

pecies

- SP1 Akkermansia muciniphila
- SP2 Bacteroides thetaiotaomicron
- SP3 Romboutsia timonensis
- SP4 Sutterella sp._str._cont1.66
- SP5 Blautia hominis
- SP6 Bifidobacterium pseudolongum
- SP8 Parasutterella excrementihominis
- SPN1 Eisenbergiella massiliensis_nov_90.734%
- SPN10 Duncaniella freteri_nov_93.774%
- SPN102 Butyrivicoccus pullicaecorum_nov_85.934%
- SPN104 Hathewayia proteolytica_nov_84.569%
- SPN105 Kineothrix alysoidea_nov_91.633%
- SPN106 Oscillibacter valericigenes_nov_90.996%
- SPN107 Blautia marasmi_nov_90.680%
- SPN108 Duncaniella freteri_nov_85.958%
- SPN109 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_90.389%
- SPN11 Eubacterium coprostanoligenes_nov_91.511%
- SPN110 Lacrimispora xylanolytica_nov_91.245%
- SPN111 Lacrimispora xylanolytica_nov_91.406%
- SPN112 Phoea massiliensis_nov_90.060%
- SPN113 Pseudoflavonifractor phocaensis_nov_90.211%
- SPN114 Kineothrix alysoidea_nov_91.451%
- SPN115 Oscillibacter valericigenes_nov_91.939%
- SPN116 Anaeroplasmabactoclasticum_nov_87.352%
- SPN117 Lachnospirillum sp._str._M62/1_nov_91.085%
- SPN118 Duncaniella freteri_nov_94.162%
- SPN119 Eisenbergiella massiliensis_nov_87.218%
- SPN12 Anaerocolumna cellulolytica_nov_90.116%
- SPN120 Turicibacter sanguinis_nov_95.635%
- SPN121 Oscillibacter valericigenes_nov_90.613%
- SPN122 Eisenbergiella massiliensis_nov_90.421%
- SPN123 Ruminococcus champanellensis_nov_92.262%
- SPN124 Bacteroides capillosus_nov_90.076%
- SPN125 Sporobacter termitidis_nov_83.168%
- SPN126 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.298%
- SPN127 Oscillibacter valericigenes_nov_94.402%
- SPN128 unclassified_Ruminococcaceae sp._str._D16_nov_95.769%
- SPN129 Duncaniella freteri_nov_93.220%
- SPN13 Blautia hominis_nov_97.773%
- SPN130 unclassified_Ruminococcaceae sp._str._D16_nov_92.115%

- SPN131 Kineothrix alysoidea_nov_88.654%
- SPN132 Acetivibrio cellulolyticus_nov_83.851%
- SPN133 Acetivibrio cellulolyticus_nov_83.090%
- SPN134 Eisenbergiella massiliensis_nov_86.346%
- SPN135 Anaerostipes sp._str._3256FAA_nov_96.899%
- SPN136 Ruminiclostridium cellulolyticum_nov_82.704%
- SPN137 Oscillibacter valericigenes_nov_92.278%
- SPN138 Eisenbergiella massiliensis_nov_90.684%
- SPN139 Hydrogenoanaerobacterium saccharovorans_nov_89.942%
- SPN14 Anaerostipes aminivorans_nov_92.184%
- SPN140 Kineothrix alysoidea_nov_87.129%
- SPN141 Oscillibacter valericigenes_nov_93.642%
- SPN142 Lachnospirillum sp._str._L2_50_nov_87.968%
- SPN143 Duncaniella freteri_nov_90.512%
- SPN144 Oscillibacter valericigenes_nov_94.981%
- SPN145 Leifsonia kafniensis_nov_84.158%
- SPN15 Kineothrix alysoidea_nov_91.473%
- SPN16 unclassified_Ruminococcaceae sp._str._D16_nov_93.077%
- SPN17 Hydrogenoanaerobacterium saccharovorans_nov_89.942%
- SPN18 Lachnospirillum [Clostridium] polysaccharolyticum_nov_93.243%
- SPN19 Oscillibacter valericigenes_nov_94.175%
- SPN2 Duncaniella freteri_nov_88.476%
- SPN20 Acetivibrio cellulolyticus_nov_85.921%
- SPN21 Oscillibacter valericigenes_nov_94.027%
- SPN22 Oscillibacter valericigenes_nov_93.822%
- SPN23 Oscillibacter valericigenes_nov_90.751%
- SPN24 Kineothrix alysoidea_nov_91.211%
- SPN25 Eisenbergiella massiliensis_nov_88.636%
- SPN26 Eisenbergiella massiliensis_nov_91.262%
- SPN27 Oscillibacter valericigenes_nov_91.954%
- SPN28 Lachnospirillum bolteaee_nov_91.683%
- SPN29 Ruminiclostridium cellulolyticum_nov_84.158%
- SPN3 Roseburia inulinivorans_nov_87.925%
- SPN30 Eisenbergiella massiliensis_nov_89.126%
- SPN31 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_88.115%
- SPN32 Duncaniella freteri_nov_93.208%
- SPN33 Eisenbergiella massiliensis_nov_85.389%
- SPN34 Kineothrix alysoidea_nov_92.636%
- SPN35 Oscillibacter valericigenes_nov_93.295%
- SPN36 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.057%

- SPN37 Faecalimonas umbilicata_nov_94.798%
- SPN38 Christensenella massiliensis_nov_84.571%
- SPN39 Oscillibacter valericigenes_nov_93.605%
- SPN4 Mobilitalea sibirica_nov_87.795%
- SPN40 Phoea massiliensis_nov_90.297%
- SPN41 Romboutsia timonensis_nov_97.951%
- SPN42 Lactobacillus amylovorus_nov_86.364%
- SPN43 Eisenbergiella massiliensis_nov_88.123%
- SPN44 Pseudoflavonifractor phocaensis_nov_86.122%
- SPN45 Kineothrix alysoidea_nov_91.054%
- SPN46 Marvinbryantia formatexigens_nov_91.942%
- SPN47 Oscillibacter valericigenes_nov_92.308%
- SPN48 Anaeromassilibacillus senegalensis_nov_92.460%
- SPN49 Ruminiclostridium cellulolyticum_nov_83.300%
- SPN5 unclassified_Ruminococcaceae sp._str._D16_nov_96.132%
- SPN50 Sporobacter termitidis_nov_87.897%
- SPN51 Butyrivibrio proteoclasticus_nov_85.714%
- SPN52 Tyzzerella [Clostridium] colinum_nov_88.655%
- SPN53 Lachnospirillum bolteaee_nov_95.146%
- SPN54 Bacteroides capillosus_nov_90.613%
- SPN55 Lacrimispora xylanolytica_nov_88.593%
- SPN6 Lacrimispora saccharolytica_nov_89.981%
- SPN60 Duncaniella freteri_nov_90.262%
- SPN64 unclassified_Ruminococcaceae sp._str._D16_nov_91.571%
- SPN7 Lachnospirillum symbiosum_nov_95.146%
- SPN70 Duncaniella freteri_nov_88.598%
- SPN75 Kineothrix alysoidea_nov_90.559%
- SPN8 Acutalibacter muris_nov_88.359%
- SPN86 Eisenbergiella massiliensis_nov_90.267%
- SPN9 Anaerotruncus rubiinfantis_nov_83.179%
- SPN91 Lachnospirillum [Clostridium] polysaccharolyticum_nov_86.320%
- SPN96 Duncaniella freteri_nov_88.224%
- SPPN1 multigenus multispecies_sppn1_2_nov_87.739%
- SPPN2 multigenus multispecies_sppn2_2_nov_90.979%
- SPPN3 multigenus multispecies_sppn3_2_nov_84.557%
- SPPN4 Bacillus multispecies_sppn4_2_nov_83.299%
- SPPN6 multigenus multispecies_sppn6_2_nov_92.456%
- SPPN7 Lachnospirillum bolteaee