

SPN98 Streptococcus thermophilus	SPN76 Prevotella sp._HMT_305_nov_97.619%
SPN1 Kiliella Kiliella laminaria_nov_87.416%	SPN77 Oscillibacter ruminantium_nov_94.344%
SPN10 Oscillibacter ruminantium_nov_88.367%	SPN78 Alistipes indistinctus_nov_96.529%
SPN100 Breznakia pachnodae_nov_87.821%	SPN79 Bacteroidetes_[G-3] bacterium_HMT_365_nov_89.655%
SPN101 Eubacterium xylanophilum_nov_97.727%	SPN8 Coprococcus catus_nov_96.154%
SPN102 Ruminococcus champanellensis_nov_95.928%	SPN80 Anaerotruncus colihominis_nov_93.850%
SPN106 Alkalibaculum bacchi_nov_83.108%	SPN81 Acetivibrio clariflavus_nov_90.745%
SPN109 Acetanaerobacterium elongatum_nov_92.081%	SPN82 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_92.411%
SPN11 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_94.369%	SPN83 Calycanthus floridus_Oral_Taxon_D07_nov_97.297%
SPN116 Anaerocolumna cellulositytica_nov_96.833%	SPN84 Lachnospira eligens_nov_97.959%
SPN119 Oscillibacter ruminantium_nov_94.144%	SPN85 Bacteroides zoogeleoformans_nov_92.625%
SPN12 Agathobaculum desmolans_nov_97.279%	SPN86 Fournierella massiliensis_nov_95.455%
SPN120 Oscillibacter valericigenes_nov_94.344%	SPN87 Agathobaculum desmolans_nov_97.738%
SPN126 Erysipelotrichaceae_[G-1] bacterium_HMT_905_nov_92.625%	SPN88 Butyricoccus pullicaecorum_nov_94.382%
SPN129 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_93.946%	SPN89 Desulfovibrio sp._HMT_040_nov_92.521%
SPN13 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_92.777%	SPN9 Firmicutes_[G] sp._Oral_Taxon_A55_nov_91.880%
SPN134 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.011%	SPN90 Muribaculum intestinale_nov_90.043%
SPN14 unclassified_Ruminococcaceae sp._str._D16_nov_89.414%	SPN91 Hespelia porcina_nov_95.238%
SPN140 Prevotella stercora_nov_97.403%	SPN92 Breznakia pachnodae_nov_83.512%
SPN144 Ruminococcus callidus_nov_94.369%	SPN93 Desulfohalotomaculum tongense_nov_87.696%
SPN149 Eubacterium coprostanoligenes_nov_97.059%	SPN94 Prevotella copri_nov_97.174%
SPN15 Treponema denticola_nov_90.000%	SPN95 Christensenella massiliensis_nov_88.739%
SPN152 Lachnobacterium bovis_nov_95.711%	SPN96 Oscillibacter ruminantium_nov_92.601%
SPN16 Succinivibrio dextrinosolvens_nov_96.372%	SPN97 Coprococcus catus_nov_96.136%
SPN161 Mediterranea massiliensis_nov_92.641%	SPN98 Holdemanella biformis_nov_97.854%
SPN17 Eubacterium coprostanoligenes_nov_97.732%	SPN99 Ihubacter massiliensis_nov_96.606%
SPN170 Kineothrix alysoides_nov_96.825%	SPP1 Bacteroides sp._str._4136_uniformis
SPN172 Tyzzerella [Clostridium] colinum_nov_95.955%	SPP103 Bacteroides cellulosityticus_sp._str._VH2
SPN18 Anaeromassilibacillus senegalensis_nov_88.435%	SPP104 Anaerostipes_Lacnociostridium_unclassified_Lachnospiraceae_hadrus_sp._SSC/2_sp._str._cont1.81
SPN182 Faecalibacterium prausnitzii_nov_97.964%	SPP105 Lactobacillus_Ligilactobacillus_ruminis
SPN183 Lachnociostridium sp._str._SS21_nov_96.364%	SPP109 Neisseria flavescens_flavescens[subflava
SPN19 Kineothrix alysoides_nov_95.918%	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
SPN192 Sporobacter termitidis_nov_93.034%	SPP111 Neisseria flavescens[subflava_perflava
SPN193 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_89.038%	SPP113 Streptococcus sanguinis_sp._HMT_074
SPN194 Oscillibacter ruminantium_nov_93.919%	SPP115 Bacteroides_Phocaeicola_coprocola
SPN2 Alloprevotella rava_nov_92.857%	SPP117 Blautia_Mediterraneibacter [Ruminococcus] lactaris_lactaris
SPN20 Saccharofermentans acetigenes_nov_89.462%	SPP118 Atopobium_Lancefieldella parvula_parvulum
SPN203 Ruminoclostridium leptum_nov_95.011%	SPP12 Roseburia faecalis_faecis
SPN205 Lachnospira pectinoschiza_nov_96.372%	SPP127 Streptococcus parasanguinis_II_parasanguinis_clade_411_sp._HMT_057
SPN21 Parasporobacterium paucivorans_nov_95.455%	SPP134 Granulicatella adiacens_padiacens
SPN213 Flavonifractor plautii_nov_95.711%	SPP135 Actinomyces_Schaalia odontolytica_odontolyticus
SPN216 Prevotella copri_nov_97.397%	SPP136 Bacteroides sp._str._3140A_sp._str._4347FAA
SPN22 Oscillibacter ruminantium_nov_93.243%	SPP137 Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
SPN222 Anaerobacterium chartisolvans_nov_89.462%	SPP138 Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torques
SPN226 Lachnospira eligens_nov_94.785%	SPP14 Lachnociostridium sp._SM4/1_sp._str._M62/1
SPN23 unclassified_Ruminococcaceae sp._str._D16_nov_93.243%	SPP15 Bacteroides_Phocaeicola_dorei_sp._Oral_Taxon_D25_sp._str._219_sp._str._3133FAA
SPN236 Eubacterium ruminantium_nov_97.279%	SPP17 Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibularis
SPN24 Eubacterium coprostanoligenes_nov_95.270%	SPP18 Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
SPN241 Christensenella hongkongensis_nov_87.220%	SPP19 Bifidobacterium adolescentis_faecale
SPN246 Butyrivibrio sp._HMT_090_nov_92.795%	SPP2 Streptococcus gordonii_sp._str._2136FAA
SPN249 Sporobacter termitidis_nov_90.562%	SPP20 Porphyromonas catoniae_sp._HMT_284
SPN25 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_93.905%	SPP21 Porphyromonas catoniae_sp._HMT_275_sp._HMT_284
SPN255 Paraprevotella clara_nov_97.609%	SPP25 Streptococcus sanguinis_sp._Oral_Taxon_B66
SPN257 Adlercreutzia muris_nov_94.796%	SPP26 Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
SPN26 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_94.369%	SPP27 Gemella morbillorum_morbillum
SPN264 Prevotella colorans_nov_93.492%	SPP28 Bacteroides_Phocaeicola coprophilus
SPN266 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_91.216%	SPP29 Blautia obeum_wexlerae
SPN27 Dialister succinatiphilus_nov_95.931%	SPP3 Streptococcus anginosus_constellatus
SPN275 Sporobacter termitidis_nov_93.243%	SPP30 Prevotella histicola_melaninogenica
SPN276 Prevotella copri_nov_97.831%	SPP39 Neisseria flava_macacae_mucosa_sicca_sp._TM101
SPN28 Anaerovibrio lipolyticus_nov_93.548%	SPP4 Actinomyces_Schaalia odontolyticus_sp._HMT_180
SPN287 Clostridium sp._str._NML04A032_nov_92.793%	SPP41 Enterobacter asburiae_cancerogenus_cloacae_hormaechei
SPN288 Prevotella copri_nov_97.835%	SPP42 Enterobacter_Leclercia_Pantoea adecarboxylata_agglomerans_cloacae_ludwigii
SPN289 Christensenella massiliensis_nov_90.766%	SPP43 Bacteroides sp._str._XB1A_xylanisolvans
SPN29 Prevotella sp._HMT_396_nov_97.614%	SPP46 Fusobacterium naviforme_nucleatum_nucleatum_ss_vincentii_nucleatum_subsp._vincentii
SPN299 Acetivibrio ethanolignens_nov_95.909%	SPP47 Fusobacterium nucleatum_nucleatum_ss_nucleatum_nucleatum_subsp._nucleatum
SPN3 Clostridium sp._str._NML04A032_nov_88.814%	SPP48 Fusobacterium nucleatum_nucleatum_ss_nucleatum_nucleatum_subsp._nucleatum_simiae
SPN30 Fusicatenibacter saccharivorans_nov_96.599%	SPP49 Fusobacterium nucleatum_nucleatum_subsp._vincentii
SPN301 Acutalibacter muris_nov_96.599%	SPP5 Actinomyces_Schaalia odontolyticus_sp._HMT_180
SPN31 Ruminococcus bromii_nov_95.475%	SPP50 Fusobacterium nucleatum_nucleatum_ss_animalis