

viensis	●	SPN2163	Mucispirillum schaedleri_nov_96.833%	●	SPN584	Papillibacter cinnamivorans_nov_93.795%
ohnsonii	●	SPN2174	Aminipila butyrica_nov_91.627%	●	SPN596	Lachnospiraceae_[G-11]_bacterium_MOT-178_nov_92.089%
ae	●	SPN218	Odoribacter splanchnicus_nov_92.562%	●	SPN60	Actinomarinicola tropica_nov_91.607%
intestinalis	●	SPN2187	Eubacteriales_[G-1]_bacterium_MOT-158_nov_95.204%	●	SPN606	Harryflintia acetispora_nov_93.079%
oides	●	SPN2207	Eubacterium ventriosum_nov_97.108%	●	SPN617	Anaerocolumna cellulositytica_nov_90.000%
ioparus	●	SPN2212	Clostridiales_[F-1][G-1]_bacterium_HMT_093_nov_87.381%	●	SPN628	Lachnospiraceae_[G-12]_bacterium_MOT-179_nov_94.501%
[G-4]_bacterium_MOT-164	●	SPN2218	Holdemania massiliensis_nov_92.483%	●	SPN640	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_93.333%
monticola	●	SPN2230	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_92.414%	●	SPN649	Lachnospiraceae_[G-3]_bacterium_MOT-168_nov_95.208%
er asaccharolyticus	●	SPN2231	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_91.724%	●	SPN660	Marvinbryantia formatexigens_nov_91.435%
n pseudolongum	●	SPN2252	Desulfovibrio fairfieldensis_nov_89.379%	●	SPN672	Povalibacter uvarum_nov_94.104%
sartorii	●	SPN2262	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_92.449%	●	SPN681	Alistipes putredinis_nov_96.782%
aeae_[G-12]_bacterium_MOT-179	●	SPN2275	Nitrosospira multiformis_nov_93.864%	●	SPN683	Roseburia faecis_nov_94.231%
aeae_[G-2]_bacterium_MOT-104	●	SPN228	Caproiciproducens galactitolivorans_nov_89.737%	●	SPN696	Eisenbergiella massiliensis_nov_94.286%
edinis	●	SPN2287	Culturomica massiliensis_nov_91.055%	●	SPN705	Muribaculum intestinale_nov_94.533%
aeae_[G-3]_bacterium_MOT-168	●	SPN2297	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_90.800%	●	SPN715	Piscinibacter defluvii_nov_97.500%
ium [Clostridium] scindens	●	SPN2309	Muribaculaceae_[G-1]_bacterium_MOT-129_nov_86.089%	●	SPN72	Lachnospiraceae_[G-6]_bacterium_MOT-171_nov_95.188%
autersii	●	SPN2326	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_92.874%	●	SPN726	Aggregatilinea lenta_nov_87.081%
ostridium [Clostridium] innocuum	●	SPN2331	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_92.922%	●	SPN735	Adlercreutzia mucosicola_nov_97.826%
rium lactatiformans	●	SPN2337	Eubacteriales_[G-4]_bacterium_MOT-165_nov_97.356%	●	SPN746	Lachnospiraceae_[G-11]_bacterium_MOT-176_nov_97.608%
ostridium [Clostridium] cocleatum	●	SPN2343	Muribaculaceae_[G-1]_bacterium_MOT-129_nov_89.431%	●	SPN747	Luteitalea pratensis_nov_92.135%
aceae_[G-1]_bacterium_MOT-198	●	SPN2354	Eubacteriales_[G-3]_bacterium_MOT-163_nov_93.765%	●	SPN757	Lacrimispora xylanolytica_nov_93.789%
us murinus	●	SPN2365	Povalibacter uvarum_nov_90.909%	●	SPN767	Hydrogenoanaerobacterium saccharovorans_nov_93.735%
mucosicola	●	SPN2376	Lachnospiraceae_[G-11]_bacterium_MOT-176_nov_94.523%	●	SPN779	Paludibaculum fermentans_nov_91.388%
acidifaciens	●	SPN238	Eubacteriales_[G-3]_bacterium_MOT-163_nov_81.065%	●	SPN789	Coprococcus catus_nov_95.192%
MOT-127	●	SPN2388	Kineothrix alysoides_nov_96.154%	●	SPN791	Lacrimispora indolis_nov_90.625%
a orotica	●	SPN2400	Muribaculaceae_[G-1]_bacterium_MOT-129_nov_90.816%	●	SPN800	Lachnoclostridium [Clostridium] populeti_nov_96.386%
eris	●	SPN2410	Ruminococcus albus_nov_92.530%	●	SPN809	Adlercreutzia muris_nov_88.745%
caccae	●	SPN2422	Duncaniella freteri_nov_87.398%	●	SPN819	Pelotomaculum terephthalicum_nov_86.667%
as sediminicola	●	SPN2433	Lachnospiraceae_[G-12]_bacterium_MOT-180_nov_94.231%	●	SPN830	Lachnospiraceae_[G-11]_bacterium_MOT-177_nov_94.606%
n psychrotolerans	●	SPN244	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_93.578%	●	SPN84	Butyrivibrio crossotus_nov_97.108%
aceae_[G-1]_bacterium_MOT-189	●	SPN2445	Duncaniella freteri_nov_90.612%	●	SPN841	Vitreimonas flagellata_nov_95.465%
idenbachensis	●	SPN2448	Amedibacillus dolichus_nov_92.290%	●	SPN852	Saccharibacteria_(TM7)_[G-3]_bacterium_HMT_351_nov_97.122%
aeae_[G-11]_bacterium_MOT-177	●	SPN2465	Ruminiclostridium cellulolyticum_nov_89.499%	●	SPN853	Prevotella sp._MOT-128_nov_91.954%
endophyticus	●	SPN2474	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_91.264%	●	SPN864	Lachnospiraceae_[G-6]_bacterium_MOT-171_nov_91.213%
ibbensis	●	SPN2478	Odoribacter splanchnicus_nov_93.807%	●	SPN876	Duncaniella freteri_nov_89.474%
raquae	●	SPN248	Oscillospiraceae_[G-4]_bacterium_MOT-151_nov_96.651%	●	SPN887	Adlercreutzia caecimuris_nov_92.291%
ae_[G-7]_bacterium_MOT-154	●	SPN2486	Lachnospiraceae_[G-12]_bacterium_MOT-179_nov_92.548%	●	SPN899	Lachnospiraceae_[G-14]_bacterium_MOT-184_nov_94.820%
_[G-2]_bacterium_MOT-162	●	SPN2498	Bacteroides uniformis_nov_95.893%	●	SPN900	Anaerotignum lactatifermentans_nov_97.847%
aeae_[G-11]_bacterium_MOT-178	●	SPN2509	Muribaculaceae_[G-1]_bacterium_MOT-129_nov_86.640%	●	SPN923	Culturomica massiliensis_nov_89.613%
aeae_[G-3]_bacterium_MOT-150	●	SPN2511	Gluciribacter canis_nov_93.501%	●	SPN932	Dongia mobilis_nov_96.145%
is bromii	●	SPN2521	Longibaculum muris_nov_92.308%	●	SPN943	Lachnospiraceae_[G-10]_bacterium_MOT-175_nov_90.612%
ae_[G-1]_bacterium_MOT-129	●	SPN2532	Enterocloster asparagiformis_nov_96.875%	●	SPN95	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_91.284%
aeae_[G-14]_bacterium_MOT-184	●	SPN2543	Butyricicoccus pullicaecorum_nov_94.737%	●	SPN953	Chryseolinea soli_nov_95.392%
ium mastitidis	●	SPN2555	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_91.034%	●	SPN962	Muribaculum intestinale_nov_93.822%
ae_[G-2]_bacterium_MOT-167	●	SPN2556	Paraburkholderia hospita_nov_91.667%	●	SPN963	Arsenicibacter rosenii_nov_97.936%
pneumotropicus	●	SPN2566	Eubacteriales_[G-4]_bacterium_MOT-165_nov_94.724%	●	SPN974	Bacteroidetes_[G-3]_bacterium_HMT_436_nov_85.223%
ae_[G-14]_bacterium_MOT-185	●	SPN2577	Parasutterella excrementihominis_nov_94.578%	●	SPN98	Kineothrix alysoides_nov_97.837%
fairfieldensis	●	SPN2588	Kiloniella majae_nov_88.836%	●	SPN985	Duncaniella freteri_nov_90.283%
aeae_[G-6]_bacterium_MOT-153	●	SPN259	Eubacteriales_[G-1]_bacterium_MOT-158_nov_96.635%	●	SPN989	Muribaculaceae_[G-1]_bacterium_MOT-129_nov_91.705%
as contaminans	●	SPN2594	Parabacteroides merdae_nov_93.265%	●	SPN99	Anaeroplasmabactoclasticum_nov_92.874%
aeae_[G-11]_bacterium_MOT-176	●	SPN2600	Holophaga foetida_nov_85.102%	●	SPN997	Eubacteriales_[G-3]_bacterium_MOT-163_nov_94.005%
s lanuginosus_nov_93.636%	●	SPN2610	Erysipelatoclostridium [Clostridium] saccharogumia_nov_93.213%	●	SPP102	Staphylococcus capitis_caprae_epidermidis
enegalensis_nov_93.686%	●	SPN2621	Anaerotruncus colihominis_nov_95.904%	●	SPP110	Afipia_Bradyrhizobium archetypum_austaliense_broomeae_elkanii_embra_pens...(9 spec
zia equolifaciens_nov_92.029%	●	SPN2633	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_92.414%	●	SPP115	Lachnospiraceae_[G-12]_bacterium_MOT-179_bacterium_MOT-180
zia equolifaciens_nov_94.498%	●	SPN2643	Lachnospiraceae_[G-10]_bacterium_MOT-175_nov_93.525%	●	SPP12	Arthrobacter_Pseudarthrobacter_humicola_oryzae_oxydans_pascens_phenanthreniv
butyrica_nov_88.544%	●	SPN2655	Lachnospiraceae_[G-7]_bacterium_MOT-172_nov_91.925%	●	SPP122	Pasteurella_Rodentibacter_caecimuris_heylii
zia caecimuris_nov_94.156%	●	SPN2665	Alistipes putredinis_nov_97.471%	●	SPP147	Ligilactobacillus animalis_apodemimurinus
aceae_[G-2]_bacterium_MOT-104_nov_89.800%	●	SPN2666	Odoribacter splanchnicus_nov_92.184%	●	SPP148	Bacteroides acidifaciens_acidofaciens
acter fermentans_nov_89.895%	●	SPN2676	Duncaniella freteri_nov_93.293%	●	SPP159	Sphingomonas aquatilis_melonis
er sanguinis_nov_95.923%	●	SPN2688	Paludibaculum fermentans_nov_92.048%	●	SPP164	Chryseobacterium_Epilithonimonas_ginsenosidimitans_zeae
etes_[G-3]_bacterium_HMT_436_nov_85.540%	●	SPN2694	Duncaniella freteri_nov_87.375%	●	SPP166	Blautia_hansenii_hominis_marasmi
us mobilis_nov_90.271%	●	SPN2698	Muribaculaceae_[G-2]_bacterium_MOT-104_nov_92.449%	●	SPP178	Escherichia_Pseudescherichia_Shigella_coli_fergusonii_flexneri_sonnei_vulneris
learius_nov_91.591%	●	SPN2708	Kineothrix alysoides_nov_97.356%	●	SPP188	Methyloburbum_aminovorans_extorquens_podarium_populi_pseudosasa...(9 spec
nella hongkongensis_nov_93.510%	●	SPN271	Dokdonella fugitiva_nov_97.955%	●	SPP189	Pseudomonas_arsenicoxydans_mandeliiprosekii_silesiensis
etes_[G-3]_bacterium_HMT_436_nov_86.585%	●	SPN2719	Lachnospiraceae_[G-10]_bacterium_MOT-175_nov_96.394%	●	SPP195	Enterocloster boltea_clostridioformis
aceae_[G-2]_bacterium_MOT-104_nov_88.761%	●	SPN2731	Muricomes intestini_nov_87.831%	●	SPP2	Hungatella_effluvii_hathewayi
aceae_[G-1]_bacterium_MOT-147_nov_96.466%	●	SPN2742	Lachnoclostridium [Clostridium] herbivorans_nov_92.917%	●	SPP209	Clostridium bowmanii_estertheticum