

Species

● SP10 Bacteroides intestinalis	● SP8 Romboutsia ilealis	● SPN423 Pseudoflavonifractor phocaeensis_nov_94.191%
● SP107 Bacteroides thetaiotaomicron	● SP80 Parabacteroides goldsteinii	● SPN43 Pseudoflavonifractor phocaeensis_nov_92.340%
● SP109 Atopostipes sp._MOT-200	● SP83 Staphylococcus haemolyticus	● SPN434 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.116%
● SP11 Haemophilus parainfluenzae	● SP87 Akkermansia muciniphila	● SPN447 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.154%
● SP115 Kosakonia sacchari	● SP91 Adlercreutzia equolifaciens	● SPN456 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_90.702%
● SP119 Corynebacterium marinum	● SP93 Monoglobus pectinilyticus	● SPN467 Oscillibacter valericigenes_nov_94.595%
● SP12 Clostridium disporicum	● SP99 Bacteroides uniformis	● SPN47 Coprococcus eutactus_nov_95.085%
● SP126 Microbacterium hydrocarbonoxydans	● SPN1 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.154%	● SPN478 Lachnospiraceae_[G-4] bacterium_MOT-169_nov_97.065%
● SP129 Corynebacterium amycolatum	● SPN107 Niveispirillum fermenti_nov_85.242%	● SPN479 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.340%
● SP13 Enterococcus avium	● SPN110 Rothia nasimurium_nov_97.228%	● SPN489 Mediterraneibacter[Ruminococcus] gnavus_nov_94.990%
● SP138 Odoribacter laneus	● SPN12 Streptococcus sp._MOT-045_nov_97.764%	● SPN498 Lachnospiraceae_[G-8] bacterium_MOT-173_nov_89.189%
● SP14 Collinsella aerofaciens	● SPN120 Anaerotignum aminivorans_nov_92.812%	● SPN509 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.304%
● SP147 Erysipelatoclostridium[Clostridium] saccharogumia	● SPN13 Kiloniella majae_nov_85.144%	● SPN519 Caproiciproducens galactitolivorans_nov_91.083%
● SP15 Phocaeicola vulgatus	● SPN132 Blautia luti_nov_96.008%	● SPN523 Arthrosira platensis_nov_90.531%
● SP17 Enterobacter cancerogenus	● SPN133 Frisingicoccus caecimuris_nov_96.451%	● SPN531 Lachnospira multipara_nov_84.298%
● SP18 Shigella sonnei	● SPN144 Blautia glucerasea_nov_97.899%	● SPN541 Lacrimispora xylanolytica_nov_96.436%
● SP19 Shigella flexneri	● SPN156 Eubacterium xylanophilum_nov_91.313%	● SPN551 Propioniferax innocua_nov_95.207%
● SP22 Escherichia coli	● SPN164 Turcibacter sanguinis_nov_96.137%	● SPN562 Blautia hydrogenotrophica_nov_93.582%
● SP23 Ligilactobacillus murinus	● SPN168 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.130%	● SPN572 Oscillospiraceae_[G-7] bacterium_MOT-154_nov_94.612%
● SP25 Blautia glucerasea	● SPN180 Eubacterium coprostanoligenes_nov_91.379%	● SPN580 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.228%
● SP27 Cutibacterium acnes	● SPN202 Coprococcus eutactus_nov_92.704%	● SPN581 Faecalicatena fissicatena_nov_96.042%
● SP31 Granulicatella adiacens	● SPN213 Agathobaculum desmolans_nov_86.345%	● SPN582 Eubacterium coprostanoligenes_nov_91.845%
● SP36 Bacteroides xylanisolvens	● SPN218 Butyricicoccus pullicaecorum_nov_87.216%	● SPN583 Lachnospiraceae_[G-1] bacterium_MOT-166_nov_93.983%
● SP37 Bifidobacterium pseudolongum	● SPN226 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_97.831%	● SPN584 Lacrimispora saccharolytica_nov_90.625%
● SP39 Rothia kristinae	● SPN238 Longibaculum muris_nov_92.754%	● SPN585 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_96.646%
● SP4 Longicatena caecimuris	● SPN25 Glucerbacter canis_nov_95.388%	● SPN586 Glucerbacter canis_nov_93.946%
● SP40 Geobacillus stearothermophilus	● SPN250 Blautia faecicola_nov_93.111%	● SPN588 Eisenbergiella massiliensis_nov_86.486%
● SP41 Lactobacillus intestinalis	● SPN252 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_90.496%	● SPN589 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_88.017%
● SP42 Shigella dysenteriae	● SPN264 Lachnoclostridium[Clostridium] herbivorans_nov_92.723%	● SPN59 Blautia schinkii_nov_94.340%
● SP44 Rothia aerolata	● SPN275 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_93.750%	● SPN590 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.347%
● SP45 Eisenbergiella massiliensis	● SPN284 Ruminococcus albus_nov_91.045%	● SPN591 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_92.500%
● SP46 Triticum aestivum	● SPN286 Lawsonibacter asaccharolyticus_nov_90.909%	● SPN592 Duncaniella freteri_nov_89.024%
● SP5 Staphylococcus ureilyticus	● SPN298 Acutalibacter muris_nov_95.661%	● SPN593 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_96.226%
● SP51 Actinidia eriantha	● SPN309 Barnesiella viscericola_nov_90.447%	● SPN603 Alistipes finegoldii_nov_94.835%
● SP53 Fusicatenibacter saccharivorans	● SPN321 Fusicatenibacter saccharivorans_nov_89.474%	● SPN616 Blautia producta_nov_96.017%
● SP54 Parasutterella excrementihominis	● SPN334 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_95.218%	● SPN627 Ruminococcus albus_nov_92.225%
● SP55 Ruthenibacterium lactatiformans	● SPN343 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.855%	● SPN639 Chryseobacterium culicis_nov_97.280%
● SP57 Stenotrophomonas pavanii	● SPN355 Erysipelatoclostridium[Clostridium] saccharogumia_nov_96.042%	● SPN71 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.473%
● SP58 Streptococcus agalactiae	● SPN36 Anaerostipes hadrus_nov_92.754%	● SPN83 Blautia producta_nov_95.178%
● SP61 Veillonella parvula	● SPN364 Fusicatenibacter saccharivorans_nov_97.479%	● SPN95 Sphingobacter marinum_nov_80.142%
● SP64 Enterococcus faecalis	● SPN365 Faecalimonas umblicata_nov_96.033%	● SPP4 Staphylococcus argenteus_aureus_roterodami
● SP66 Bilophia wadsworthia	● SPN377 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.929%	● SPP5 Streptococcus parasanguinis_parasanguinis_clade_721
● SP69 Streptococcus sp._HMT_064	● SPN388 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.946%	● SPP7 Veillonella dispar_parvula
● SP7 Streptococcus mutans	● SPN400 Anaerotruncus rubiinfantis_nov_88.820%	● SPP8 Staphylococcus saprophyticus_xylosus
● SP75 Mammalicoccus sciuri	● SPN405 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.097%	● SPPN3 Kordiimonas multispecies_sppn3_2_nov_79.121%
● SP79 Alistipes finegoldii	● SPN411 Zhongshania marina_nov_79.765%	