

pecies

- SP1 Bacteroides thetaiotaomicron
- SP10 Lachnospiraceae_[G-1] bacterium_MOT-166
- SP2 Akkermansia muciniphila
- SP3 Lachnospiraceae_[G-14] bacterium_MOT-184
- SP4 Parasutterella excrementihominis
- SP5 Blautia hominis
- SP6 Eubacteriales_[G-1] bacterium_MOT-159
- SP7 Lachnospiraceae_[G-11] bacterium_MOT-177
- SP8 Oscillospiraceae_[G-7] bacterium_MOT-154
- SP9 Romboutsia ilealis
- SPN1 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.069%
- SPN10 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_91.506%
- SPN100 Blautia faecicola_nov_89.749%
- SPN101 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.549%
- SPN102 Duncaniella freteri_nov_94.162%
- SPN103 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.478%
- SPN104 Acetivibrio cellulolyticus_nov_83.851%
- SPN105 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_91.085%
- SPN106 Sporobacter termitidis_nov_83.168%
- SPN107 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.644%
- SPN108 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.091%
- SPN109 Lawsonibacter asaccharolyticus_nov_94.798%
- SPN110 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.450%
- SPN111 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.899%
- SPN112 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.990%
- SPN113 Acetivibrio cellulolyticus_nov_83.090%
- SPN114 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_87.033%
- SPN115 Anaerostipes caccae_nov_96.058%
- SPN116 Duncaniella freteri_nov_90.323%
- SPN117 Ruminiclostridium cellulolyticum_nov_82.704%
- SPN118 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.040%
- SPN119 Hydrogenoanaerobacterium saccharovorans_nov_89.942%
- SPN12 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_95.335%
- SPN120 Kineothrix alysoides_nov_87.129%
- SPN121 Oscillibacter valericigenes_nov_93.642%
- SPN122 Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.778%
- SPN123 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_87.452%
- SPN124 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.098%
- SPN125 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.850%
- SPN126 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.805%
- SPN127 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_92.293%
- SPN128 Duncaniella freteri_nov_93.774%
- SPN13 Turicibacter sanguinis_nov_95.635%
- SPN130 Pseudoflavonifractor phocaeensis_nov_95.568%
- SPN131 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_91.602%
- SPN132 Enterocloster bolteae_nov_94.951%
- SPN133 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_95.183%
- SPN134 Acutalibacter muris_nov_88.359%
- SPN135 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_92.731%
- SPN136 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_90.430%
- SPN137 Blautia hominis_nov_97.773%
- SPN138 Anaerotignum aminivorans_nov_92.184%
- SPN139 Parasutterella excrementihominis_nov_97.852%
- SPN14 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.849%
- SPN15 Faecalimonas umbilicata_nov_91.538%
- SPN16 Ruminiclostridium cellulolyticum_nov_84.158%
- SPN17 Eisenbergiella massiliensis_nov_89.126%
- SPN18 Eubacteriales_[G-1] bacterium_MOT-158_nov_90.946%
- SPN19 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_85.769%
- SPN2 Eubacteriales_[G-3] bacterium_MOT-163_nov_89.695%
- SPN20 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.071%
- SPN21 Faecalimonas umbilicata_nov_94.798%
- SPN23 Christensenella massiliensis_nov_84.571%
- SPN24 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_93.064%
- SPN25 Lachnoclostridium [Clostridium] scindens_nov_90.979%
- SPN26 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.427%
- SPN27 Eisenbergiella massiliensis_nov_88.123%
- SPN28 Phocaea massiliensis_nov_90.297%
- SPN29 Pseudoflavonifractor phocaeensis_nov_86.122%
- SPN3 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.851%
- SPN30 Marvinbryantia formatexigens_nov_91.942%
- SPN31 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_92.941%
- SPN32 Duncaniella freteri_nov_88.598%
- SPN33 Ruminiclostridium cellulolyticum_nov_83.300%
- SPN34 Anaeromassilibacillus senegalensis_nov_92.460%
- SPN35 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_87.549%
- SPN36 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.103%
- SPN37 Sporobacter termitidis_nov_87.897%
- SPN38 Tyzzerella [Clostridium] colinum_nov_88.655%
- SPN39 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.155%
- SPN4 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_92.293%
- SPN40 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_91.765%
- SPN41 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_86.111%
- SPN42 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_93.254%
- SPN44 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_90.361%
- SPN45 Butyricoccus pullicaecorum_nov_85.934%
- SPN46 Lawsonibacter asaccharolyticus_nov_90.421%
- SPN47 Lawsonibacter asaccharolyticus_nov_90.038%
- SPN5 Hydrogenoanaerobacterium saccharovorans_nov_89.942%
- SPN52 Duncaniella freteri_nov_90.262%
- SPN56 Lawsonibacter asaccharolyticus_nov_91.762%
- SPN6 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.243%
- SPN63 Duncaniella freteri_nov_93.208%
- SPN68 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.778%
- SPN7 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.245%
- SPN74 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.653%
- SPN77 Hathewayia proteolytica_nov_84.569%
- SPN8 Lachnoclostridium [Clostridium] scindens_nov_87.739%
- SPN85 Leifsonia kafniensis_nov_84.158%
- SPN87 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.697%
- SPN88 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_97.154%
- SPN89 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.267%
- SPN9 Acetivibrio cellulolyticus_nov_85.921%
- SPN90 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.886%
- SPN91 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_93.922%
- SPN92 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.590%
- SPN93 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_92.549%
- SPN94 Duncaniella freteri_nov_93.208%
- SPN95 Phocaea massiliensis_nov_90.060%
- SPN96 Eubacterium coprostanoligenes_nov_91.511%
- SPN97 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_94.320%
- SPN98 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.713%
- SPN99 Anaeroplasma abactoclasticum_nov_87.352%
- SPPN1 Eubacteriales_[G-4] multispecies_sppn1_2_nov_94.769%