Species

SP236 Campylobacter concisus
SP238 Parabacteroides distasonis
SP239 Prevotella sp._HMT_443
SP24 Ruminococcus callidus
SP240 Parabacteroides merdae
SP241 Actinomyces graevenitzii
SP242 Mycoplasma orale
SP243 Alistipes putredinis
SP248 Porphyromonas endodontalis
SP249 Veillonellaceae_[G−1] bacterium_HMT_135
SP25 Rothia mucilaginosa
SP250 Selenomonas sputigena
SP251 Bifidobacterium dentium
SP252 Porphyromonas sp._HMT_285
SP254 Mediterraneibacter faecis
SP255 Slackia exigua
SP256 Lachnoanaerobaculum orale
SP258 Selenomonas sp._HMT_136
SP259 Lactobacillus rogosae
SP26 Eubacterium rectale
SP260 Bacteroidetes_[G−3] bacterium_HMT_365
SP261 Saccharibacteria_(TM7)_[G−4] bacterium_HMT_355
SP262 Bacteroidetes_[G−3] bacterium_HMT_280
SP263 Romboutsia timonensis
SP266 Coprococcus eutactus
SP267 Treponema sp._HMT_237
SP268 Capnocytophaga gingivalis
SP269 Ruminococcus champanellensis
SP27 Prevotella copri
SP270 Bilophila wadsworthia
SP272 Prevotella sp._HMT_313
SP273 Dorea formicigenerans
SP274 Mollicutes_[G−2] bacterium_HMT_906
SP275 Gemella haemolysans
SP279 Treponema socranskii
SP28 Blautia luti
SP280 Corynebacterium durum
SP283 Stomatobaculum sp._HMT_097
SP284 Veillonellaceae_[G−1] bacterium_HMT_129
SP288 Alistipes shahii
SP29 Lachnospira eligens
SP295 Lacnoclostridium sp._SS3/4
SP296 Stomatobaculum longum
SP297 Coprococcus comes
SP30 Prevotella pallens
SP301 Ruminococcus bromii
SP303 Mitsuokella jalaludinii
SP304 Prevotella sp._HMT_304
SP305 Slackia isoflavoniconvertens
SP306 Kineothrix alysoides
SP307 Roseburia inulinivorans
SP308 Porphyromonas gingivalis
SP309 Catenibacterium mitsuokai
SP310 Prevotella sp._HMT_305
SP312 Schaalia cardiffensis
SP313 Alloprevotella tannerae
SP318 Catonella morbi
SP321 Firmicutes_[G] sp._Oral_Taxon_A55
SP322 Porphyromonas sp._Oral_Taxon_B43
SP323 Capnocytophaga leadbetteri
SP325 Peptostreptococcaceae_[XI][G−4] bacterium_HMT_369
SP326 Mollicutes_[G−1] bacterium_HMT_504
SP327 Peptostreptococcaceae_[XI][G−5] [Eubacterium]_saphenum
SP328 Alloscardovia omnicolens
SP329 Absconditabacteria_(SR1)_[G−1] bacterium_HMT_874
SP330 Adlercreutzia equolifaciens
SP332 Treponema sp._HMT_257
SP333 Prevotella sp._HMT_306
SP334 Veillonella parvula
SP338 Eggerthella sp._str._1356FAA
SP98 Streptococcus thermophilus
SPN1 Kiloniella Kiliella laminaria_nov_87.416%
SPN10 Oscillibacter ruminantium_nov_88.367%
SPN100 Breznakia pachnodae_nov_87.821%
SPN101 Eubacterium xylanophilum_nov_97.727%
SPN102 Ruminococcus champanellensis_nov_95.928%
SPN106 Alkalibaculum bacchi_nov_83.108%
SPN109 Acetanaerobacterium elongatum_nov_92.081%
SPN11 Ruminococcaceae_[G−2] bacterium_HMT_085_nov_94.369%
SPN116 Anaerocolumna cellulosilytica_nov_96.833%
SPN119 Oscillibacter ruminantium_nov_94.144%
SPN12 Agathobaculum desmolans_nov_97.279%
SPN120 Oscillibacter valericigenes_nov_94.344%
SPN126 Erysipelotrichaceae_[G−1] bacterium_HMT_905_nov_92.625%
SPN129 Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.946%
SPN13 Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_92.777%
SPN134 Ruminococcaceae_[G−2] bacterium_HMT_085_nov_91.011%
SPN14 unclassified_Ruminococcaceae sp._str._D16_nov_89.414%
SPN140 Prevotella stercorea_nov_97.403%
SPN144 Ruminococcus callidus_nov_94.369%
SPN149 Eubacterium coprostanoligenes_nov_97.059%
SPN15 Treponema denticola_nov_90.000%
SPN152 Lachnobacterium bovis_nov_95.711%
SPN16 Succinivibrio dextrinosolvens_nov_96.372%
SPN161 Mediterranea massiliensis_nov_92.641%
SPN17 Eubacterium coprostanoligenes_nov_97.732%
SPN170 Kineothrix alysoides_nov_96.825%
SPN172 Tyzzerella [Clostridium] colinum_nov_95.955%
SPN18 Anaeromassilibacillus senegalensis_nov_88.435%
SPN182 Faecalibacterium prausnitzii_nov_97.964%
SPN183 Lacnoclostridium sp._str._SS2/1_nov_96.364%
SPN19 Kineothrix alysoides_nov_95.918%
SPN192 Sporobacter termitidis_nov_93.034%
SPN193 Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
SPN194 Oscillibacter ruminantium_nov_93.919%
SPN2 Alloprevotella rava_nov_92.857%
SPN20 Saccharofermentans acetigenes_nov_89.462%
SPN203 Ruminoclostridium leptum_nov_95.011%
SPN205 Lachnospira pectinoschiza_nov_96.372%
SPN21 Parasporobacterium paucivorans_nov_95.455%
SPN213 Flavonifractor plautii_nov_95.711%
SPN216 Prevotella copri_nov_97.397%
SPN22 Oscillibacter ruminantium_nov_93.243%
SPN222 Anaerobacterium chartisolvens_nov_89.462%
SPN226 Lachnospira eligens_nov_94.785%
SPN23 unclassified_Ruminococcaceae sp._str._D16_nov_93.243%
SPN236 Eubacterium ruminantium_nov_97.279%
SPN24 Eubacterium coprostanoligenes_nov_95.270%
SPN241 Christensenella hongkongensis_nov_87.220%
SPN246 Butyrivibrio sp._HMT_090_nov_92.795%
SPN249 Sporobacter termitidis_nov_90.562%
SPN25 Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.905%
SPN255 Paraprevotella clara_nov_97.609%
SPN257 Adlercreutzia muris_nov_94.796%
SPN26 Ruminococcaceae_[G−2] bacterium_HMT_085_nov_94.369%
SPN264 Prevotella colorans_nov_93.492%
SPN266 Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_91.216%
SPN27 Dialister succinatiphilus_nov_95.931%
SPN275 Sporobacter termitidis_nov_93.243%
SPN276 Prevotella copri_nov_97.831%
SPN28 Anaerovibrio lipolyticus_nov_93.548%
SPN287 Clostridium sp._str._NML04A032_nov_92.793%
SPN288 Prevotella copri_nov_97.835%
SPN289 Christensenella massiliensis_nov_90.766%
SPN29 Prevotella sp._HMT_396_nov_97.614%
SPN299 Acetivibrio ethanolgignens_nov_95.909%
SPN3 Clostridium sp._str._NML04A032_nov_88.814%
SPN30 Fusicatenibacter saccharivorans_nov_96.599%
SPN301 Acutalibacter muris_nov_96.599%
SPN31 Ruminococcus bromii_nov_95.475%
SPN76 Prevotella sp._HMT_305_nov_97.619%
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 zoogleoformans_nov_92.625%
SPN86 Fournierella massiliensis_nov_95.455%
SPN87 Agathobaculum desmolans_nov_97.738%
SPN88 Butyricicoccus 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 pachnodae_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 biformis_nov_97.854%
SPN99 Ihubacter massiliensis_nov_96.606%
SPP1 Bacteroides sp._str._4136_uniformis
SPP103 Bacteroides cellulosilyticus_sp._str._WH2
SPP104 Anaerostipes_Lacnoclostridium_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_ ...(20 species)
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 Lacnoclostridium sp._SM4/1_sp._str._M62/1
SPP15 Bacteroides_Phocaeicola dorei_sp._Oral_Taxon_D25_sp._str._219_sp._str._313 ...(4 species)
SPP17 Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibulari ...(4 species)
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_macacae_mucosa_sicca_sp._TM101
SPP4 Actinomyces_Schaalia odontolyticus_sp._HMT_180
SPP41 Enterobacter asburiae_cancerogenus_cloacae_hormaechei
SPP42 Enterobacter_Leclercia_Pantoea adecarboxylata_agglomerans_cloacae_ludwigii
SPP43 Bacteroides sp._str._XB1A_xylanisolvens
SPP46 Fusobacterium naviforme_nucleatum_nucleatum_ss_vincentii_nucleat ...(4 species)
SPP47 Fusobacterium nucleatum_nucleatum_ss_nucleatum_nucleatum_subsp._nucleatum
SPP48 Fusobacterium nucleatum_nucleatum_ss_nucleatum_nucleatum_subsp._ ...(4 species)
SPP49 Fusobacterium nucleatum_nucleatum_subsp._vincentii
SPP5 Actinomyces_Schaalia odontolyticus_sp._HMT_180
SPP50 Fusobacterium nucleatum_nucleatum_ss_animalis