

Species

SP1 Duncaniella freteri
SP10 Oscillospiraceae_[G-7] bacterium_MOT-154
SP11 Peptococcaceae_[G-1] bacterium_MOT-146
SP12 Lactobacillus johnsonii
SP14 Eubacteriales_[G-4] bacterium_MOT-164
SP15 Erysipelatoclostridium [Clostridium] coeleatum
SP16 Ileibacterium valens
SP17 Lachnospiraceae_[G-7] bacterium_MOT-172
SP18 Eubacteriales_[G-1] bacterium_MOT-144
SP19 Parvibacter caecicola
SP2 Erysipelotrichaceae_[G-1] bacterium_MOT-189
SP20 Lactobacillus intestinalis
SP21 Faecalibaculum rodentium
SP22 Bacteroides acidifaciens
SP23 Lachnospiraceae_[G-14] bacterium_MOT-185
SP24 Eubacteriales_[G-2] bacterium_MOT-162
SP25 Lactobacillus gasseri
SP26 Akkermansia muciniphila
SP27 Adlercreutzia muris
SP28 Helicobacter ganmani
SP29 Phocaecicola sartorii
SP3 Prevotella sp._MOT-128
SP30 Alistipes sp._MOT-127
SP31 Oscillospiraceae_[G-3] bacterium_MOT-150
SP32 Acutalibacter muris
SP4 Staphylococcus saprophyticus
SP5 Muribaculaceae_[G-1] bacterium_MOT-129
SP6 Mammalicoccus lentus
SP7 Ligilactobacillus murinus
SP8 Limosilactobacillus reuteri
SP9 Lachnospiraceae_[G-11] bacterium_MOT-178
SPN100 Kineothrix alysoides_nov_90.079%
SPN101 Lacrimispora xylanolytica_nov_93.969%
SPN102 Kiloniella majae_nov_85.124%
SPN103 Duncaniella freteri_nov_90.772%
SPN104 Eubacterium ramulus_nov_87.814%
SPN105 Longibaculum muris_nov_90.820%
SPN106 Adlercreutzia equolifaciens_nov_90.020%
SPN107 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.552%
SPN108 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_93.642%
SPN109 Culturomica massiliensis_nov_89.792%
SPN11 Kiloniella majae_nov_87.397%
SPN110 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.961%
SPN111 Duncaniella freteri_nov_89.552%
SPN112 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.990%
SPN113 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.172%
SPN114 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.174%
SPN115 Eubacteriales_[G-3] bacterium_MOT-163_nov_81.321%
SPN116 Eubacteriales_[G-1] bacterium_MOT-159_nov_92.323%
SPN117 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_96.040%
SPN118 Duncaniella freteri_nov_86.278%
SPN119 Duncaniella freteri_nov_86.456%
SPN120 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.115%
SPN121 Pseudoflavonifractor phocaeensis_nov_93.077%
SPN122 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.333%
SPN123 Duncaniella freteri_nov_95.057%
SPN124 Muribaculaceae_[G-2] bacterium_MOT-104_nov_85.741%
SPN125 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_97.012%
SPN126 Duncaniella freteri_nov_88.180%
SPN127 Eubacteriales_[G-4] bacterium_MOT-165_nov_93.763%
SPN128 Agathobaculum desmolans_nov_91.262%
SPN129 Magnetovibrio blakemorei_nov_83.438%
SPN130 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.618%
SPN131 Parvularcula lutaonensis_nov_83.610%
SPN132 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.981%
SPN133 Algimonas porphyrae_nov_79.733%
SPN134 Adlercreutzia muris_nov_92.245%
SPN135 Faecalimonas umbilicata_nov_95.358%

SPN136 Devosia geojensis_nov_84.440%
SPN137 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_92.717%
SPN138 Anaeromassilibacillus senegalensis_nov_85.602%
SPN139 Anaerocolumna cellulositlytica_nov_90.504%
SPN14 Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.429%
SPN140 Lawsonibacter asaccharolyticus_nov_91.762%
SPN141 Sporosolibacterium tautonense_nov_82.852%
SPN142 Duncaniella freteri_nov_87.006%
SPN143 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.874%
SPN144 Muribaculaceae_[G-2] bacterium_MOT-104_nov_87.262%
SPN145 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_94.094%
SPN146 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.688%
SPN147 Erysipelothrix rhusiopathiae_nov_85.902%
SPN148 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.280%
SPN149 Acutalibacter muris_nov_94.828%
SPN15 Faecalicatena fissicatena_nov_94.542%
SPN150 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.455%
SPN151 Marvinbryantia formatexigens_nov_91.358%
SPN152 Eisenbergiella massiliensis_nov_88.100%
SPN153 Duncaniella freteri_nov_87.453%
SPN154 Eisenbergiella massiliensis_nov_87.838%
SPN155 Phocaea massiliensis_nov_86.680%
SPN156 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.504%
SPN157 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_91.456%
SPN158 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_97.614%
SPN159 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_97.642%
SPN160 Adlercreutzia caecimuris_nov_94.177%
SPN161 Adlercreutzia caecimuris_nov_89.379%
SPN162 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.347%
SPN163 Hydrogenoanaerobacterium saccharovorans_nov_88.781%
SPN164 Adlercreutzia caecimuris_nov_92.644%
SPN165 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.602%
SPN166 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.446%
SPN167 Caproicobacter fermentans_nov_87.354%
SPN168 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.333%
SPN169 Roseburia intestinalis_nov_92.941%
SPN170 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.257%
SPN171 Culturomica massiliensis_nov_89.792%
SPN172 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.091%
SPN173 Alpinimonas psychrophila_nov_82.549%
SPN174 Eubacteriales_[G-3] bacterium_MOT-163_nov_81.921%
SPN175 Mollicutes_[G-2] bacterium_MOT-187_nov_93.246%
SPN176 Duncaniella freteri_nov_88.827%
SPN177 Anaerotignum lactatifermentans_nov_96.139%
SPN178 Faecalicatena orotica_nov_92.218%
SPN179 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_88.610%
SPN180 Odoribacter splanchnicus_nov_92.146%
SPN181 Mailhella massiliensis_nov_89.888%
SPN182 Eggerthella timonensis_nov_89.222%
SPN183 Phocaea massiliensis_nov_90.239%
SPN190 Alistipes putredinis_nov_94.847%
SPN191 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.354%
SPN2 Duncaniella freteri_nov_86.679%
SPN201 Duncaniella freteri_nov_96.591%
SPN21 Duncaniella freteri_nov_90.038%
SPN22 Bacteroidetes_[G-3] bacterium_HMT_436_nov_86.078%
SPN23 Duncaniella freteri_nov_87.759%
SPN24 Parasutterella excrementihominis_nov_94.778%
SPN25 Alistipes senegalensis_nov_93.846%
SPN26 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.870%
SPN27 Duncaniella freteri_nov_88.972%
SPN28 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_94.757%
SPN29 Duncaniella freteri_nov_87.709%
SPN30 Duncaniella freteri_nov_88.368%
SPN31 Duncaniella freteri_nov_90.530%
SPN32 Olsenella phocaeensis_nov_92.172%
SPN33 Odoribacter splanchnicus_nov_92.337%
SPN34 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_94.929%

SPN35 Duncaniella freteri_nov_93.573%
SPN36 Turicibacter sanguinis_nov_95.833%
SPN37 Tidjanibacter massiliensis_nov_89.583%
SPN38 Parabacteroides merdae_nov_93.182%
SPN39 Mucispirillum schaedleri_nov_93.124%
SPN40 Saccharibacteria (TM7)_[G-3] bacterium_HMT_351_nov_93.617%
SPN41 Parabacteroides distasonis_nov_97.706%
SPN42 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.651%
SPN43 Duncaniella freteri_nov_87.290%
SPN44 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.398%
SPN45 Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.114%
SPN46 Duncaniella freteri_nov_89.366%
SPN47 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.838%
SPN48 Desulfovibrio fairfieldensis_nov_89.555%
SPN49 Clostridium collagenovorans_nov_81.460%
SPN50 Kineothrix alysoides_nov_87.327%
SPN51 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.522%
SPN52 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.060%
SPN53 Ihubacter massiliensis_nov_94.788%
SPN54 Longibaculum muris_nov_87.574%
SPN55 Alistipes putredinis_nov_92.776%
SPN56 Muribaculaceae_[G-1] bacterium_MOT-129_nov_91.279%
SPN57 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.792%
SPN58 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.455%
SPN59 Eubacteriales_[G-1] bacterium_MOT-161_nov_93.333%
SPN60 Mucispirillum schaedleri_nov_94.862%
SPN61 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_95.935%
SPN62 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.797%
SPN63 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.862%
SPN64 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_97.624%
SPN65 Adlercreutzia caecimuris_nov_89.442%
SPN66 Pseudoflavonifractor phocaeensis_nov_86.513%
SPN67 Bacteroidetes_[G-3] bacterium_HMT_436_nov_86.275%
SPN68 Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.114%
SPN69 Bacteroides uniformis_nov_95.785%
SPN70 Neglectibacter timonensis_nov_95.076%
SPN71 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.610%
SPN72 Duncaniella freteri_nov_89.700%
SPN73 Duncaniella freteri_nov_87.030%
SPN74 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.070%
SPN75 Duncaniella freteri_nov_88.213%
SPN76 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.267%
SPN77 Saccharibacteria (TM7)_[G-3] bacterium_HMT_351_nov_95.358%
SPN78 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_89.555%
SPN79 Ruminiclostridium cellulolyticum_nov_83.433%
SPN80 Duncaniella freteri_nov_89.434%
SPN81 Longibaculum muris_nov_90.588%
SPN82 Mailhella massiliensis_nov_90.377%
SPN83 Lacrimispora saccharolytica_nov_92.621%
SPN84 Eubacterium coprostanoligenes_nov_91.892%
SPN85 Alistipes senegalensis_nov_93.690%
SPN86 Bacteroidetes_[G-3] bacterium_HMT_436_nov_85.938%
SPN87 Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.971%
SPN88 Faecalicatena fissicatena_nov_93.933%
SPN89 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.617%
SPN90 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_90.669%
SPN91 Alistipes finegoldii_nov_94.073%
SPN92 Alloprevotella sp._HMT_473_nov_90.215%
SPN93 Eisenbergiella massiliensis_nov_87.548%
SPN94 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_91.602%
SPN95 Eisenbergiella massiliensis_nov_90.805%
SPN96 Mollicutes_[G-2] bacterium_MOT-187_nov_95.701%
SPN97 Butyrivibrio pullicaeorum_nov_86.320%
SPN98 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_85.393%
SPN99 Anaerotignum lactatifermentans_nov_95.560%
SPPN1 Gloeobacter multispecies_sppn1_2_nov_82.056%
SPPN2 Prevotella multispecies_sppn2_2_nov_89.792%
SPPN3 multigenus multispecies_sppn3_3_nov_93.798%