

- SP7 Oscillospiraceae [G-6] bacterium_MOT-153
- SP8 Lactobacillus johnsonii
- SP9 Eubacteriales [G-4] bacterium_MOT-164
- SPN10 Muribaculaceae [G-1] bacterium_MOT-129_nov_85.887%
- SPN100 Eubacteriales [G-1] bacterium_MOT-161_nov_93.843%
- SPN101 Eubacteriales [G-4] bacterium_MOT-164_nov_97.228%
- SPN102 Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.188%
- SPN103 Oscillospiraceae [G-2] bacterium_MOT-149_nov_94.154%
- SPN104 Lachnospiraceae [G-11] bacterium_MOT-177_nov_96.066%
- SPN105 Lachnospiraceae [G-3] bacterium_MOT-168_nov_89.353%
- SPN106 Oscillospiraceae [G-4] bacterium_MOT-151_nov_92.931%
- SPN107 Lachnospiraceae [G-7] bacterium_MOT-172_nov_91.718%
- SPN108 Acetivibrio cellulolyticus_nov_83.153%
- SPN109 Eisenbergiella massiliensis_nov_87.578%
- SPN11 Lachnospiraceae [G-12] bacterium_MOT-180_nov_91.093%
- SPN110 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.528%
- SPN111 Oscillospiraceae [G-2] bacterium_MOT-149_nov_85.773%
- SPN112 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.776%
- SPN113 Lachnospiraceae [G-3] bacterium_MOT-168_nov_94.572%
- SPN114 Eisenbergiella massiliensis_nov_88.174%
- SPN115 Acutalibacter muris_nov_94.227%
- SPN116 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.644%
- SPN117 Anaerotignum faecicola_nov_85.287%
- SPN118 Acetatifactor muris_nov_95.652%
- SPN119 Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.971%
- SPN12 Lachnoclostridium phytofermentans_nov_91.458%
- SPN120 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.351%
- SPN121 Oscillospiraceae [G-2] bacterium_MOT-149_nov_92.484%
- SPN122 Oscillospiraceae [G-4] bacterium_MOT-151_nov_96.050%
- SPN123 Oscillibacter valericigenes_nov_96.042%
- SPN124 Lacrimispora xylanolytica_nov_94.363%
- SPN125 Lachnospiraceae [G-13] bacterium_MOT-181_nov_91.189%
- SPN126 Blautia producta_nov_95.833%
- SPN127 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_88.820%
- SPN128 Oscillospiraceae [G-3] bacterium_MOT-150_nov_91.925%
- SPN129 Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.466%
- SPN130 Kineothrix alysoides_nov_86.848%
- SPN131 Oscillospiraceae [G-4] bacterium_MOT-151_nov_96.875%
- SPN132 Oscillibacter valericigenes_nov_93.595%

- SPN151 Flavonifractor plautii_nov_92.516%
- SPN152 Phoceia massiliensis_nov_88.034%
- SPN153 Clostridiales [F-1][G-1] bacterium_HMT_093_nov_91.775%
- SPN154 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.375%
- SPN155 Roseburia hominis_nov_91.476%
- SPN156 Lachnospiraceae [G-6] bacterium_MOT-171_nov_92.484%
- SPN157 Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.851%
- SPN158 Oscillospiraceae [G-2] bacterium_MOT-149_nov_94.363%
- SPN159 Eubacteriales [G-1] bacterium_MOT-159_nov_92.161%
- SPN160 Paludicola psychrotolerans_nov_87.759%
- SPN161 Lachnospiraceae [G-13] bacterium_MOT-181_nov_87.225%
- SPN162 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_89.855%
- SPN163 Oscillospiraceae [G-4] bacterium_MOT-151_nov_86.722%
- SPN164 Lacrimispora saccharolytica_nov_93.082%
- SPN165 Acetivibrio cellulolyticus_nov_82.189%
- SPN166 Eubacterium ramulus_nov_91.232%
- SPN167 Lachnospiraceae [G-5] bacterium_MOT-170_nov_97.904%
- SPN168 Eisenbergiella massiliensis_nov_85.655%
- SPN169 Lachnospiraceae [G-14] bacterium_MOT-182_nov_88.608%
- SPN170 Dehalobacter restrictus_nov_85.221%
- SPN171 Lachnospiraceae [G-14] bacterium_MOT-182_nov_93.348%
- SPN172 Oscillospiraceae [G-4] bacterium_MOT-151_nov_93.971%
- SPN173 Lachnospiraceae [G-11] bacterium_MOT-176_nov_89.858%
- SPN174 Lachnospiraceae [G-6] bacterium_MOT-171_nov_90.586%
- SPN175 Anaerotruncus rubiinfantis_nov_92.708%
- SPN176 Longibaculum muris_nov_93.154%
- SPN177 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.489%
- SPN178 Faecalicoccus acidiformans_nov_89.600%
- SPN179 Anaerotruncus rubiinfantis_nov_88.223%
- SPN180 Pseudoflavonifractor phocaeensis_nov_93.776%
- SPN181 Roseburia inulinivorans_nov_91.476%
- SPN182 Oscillospiraceae [G-2] bacterium_MOT-149_nov_92.531%
- SPN183 Lacrimispora amygdalina_nov_93.082%
- SPN184 Oscillospiraceae [G-3] bacterium_MOT-150_nov_92.324%
- SPN185 Anaeromassilibacillus senegalensis_nov_92.489%
- SPN186 Lachnospiraceae [G-3] bacterium_MOT-168_nov_93.319%
- SPN187 Fusicatenibacter saccharivorans_nov_87.992%
- SPN188 Phoceia massiliensis_nov_87.660%
- SPN189 Turicibacter sanguinis_nov_95.923%

- SPN56 Hathewayia proteolytica_nov_83.297%
- SPN57 Marvinbryantia formatexigens_nov_91.435%
- SPN58 Pseudoflavonifractor capillosus_nov_89.897%
- SPN59 Lachnospiraceae [G-14] bacterium_MOT-184_nov_92.584%
- SPN60 Acetivibrio cellulolyticus_nov_83.801%
- SPN61 Lachnospiraceae [G-12] bacterium_MOT-180_nov_89.613%
- SPN62 Butyrivibrio fibrisolvens_nov_86.831%
- SPN63 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.182%
- SPN64 Blautia faecicola_nov_91.170%
- SPN65 Oscillospiraceae [G-4] bacterium_MOT-151_nov_95.634%
- SPN66 Christensenella hongkongensis_nov_85.294%
- SPN67 Oscillospiraceae [G-3] bacterium_MOT-150_nov_91.736%
- SPN68 Hathewayia proteolytica_nov_84.233%
- SPN69 Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.041%
- SPN70 Lachnospiraceae [G-12] bacterium_MOT-179_nov_94.501%
- SPN71 Eubacteriales [G-3] bacterium_MOT-163_nov_85.944%
- SPN72 Eubacteriales [G-1] bacterium_MOT-159_nov_94.055%
- SPN73 Oscillospiraceae [G-4] bacterium_MOT-151_nov_96.881%
- SPN74 Oscillospiraceae [G-2] bacterium_MOT-149_nov_94.375%
- SPN75 Faecalimonas umbilicata_nov_91.286%
- SPN76 Eubacteriales [G-4] bacterium_MOT-164_nov_94.292%
- SPN77 Lachnoclostridium [Clostridium] scindens_nov_89.648%
- SPN78 Fusicatenibacter saccharivorans_nov_90.041%
- SPN79 Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.096%
- SPN80 Lachnospiraceae [G-14] bacterium_MOT-183_nov_93.348%
- SPN81 Lachnospiraceae [G-11] bacterium_MOT-176_nov_94.726%
- SPN82 Lachnospiraceae [G-11] bacterium_MOT-178_nov_91.886%
- SPN83 Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.388%
- SPN84 Hydrogenoanaerobacterium saccharovorans_nov_88.589%
- SPN85 Eisenbergiella massiliensis_nov_87.164%
- SPN86 Christensenella hongkongensis_nov_85.645%
- SPN87 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.351%
- SPN88 Hydrogenoanaerobacterium saccharovorans_nov_90.041%
- SPN89 Oscillospiraceae [G-3] bacterium_MOT-150_nov_91.511%
- SPN90 Eubacterium xylanophilum_nov_91.075%
- SPN91 Kineothrix alysoides_nov_87.474%
- SPN92 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.719%
- SPN93 Oscillospiraceae [G-3] bacterium_MOT-150_nov_93.582%
- SPN94 Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.451%