

Species

●	SPN109 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.555%
●	SPN11 Eubacteriales_[G-4] bacterium_MOT-164_nov_94.904%
●	SPN110 Robinsoniella peoriensis_nov_94.363%
●	SPN111 Mediterraneibacter [Ruminococcus] gnavus_nov_93.528%
●	SPN112 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.375%
●	SPN113 Anaerostipes hadrus_nov_87.578%
●	SPN114 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_97.077%
●	SPN115 Agathobaculum desmolans_nov_91.649%
●	SPN116 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_94.200%
●	SPN117 Blautia producta_nov_95.407%
●	SPN118 Christensenella hongkongensis_nov_86.275%
●	SPN119 Magnetovibrio blakemorei_nov_83.146%
●	SPN12 Anaerotignum aminivorans_nov_92.405%
●	SPN120 Oscillibacter valericigenes_nov_93.595%
●	SPN121 Roseburia faecis_nov_94.167%
●	SPN122 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.358%
●	SPN123 Parvibacter caecicola_nov_87.124%
●	SPN124 Eubacterium ramulus_nov_87.265%
●	SPN125 Ihubacter massiliensis_nov_90.644%
●	SPN126 Phoceia massiliensis_nov_90.426%
●	SPN127 Paludicola psychrotolerans_nov_87.759%
●	SPN128 Alistipes senegalensis_nov_93.443%
●	SPN129 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.132%
●	SPN13 Duncaniella freteri_nov_89.069%
●	SPN130 Emergencia timonensis_nov_90.188%
●	SPN131 Duncaniella freteri_nov_85.685%
●	SPN135 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.640%
●	SPN14 Marvinbryantia formatexigens_nov_91.649%
●	SPN141 Lachnoclostridium [Clostridium] scindens_nov_87.942%
●	SPN146 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.305%
●	SPN15 Mediterraneibacter [Ruminococcus] torques_nov_94.969%
●	SPN153 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_92.713%
●	SPN157 Lachnoclostridium [Clostridium] herbivorans_nov_92.917%
●	SPN16 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.133%
●	SPN165 Prevotella shahii_nov_87.602%
●	SPN168 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.528%
●	SPN17 Kineothrix alysoides_nov_87.474%
●	SPN177 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_92.793%
●	SPN179 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_92.693%
●	SPN18 Lachnospira eligens_nov_90.021%

●	SPN326 Lachnospiraceae_[Clostridium] aminophilum_nov_92.276%
●	SPN327 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.781%
●	SPN328 Lachnoclostridium [Clostridium] aminophilum_nov_87.318%
●	SPN329 Duncaniella freteri_nov_86.061%
●	SPN33 Adlercreutzia caecimuris_nov_92.688%
●	SPN330 Parasutterella excrementihominis_nov_94.578%
●	SPN331 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.718%
●	SPN332 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_90.586%
●	SPN333 Mailhella massiliensis_nov_90.244%
●	SPN334 Lachnoclostridium phytofermentans_nov_91.268%
●	SPN335 Lacrimispora xylanolytica_nov_92.276%
●	SPN336 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.489%
●	SPN337 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_87.083%
●	SPN338 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
●	SPN339 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.833%
●	SPN34 Eubacteriales_[G-1] bacterium_MOT-161_nov_93.843%
●	SPN340 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.442%
●	SPN341 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.347%
●	SPN342 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.770%
●	SPN343 Gordonibacter urolithinfaciens_nov_89.633%
●	SPN344 Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.816%
●	SPN345 Anaerotignum aminivorans_nov_93.008%
●	SPN346 Eubacteriales_[G-1] bacterium_MOT-159_nov_94.268%
●	SPN347 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_89.613%
●	SPN348 Hathewayia proteolytica_nov_83.514%
●	SPN349 Roseburia hominis_nov_92.547%
●	SPN35 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.466%
●	SPN350 Oribacterium parvum_nov_89.979%
●	SPN351 Muribaculaceae_[G-2] bacterium_MOT-104_nov_87.800%
●	SPN352 Lacrimispora xylanolytica_nov_93.789%
●	SPN353 Duncaniella freteri_nov_86.061%
●	SPN354 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_94.606%
●	SPN355 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.956%
●	SPN356 Enterocloster bolteae_nov_95.388%
●	SPN357 Lacrimispora saccharolytica_nov_93.082%
●	SPN358 Duncaniella freteri_nov_89.113%
●	SPN359 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.451%
●	SPN36 Eubacteriales_[G-1] bacterium_MOT-161_nov_97.868%
●	SPN360 Eisenbergiella massiliensis_nov_85.655%
●	SPN361 Anaerotruncus rubiinfantis_nov_92.708%

●	SPN423 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.754%
●	SPN424 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.102%
●	SPN43 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.347%
●	SPN44 Eubacteriales_[G-1] bacterium_MOT-159_nov_92.389%
●	SPN45 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.628%
●	SPN46 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.572%
●	SPN47 Duncaniella freteri_nov_87.375%
●	SPN48 Roseburia inulinivorans_nov_93.776%
●	SPN49 Mollicutes_[G-2] bacterium_MOT-187_nov_82.890%
●	SPN5 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_93.528%
●	SPN50 Kineothrix alysoides_nov_87.885%
●	SPN51 Erysipelatoclostridium [Clostridium] innocuum_nov_88.270%
●	SPN52 Hydrogenoanaerobacterium saccharovorans_nov_88.773%
●	SPN53 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_93.348%
●	SPN54 Hydrogenoanaerobacterium saccharovorans_nov_88.589%
●	SPN55 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.182%
●	SPN56 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.451%
●	SPN57 Pseudoflavonifractor phocaensis_nov_93.776%
●	SPN58 Duncaniella freteri_nov_91.002%
●	SPN59 Duncaniella freteri_nov_86.028%
●	SPN6 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.711%
●	SPN60 Drancourtella massiliensis_nov_90.229%
●	SPN61 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.634%
●	SPN62 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.200%
●	SPN63 Blautia faecicola_nov_91.170%
●	SPN64 Peptostreptococcaceae_[G-5] bacterium_HMT_493_nov_90.000%
●	SPN65 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.992%
●	SPN66 Faecalimonas umbilicata_nov_95.625%
●	SPN67 Adlercreutzia muris_nov_91.845%
●	SPN68 Duncaniella freteri_nov_86.747%
●	SPN69 Hathewayia proteolytica_nov_83.297%
●	SPN7 Butyrivacter intestini_nov_90.265%
●	SPN70 Duncaniella freteri_nov_93.293%
●	SPN71 Anaerotignum lactatifermentans_nov_95.464%
●	SPN72 Alloprevotella sp._HMT_473_nov_89.431%
●	SPN73 Neglectibacter timonensis_nov_95.132%
●	SPN74 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_85.773%
●	SPN75 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.125%
●	SPN76 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.351%
●	SPN77 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_87.657%