

Species

SP39 Ralstonia sp._Oral_Taxon_C91
SP4 Lactobacillus murinus
SP40 Haemophilus parainfluenzae
SP41 Bradyrhizobium lupini
SP42 Pseudomonas oryzaehabitans
SP43 Oxalobacteraceae_[G] sp._Oral_Taxon_C91
SP44 Turicella otitidis
SP46 Roseburia inulinivorans
SP5 Streptococcus thoraltensis
SP50 Streptococcus infantis
SP51 Paenibacillus campinansensis
SP52 Mycobacterium cookii
SP53 Acinetobacter pittii
SP56 Capnocytophaga sputigena
SP57 Bifidobacterium longum
SP6 Bifidobacterium pseudolongum
SP60 Streptococcus mitis
SP61 Staphylococcus auricularis
SP62 Sphingomonas yabuuchiae
SP65 Methylobacterium goesingense
SP66 Cutibacterium namnetense
SP67 Haemophilus haemolyticus
SP68 Parasutterella excrementihominis
SP69 Lactobacillus ruminis
SP7 Escherichia coli
SP74 Clostridium disporicum
SP75 Microbacterium lacticum
SP76 Corynebacterium pyruviciproducens
SP77 Clostridium saudiense
SP78 Staphylococcus hominis
SP8 Sediminibacterium aquarii
SP81 Sphingomonas panni
SP82 Pseudomonas putida
SP83 Odoribacter splanchnicus
SP84 Finegoldia magna
SP9 Enterococcus sp._str._F95
SPN1 Lachnoclostridium polysaccharolyticum nov_91.522%
SPN10 Lachnoclostridium polysaccharolyticum nov_92.829%
SPN100 Eisenbergiella massiliensis nov_90.385%
SPN101 Blautia marasmi nov_91.650%
SPN102 Acetivibrio cellulolyticus nov_84.189%
SPN103 Gabonia massiliensis nov_84.877%
SPN104 Bacteroidetes_[G-7] bacterium_HMT_911 nov_85.164%
SPN105 Lachnoclostridium bolteae nov_92.621%
SPN106 Bacteroides thetaiotaomicron nov_93.762%
SPN107 Blautia schinkii nov_93.548%
SPN108 Eubacterium hallii nov_90.751%
SPN109 Ruthenibacterium lactatiformans nov_95.769%
SPN11 Bacteroidetes_[G-7] bacterium_HMT_911 nov_85.910%
SPN110 Enterorhabdus muris nov_92.245%
SPN111 Terrimicrobium sacchariphilum nov_81.224%
SPN112 Bacteroidetes_[G-7] bacterium_HMT_911 nov_84.854%
SPN113 Massilia yuzhufengensis nov_97.395%
SPN114 Colidextribacter massiliensis nov_87.908%
SPN115 Eisenbergiella massiliensis nov_90.133%
SPN116 Neglecta timonensis nov_94.497%
SPN117 Ruminococcaceae_[G-2] bacterium_HMT_085 nov_85.657%
SPN118 Herbaspirillum huttiense nov_83.151%
SPN119 Roseburia inulinivorans nov_93.282%
SPN12 Sediminibacterium aquarii nov_92.460%
SPN120 Erysipelatoclostridium sp._str._HGF2 nov_88.971%
SPN121 Oscillibacter valericigenes nov_93.295%
SPN122 Turicella otitidis nov_96.071%
SPN123 Bacteroides capillosus nov_91.379%
SPN124 Butyrivibrio proteoclasticus nov_78.519%
SPN125 Phoea massiliensis nov_87.354%
SPN126 Hydrogenoanaerobacterium saccharovorans nov_87.379%
SPN127 Bacteroides massiliensis nov_96.522%
SPN128 Saccharibacteria_(TM7)[G-3] bacterium_HMT_351 nov_92.706%
SPN129 unclassified_Ruminococcaceae sp._str._D16 nov_85.851%
SPN13 Butyrivibrio proteoclasticus nov_91.628%

SPN174 Sutterella sp._str._cont1.06 nov_97.744%
SPN175 Terrimonas pekingensis nov_90.822%
SPN176 Lachnoclostridium polysaccharolyticum nov_89.555%
SPN177 unclassified_Lachnospiraceae sp._str._cont1.79 nov_92.039%
SPN178 Agathobaculum desmolans nov_91.845%
SPN179 Phoea massiliensis nov_90.748%
SPN18 Dorea longicatena nov_94.563%
SPN180 unclassified_Ruminococcaceae sp._str._D16 nov_92.898%
SPN181 Lactobacillus murinus nov_94.528%
SPN182 unclassified_Ruminococcaceae sp._str._D16 nov_90.249%
SPN183 Anaeroplasmabactoclasticum nov_86.957%
SPN184 Marvinbryantia formatexigens nov_91.975%
SPN185 Phoea massiliensis nov_89.286%
SPN186 Bacteroidetes_[G-7] bacterium_HMT_911 nov_85.352%
SPN187 Hydrogenoanaerobacterium saccharovorans nov_89.341%
SPN188 Lachnoclostridium asparagiforme nov_94.347%
SPN189 Anaerocolumna cellulolytica nov_90.522%
SPN19 Lachnoclostridium bolteae nov_94.951%
SPN190 Alistipes putredinis nov_93.156%
SPN191 Eubacterium siraeum nov_91.849%
SPN192 Blautia producta nov_94.574%
SPN193 Alistipes ihumii nov_93.905%
SPN194 Eubacterium siraeum nov_91.984%
SPN195 Anaerocolumna cellulolytica nov_86.863%
SPN196 Anaerostipes sp._str._3256FAA nov_89.017%
SPN197 Eisenbergiella massiliensis nov_86.867%
SPN198 Acinetobacter johnsonii nov_97.697%
SPN199 Enterorhabdus muris nov_89.344%
SPN2 Bacteroidetes_[G-7] bacterium_HMT_911 nov_85.742%
SPN20 Prevotella fusca nov_82.772%
SPN200 unclassified_Ruminococcaceae sp._str._D16 nov_94.423%
SPN201 Anaeroplasmabactoclasticum nov_87.059%
SPN202 Agathobaculum desmolans nov_91.794%
SPN203 Oscillibacter valericigenes nov_93.798%
SPN204 Dorea longicatena nov_94.563%
SPN205 Gabonia massiliensis nov_86.704%
SPN206 Blautia producta nov_94.778%
SPN207 Ruminiclostridium cellulolyticum nov_82.505%
SPN208 Faecalibacillus acidiformans nov_90.075%
SPN209 Muricomes intestini nov_88.998%
SPN21 Anaerotruncus rubiinfantis nov_82.560%
SPN210 Bacteroidetes_[G-7] bacterium_HMT_911 nov_85.659%
SPN211 Alistipes senegalensis nov_96.905%
SPN212 Longibaculum muris nov_93.517%
SPN213 Barnesiella viscericola nov_83.673%
SPN214 unclassified_Ruminococcaceae sp._str._D16 nov_95.192%
SPN215 Streptococcus troglodytidis nov_90.257%
SPN216 Clostridiales_[F-1][G-1] bacterium_HMT_093 nov_92.067%
SPN217 Bacteroides capillosus nov_90.421%
SPN218 Anaerotruncus rubiinfantis nov_81.262%
SPN219 Oscillibacter valericigenes nov_93.617%
SPN22 Anaerotruncus rubiinfantis nov_80.705%
SPN220 Anaeromassilibacillus senegalensis nov_92.063%
SPN221 Deinococcus aerophilus nov_93.097%
SPN222 Ruminococcaceae_[G-2] bacterium_HMT_085 nov_88.798%
SPN223 Bacteroidetes_[G-7] bacterium_HMT_911 nov_86.127%
SPN224 Faecalicatena fissicatena nov_85.660%
SPN225 Ihubacter massiliensis nov_90.751%
SPN226 unclassified_Ruminococcaceae sp._str._D16 nov_90.440%
SPN227 Streptococcus acidominimus nov_94.163%
SPN228 unclassified_Ruminococcaceae sp._str._D16 nov_92.692%
SPN229 Eisenbergiella massiliensis nov_90.267%
SPN23 Roseburia hominis nov_90.485%
SPN230 Eisenbergiella massiliensis nov_91.082%
SPN231 Neglecta timonensis nov_94.677%
SPN232 Ruminococcus champanellensis nov_92.063%
SPN233 Bacteroidetes_[G-7] bacterium_HMT_911 nov_83.172%
SPN234 Alistipes senegalensis nov_93.846%
SPN235 Eubacterium xylanophilum nov_92.075%
SPN236 Kiloniella majae nov_87.759%
SPN237 Streptococcus thermophilus nov_96.505%

SPN2 Lactobacillus roridimorphus nov_92.063%
SPN200 Alistipes finegoldii nov_93.690%
SPN41 Bacteroides capillosus nov_92.871%
SPN42 Bacteroidetes_[G-7] bacterium_HMT_911 nov_87.452%
SPN43 Phoea massiliensis nov_91.519%
SPN44 Faecalibaculum rodentium nov_84.952%
SPN45 Eubacterium siraeum nov_91.566%
SPN46 Phoea massiliensis nov_90.675%
SPN47 Eubacterium coprostanoligenes nov_91.546%
SPN48 Bacteroidetes_[G-7] bacterium_HMT_911 nov_87.476%
SPN49 Ruminococcaceae_[G-2] bacterium_HMT_085 nov_91.616%
SPN5 Lachnoclostridium sp._str._M62/1 nov_94.175%
SPN50 Oscillibacter valericigenes nov_94.208%
SPN51 Streptococcus danieliae nov_96.516%
SPN52 Fusicatenibacter saccharivorans nov_90.138%
SPN53 Lachnoclostridium polysaccharolyticum nov_92.442%
SPN54 Colidextribacter massiliensis nov_89.382%
SPN55 Enterococcus sp._str._F95 nov_95.446%
SPN56 Oscillibacter valericigenes nov_91.908%
SPN57 Barnesiella viscericola nov_82.900%
SPN58 unclassified_Ruminococcaceae sp._str._D16 nov_93.269%
SPN59 Bacteroidetes_[G-7] bacterium_HMT_911 nov_82.752%
SPN6 Negativibacillus massiliensis nov_86.914%
SPN60 Lactobacillus oeni nov_87.778%
SPN61 Parabacteroides distasonis nov_83.426%
SPN62 Eisenbergiella massiliensis nov_86.705%
SPN63 Sporobacter termitidis nov_88.492%
SPN64 Eisenbergiella massiliensis nov_90.631%
SPN65 NA rectale nov_92.816%
SPN66 Enterorhabdus caecimuris nov_93.800%
SPN67 Eubacterium xylanophilum nov_91.149%
SPN68 Roseburia hominis nov_92.100%
SPN69 Bacteroidetes_[G-7] bacterium_HMT_911 nov_84.423%
SPN7 Algimonas arctica nov_79.276%
SPN70 Acetivibrio cellulolyticus nov_83.125%
SPN71 Leifsonia kafiensis nov_84.692%
SPN72 Clostridium merdae nov_91.321%
SPN73 Barnesiella intestinhominis nov_83.554%
SPN74 Absiella tortuosus nov_86.716%
SPN75 Neglecta timonensis nov_95.472%
SPN76 Ruminococcaceae_[G-2] bacterium_HMT_085 nov_91.207%
SPN77 Streptococcus acidominimus nov_94.788%
SPN78 Ruminococcaceae_[G-2] bacterium_HMT_085 nov_91.207%
SPN79 Oscillibacter valericigenes nov_92.337%
SPN8 Intestinimonas butyriciproducens nov_95.775%
SPN80 Ruminococcaceae_[G-2] bacterium_HMT_085 nov_87.018%
SPN81 Eisenbergiella massiliensis nov_92.008%
SPN82 Anaerotruncus rubiinfantis nov_81.227%
SPN83 Hathewayia proteolytica nov_83.768%
SPN84 Anaerotruncus rubiinfantis nov_93.050%
SPN85 Tyzzerella colinum nov_87.579%
SPN86 Eubacterium ventriosum nov_93.217%
SPN87 Saccharibacteria_(TM7)[G-3] bacterium_HMT_351 nov_93.050%
SPN88 Ruminococcaceae_[G-2] bacterium_HMT_085 nov_90.650%
SPN89 Prevotella oris nov_82.772%
SPN9 Lachnoclostridium herbivorans nov_91.346%
SPN90 Alistipes senegalensis nov_94.073%
SPN91 Vampirovibrio cholerae nov_81.520%
SPN92 Enterorhabdus caecimuris nov_95.382%
SPN93 Leifsonia kafiensis nov_84.554%
SPN94 Alloabaculum stercoricanis nov_86.139%
SPN95 Roseburia faecalis nov_94.197%
SPN96 unclassified_Lachnospiraceae sp._str._cont1.79 nov_91.489%
SPN97 unclassified_Lachnospiraceae sp._str._cont1.79 nov_95.155%
SPN98 Hydrogenoanaerobacterium saccharovorans nov_88.395%
SPN99 Intestinimonas butyriciproducens nov_93.812%
SPP1 Cutibacterium_Propionibacterium_acnes
SPP10 Burkholderia cenocepacia_lata_multivorans.sp._str._383
SPP11 Kocuria rhizophila.sp._Oral_Taxon_B56
SPP2 Pseudomonas kuykendallii_selenipraecipitans
SPN3 Roseburia faecalis Bifidobacterium longum Bifidobacterium