

SP40	Haemophilus parainfluenzae	SPN176	Lachnoclostridium polysaccharolyticum nov_89.555%	SPN41	Bacteroides capillosus nov_92.871%
SP41	Bradyrhizobium lupini	SPN177	unclassified_Lachnospiraceae sp._str._cont1.79_nov_92.039%	SPN42	Bacteroidetes_[G-7] bacterium_HMT_911_nov_87.452%
SP42	Pseudomonas oryzaehabitans	SPN178	Agathobaculum desmolans nov_91.845%	SPN43	Phoea massiliensis nov_91.519%
SP43	Oxalobacteraceae_[G] sp._Oral_Taxon_C91	SPN179	Phoea massiliensis nov_90.748%	SPN44	Faecalibaculum rodentium nov_84.952%
SP44	Turicella otitidis	SPN180	Dorea longicatena nov_94.563%	SPN45	Eubacterium siraeum nov_91.566%
SP46	Roseburia inulinivorans	SPN181	unclassified_Ruminococcaceae sp._str._D16_nov_92.898%	SPN46	Phoea massiliensis nov_90.675%
SP5	Streptococcus thoraltensis	SPN182	Lactobacillus murinus nov_94.528%	SPN47	Eubacterium coprostanoligenes nov_91.546%
SP50	Streptococcus infantis	SPN182	unclassified_Ruminococcaceae sp._str._D16_nov_90.249%	SPN48	Bacteroidetes_[G-7] bacterium_HMT_911_nov_87.476%
SP51	Paenibacillus campinasensis	SPN183	Anaeroplasma abactoclasticum nov_86.957%	SPN49	Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.616%
SP52	Mycobacterium cookii	SPN184	Marvinbryantia formatexigens nov_91.975%	SPN5	Lachnoclostridium sp._str._M62/1_nov_94.175%
SP53	Acinetobacter pittii	SPN185	Phoea massiliensis nov_89.286%	SPN50	Oscillibacter valericigenes nov_94.208%
SP56	Capnocytophaga sputigena	SPN186	Bacteroidetes_[G-7] bacterium_HMT_911_nov_85.352%	SPN51	Streptococcus danieliae nov_96.516%
SP57	Bifidobacterium longum	SPN187	Hydrogenoanaerobacterium saccharovorans nov_89.341%	SPN52	Fusicatenibacter saccharivorans nov_90.138%
SP6	Bifidobacterium pseudolongum	SPN188	Lachnoclostridium asparagiforme nov_94.347%	SPN53	Lachnoclostridium polysaccharolyticum nov_92.442%
SP60	Streptococcus mitis	SPN189	Anaerocolumna cellulositlytica nov_90.522%	SPN54	Colidextribacter massiliensis nov_89.382%
SP61	Staphylococcus auricularis	SPN19	Lachnoclostridium bolteae nov_94.951%	SPN55	Enterococcus sp._str._F95_nov_95.446%
SP62	Sphingomonas yabuuchiae	SPN190	Alistipes putredinis nov_93.156%	SPN56	Oscillibacter valericigenes nov_91.908%
SP65	Methylobacterium goeisingense	SPN191	Eubacterium siraeum nov_91.849%	SPN57	Barnesiella viscericola nov_82.900%
SP66	Cutibacterium namnetense	SPN192	Blautia producta nov_94.574%	SPN58	unclassified_Ruminococcaceae sp._str._D16_nov_93.269%
SP67	Haemophilus haemolyticus	SPN193	Alistipes ihumii nov_93.905%	SPN59	Bacteroidetes_[G-7] bacterium_HMT_911_nov_82.752%
SP68	Parasutterella excrementihominis	SPN194	Eubacterium siraeum nov_91.984%	SPN6	Negativibacillus massiliensis nov_86.914%
SP69	Lactobacillus ruminis	SPN195	Anaerocolumna cellulositlytica nov_86.863%	SPN60	Lactobacillus oeni nov_87.778%
SP7	Escherichia coli	SPN196	Anaerostipes sp._str._3256FAA_nov_89.017%	SPN61	Parabacteroides distasonis nov_83.426%
SP74	Clostridium disporicum	SPN197	Eisenbergiella massiliensis nov_86.867%	SPN62	Eisenbergiella massiliensis nov_86.705%
SP75	Microbacterium lacticum	SPN198	Acinetobacter johnsonii nov_97.697%	SPN63	Sporobacter termitidis nov_88.492%
SP76	Corynebacterium pyruviciproducens	SPN199	Enterorhabdus muris nov_89.344%	SPN64	Eisenbergiella massiliensis nov_90.631%
SP77	Clostridium saudiense	SPN2	Bacteroidetes_[G-7] bacterium_HMT_911_nov_85.742%	SPN65	NA rectale nov_92.816%
SP78	Staphylococcus hominis	SPN20	Prevotella fusca nov_82.772%	SPN66	Enterorhabdus caecimuris nov_93.800%
SP8	Sediminibacterium aquarii	SPN200	unclassified_Ruminococcaceae sp._str._D16_nov_94.423%	SPN67	Eubacterium xylanophilum nov_91.149%
SP81	Sphingomonas panni	SPN201	Anaeroplasma abactoclasticum nov_87.059%	SPN68	Roseburia hominis nov_92.100%
SP82	Pseudomonas putida	SPN202	Agathobaculum desmolans nov_91.794%	SPN69	Bacteroidetes_[G-7] bacterium_HMT_911_nov_84.423%
SP83	Odoribacter splanchnicus	SPN203	Oscillibacter valericigenes nov_93.798%	SPN7	Algimonas arctica nov_79.276%
SP84	Finegoldia magna	SPN204	Dorea longicatena nov_94.563%	SPN70	Acetivibrio cellulolyticus nov_83.125%
SP9	Enterococcus sp._str._F95	SPN205	Gabonia massiliensis nov_86.704%	SPN71	Leifsonia kafniensis nov_84.692%
SPN1	Lachnoclostridium polysaccharolyticum nov_91.522%	SPN206	Blautia producta nov_94.778%	SPN72	Clostridium merdae nov_91.321%
SPN10	Lachnoclostridium polysaccharolyticum nov_92.829%	SPN207	Ruminiclostridium cellulolyticum nov_82.505%	SPN73	Barnesiella intestinihominis nov_83.554%
SPN100	Eisenbergiella massiliensis nov_90.385%	SPN208	Faecalicoccus acidiformans nov_90.075%	SPN74	Abssiella tortuosum nov_86.716%
SPN101	Blautia marasmi nov_91.650%	SPN209	Muricomes intestini nov_88.998%	SPN75	Neglecta timonensis nov_95.472%
SPN102	Acetivibrio cellulolyticus nov_84.189%	SPN21	Anaerotruncus rubiinfantis nov_82.560%	SPN76	Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.207%
SPN103	Gabonia massiliensis nov_84.877%	SPN210	Bacteroidetes_[G-7] bacterium_HMT_911_nov_85.659%	SPN77	Streptococcus acidominimus nov_94.788%
SPN104	Bacteroidetes_[G-7] bacterium_HMT_911_nov_85.164%	SPN211	Alistipes senegalensis nov_96.905%	SPN78	Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.207%
SPN105	Lachnoclostridium bolteae nov_92.621%	SPN212	Longibaculum muris nov_93.517%	SPN79	Oscillibacter valericigenes nov_92.337%
SPN106	Bacteroides thetaiotaomicron nov_93.762%	SPN213	Barnesiella viscericola nov_83.673%	SPN8	Intestinimonas butyriciproducens nov_95.775%
SPN107	Blautia schinkii nov_93.548%	SPN214	unclassified_Ruminococcaceae sp._str._D16_nov_95.192%	SPN80	Ruminococcaceae_[G-2] bacterium_HMT_085_nov_87.018%
SPN108	Eubacterium hallii nov_90.751%	SPN215	Streptococcus troglodytidis nov_90.257%	SPN81	Eisenbergiella massiliensis nov_92.008%
SPN109	Ruthenibacterium lactaiformans nov_95.769%	SPN216	Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_92.067%	SPN82	Anaerotruncus rubiinfantis nov_81.227%
SPN11	Bacteroidetes_[G-7] bacterium_HMT_911_nov_85.910%	SPN217	Bacteroides capillosus nov_90.421%	SPN83	Hathewayia proteolytica nov_83.768%
SPN110	Enterorhabdus muris nov_92.245%	SPN218	Anaerotruncus rubiinfantis nov_81.262%	SPN84	Anaerotruncus rubiinfantis nov_93.050%
SPN111	Terrimicrobium sacchariphilum nov_81.224%	SPN219	Oscillibacter valericigenes nov_93.617%	SPN85	Tyzzerella colinum nov_87.579%
SPN112	Bacteroidetes_[G-7] bacterium_HMT_911_nov_84.854%	SPN22	Anaerotruncus rubiinfantis nov_80.705%	SPN86	Eubacterium ventriosum nov_93.217%
SPN113	Massilia yuzhufengensis nov_97.395%	SPN220	Anaeromassilibacillus senegalensis nov_92.063%	SPN87	Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_93.050%
SPN114	Colidextribacter massiliensis nov_87.908%	SPN221	Deinococcus aerophilus nov_93.097%	SPN88	Ruminococcaceae_[G-2] bacterium_HMT_085_nov_90.650%
SPN115	Eisenbergiella massiliensis nov_90.133%	SPN222	Ruminococcaceae_[G-2] bacterium_HMT_085_nov_88.798%	SPN89	Prevotella oris nov_82.772%
SPN116	Neglecta timonensis nov_94.497%	SPN223	Bacteroidetes_[G-7] bacterium_HMT_911_nov_86.127%	SPN9	Lachnoclostridium herbivorans nov_91.346%
SPN117	Ruminococcaceae_[G-2] bacterium_HMT_085_nov_85.657%	SPN224	Faecalicatena fissicatena nov_85.660%	SPN90	Alistipes senegalensis nov_94.073%
SPN118	Herbaspirillum huttiense nov_83.151%	SPN225	lhubacter massiliensis nov_90.751%	SPN91	Vampirovibrio chlorellavorus nov_81.520%
SPN119	Roseburia inulinivorans nov_93.282%	SPN226	unclassified_Ruminococcaceae sp._str._D16_nov_90.440%	SPN92	Enterorhabdus caecimuris nov_95.382%
SPN12	Sediminibacterium aquarii nov_92.460%	SPN227	Streptococcus acidominimus nov_94.163%	SPN93	Leifsonia kafniensis nov_84.554%
SPN120	Erysipelatoclostridium sp._str._HGF2_nov_88.971%	SPN228	unclassified_Ruminococcaceae sp._str._D16_nov_92.692%	SPN94	Allobaculum stercoricanis nov_86.139%
SPN121	Oscillibacter valericigenes nov_93.295%	SPN229	Eisenbergiella massiliensis nov_90.267%	SPN95	Roseburia faecalis nov_94.197%
SPN122	Turicella otitidis nov_96.071%	SPN23	Roseburia hominis nov_90.485%	SPN96	unclassified_Lachnospiraceae sp._str._cont1.79_nov_91.489%
SPN123	Bacteroides capillosus nov_91.379%	SPN230	Eisenbergiella massiliensis nov_91.082%	SPN97	unclassified_Lachnospiraceae sp._str._cont1.79_nov_95.155%
SPN124	Butyrivibrio proteoclasticus nov_78.519%	SPN231	Neglecta timonensis nov_94.677%	SPN98	Hydrogenoanaerobacterium saccharovorans nov_88.395%
SPN125	Phoea massiliensis nov_87.354%	SPN232	Ruminococcus champanelensis nov_92.063%	SPN99	Intestinimonas butyriciproducens nov_93.812%
SPN126	Hydrogenoanaerobacterium saccharovorans nov_87.379%	SPN233	Bacteroidetes_[G-7] bacterium_HMT_911_nov_83.172%	SPP1	Cutibacterium_Propionibacterium_acnes
SPN127	Bacteroides massiliensis nov_96.522%	SPN234	Alistipes senegalensis nov_93.846%	SPP10	Burkholderia_cenoccepacia_lata,multivorans,sp._str._383
SPN128	Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_92.706%	SPN235	Eubacterium xylanophilum nov_92.075%	SPP11	Kocuria_rhizophila,sp._Oral_Taxon_B56