

● SPN77 Oscillibacter ruminantium nov_94.344%
● SPN78 Alistipes indistinctus nov_96.529%
● SPN79 Bacteroidetes [G-3] bacterium HMT_365_nov_89.655%
● SPN8 Coprococcus catus nov_96.154%
● SPN80 Anaerotruncus colihominis nov_93.850%
● SPN81 Acetivibrio clariflavus nov_90.745%
● SPN82 Ruminococcaceae [G-2] bacterium HMT_085_nov_92.411%
● SPN83 Calycanthus floridus Oral_Taxon_D07_nov_97.297%
● SPN84 Lachnospira eligens nov_97.959%
● SPN85 Bacteroides zooglyphiformans nov_92.625%
● SPN86 Fournierella massiliensis nov_95.455%
● SPN87 Agathobaculum desmolaens nov_97.738%
● SPN88 Butyrivibrio pullicaecorum nov_94.382%
● SPN89 Desulfovibrio sp. HMT_040_nov_92.521%
● SPN9 Firmicutes [G] sp. Oral_Taxon_A55_nov_91.880%
● SPN90 Muribaculum intestinale nov_90.043%
● SPN91 Hespellia porcina nov_95.238%
● SPN92 Breznakia pachnoda nov_83.512%
● SPN93 Desulfohalotomaculum tongense nov_87.696%
● SPN94 Prevotella copri nov_97.174%
● SPN95 Christensenella massiliensis nov_88.739%
● SPN96 Oscillibacter ruminantium nov_92.601%
● SPN97 Coprococcus catus nov_96.136%
● SPN98 Holdemanella bififormis nov_97.854%
● SPN99 Lhubacter massiliensis nov_96.606%
● SPP1 Bacteroides sp. str. 4136_uniformis
● SPP103 Bacteroides cellulosilyticus sp. str. WH2
● SPP104 Anaerostipes Lachnospiraceae 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
● SPP111 Neisseria flavescens|subflava_perflava
● SPP113 Streptococcus sanguinis sp. HMT_074
● SPP115 Bacteroides Phocaeicola coprocola
● SPP117 Blautia Mediterraneibacter [Ruminococcus] lactaris_lactaris
● SPP118 Atopobium Lancefieldella parvula_parvulum
● SPP12 Roseburia faecalis faecis
● SPP127 Streptococcus parasanguinis II parasanguinis_clade_411_sp._HMT_057
● SPP134 Granulicatella adiacens paradiacens
● SPP135 Actinomyces Schaalia odontolytica odontolyticus
● SPP136 Bacteroides sp. str. 3140A sp. str. 4347FAA
● SPP137 Blautia Mediterraneibacter [Ruminococcus] gnavus_gnavus
● SPP138 Blautia unclassified_Lachnospiraceae sp. str. cont1.79_torques
● SPP14 Lachnospiraceae sp. str. SM4/1 sp. str. M62/I
● 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
● SPP2 Streptococcus gordonii sp. str. 2136FAA
● SPP20 Porphyromonas catoniae sp. HMT_284
● SPP21 Porphyromonas catoniae sp. HMT_275 sp. HMT_284
● SPP25 Streptococcus sanguinis sp. Oral_Taxon_B66
● SPP26 Escherichia Shigella coli dysenteriae_fergusonii_flexneri_sonnei
● SPP27 Gemella morbillorum moribillum
● SPP28 Bacteroides Phocaeicola coprophilus
● SPP29 Blautia obeum_wexlerae
● SPP3 Streptococcus anginosus_constellatus
● SPP30 Prevotella histicola_melaninogenica
● SPP39 Neisseria flava_mucosae_mucosa_sicca_sp._TM101
● SPP4 Actinomyces Schaalia odontolyticus sp. HMT_180
● SPP41 Enterobacter asburiae_cancerogenus_cloacae_hormaechei
● SPP42 Enterobacter Leclercia_Pantoea_adeocarboxylata_agglomerans_cloacae_ludwigii
● SPP43 Bacteroides sp. str. XB1A_xylanisolvens
● SPP46 Fusobacterium naviforme_nucleatum_nucleatum_ss_vincentii_nucleatum_subsp._vincentii
● SPP47 Fusobacterium nucleatum_nucleatum_ss_nucleatum_nucleatum_subsp._nucleatum
● SPP48 Fusobacterium nucleatum_nucleatum_ss_nucleatum_nucleatum_subsp._nucleatum_simiae
● SPP49 Fusobacterium nucleatum_nucleatum_subsp._vincentii
● SPP5 Actinomyces Schaalia odontolyticus sp. HMT_180