

● SPN109 Oscillospiraceae [G-4] bacterium_MOT-151_nov_93.553%
● 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 Phocaea 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 Lachnospiraceae [G-3] bacterium_MOT-168_nov_93.528%

● SPN326 Lachnoclostridium [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 bolteaee_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 Anaerotignum urolithinfaciens_nov_89.708%

● SPN423 Oscillospiraceae [G-3] bacterium_MOT-150_nov_92.734%
● 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 phocaeensis_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_90.0
● 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 Butyribacter 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_93.657%