

Species

SP21 Peptococcaceae_[G-1] bacterium_MOT-146
SP22 Duncaniella freteri
SP23 Oscillospiraceae_[G-6] bacterium_MOT-153
SP24 Sphingobium limneticum
SP25 Actinidia eriantha
SP26 Lactococcus cremoris
SP27 Microbacterium foliorum
SP28 Lactococcus lactis
SP29 Moraxella osloensis
SP3 Ligactobacillus murinus
SP30 Leclercia adecarboxylata
SP31 Parabacteroides distasonis
SP32 Adlercreutzia mucosicola
SP33 Acinetobacter lwofii
SP34 Streptococcus danieliae
SP35 Burkholderia lata
SP36 Phocaeicola sartorii
SP37 Adlercreutzia muris
SP38 Erwinia billingiae
SP39 Sphingomonas carotinfaciens
SP4 Limosilactobacillus reuteri
SP40 Clostridium disporicum
SP41 Yokenella regensburgei
SP42 Phocaeicola vulgatus
SP43 Pseudomonas veronii
SP44 Lachnospiraceae_[G-14] bacterium_MOT-185
SP45 Eubacteriales_[G-4] bacterium_MOT-164
SP46 Eubacteriales_[G-2] bacterium_MOT-162
SP47 Blastococcus aggregatus
SP48 Acinetobacter johnsonii
SP49 Massilia arenae
SP5 Kocuria indica
SP50 Bacteroides acidifaciens
SP51 Pseudomonas cedrina
SP52 Siccibacter turicensis
SP53 Pelomonas saccharophila
SP54 Parasutterella excrementihominis
SP55 Eubacteriales_[G-1] bacterium_MOT-159
SP59 Parabacteroides goldsteini
SP6 Eubacteriales_[G-3] bacterium_MOT-163
SP60 Pedobacter quisquiliarium
SP61 Helicobacter ganmani
SP62 Lachnospiraceae_[G-1] bacterium_MOT-166
SP63 Parvibacter caecicola
SP64 Blastococcus saxosidens
SP66 Sphingomonas echinoides
SP67 Pseudomonas helleri
SP68 Afipia broomeae
SP69 Lachnospiraceae_[G-3] bacterium_MOT-168
SP7 Erysipelatoclostridium [Clostridium] cocleatum
SP70 Pseudomonas extremaustralis
SP71 Klebsiella michiganensis
SP72 Microbacterium kitamiense
SP73 Enterobacter cloacae
SP74 Oscillospiraceae_[G-7] bacterium_MOT-154
SP75 Pararheinheimera mesophila
SP76 Enterobacter hormaechei
SP77 Streptococcus sanguinis
SP8 Lactobacillus johnsonii
SP9 Lactobacillus gasseri
SPN1 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.307%
SPN10 Oscillibacter valericigenes_nov_93.654%
SPN100 Lachnospiraceae_[G-1] bacterium_MOT-166_nov_95.661%
SPN101 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.543%
SPN102 Bacteroidetes_[G-3] bacterium_HMT_436_nov_85.575%
SPN103 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.059%
SPN104 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.105%
SPN105 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.644%
SPN106 Prevotella shahii_nov_87.242%
SPN107 Faecalicatena orotica_nov_94.553%

SPN131 Mailhella massiliensis_nov_90.377%
SPN132 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.945%
SPN133 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.204%
SPN134 Duncaniella freteri_nov_90.152%
SPN135 Duncaniella freteri_nov_87.896%
SPN136 Parasutterella excrementihominis_nov_94.584%
SPN137 Phocaea massiliensis_nov_90.297%
SPN138 Pseudoflavonifractor phocaeensis_nov_95.761%
SPN139 Anaerotignum aminovorans_nov_93.173%
SPN14 Faecalicatena fissicatena_nov_93.580%
SPN140 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.308%
SPN141 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_86.957%
SPN142 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_92.885%
SPN143 Ihubacter massiliensis_nov_94.767%
SPN144 Tyzzerella [Clostridium] colinum_nov_88.494%
SPN145 Planococcus massiliensis_nov_96.992%
SPN146 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_86.508%
SPN147 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_89.942%
SPN148 Eubacterium coprostanoligenes_nov_91.892%
SPN149 Duncaniella freteri_nov_87.759%
SPN15 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.241%
SPN150 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_92.843%
SPN151 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.050%
SPN152 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_86.047%
SPN153 Mediterraneibacter [Ruminococcus] torques_nov_95.200%
SPN154 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_89.349%
SPN155 Caproicibacter fermentans_nov_89.824%
SPN156 Fusicatenibacter saccharivorans_nov_90.514%
SPN157 Erwinia billingiae_nov_97.519%
SPN158 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.059%
SPN159 Lachnoclostridium [Clostridium] scindens_nov_89.827%
SPN16 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.220%
SPN160 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_89.200%
SPN161 Eubacterium oxidoreducens_nov_88.846%
SPN162 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.798%
SPN163 Saccharibacillus kuerlensis_nov_95.336%
SPN164 Acutalibacter muris_nov_94.264%
SPN165 Phocaea massiliensis_nov_90.239%
SPN166 Duncaniella freteri_nov_89.077%
SPN167 Fusicatenibacter saccharivorans_nov_91.018%
SPN168 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_90.234%
SPN169 Adlercreutzia caecimuris_nov_95.382%
SPN17 Breznakia pachnodae_nov_81.284%
SPN170 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.491%
SPN171 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.174%
SPN172 Faecalicatena fissicatena_nov_94.521%
SPN173 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.664%
SPN174 Saccharibacillus kuerlensis_nov_95.709%
SPN175 Clostridium collagenovorans_nov_83.466%
SPN176 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_91.634%
SPN177 Sporosolibacterium tautonense_nov_82.659%
SPN178 Absiella tortuosum_nov_88.725%
SPN179 Adlercreutzia caecimuris_nov_92.277%
SPN18 Lawsonibacter asaccharolyticus_nov_92.514%
SPN180 Oscillibacter valericigenes_nov_95.164%
SPN181 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_97.967%
SPN182 Breznakia pachnodae_nov_83.181%
SPN183 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_88.462%
SPN184 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.050%
SPN185 Parabacteroides merdae_nov_93.182%
SPN186 Hathewayia proteolytica_nov_84.970%
SPN187 Eubacterium ramulus_nov_89.362%
SPN188 Eisenbergiella massiliensis_nov_86.127%
SPN189 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.751%
SPN19 Blautia producta_nov_95.164%
SPN190 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_94.831%
SPN191 Deschloromonas denitrificans_nov_97.446%
SPN192 Blautia producta_nov_96.132%
SPN193 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_96.252%
SPN194 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.905%

SPN218 Duncaniella freteri_nov_87.453%
SPN219 Butyrivibrio sp._HMT_455_nov_83.556%
SPN22 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.480%
SPN220 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.544%
SPN221 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.640%
SPN222 Sporobacter termitidis_nov_88.294%
SPN223 Lacrimispora xylanolytica_nov_93.969%
SPN224 Eubacteriales_[G-1] bacterium_MOT-159_nov_88.550%
SPN225 Acidovorax defluviu_nov_95.946%
SPN226 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.585%
SPN227 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.276%
SPN228 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.452%
SPN229 Adlercreutzia equolifaciens_nov_94.389%
SPN23 Adlercreutzia equolifaciens_nov_96.593%
SPN230 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.980%
SPN231 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_89.981%
SPN232 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_89.443%
SPN233 Mollicutes_[G-2] bacterium_MOT-187_nov_93.284%
SPN234 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_91.732%
SPN235 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.477%
SPN236 Faecalibaculum rodentium_nov_96.571%
SPN237 Duncaniella freteri_nov_83.895%
SPN238 Longibaculum muris_nov_90.392%
SPN239 Adlercreutzia muris_nov_89.506%
SPN24 Cuneatibacter caecimuris_nov_92.486%
SPN240 Roseburia hominis_nov_91.715%
SPN241 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_97.018%
SPN242 Acetivibrio cellulolyticus_nov_83.644%
SPN243 Mailhella massiliensis_nov_89.888%
SPN244 Faecalimonas umblicata_nov_92.692%
SPN245 Anaerotignum aminovorans_nov_92.600%
SPN246 Blautia producta_nov_95.174%
SPN247 Pseudoflavonifractor phocaeensis_nov_92.131%
SPN25 Acetivibrio cellulolyticus_nov_84.058%
SPN26 Blautia faecicola_nov_89.709%
SPN27 Eisenbergiella massiliensis_nov_92.969%
SPN28 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.333%
SPN29 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_87.160%
SPN3 Alloprevotella sp._HMT_473_nov_90.177%
SPN30 Lactobacillus gasseri_nov_93.345%
SPN31 Eubacteriales_[G-1] bacterium_MOT-159_nov_93.976%
SPN32 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_97.614%
SPN33 Eisenbergiella massiliensis_nov_87.669%
SPN34 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.858%
SPN38 Lacrimispora xylanolytica_nov_94.314%
SPN4 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_91.715%
SPN43 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.768%
SPN49 Bacteroides uniformis_nov_95.594%
SPN5 Mollicutes_[G-2] bacterium_MOT-187_nov_90.841%
SPN54 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_93.064%
SPN6 Eisenbergiella massiliensis_nov_87.308%
SPN60 Lacrimispora xylanolytica_nov_91.992%
SPN64 Massilia arenae_nov_97.885%
SPN7 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.235%
SPN72 Eisenbergiella massiliensis_nov_90.805%
SPN75 Lachnoclostridium [Clostridium] aminophilum_nov_87.476%
SPN79 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.643%
SPN8 Lacrimispora xylanolytica_nov_92.308%
SPN80 Flavobacterium branchiicola_nov_96.282%
SPN81 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.050%
SPN82 Anaeroplasmata abactoclasticum_nov_87.352%
SPN83 Faecalicatena orotica_nov_92.218%
SPN84 Lachnoclostridium [Clostridium] scindens_nov_88.247%
SPN85 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.910%
SPN86 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.517%
SPN87 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.050%
SPN88 Olsenella phocaeensis_nov_92.172%
SPN89 Eisenbergiella massiliensis_nov_88.292%
SPN9 Muricomes intestini_nov_89.921%
SPN90 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_87.692%