

colifaciens_nov_94.444%	● SPN556 Muribaculum intestinale_nov_92.520%	● SPP10 Buttiauxella_Enterobacter_Erwinia_Klebsiella_Kluuyv ...(10 species) a decarboxylata_aerogenes_a...
ov_97.638%	● SPN557 Lachnoclostridium pacaense_nov_85.659%	● SPP11 Ligilactobacillus animalis_apodemii_faecis_murinus
ris_nov_96.471%	● SPN558 Pseudoflavonifractor capillosus_nov_94.191%	● SPP12 Bacteroides faecichinchillae_faecis_thetaiotaomicron
G-2] bacterium_MOT-162_nov_96.457%	● SPN559 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.764%	● SPP13 Acinetobacter populi_radioresistens
rotolerans_nov_91.732%	● SPN56 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.701%	● SPP14 Agrobacterium_Rhizobium fabrum_glycinendophyticum
[G-2] bacterium_MOT-149_nov_90.980%	● SPN560 Nocardiosis coralliicola_nov_96.850%	● SPP15 Capnocytophaga canimorsus_cynodegmi
[G-1] bacterium_MOT-146_nov_94.094%	● SPN561 Muribaculum intestinale_nov_94.071%	● SPP16 Thermobacillus composti_xylanilyticus
nitidis_nov_94.510%	● SPN562 Eubacterium coprostanoligenes_nov_94.902%	● SPP17 Acidovorax avenae_cattleyae_citrulli_oryzae
ii_nov_96.850%	● SPN563 Kineothrix alysoides_nov_86.614%	● SPP18 Methylobacterium bullatum_marchantiae
odentium_nov_94.118%	● SPN564 Tindallia texcoconensis_nov_82.031%	● SPP19 Lactobacillus gasseri_hominis_johnsonii_taiwanensis
spora_nov_87.843%	● SPN565 Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.157%	● SPP2 Clostridium beijerinckii_diolis_puniceum_saccharoperbutylaceto ...(4 species)
er [Ruminococcus] lactaris_nov_92.157%	● SPN566 Eubacteriales_[G-2] bacterium_MOT-162_nov_90.588%	● SPP20 Clostridium disporicum_saudiense
ae_nov_88.583%	● SPN567 Muribaculum intestinale_nov_86.275%	● SPP21 Bifidobacterium choerinum_pseudolongum
spora_nov_93.281%	● SPN568 Neglectibacter timonensis_nov_93.333%	● SPP22 Lacrimispora algidixylanolytica_xylanolytica
[G-1] bacterium_MOT-129_nov_94.882%	● SPN569 Ethanolgenens harbinense_nov_85.603%	● SPP23 Staphylococcus casei_cohnii_cohnii_subsp._cohnii_edaphicus_gallin ...(9 species)
bacterium_MOT-187_nov_90.119%	● SPN57 Muribaculum intestinale_nov_86.381%	● SPP24 Blautia hansenii_hominis_marasmi
-2] bacterium_MOT-162_nov_87.059%	● SPN570 Acetatifactor muris_nov_87.843%	● SPP25 Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexneri_marmotae_sonnei_vu ...(
e_[G-11] bacterium_MOT-176_nov_96.457%	● SPN571 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_91.339%	● SPP26 Adlercreutzia caecimuris_mucosicola
ens_nov_96.063%	● SPN572 Muribaculum intestinale_nov_90.945%	● SPP27 Enterococcus canintestini_canis_dispar_durans_faecalis_faecium_...(15 species)
[G-2] bacterium_MOT-104_nov_85.375%	● SPN573 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.095%	● SPP3 Limosilactobacillus antri_caviae_fruventi_oris_panis_reuteri_reuteri_c ...(8 species)
ctatifermentans_nov_96.863%	● SPN574 Faecalimonas umbilicata_nov_93.750%	● SPP4 Massilia alkalitolerans_brevitalea_jejuensis_niabensis_suwo ...(6 species)
ae_[G-1] bacterium_MOT-189_nov_90.551%	● SPN575 Desulfolucianica intricata_nov_85.827%	● SPP5 Lactobacillus amylovorus_ultunensis
uvativorans_nov_90.157%	● SPN576 Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.119%	● SPP6 Afipia Bradyrhizobium americanum_archetypum_australiense_betae_broomeae_...(37 species)
[G-4] bacterium_MOT-151_nov_95.669%	● SPN577 Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.119%	● SPP7 Eubacteriales_[G-4] bacterium_MOT-164_bacterium_MOT-165
e_[G-14] bacterium_MOT-185_nov_96.850%	● SPN578 Erysipelotrichaceae_[G-1] bacterium_MOT-189_nov_84.496%	● SPP8 Enterocloster bolteae_clostridioformis
tinilyticus_nov_91.732%	● SPN579 Alistipes fingoldii_nov_90.272%	● SPP9 Pseudomonas antarctica_azotoformans_canadensis_costantini_ext ...(14 species)
ris_nov_95.276%	● SPN58 Muricomes intestini_nov_84.496%	● SPPN1 Oscillospiraceae_multigenus multispecies_sppn1_2_nov_93.676%
otus_nov_91.870%	● SPN580 Papillibacter cinnamivorans_nov_88.976%	● SPPN10 Lachnospiraceae_multigenus multispecies_sppn10_2_nov_88.980%
[G-1] bacterium_MOT-129_nov_92.913%	● SPN581 Faecalimonas umbilicata_nov_94.215%	● SPPN100 Ligilactobacillus multispecies_sppn100_4_nov_84.706%
[G-2] bacterium_MOT-104_nov_93.281%	● SPN582 Anaerotruncus colihominis_nov_94.466%	● SPPN101 Prolixibacter multispecies_sppn101_2_nov_83.594%
ssiliensis_nov_85.000%	● SPN583 Muribaculum intestinale_nov_90.157%	● SPPN102 Lachnospiraceae_multigenus multispecies_sppn102_4_nov_94.510%
accharolyticum_nov_92.126%	● SPN584 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_88.583%	● SPPN103 Blautia multispecies_sppn103_3_nov_90.551%
e_[G-11] bacterium_MOT-178_nov_84.836%	● SPN585 Clostridium prolinivorans_nov_86.667%	● SPPN104 Lactobacillus multispecies_sppn104_5_nov_93.333%
e_[G-11] bacterium_MOT-177_nov_88.627%	● SPN586 Kineothrix alysoides_nov_86.220%	● SPPN105 Clostridium multispecies_sppn105_2_nov_88.672%
is_nov_90.945%	● SPN587 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_84.314%	● SPPN106 Lachnospiraceae_multigenus multispecies_sppn106_2_nov_94.882%
ris_nov_93.307%	● SPN588 Blautia schinkii_nov_87.549%	● SPPN107 Lactobacillus multispecies_sppn107_4_nov_93.333%
ciniphila_nov_95.257%	● SPN589 Muribaculaceae_[G-1] bacterium_MOT-129_nov_93.701%	● SPPN108 Lachnospiraceae_multigenus multispecies_sppn108_3_nov_87.451%
e_[G-14] bacterium_MOT-183_nov_96.063%	● SPN59 Christensenella minuta_nov_86.328%	● SPPN109 Lachnospiraceae_multigenus multispecies_sppn109_2_nov_93.388%
[G-1] bacterium_MOT-129_nov_86.667%	● SPN590 Muribaculaceae_[G-1] bacterium_MOT-129_nov_91.304%	● SPPN11 Lactobacillus multispecies_sppn11_4_nov_86.719%