

Species

SP17 Lactococcus lactis	SPN144 Lachnospiraceae [G-14] bacterium_MOT-182_nov_89.200%	SPN219 Adlercreutzia muris_nov_89.506%
SP18 Parabacteroides goldsteinii	SPN145 Eubacterium oxioreducens_nov_88.846%	SPN222 Lachnospiraceae [G-11] bacterium_MOT-176_nov_94.981%
SP19 Eubacteriales [G-1] bacterium_MOT-159	SPN146 Duncaniella freteri_nov_89.077%	SPN220 Lachnospiraceae [G-2] bacterium_MOT-167_nov_97.018%
SP2 Erysipelatoclostridium [Clostridium] coeleatum	SPN147 Acutalibacter muris_nov_94.264%	SPN221 Roseburia hominis_nov_91.715%
SP20 Eubacteriales [G-3] bacterium_MOT-163	SPN148 Phoea massiliensis_nov_90.239%	SPN222 Acetivibrio cellulolyticus_nov_83.644%
SP21 Eubacteriales [G-2] bacterium_MOT-162	SPN149 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_88.462%	SPN223 Mailhella massiliensis_nov_89.888%
SP22 Streptococcus danieliae	SPN15 Oscillospiraceae [G-3] bacterium_MOT-150_nov_93.910%	SPN224 Faecalimonas umbilicata_nov_92.692%
SP23 Lactococcus cremoris	SPN150 Lachnospiraceae [G-13] bacterium_MOT-181_nov_91.634%	SPN225 Anaerostignum aminivorans_nov_92.600%
SP24 Bacteroides acidifaciens	SPN151 Muribaculaceae [G-2] bacterium_MOT-104_nov_89.905%	SPN226 Blautia producta_nov_95.174%
SP25 Duncaniella freteri	SPN152 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.841%	SPN227 Lachnospiraceae [G-11] bacterium_MOT-178_nov_93.064%
SP26 Lactobacillus gasseri	SPN153 Anaerostignum aminivorans_nov_92.585%	SPN228 Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.307%
SP27 Microbacterium kitamiense	SPN154 Faecalicatena orotica_nov_94.553%	SPN229 Mollicutes [G-2] bacterium_MOT-187_nov_90.841%
SP28 Enterococcus gallinarum	SPN155 Lhubacter massiliensis_nov_94.402%	SPN23 Anaerotruncus rubifantans_nov_87.931%
SP29 Parvibacter caecicola	SPN156 Clostridium collagenovorans_nov_83.466%	SPN230 Lachnospiraceae [G-11] bacterium_MOT-178_nov_91.715%
SP3 Ligilactobacillus murinus	SPN157 Sporosalibacterium tautonense_nov_82.659%	SPN231 Alloprevotella sp._HMT_473_nov_90.177%
SP30 Helicobacter ganmani	SPN158 Absiella tortuosum_nov_88.725%	SPN232 Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.254%
SP31 Lachnospiraceae [G-3] bacterium_MOT-168	SPN159 Adlercreutzia caecimuris_nov_92.277%	SPN233 Eisenbergiella massiliensis_nov_87.308%
SP32 Lachnospiraceae [G-14] bacterium_MOT-185	SPN16 Oscillospiraceae [G-4] bacterium_MOT-151_nov_95.858%	SPN234 Lacrimispora xylanolytica_nov_92.308%
SP33 Lachnospiraceae [G-1] bacterium_MOT-166	SPN160 Oscillibacter valericigenes_nov_95.164%	SPN235 Muricomes intestini_nov_89.921%
SP34 Eubacteriales [G-4] bacterium_MOT-164	SPN161 Faecalicatena fissicatena_nov_94.521%	SPN236 Oscillibacter valericigenes_nov_93.654%
SP35 Oscillospiraceae [G-7] bacterium_MOT-154	SPN162 Lachnospiraceae [G-14] bacterium_MOT-183_nov_97.967%	SPN237 Kineothrix alysoides_nov_87.352%
SP36 Phocaeicola vulgatus	SPN163 Lachnospiraceae [G-3] bacterium_MOT-168_nov_95.050%	SPN238 Lachnoclostridium [Clostridium] aminophilum_nov_87.476%
SP37 Adlercreutzia mucosicola	SPN164 Faecalicatena fissicatena_nov_94.129%	SPN24 Hydrogenoanaerobacterium saccharovorans_nov_88.439%
SP4 Romboutsia ilealis	SPN165 Breznakia pachnodae_nov_83.181%	SPN26 Blautia faecicola_nov_89.709%
SP40 Microbacterium foliorum	SPN166 Hathewayia proteolytica_nov_84.970%	SPN3 Lachnospiraceae [G-11] bacterium_MOT-178_nov_94.220%
SP41 Oscillospiraceae [G-6] bacterium_MOT-153	SPN167 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.751%	SPN32 Lawsonibacter asaccharolyticus_nov_90.421%
SP42 Acetatifactor muris	SPN168 Parabacteroides merdae_nov_93.182%	SPN38 Anaeroplasmata abactoclasticum_nov_87.352%
SP43 Streptococcus sp._MOT-012	SPN169 Eubacterium ramulus_nov_89.362%	SPN4 Breznakia pachnodae_nov_81.284%
SP5 Adlercreutzia muris	SPN17 Lactobacillus gasseri_nov_93.345%	SPN42 Lachnoclostridium [Clostridium] aminophilum_nov_89.961%
SP6 Lactobacillus johnsonii	SPN170 Eisenbergiella massiliensis_nov_86.127%	SPN49 Bacteroides uniformis_nov_95.594%
SP7 Adlercreutzia caecimuris	SPN171 Lachnospiraceae [G-7] bacterium_MOT-172_nov_94.831%	SPN5 Lawsonibacter asaccharolyticus_nov_92.514%
SP8 Limosilactobacillus reuteri	SPN172 Faecalibaculum rodentium_nov_96.571%	SPN53 Hydrogenoanaerobacterium saccharovorans_nov_88.395%
SP9 Parabacteroides distasonis	SPN173 Lachnospiraceae [G-3] bacterium_MOT-168_nov_96.252%	SPN6 Cuneatibacter caecimuris_nov_92.486%
SPN1 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.241%	SPN174 Lachnospiraceae [G-9] bacterium_MOT-174_nov_92.216%	SPN60 Lacrimispora xylanolytica_nov_91.992%
SPN10 Lachnospiraceae [G-14] bacterium_MOT-185_nov_93.333%	SPN175 Adlercreutzia caecimuris_nov_92.644%	SPN63 Lachnospiraceae [G-12] bacterium_MOT-179_nov_94.737%
SPN100 Eubacterium ramulus_nov_91.296%	SPN176 Lachnoclostridium [Clostridium] scindens_nov_90.211%	SPN66 Eisenbergiella massiliensis_nov_90.805%
SPN101 Lachnospiraceae [G-3] bacterium_MOT-168_nov_94.653%	SPN177 Duncaniella freteri_nov_88.598%	SPN67 Lacrimispora xylanolytica_nov_94.314%
SPN102 Lawsonibacter asaccharolyticus_nov_91.571%	SPN178 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.636%	SPN68 Lachnospiraceae [G-11] bacterium_MOT-177_nov_95.874%
SPN103 Adlercreutzia caecimuris_nov_94.000%	SPN179 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.083%	SPN69 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.643%
SPN104 Lancefieldella parvula_nov_91.552%	SPN18 Lachnospiraceae [G-5] bacterium_MOT-170_nov_97.614%	SPN7 Acetivibrio cellulolyticus_nov_84.058%
SPN105 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.861%	SPN180 Lachnospiraceae [G-11] bacterium_MOT-178_nov_95.358%	SPN70 Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.050%
SPN106 Lachnospiraceae [G-9] bacterium_MOT-174_nov_90.234%	SPN181 Faecalicatena fissicatena_nov_95.499%	SPN71 Eisenbergiella massiliensis_nov_88.697%
SPN107 Faecalibaculum rodentium_nov_96.571%	SPN182 Lachnospiraceae [G-14] bacterium_MOT-185_nov_94.726%	SPN72 Bacteroidetes [G-3] bacterium_HMT_436_nov_85.575%
SPN108 Prevotella shahii_nov_87.054%	SPN183 Pseudoflavonifractor phocaeensis_nov_92.131%	SPN73 Lachnoclostridium [Clostridium] scindens_nov_88.247%
SPN109 Meditteraneibacter [Ruminococcus] torques_nov_94.212%	SPN184 Hydrogenoanaerobacterium saccharovorans_nov_90.522%	SPN74 Muribaculaceae [G-1] bacterium_MOT-129_nov_87.308%
SPN11 Lachnospiraceae [G-14] bacterium_MOT-185_nov_93.517%	SPN185 Pseudoflavonifractor phocaeensis_nov_86.756%	SPN75 Lachnospiraceae [G-11] bacterium_MOT-176_nov_95.155%
SPN110 Anaerotruncus rubifantans_nov_91.506%	SPN186 Phoea massiliensis_nov_90.060%	SPN76 Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.050%
SPN111 Sporobacter termitidis_nov_82.970%	SPN187 Lachnoclostridium phytofermentans_nov_89.057%	SPN77 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_87.308%
SPN112 Anaerostignum lactatifermentans_nov_95.577%	SPN188 Eubacterium coprostanoligenes_nov_90.476%	SPN78 Olsenella phocaeensis_nov_92.172%
SPN113 Leifsonia kafniensis_nov_84.158%	SPN189 Duncaniella freteri_nov_89.013%	SPN79 Roseburia hominis_nov_92.471%
SPN114 Lachnospiraceae [G-7] bacterium_MOT-172_nov_93.204%	SPN19 Eubacteriales [G-1] bacterium_MOT-159_nov_93.976%	SPN8 Roseburia inulinivorans_nov_93.786%
SPN115 Lachnospiraceae [G-7] bacterium_MOT-172_nov_91.945%	SPN190 Clostridiales [F-1][G-1] bacterium_HMT_093_nov_84.298%	SPN80 Parabacteroides distasonis_nov_97.514%
SPN116 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.353%	SPN191 Oscillospiraceae [G-4] bacterium_MOT-151_nov_94.477%	SPN81 Lachnospiraceae [G-14] bacterium_MOT-182_nov_92.245%
SPN117 Adlercreutzia caecimuris_nov_95.382%	SPN192 Clostridiales [F-1][G-1] bacterium_HMT_093_nov_91.892%	SPN82 Faecalicatena orotica_nov_92.218%
SPN118 Duncaniella freteri_nov_90.152%	SPN193 Neglectibacter timonensis_nov_94.318%	SPN83 Eubacterium xylanophilum_nov_91.149%
SPN119 Mailhella massiliensis_nov_90.377%	SPN194 Eisenbergiella massiliensis_nov_87.669%	SPN84 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.050%
SPN12 Eisenbergiella massiliensis_nov_92.969%	SPN195 Duncaniella freteri_nov_87.453%	SPN85 Lachnospiraceae [G-1] bacterium_MOT-166_nov_95.661%
SPN120 Duncaniella freteri_nov_87.896%	SPN196 Saccharibacteria (TM7) [G-3] bacterium_HMT_351_nov_95.551%	SPN86 Longibaculum muris_nov_93.910%
SPN121 Parasutterella excrementihominis_nov_94.584%	SPN197 Butyrivibrio sp._HMT_455_nov_83.556%	SPN87 Parabacteroides distasonis_nov_97.323%
SPN122 Phoea massiliensis_nov_90.297%	SPN198 Muribaculaceae [G-2] bacterium_MOT-104_nov_89.544%	SPN88 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.105%
SPN123 Pseudoflavonifractor phocaeensis_nov_95.761%	SPN199 Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.640%	SPN89 Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.644%
SPN124 Anaerostignum aminivorans_nov_93.173%	SPN2 Lachnospiraceae [G-11] bacterium_MOT-176_nov_94.798%	SPN9 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.097%
SPN125 Blautia producta_nov_96.132%	SPN20 Anaerotruncus rubifantans_nov_92.636%	SPN90 Turicibacter sanguinis_nov_95.817%
SPN126 Lachnospiraceae [G-9] bacterium_MOT-174_nov_86.957%	SPN200 Faecalicatena fissicatena_nov_93.810%	SPN91 Bariatricus massiliensis_nov_93.037%
SPN127 Lachnospiraceae [G-11] bacterium_MOT-176_nov_92.885%	SPN201 Sporobacter termitidis_nov_88.294%	SPN92 Clostridium collagenovorans_nov_80.952%
SPN128 Oscillospiraceae [G-4] bacterium_MOT-151_nov_93.491%	SPN202 Eubacteriales [G-1] bacterium_MOT-159_nov_88.550%	SPN93 Eubacterium ventriosum_nov_92.843%
SPN129 Tyzzerella [Clostridium] colinum_nov_88.494%	SPN203 Lacrimispora xylanolytica_nov_93.969%	SPN94 Mucispirillum schaedleri_nov_93.110%