

Species

●	SP25 Eubacteriales_[G-4] bacterium_MOT-164
●	SP253 Mesosutterella multiformis
●	SP254 Parabacteroides johnsonii
●	SP255 Methanobrevibacter smithii
●	SP258 Fusicatenibacter saccharivorans
●	SP259 Megasphaera massiliensis
●	SP260 Phocaeicola faecicola
●	SP269 Ligilactobacillus ruminis
●	SP275 Bacteroides fragilis
●	SP276 Bacteroides intestinalis
●	SP279 Blautia luti
●	SP282 Clostridium chartatabidum
●	SP283 Lachnospira pectinoschiza
●	SP29 Catenibacterium mitsuokai
●	SP290 Haemophilus parainfluenzae
●	SP291 Blautia obeum
●	SP293 Eubacterium coprostanoligenes
●	SP294 Ruminococcus flavefaciens
●	SP295 Mediterraneibacter [Ruminococcus] torques
●	SP296 Hungatella xylanolytica
●	SP298 Anaerotruncus colihominis
●	SP31 Ruminococcus champanellensis
●	SP32 Alistipes onderdonkii
●	SP33 Blautia wexlerae
●	SP338 Bacteroides caccae
●	SP34 Butyricimonas virosa
●	SP343 Holdemania filiformis
●	SP344 Alistipes indistinctus
●	SP345 Duodenibacillus massiliensis
●	SP351 Sutterella wadsworthensis
●	SP352 Alistipes communis
●	SP37 Coprococcus catus
●	SP38 Akkermansia muciniphila
●	SP380 Frisingioccocus caecimuris
●	SP385 Ruminococcus bromii
●	SP387 Alistipes putredinis
●	SP389 Flavonifractor plautii
●	SP39 Anaerostipes hadrus
●	SP390 Kineothrix alysoides
●	SP391 Eubacterium ventriosum

●	SPN162 Pseudoprevotella muciniphila_nov_87.215%
●	SPN164 Paraprobacterium paucivorans_nov_96.279%
●	SPN165 Acetitomaculum ruminis_nov_97.209%
●	SPN166 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_92.558%
●	SPN167 Ihubacter massiliensis_nov_93.925%
●	SPN168 Muribaculaceae_[G-2] bacterium_MOT-104_nov_97.664%
●	SPN169 Tepidibaculum saccharolyticum_nov_86.512%
●	SPN170 Caproicibacterium amylolyticum_nov_83.645%
●	SPN171 Eubacterium ventriosum_nov_97.209%
●	SPN172 Sporobacter termitidis_nov_93.488%
●	SPN174 Muribaculum intestinale_nov_85.116%
●	SPN175 Prevotella stercorea_nov_97.209%
●	SPN176 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_94.931%
●	SPN177 Acetivibrio straminisolvans_nov_93.488%
●	SPN178 Lutispora thermophila_nov_91.628%
●	SPN179 Aestuariaispira insulae_nov_87.037%
●	SPN18 Mollicutes_[G-2] bacterium_MOT-187_nov_86.916%
●	SPN180 Eubacteriales_[G-1] bacterium_MOT-144_nov_93.488%
●	SPN182 Peptococcus simiae_nov_89.302%
●	SPN183 Coprobacter fastidiosus_nov_88.940%
●	SPN184 Aestuariaispira insulae_nov_85.000%
●	SPN185 Paludicola psychrotolerans_nov_93.925%
●	SPN186 Faecalibacterium prausnitzii_nov_85.922%
●	SPN187 Mollicutes_[G-2] bacterium_MOT-187_nov_85.981%
●	SPN188 Caproicibacterium amylolyticum_nov_86.512%
●	SPN189 Peptoniphilaceae_[G-3] bacterium_HMT_929_nov_81.395%
●	SPN190 Peptostreptococcaceae_[G-5] bacterium_HMT_493_nov_93.953%
●	SPN191 Prevotella stercorea_nov_97.674%
●	SPN192 Muribaculum intestinale_nov_88.073%
●	SPN193 Coprococcus eutactus_nov_96.279%
●	SPN194 Luteimonas terricola_nov_80.930%
●	SPN195 Fournierella massiliensis_nov_95.349%
●	SPN196 Dysosmobacter welbionis_nov_95.349%
●	SPN197 Ruminococcus callidus_nov_87.442%
●	SPN198 Harryflintia acetispora_nov_96.262%
●	SPN199 Holdemania biformis_nov_97.209%
●	SPN200 Faecalicoccus acidiformans_nov_95.327%
●	SPN201 Pacificispira spongicola_nov_84.862%
●	SPN202 Oscillibacter ruminantium_nov_94.884%
●	SPN203 Sporobacter termitidis_nov_92.991%

●	SPN851 Anaerobutyricum hallii_nov_97.674%
●	SPN858 Eubacterium coprostanoligenes_nov_93.981%
●	SPN859 Eubacterium ramulus_nov_97.642%
●	SPN86 Duodenibacillus massiliensis_nov_97.209%
●	SPN868 Ruminococcus flavefaciens_nov_85.648%
●	SPN869 Dysosmobacter welbionis_nov_93.953%
●	SPN87 Lawsonibacter asaccharolyticus_nov_96.279%
●	SPN870 Papillibacter cinnamivorans_nov_92.593%
●	SPN877 Kineosporia aurantiaca_nov_81.106%
●	SPN88 Phocaeicola coprocola_nov_94.884%
●	SPN881 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.636%
●	SPN887 Mollicutes_[G-1] bacterium_MOT-186_nov_91.204%
●	SPN89 Kiloniella antarctica_nov_87.097%
●	SPN890 Hungatella effluvii_nov_95.349%
●	SPN900 Eubacteriales_[G-3] bacterium_MOT-163_nov_94.419%
●	SPN904 Ruminiclostridium cellobioparum_nov_89.352%
●	SPN91 Oscillospiraceae_[G-7] bacterium_MOT-154_nov_93.023%
●	SPN92 Hungatella effluvii_nov_96.744%
●	SPN93 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_83.333%
●	SPN94 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.931%
●	SPN95 Lachnospira multipara_nov_90.741%
●	SPN96 Hungatella effluvii_nov_96.744%
●	SPN97 Papillibacter cinnamivorans_nov_91.163%
●	SPN98 Mollicutes_[G-2] bacterium_MOT-188_nov_90.099%
●	SPN99 Oscillibacter ruminantium_nov_97.209%
●	SPP11 Bacteroides faecichinchillae_faecis_thetaiotaomicon
●	SPP13 Bifidobacterium breve_cebidarum_longum
●	SPP15 Lactocaseibacillus casei_rhamnosus
●	SPP2 Escherichia_Pseudesccherichia_Shigella_boydii_coli_fergusonii_flexneri_marmotae_sonnei_vu ...(7 species)
●	SPP20 Butyricimonas faecihominis_paraviroa
●	SPP21 Streptococcus hongkongensis_salivarius_thermophilus_vestibularis
●	SPP27 Faecalibacillus faecis_intestinalis
●	SPP3 Sutterella massiliensis_stercoricanis
●	SPP31 Eubacteriales_[G-4] bacterium_MOT-164_bacterium_MOT-165
●	SPP32 Streptococcus australis_koreensis_parasanguinis_parasanguinis_cl ...(8 species)
●	SPP33 Bifidobacterium adolescentis_faecale
●	SPP35 Buttiiauxella_Enterobacter_Erwinia_Klebsiella_Kluyv ...(10 species) adecarboxylata_aerogenes_agglomerans_agrestis_amni ...(30 species)
●	SPP37 Veillonella atypica_dispar
●	SPP39 Fusobacterium ulcerans_varium
●	SPP45 Bacteroides koreensis_kribbi