

Species

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 boltae_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 Butyrivibrio 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 kafiensis_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%