaceae_sp._str._D16_nov_93.243%	SPP70 Veillonella dispar_parvula
SPN236 Eubacterium ruminantium_nov_97.279%	SPP71 Veillonella parvula_tobetsuensis
SPN246 Butyrivibrio sp._HMT_090_nov_92.795%	SPP74 Bacteroides_Phocaeicola plebeius
SPN249 Sporobacter termitidis_nov_90.562%	SPP79 Oribacterium parvum_sp._str._ACB1
SPN255 Paraprevotella clara_nov_97.609%	SPP8 Mogibacterium diversum_neglectum_vescum
SPN264 Prevotella colorans_nov_93.492%	SPP87 Actinomyces sp._HMT_169_sp._str._ChDCB197
SPN275 Sporobacter termitidis_nov_93.243%	SPP95 Streptococcus parasanguinis_parasanguinis_I_parasanguinis_clade_721_sp._HMT_066
SPN289 Christensenella massiliensis_nov_90.766%	SPPN1 Ruminococcus multispecies_sppn1_2_nov_95.023%
SPN301 Acutalibacter muris_nov_96.599%	SPPN106 Roseburia multispecies_sppn106_3_nov_96.833%
SPN322 Prevotellamassilia timonensis_nov_97.397%	SPPN2 Christensenella multispecies_sppn2_2_nov_91.216%
SPN334 Eubacterium coprostanoligenes_nov_92.777%	SPPN27 Clostridium multispecies_sppn27_2_nov_97.959%
SPN335 Kineothrix alysoides_nov_96.614%	SPPN34 Mobiluncus multispecies_sppn34_2_nov_97.817%
SPN346 Asteroleplasma anaerobium_nov_80.300%	SPPN46 multigenus multispecies_sppn46_2_nov_92.568%
SPN35 Succinivibrio dextrinosolvens_nov_97.079%	SPPN52 Roseburia multispecies_sppn52_2_nov_97.072%
SPN358 Eubacterium coprostanoligenes_nov_95.711%	SPPN68 Lachnospirillum multispecies_sppn68_2_nov_97.279%
SPN366 Prevotella oralis_nov_92.641%	