

Species

SP10 Oscillospiraceae [G-7] bacterium_MOT-154	SPN136 Devosia geojerensis nov. 84.440%	SPN35 Duncaniella freteri nov. 93.573%
SP11 Peptococcaceae [G-1] bacterium_MOT-146	SPN137 Lachnospiraceae [G-11] bacterium_MOT-177 nov. 92.717%	SPN36 Turicibacter sanguinis nov. 95.833%
SP12 Lactobacillus johnsonii	SPN138 Anaeromassilibacillus senegalensis nov. 85.602%	SPN37 Tidjanibacter massiliensis nov. 89.583%
SP14 Eubacteriales [G-4] bacterium_MOT-164	SPN139 Anaerocolumna cellulositytica nov. 90.504%	SPN38 Parabacteroides merdae nov. 93.182%
SP15 Erysipelatoclostridium [Clostridium] cocleatum	SPN14 Muribaculaceae [G-2] bacterium_MOT-104 nov. 91.429%	SPN39 Mucispirillum schaedleri nov. 93.124%
SP16 Ileibacterium valens	SPN140 Lawsonibacter asaccharolyticus nov. 91.762%	SPN40 Saccharibacteria (TM7) [G-3] bacterium_HMT_351 nov. 93.617%
SP17 Lachnospiraceae [G-7] bacterium_MOT-172	SPN141 Sporosalibacterium tautonense nov. 82.852%	SPN41 Parabacteroides distasonis nov. 97.706%
SP18 Eubacteriales [G-1] bacterium_MOT-144	SPN142 Duncaniella freteri nov. 87.006%	SPN42 Lachnospiraceae [G-6] bacterium_MOT-171 nov. 93.651%
SP19 Parvibacter caecicola	SPN143 Oscillospiraceae [G-2] bacterium_MOT-149 nov. 93.874%	SPN43 Duncaniella freteri nov. 87.290%
SP2 Erysipelotrichaceae [G-1] bacterium_MOT-189	SPN144 Muribaculaceae [G-2] bacterium_MOT-104 nov. 87.262%	SPN44 Muribaculaceae [G-1] bacterium_MOT-129 nov. 86.398%
SP20 Lactobacillus intestinalis	SPN145 Lachnospiraceae [G-11] bacterium_MOT-177 nov. 94.094%	SPN45 Muribaculaceae [G-2] bacterium_MOT-104 nov. 90.114%
SP21 Faecalibaculum rodentium	SPN146 Oscillospiraceae [G-4] bacterium_MOT-151 nov. 93.688%	SPN46 Duncaniella freteri nov. 89.366%
SP22 Bacteroides acidifaciens	SPN147 Erysipelothrix rhusiopathiae nov. 85.902%	SPN47 Muribaculaceae [G-1] bacterium_MOT-129 nov. 87.838%
SP23 Lachnospiraceae [G-14] bacterium_MOT-185	SPN148 Lachnospiraceae [G-6] bacterium_MOT-171 nov. 94.280%	SPN48 Desulfovibrio fairfieldensis nov. 89.555%
SP24 Eubacteriales [G-2] bacterium_MOT-162	SPN149 Acutalibacter muris nov. 94.828%	SPN49 Clostridium collagenovorans nov. 81.460%
SP25 Lactobacillus gasseri	SPN15 Faecalicatena fissicatena nov. 94.542%	SPN50 Kineothrix alysoides nov. 87.327%
SP26 Akkermansia muciniphila	SPN150 Oscillospiraceae [G-2] bacterium_MOT-149 nov. 94.455%	SPN51 Lachnoclostridium [Clostridium] polysaccharolyticum nov. 90.522%
SP27 Adlercreutzia muris	SPN151 Marvinbryantia formatexigens nov. 91.358%	SPN52 Muribaculaceae [G-2] bacterium_MOT-104 nov. 89.060%
SP28 Helicobacter ganmani	SPN152 Eisenbergiella massiliensis nov. 88.100%	SPN53 Ihubacter massiliensis nov. 94.788%
SP29 Phocaecicola sartorii	SPN153 Duncaniella freteri nov. 87.453%	SPN54 Longibaculum muris nov. 87.574%
SP3 Prevotella sp. MOT-128	SPN154 Eisenbergiella massiliensis nov. 87.838%	SPN55 Alistipes putredinis nov. 92.776%
SP30 Alistipes sp. MOT-127	SPN155 Phocaea massiliensis nov. 86.680%	SPN56 Muribaculaceae [G-1] bacterium_MOT-129 nov. 91.279%
SP31 Oscillospiraceae [G-3] bacterium_MOT-150	SPN156 Muribaculaceae [G-2] bacterium_MOT-104 nov. 89.504%	SPN57 Muribaculaceae [G-2] bacterium_MOT-104 nov. 89.792%
SP32 Acutalibacter muris	SPN157 Lachnospiraceae [G-12] bacterium_MOT-180 nov. 91.456%	SPN58 Lachnospiraceae [G-3] bacterium_MOT-168 nov. 95.455%
SP4 Staphylococcus saprophyticus	SPN158 Lachnospiraceae [G-5] bacterium_MOT-170 nov. 97.614%	SPN59 Eubacteriales [G-1] bacterium_MOT-161 nov. 93.333%
SP5 Muribaculaceae [G-1] bacterium_MOT-129	SPN159 Lachnospiraceae [G-11] bacterium_MOT-177 nov. 97.642%	SPN60 Mucispirillum schaedleri nov. 94.862%
SP6 Mammaliicoccus lentus	SPN160 Adlercreutzia caecimuris nov. 94.177%	SPN61 Lachnospiraceae [G-14] bacterium_MOT-185 nov. 95.935%
SP7 Ligilactobacillus murinus	SPN161 Adlercreutzia caecimuris nov. 89.379%	SPN62 Muribaculaceae [G-1] bacterium_MOT-129 nov. 85.797%
SP8 Limosilactobacillus reuteri	SPN162 Lachnoclostridium [Clostridium] polysaccharolyticum nov. 90.347%	SPN63 Lachnospiraceae [G-3] bacterium_MOT-168 nov. 94.862%
SP9 Lachnospiraceae [G-11] bacterium_MOT-178	SPN163 Hydrogenoanaerobacterium saccharovorans nov. 88.781%	SPN64 Lachnospiraceae [G-3] bacterium_MOT-168 nov. 97.624%
SPN100 Kineothrix alysoides nov. 90.079%	SPN164 Adlercreutzia caecimuris nov. 92.644%	SPN65 Adlercreutzia caecimuris nov. 89.442%
SPN101 Lacrimispora xylanolytica nov. 93.969%	SPN165 Oscillospiraceae [G-3] bacterium_MOT-150 nov. 91.602%	SPN66 Pseudoflavonifractor phocaeensis nov. 86.513%
SPN102 Kiloniella majae nov. 85.124%	SPN166 Lachnospiraceae [G-6] bacterium_MOT-171 nov. 95.446%	SPN67 Bacteroidetes [G-3] bacterium_HMT_436 nov. 86.275%
SPN103 Duncaniella freteri nov. 90.772%	SPN167 Caproicibacter fermentans nov. 87.354%	SPN68 Muribaculaceae [G-2] bacterium_MOT-104 nov. 90.114%
SPN104 Eubacterium ramulus nov. 87.814%	SPN168 Oscillospiraceae [G-2] bacterium_MOT-149 nov. 93.333%	SPN69 Bacteroides uniformis nov. 95.785%
SPN105 Longibaculum muris nov. 90.820%	SPN169 Roseburia intestinalis nov. 92.941%	SPN70 Neglectibacter timonensis nov. 95.076%
SPN106 Adlercreutzia equolifaciens nov. 90.020%	SPN170 Oscillospiraceae [G-2] bacterium_MOT-149 nov. 94.257%	SPN71 Muribaculaceae [G-1] bacterium_MOT-129 nov. 88.610%
SPN107 Oscillospiraceae [G-3] bacterium_MOT-150 nov. 91.552%	SPN171 Clostridiales [F-1][G-1] bacterium_HMT_093 nov. 84.091%	SPN72 Duncaniella freteri nov. 89.700%
SPN108 Lachnospiraceae [G-11] bacterium_MOT-178 nov. 93.642%	SPN172 Clostridiales [F-1][G-1] bacterium_HMT_093 nov. 84.091%	SPN73 Duncaniella freteri nov. 87.030%
SPN109 Culturomicrobia massiliensis nov. 89.792%	SPN173 Alpinimonas psychrophila nov. 82.549%	SPN74 Lachnoclostridium [Clostridium] polysaccharolyticum nov. 92.070%
SPN11 Kiloniella majae nov. 87.397%	SPN174 Eubacteriales [G-3] bacterium_MOT-163 nov. 81.921%	SPN75 Duncaniella freteri nov. 88.213%
SPN110 Muribaculaceae [G-1] bacterium_MOT-129 nov. 89.961%	SPN175 Mollicutes [G-2] bacterium_MOT-187 nov. 93.246%	SPN76 Oscillospiraceae [G-3] bacterium_MOT-150 nov. 93.267%
SPN111 Duncaniella freteri nov. 89.552%	SPN176 Duncaniella freteri nov. 88.827%	SPN77 Saccharibacteria (TM7) [G-3] bacterium_HMT_351 nov. 95.351%
SPN112 Lachnospiraceae [G-11] bacterium_MOT-176 nov. 94.990%	SPN177 Anaerotignum lactatifermentans nov. 96.139%	SPN78 Lachnoclostridium [Clostridium] polysaccharolyticum nov. 89.555%
SPN113 Eubacteriales [G-4] bacterium_MOT-164 nov. 97.172%	SPN178 Faecalicatena otitica nov. 92.218%	SPN79 Ruminoclostridium cellulolyticum nov. 83.433%
SPN114 Lachnospiraceae [G-11] bacterium_MOT-176 nov. 95.174%	SPN179 Lachnoclostridium [Clostridium] polysaccharolyticum nov. 88.610%	SPN80 Duncaniella freteri nov. 89.434%
SPN115 Eubacteriales [G-3] bacterium_MOT-163 nov. 81.321%	SPN180 Odoribacter splanchnicus nov. 92.146%	SPN81 Longibaculum muris nov. 90.588%
SPN116 Eubacteriales [G-1] bacterium_MOT-159 nov. 92.323%	SPN181 Mailhella massiliensis nov. 89.888%	SPN82 Mailhella massiliensis nov. 90.377%
SPN117 Oscillospiraceae [G-4] bacterium_MOT-151 nov. 96.040%	SPN182 Eggerthella timonensis nov. 89.222%	SPN83 Lacrimispora saccharolytica nov. 92.621%
SPN118 Duncaniella freteri nov. 86.278%	SPN183 Phocaea massiliensis nov. 90.239%	SPN84 Eubacterium coprostanoligenes nov. 91.892%
SPN119 Duncaniella freteri nov. 86.456%	SPN190 Alistipes putredinis nov. 94.847%	SPN85 Alistipes senegalensis nov. 93.690%
SPN120 Muribaculaceae [G-1] bacterium_MOT-129 nov. 87.115%	SPN191 Muribaculaceae [G-2] bacterium_MOT-104 nov. 89.354%	SPN86 Bacteroidetes [G-3] bacterium_HMT_436 nov. 85.938%
SPN121 Pseudoflavonifractor phocaeensis nov. 93.077%	SPN2 Duncaniella freteri nov. 86.679%	SPN87 Lachnospiraceae [G-12] bacterium_MOT-179 nov. 94.971%
SPN122 Lachnospiraceae [G-14] bacterium_MOT-185 nov. 93.333%	SPN201 Duncaniella freteri nov. 96.591%	SPN88 Faecalicatena fissicatena nov. 93.933%
SPN123 Duncaniella freteri nov. 95.057%	SPN21 Duncaniella freteri nov. 90.038%	SPN89 Lachnospiraceae [G-7] bacterium_MOT-172 nov. 93.617%
SPN124 Muribaculaceae [G-2] bacterium_MOT-104 nov. 85.741%	SPN22 Bacteroidetes [G-3] bacterium_HMT_436 nov. 86.078%	SPN90 Lachnospiraceae [G-14] bacterium_MOT-182 nov. 90.669%
SPN125 Lachnospiraceae [G-2] bacterium_MOT-167 nov. 97.012%	SPN23 Duncaniella freteri nov. 87.759%	SPN91 Alistipes finegoldii nov. 94.073%
SPN126 Duncaniella freteri nov. 88.180%	SPN24 Parasutterella excrementihominis nov. 94.778%	SPN92 Alloprevotella sp. HMT_473 nov. 90.215%
SPN127 Eubacteriales [G-4] bacterium_MOT-165 nov. 93.763%	SPN25 Alistipes senegalensis nov. 93.846%	SPN93 Eisenbergiella massiliensis nov. 87.548%
SPN128 Agathobaculum desmolans nov. 91.262%	SPN26 Oscillospiraceae [G-6] bacterium_MOT-153 nov. 91.870%	SPN94 Lachnospiraceae [G-13] bacterium_MOT-181 nov. 91.602%
SPN129 Magnetovibrio blakemorei nov. 83.438%	SPN27 Duncaniella freteri nov. 88.972%	SPN95 Eisenbergiella massiliensis nov. 90.805%
SPN130 Lachnospiraceae [G-6] bacterium_MOT-171 nov. 95.618%	SPN28 Lachnospiraceae [G-12] bacterium_MOT-180 nov. 94.757%	SPN96 Mollicutes [G-2] bacterium_MOT-187 nov. 95.701%
SPN131 Parvularcula lutaonensis nov. 83.610%	SPN29 Duncaniella freteri nov. 87.709%	SPN97 Lutyricoccus pullicacorum nov. 86.320%
SPN132 Muribaculaceae [G-2] bacterium_MOT-104 nov. 89.981%	SPN30 Duncaniella freteri nov. 88.368%	SPN98 Lachnospiraceae [G-13] bacterium_MOT-181 nov. 85.393%
SPN133 Algimona porphyrae nov. 79.733%	SPN31 Duncaniella freteri nov. 90.530%	SPN99 Anaerotignum lactatifermentans nov. 95.560%
SPN134 Adlercreutzia muris nov. 92.245%	SPN32 Olsenella phocaeensis nov. 92.172%	SPPN1 Gloeobacter multispecies sppn1_2 nov. 82.056%
SPN135 Faecalimonas umhilicata nov. 95.358%	SPN33 Odoribacter splanchnicus nov. 92.337%	SPPN2 Prevotella multispecies sppn2_2 nov. 89.792%
	SPN34 Lachnospiraceae [G-14] bacterium_MOT-185 nov. 94.920%	SPPN3 multigenus multispecies sppn3_2 nov. 93.709%