

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 Fusicatenibacter saccharivorans nov. 97.279%
SPN467 Saccharibacteria (TM7) [G-3] bacterium_HMT_351_nov_97.072%
SPN536 Lacrimispora 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 Lachnoclostridium 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_Lachnoclostridium_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_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._dentisani
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_paradiacens
SPP135 Actinomyces_Schaalia odontolytica_odontolyticus
SPP137 Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
SPP138 Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torques
SPP14 Lachnoclostridium sp._SM4/1_sp._str._M62/1
SPP15 Bacteroides_Phocaeicola_dorei_sp._Oral_Taxon_D25_sp._str._219_sp._str._3133FAA
SPP17 Streptococcus salivarius sp._str._ACS2_sp._str._C150_vestibularis
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
SPP56 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_Phocaeicola 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_721_sp._HMT_066
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 Lachnoclostridium multispecies_sppn68_2_nov_97.279%