cies

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 Phocea 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%
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 Hathewaya 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%
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 Hathewaya 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%