

SPN7 Oscillospiraceae [G-6] bacterium_MOT-153	SPN151 Flavonifractor plautii_nov_92.516%	SPN56 Hathewayia proteolytica_nov_83.297%
SPN8 Lactobacillus johnsonii	SPN152 Phocaea massiliensis_nov_88.034%	SPN57 Marvinbryantia formatexigens_nov_91.435%
SPN9 Eubacteriales [G-4] bacterium_MOT-164	SPN153 Clostridiales [F-1][G-1] bacterium_HMT_093_nov_91.775%	SPN58 Pseudoflavonifractor capillosus_nov_89.897%
SPN10 Muribaculaceae [G-1] bacterium_MOT-129_nov_85.887%	SPN154 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.375%	SPN59 Lachnospiraceae [G-14] bacterium_MOT-184_nov_92.584%
SPN100 Eubacteriales [G-1] bacterium_MOT-161_nov_93.843%	SPN155 Roseburia hominis_nov_91.476%	SPN60 Acetivibrio cellulolyticus_nov_83.801%
SPN101 Eubacteriales [G-4] bacterium_MOT-164_nov_97.228%	SPN156 Lachnospiraceae [G-6] bacterium_MOT-171_nov_92.484%	SPN61 Lachnospiraceae [G-12] bacterium_MOT-180_nov_89.613%
SPN102 Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.188%	SPN157 Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.851%	SPN62 Butyrivibrio fibrisolvens_nov_86.831%
SPN103 Oscillospiraceae [G-2] bacterium_MOT-149_nov_94.154%	SPN158 Oscillospiraceae [G-2] bacterium_MOT-149_nov_94.363%	SPN63 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.182%
SPN104 Lachnospiraceae [G-11] bacterium_MOT-177_nov_96.066%	SPN159 Eubacteriales [G-1] bacterium_MOT-159_nov_92.161%	SPN64 Blautia faecicola_nov_91.170%
SPN105 Lachnospiraceae [G-3] bacterium_MOT-168_nov_89.353%	SPN160 Paludicola psychrotolerans_nov_87.759%	SPN65 Oscillospiraceae [G-4] bacterium_MOT-151_nov_95.634%
SPN106 Oscillospiraceae [G-4] bacterium_MOT-151_nov_92.931%	SPN161 Lachnospiraceae [G-13] bacterium_MOT-181_nov_87.225%	SPN66 Christensenella hongkongensis_nov_85.294%
SPN107 Lachnospiraceae [G-7] bacterium_MOT-172_nov_91.718%	SPN162 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_89.855%	SPN67 Oscillospiraceae [G-3] bacterium_MOT-150_nov_91.736%
SPN108 Acetivibrio cellulolyticus_nov_83.153%	SPN163 Oscillospiraceae [G-4] bacterium_MOT-151_nov_86.722%	SPN68 Hathewayia proteolytica_nov_84.233%
SPN109 Eisenbergiella massiliensis_nov_87.578%	SPN164 Lacrimispora saccharolytica_nov_93.082%	SPN69 Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.041%
SPN11 Lachnospiraceae [G-12] bacterium_MOT-180_nov_91.093%	SPN165 Acetivibrio cellulolyticus_nov_82.189%	SPN70 Lachnospiraceae [G-12] bacterium_MOT-179_nov_94.501%
SPN110 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.528%	SPN166 Eubacterium ramulus_nov_91.232%	SPN71 Eubacteriales [G-3] bacterium_MOT-163_nov_85.944%
SPN111 Oscillospiraceae [G-2] bacterium_MOT-149_nov_85.773%	SPN167 Lachnospiraceae [G-5] bacterium_MOT-170_nov_97.904%	SPN72 Eubacteriales [G-1] bacterium_MOT-159_nov_94.055%
SPN112 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.776%	SPN168 Eisenbergiella massiliensis_nov_85.655%	SPN73 Oscillospiraceae [G-4] bacterium_MOT-151_nov_96.881%
SPN113 Lachnospiraceae [G-3] bacterium_MOT-168_nov_94.572%	SPN169 Lachnospiraceae [G-14] bacterium_MOT-182_nov_88.608%	SPN74 Oscillospiraceae [G-2] bacterium_MOT-149_nov_94.375%
SPN114 Eisenbergiella massiliensis_nov_88.174%	SPN170 Dehalobacter restrictus_nov_85.221%	SPN75 Faecalimonas umbilicata_nov_91.286%
SPN115 Acutalibacter muris_nov_94.227%	SPN171 Lachnospiraceae [G-14] bacterium_MOT-182_nov_93.348%	SPN76 Eubacteriales [G-4] bacterium_MOT-164_nov_94.292%
SPN116 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.644%	SPN172 Oscillospiraceae [G-4] bacterium_MOT-151_nov_93.971%	SPN77 Lachnoclostridium [Clostridium] scindens_nov_89.648%
SPN117 Anaerotrignum faecicola_nov_85.287%	SPN173 Lachnospiraceae [G-11] bacterium_MOT-176_nov_89.858%	SPN78 Fusicatenibacter saccharivorans_nov_90.041%
SPN118 Acetatifactor muris_nov_95.652%	SPN174 Lachnospiraceae [G-6] bacterium_MOT-171_nov_90.586%	SPN79 Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.096%
SPN119 Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.971%	SPN175 Anaerotruncus rubiinfantis_nov_92.708%	SPN80 Lachnospiraceae [G-14] bacterium_MOT-183_nov_93.348%
SPN12 Lachnoclostridium phytofermentans_nov_91.458%	SPN176 Longibaculum muris_nov_93.154%	SPN81 Lachnospiraceae [G-11] bacterium_MOT-176_nov_94.726%
SPN120 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.351%	SPN177 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.489%	SPN82 Lachnospiraceae [G-11] bacterium_MOT-178_nov_91.886%
SPN121 Oscillospiraceae [G-2] bacterium_MOT-149_nov_92.484%	SPN178 Faecalicoccus acidiformans_nov_89.600%	SPN83 Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.388%
SPN122 Oscillospiraceae [G-4] bacterium_MOT-151_nov_96.050%	SPN179 Anaerotruncus rubiinfantis_nov_88.223%	SPN84 Hydrogenoanaerobacterium saccharovorans_nov_88.589%
SPN123 Oscillibacter valericigenes_nov_96.042%	SPN180 Pseudoflavonifractor phocaensis_nov_93.776%	SPN85 Eisenbergiella massiliensis_nov_87.164%
SPN124 Lacrimispora xylanolytica_nov_94.363%	SPN181 Roseburia inulinivorans_nov_91.476%	SPN86 Christensenella hongkongensis_nov_85.645%
SPN125 Lachnospiraceae [G-13] bacterium_MOT-181_nov_91.189%	SPN182 Oscillospiraceae [G-2] bacterium_MOT-149_nov_92.531%	SPN87 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.351%
SPN126 Blautia producta_nov_95.833%	SPN183 Lacrimispora amygdalina_nov_93.082%	SPN88 Hydrogenoanaerobacterium saccharovorans_nov_90.041%
SPN127 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_88.820%	SPN184 Oscillospiraceae [G-3] bacterium_MOT-150_nov_92.324%	SPN89 Oscillospiraceae [G-3] bacterium_MOT-150_nov_91.511%
SPN128 Oscillospiraceae [G-3] bacterium_MOT-150_nov_91.925%	SPN185 Anaeromassilibacillus senegalensis_nov_92.489%	SPN90 Eubacterium xylanophilum_nov_91.075%
SPN129 Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.466%	SPN186 Lachnospiraceae [G-3] bacterium_MOT-168_nov_93.319%	SPN91 Kineothrix alysoides_nov_87.474%
SPN130 Kineothrix alysoides_nov_86.848%	SPN187 Fusicatenibacter saccharivorans_nov_87.992%	SPN92 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.719%
SPN131 Oscillospiraceae [G-4] bacterium_MOT-151_nov_96.875%	SPN188 Phocaea massiliensis_nov_87.660%	SPN93 Oscillospiraceae [G-3] bacterium_MOT-150_nov_93.582%
SPN132 Oscillibacter valericigenes_nov_93.595%	SPN189 Turicibacter sanguinis_nov_95.923%	SPN94 Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.451%