

Species

SP22 Duncaniella freteri	SPN132 Lachnospiraceae [G-7] bacterium MOT-172_nov_91.945%	SPN219 Butyrivibrio sp. HMT_455_nov_83.556%
SP23 Oscillospiraceae [G-6] bacterium MOT-153	SPN133 Lachnospiraceae [G-7] bacterium MOT-172_nov_93.204%	SPN22 Lachnospiraceae [G-14] bacterium MOT-182_nov_92.480%
SP24 Sphingobium limneticum	SPN134 Duncaniella freteri_nov_90.152%	SPN220 Muribaculaceae [G-2] bacterium MOT-104_nov_89.544%
SP25 Actinidia eriantha	SPN135 Duncaniella freteri_nov_87.896%	SPN221 Oscillospiraceae [G-2] bacterium MOT-149_nov_96.640%
SP26 Lactococcus cremoris	SPN136 Parasutterella excrementihominis_nov_94.584%	SPN222 Sporobacter termitidis_nov_88.294%
SP27 Microbacterium foliorum	SPN137 Phoea massiliensis_nov_90.297%	SPN223 Lacrimispora xylanolytica_nov_93.969%
SP28 Lactococcus lactis	SPN138 Pseudoflavonifractor phocaeensis_nov_95.761%	SPN224 Eubacteriales [G-1] bacterium MOT-159_nov_88.550%
SP29 Moraxella osloensis	SPN139 Anaerotignum aminivorans_nov_93.173%	SPN225 Acidovorax defluviu_nov_95.946%
SP3 Ligilactobacillus murinus	SPN14 Faecalicatena fissicatena_nov_93.580%	SPN226 Oscillospiraceae [G-3] bacterium MOT-150_nov_91.585%
SP30 Leclercia adecarboxylata	SPN140 Muribaculaceae [G-1] bacterium MOT-129_nov_87.308%	SPN227 Oscillospiraceae [G-2] bacterium MOT-149_nov_95.276%
SP31 Parabacteroides distasonis	SPN141 Lachnospiraceae [G-9] bacterium MOT-174_nov_86.957%	SPN228 Oscillospiraceae [G-2] bacterium MOT-149_nov_93.452%
SP32 Adlercreutzia mucosicola	SPN142 Lachnospiraceae [G-11] bacterium MOT-176_nov_92.885%	SPN229 Adlercreutzia equolifaciens_nov_94.389%
SP33 Acinetobacter lwoffii	SPN143 Ihubacter massiliensis_nov_94.767%	SPN23 Adlercreutzia equolifaciens_nov_96.593%
SP34 Streptococcus danieliae	SPN144 Tyzzerella [Clostridium] colinum_nov_88.494%	SPN230 Eubacteriales [G-4] bacterium MOT-164_nov_97.980%
SP35 Burkholderia lata	SPN145 Planococcus massiliensis_nov_96.992%	SPN231 Lachnospiraceae [G-11] bacterium MOT-176_nov_89.981%
SP36 Phocaeicola sartorii	SPN146 Lachnospiraceae [G-7] bacterium MOT-172_nov_86.508%	SPN232 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_89.44%
SP37 Adlercreutzia muris	SPN147 Lachnospiraceae [G-12] bacterium MOT-180_nov_89.942%	SPN233 Mollicutes [G-2] bacterium MOT-187_nov_93.284%
SP38 Erwinia billingiae	SPN148 Eubacterium coprostanoligenes_nov_91.892%	SPN234 Oscillospiraceae [G-4] bacterium MOT-151_nov_91.732%
SP39 Sphingomonas carotinifaciens	SPN149 Duncaniella freteri_nov_87.759%	SPN235 Oscillospiraceae [G-4] bacterium MOT-151_nov_94.477%
SP4 Limosilactobacillus reuteri	SPN15 Oscillospiraceae [G-2] bacterium MOT-149_nov_93.241%	SPN236 Faecalibaculum rodentium_nov_96.571%
SP40 Clostridium disporicum	SPN150 Lachnospiraceae [G-7] bacterium MOT-172_nov_92.843%	SPN237 Duncaniella freteri_nov_83.895%
SP41 Yokenella regensburgei	SPN151 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.050%	SPN238 Longibaculum muris_nov_90.392%
SP42 Phocaeicola vulgatus	SPN152 Lachnospiraceae [G-7] bacterium MOT-172_nov_86.047%	SPN239 Adlercreutzia muris_nov_89.506%
SP43 Pseudomonas veronii	SPN153 Mediterraneibacter [Ruminococcus] torques_nov_95.200%	SPN24 Cuneatibacter caecimuris_nov_92.486%
SP44 Lachnospiraceae [G-14] bacterium MOT-185	SPN154 Lachnospiraceae [G-2] bacterium MOT-167_nov_89.349%	SPN240 Roseburia hominis_nov_91.715%
SP45 Eubacteriales [G-4] bacterium MOT-164	SPN155 Caproicibacter fermentans_nov_89.824%	SPN241 Lachnospiraceae [G-2] bacterium MOT-167_nov_97.018%
SP46 Eubacteriales [G-2] bacterium MOT-162	SPN156 Fusicatenibacter saccharivorans_nov_90.514%	SPN242 Acetivibrio cellulolyticus_nov_83.644%
SP47 Blastococcus aggregatus	SPN157 Erwinia billingiae_nov_97.519%	SPN243 Mailhella massiliensis_nov_89.888%
SP48 Acinetobacter johnsonii	SPN158 Lachnospiraceae [G-3] bacterium MOT-168_nov_95.059%	SPN244 Faecalimonas umbilicata_nov_92.692%
SP49 Massilia arenae	SPN159 Lachnoclostridium [Clostridium] scindens_nov_89.827%	SPN245 Anaerotignum aminivorans_nov_92.600%
SP5 Kocuria indica	SPN16 Lachnospiraceae [G-11] bacterium MOT-178_nov_94.220%	SPN246 Blautia producta_nov_95.174%
SP50 Bacteroides acidifaciens	SPN160 Lachnospiraceae [G-14] bacterium MOT-182_nov_89.200%	SPN247 Pseudoflavonifractor phocaeensis_nov_92.131%
SP51 Pseudomonas cedrina	SPN161 Eubacterium oxidoreducens_nov_88.846%	SPN25 Acetivibrio cellulolyticus_nov_84.058%
SP52 Siccibacter turicensis	SPN162 Lachnospiraceae [G-11] bacterium MOT-176_nov_94.798%	SPN26 Blautia faecicola_nov_89.709%
SP53 Pelomonas saccharophila	SPN163 Saccharibacillus kuerlensis_nov_95.336%	SPN27 Eisenbergiella massiliensis_nov_92.969%
SP54 Parasutterella excrementihominis	SPN164 Acutalibacter muris_nov_94.264%	SPN28 Lachnospiraceae [G-14] bacterium MOT-185_nov_93.333%
SP55 Eubacteriales [G-1] bacterium MOT-159	SPN165 Phoea massiliensis_nov_90.239%	SPN29 Lachnospiraceae [G-14] bacterium MOT-182_nov_87.160%
SP59 Parabacteroides goldsteinii	SPN166 Duncaniella freteri_nov_89.077%	SPN30 Alloprevotella sp. HMT_473_nov_90.177%
SP6 Eubacteriales [G-3] bacterium MOT-163	SPN167 Fusicatenibacter saccharivorans_nov_91.018%	SPN3 Lactobacillus gasserii_nov_93.345%
SP60 Pedobacter quisquiliarum	SPN168 Lachnospiraceae [G-9] bacterium MOT-174_nov_90.234%	SPN31 Eubacteriales [G-1] bacterium MOT-159_nov_93.976%
SP61 Helicobacter ganmani	SPN169 Adlercreutzia caecimuris_nov_95.382%	SPN32 Lachnospiraceae [G-5] bacterium MOT-170_nov_97.614%
SP62 Lachnospiraceae [G-1] bacterium MOT-166	SPN17 Breznakia pachnodae_nov_81.284%	SPN33 Eisenbergiella massiliensis_nov_87.669%
SP63 Parvibacter caecicola	SPN170 Oscillospiraceae [G-4] bacterium MOT-151_nov_93.491%	SPN34 Oscillospiraceae [G-4] bacterium MOT-151_nov_95.858%
SP64 Blastococcus saxosidens	SPN171 Lachnospiraceae [G-11] bacterium MOT-176_nov_95.174%	SPN38 Lacrimispora xylanolytica_nov_94.314%
SP66 Sphingomonas echinoides	SPN172 Faecalicatena fissicatena_nov_94.521%	SPN4 Lachnospiraceae [G-11] bacterium MOT-178_nov_91.715%
SP67 Pseudomonas helleri	SPN173 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.664%	SPN43 Muribaculaceae [G-1] bacterium MOT-129_nov_89.768%
SP68 Afipia broomeae	SPN174 Saccharibacillus kuerlensis_nov_95.709%	SPN49 Bacteroides uniformis_nov_95.594%
SP69 Lachnospiraceae [G-3] bacterium MOT-168	SPN175 Clostridium collagenovorans_nov_83.466%	SPN5 Mollicutes [G-2] bacterium MOT-187_nov_90.841%
SP7 Erysipelatoclostridium [Clostridium] cocleatum	SPN176 Lachnospiraceae [G-13] bacterium MOT-181_nov_91.634%	SPN54 Lachnospiraceae [G-11] bacterium MOT-178_nov_93.064%
SP70 Pseudomonas extrem australis	SPN177 Sporosalibacterium tautonense_nov_82.659%	SPN6 Eisenbergiella massiliensis_nov_87.308%
SP71 Klebsiella michiganensis	SPN178 Absiella tortuosum_nov_88.725%	SPN60 Lacrimispora xylanolytica_nov_91.992%
SP72 Microbacterium kitamiense	SPN179 Adlercreutzia caecimuris_nov_92.277%	SPN64 Massilia arenae_nov_97.885%
SP73 Enterobacter cloacae	SPN18 Lawsonibacter asaccharolyticus_nov_92.514%	SPN7 Oscillospiraceae [G-2] bacterium MOT-149_nov_94.235%
SP74 Oscillospiraceae [G-7] bacterium MOT-154	SPN180 Oscillibacter valericigenes_nov_95.164%	SPN72 Eisenbergiella massiliensis_nov_90.805%
SP75 Pararheinheimera mesophila	SPN181 Lachnospiraceae [G-14] bacterium MOT-183_nov_97.967%	SPN75 Lachnoclostridium [Clostridium] aminophilum_nov_87.476%
SP76 Enterobacter hormaechei	SPN182 Breznakia pachnodae_nov_83.181%	SPN79 Lachnospiraceae [G-6] bacterium MOT-171_nov_94.643%
SP77 Streptococcus sanguinis	SPN183 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_88.462%	SPN8 Lacrimispora xylanolytica_nov_92.308%
SP8 Lactobacillus johnsonii	SPN184 Lachnospiraceae [G-3] bacterium MOT-168_nov_95.050%	SPN80 Flavobacterium branchiicola_nov_96.282%
SP9 Lactobacillus gasserii	SPN185 Parabacteroides merdae_nov_93.182%	SPN81 Oscillospiraceae [G-2] bacterium MOT-149_nov_95.050%
SPN1 Lachnospiraceae [G-6] bacterium MOT-171_nov_93.307%	SPN186 Hathewayia proteolytica_nov_84.970%	SPN82 Anaeroplasmata abactoclasticum_nov_87.352%
SPN10 Oscillibacter valericigenes_nov_93.654%	SPN187 Eubacterium ramulus_nov_89.362%	SPN83 Faecalicatena orotica_nov_92.218%
SPN100 Lachnospiraceae [G-1] bacterium MOT-166_nov_95.661%	SPN188 Eisenbergiella massiliensis_nov_86.127%	SPN84 Lachnoclostridium [Clostridium] scindens_nov_88.247%
SPN101 Lachnospiraceae [G-11] bacterium MOT-176_nov_95.543%	SPN189 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.751%	SPN85 Oscillospiraceae [G-3] bacterium MOT-150_nov_93.910%
SPN102 Bacteroidetes [G-3] bacterium HMT_436_nov_85.575%	SPN19 Blautia producta_nov_95.164%	SPN86 Lachnospiraceae [G-14] bacterium MOT-185_nov_93.517%
SPN103 Lachnospiraceae [G-3] bacterium MOT-168_nov_94.059%	SPN190 Lachnospiraceae [G-7] bacterium MOT-172_nov_94.831%	SPN87 Lachnospiraceae [G-6] bacterium MOT-171_nov_95.050%
SPN104 Lachnospiraceae [G-14] bacterium MOT-185_nov_92.105%	SPN191 Dechloromonas denitrificans_nov_97.446%	SPN88 Olsenella phocaeensis_nov_92.172%
SPN105 Lachnospiraceae [G-6] bacterium MOT-171_nov_95.644%	SPN192 Blautia producta_nov_96.132%	SPN89 Eisenbergiella massiliensis_nov_88.292%
SPN106 Prevotella shahii_nov_87.242%	SPN193 Lachnospiraceae [G-3] bacterium MOT-168_nov_96.252%	SPN90 Muricoccus intestinalis_nov_89.921%