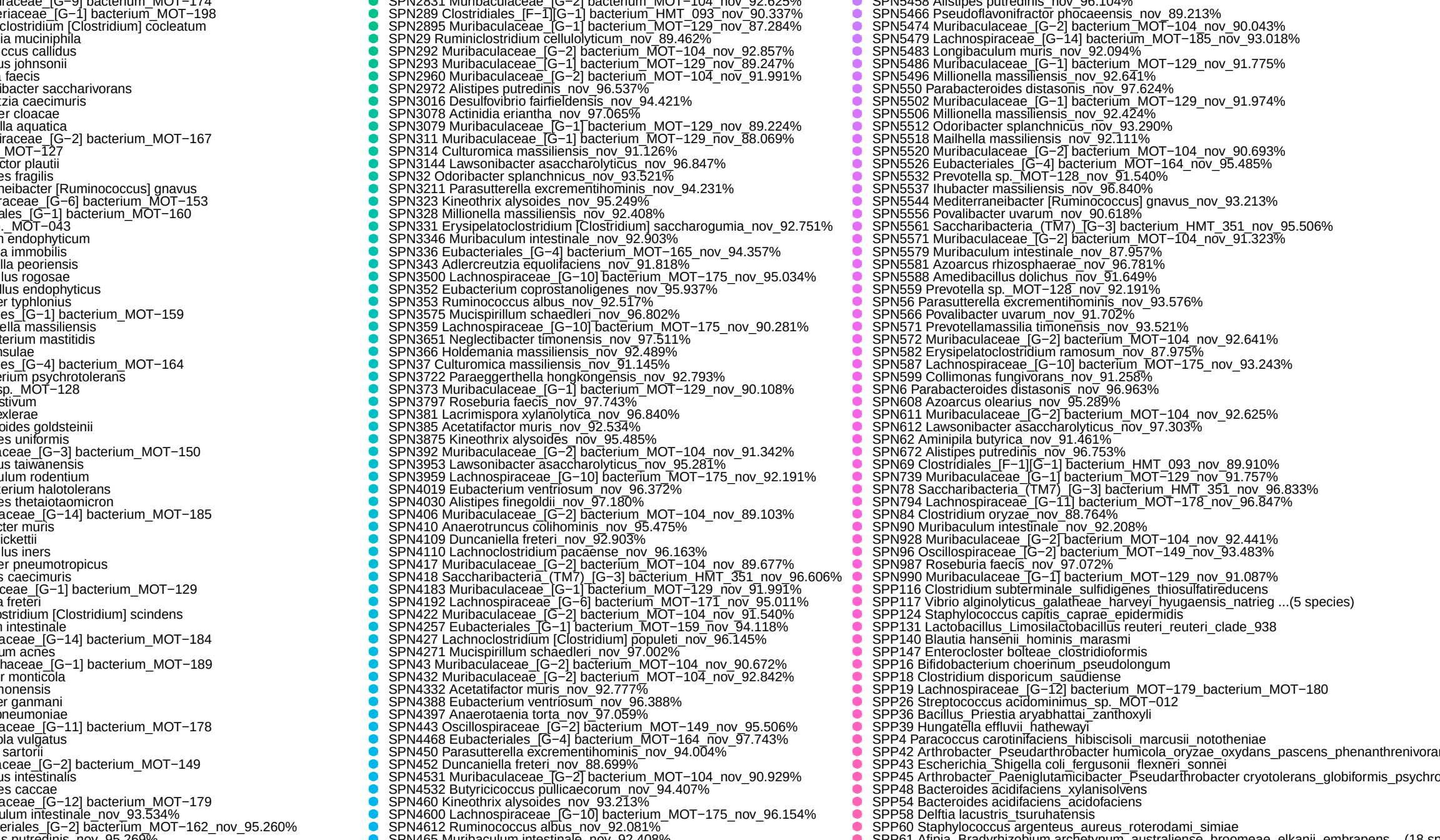
raceae_[G−9] bacterium_MOT−174
eriaceae_[G−1] bacterium_MOT−198
clostridium [Clostridium] cocleatum
ia muciniphila
ccus callidus
us johnsonii
faecis
ibacter saccharivorans
tzia caecimuris
er cloacae
lla aquatica
iraceae_[G−2] bacterium_MOT−167
_MOT−127
ctor plautii
es fragilis
neibacter [Ruminococcus] gnavus
raceae_[G−6] bacterium_MOT−153
ales_[G−1] bacterium_MOT−160
._MOT−043
n endophyticum
a immobilis
lla peoriensis
lus rogosae
llus endophyticus
er typhlonius
es_[G−1] bacterium_MOT−159
ella massiliensis
terium mastitidis
nsulae
es_[G−4] bacterium_MOT−164
erium psychrotolerans
sp._MOT−128
stivum
xlerae
oides goldsteinii
es uniformis
aceae_[G−3] bacterium_MOT−150
us taiwanensis
ulum rodentium
terium halotolerans
es thetaiotaomicron
aceae_[G−14] bacterium_MOT−185
cter muris
ickettii
lus iners
er pneumotropicus
s caecimuris
ceae_[G−1] bacterium_MOT−129
a freteri
ostridium [Clostridium] scindens
n intestinale
aceae_[G−14] bacterium_MOT−184
um acnes
haceae_[G−1] bacterium_MOT−189
r monticola
nonensis
er ganmani
pneumoniae
aceae_[G−11] bacterium_MOT−178
ola vulgatus
sartorii
aceae_[G−2] bacterium_MOT−149
us intestinalis
es caccae
aceae_[G−12] bacterium_MOT−179
ulum intestinale_nov_93.534%
eriales_[G−2] bacterium_MOT−162_nov_95.260%
SPN2831 Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.625%
SPN289 Clostridiales_[F−1][G−1] bacterium_HMT_093_nov_90.337%
SPN2895 Muribaculaceae_[G−1] bacterium_MOT−129_nov_87.284%
SPN29 Ruminiclostridium cellulolyticum_nov_89.462%
SPN292 Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.857%
SPN293 Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.247%
SPN2960 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.991%
SPN2972 Alistipes putredinis_nov_96.537%
SPN3016 Desulfovibrio fairfieldensis_nov_94.421%
SPN3078 Actinidia eriantha_nov_97.065%
SPN3079 Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.224%
SPN311 Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.069%
SPN314 Culturomica massiliensis_nov_91.126%
SPN3144 Lawsonibacter asaccharolyticus_nov_96.847%
SPN32 Odoribacter splanchnicus_nov_93.521%
SPN3211 Parasutterella excrementihominis_nov_94.231%
SPN323 Kineothrix alysoides_nov_95.249%
SPN328 Millionella massiliensis_nov_92.408%
SPN331 Erysipelatoclostridium [Clostridium] saccharogumia_nov_92.751%
SPN3346 Muribaculum intestinale_nov_92.903%
SPN336 Eubacteriales_[G−4] bacterium_MOT−165_nov_94.357%
SPN343 Adlercreutzia equolifaciens_nov_91.818%
SPN3500 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_95.034%
SPN352 Eubacterium coprostanoligenes_nov_95.937%
SPN353 Ruminococcus albus_nov_92.517%
SPN3575 Mucispirillum schaedleri_nov_96.802%
SPN359 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_90.281%
SPN3651 Neglectibacter timonensis_nov_97.511%
SPN366 Holdemania massiliensis_nov_92.489%
SPN37 Culturomica massiliensis_nov_91.145%
SPN3722 Paraeggerthella hongkongensis_nov_92.793%
SPN373 Muribaculaceae_[G−1] bacterium_MOT−129_nov_90.108%
SPN3797 Roseburia faecis_nov_97.743%
SPN381 Lacrimispora xylanolytica_nov_96.840%
SPN385 Acetatifactor muris_nov_92.534%
SPN3875 Kineothrix alysoides_nov_95.485%
SPN392 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.342%
SPN3953 Lawsonibacter asaccharolyticus_nov_95.281%
SPN3959 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_92.191%
SPN4019 Eubacterium ventriosum_nov_96.372%
SPN4030 Alistipes finegoldii_nov_97.180%
SPN406 Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.103%
SPN410 Anaerotruncus colihominis_nov_95.475%
SPN4109 Duncaniella freteri_nov_92.903%
SPN4110 Lachnoclostridium pacaense_nov_96.163%
SPN417 Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.677%
SPN418 Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351_nov_96.606%
SPN4183 Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.991%
SPN4192 Lachnospiraceae_[G−6] bacterium_MOT−171_nov_95.011%
SPN422 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.540%
SPN4257 Eubacteriales_[G−1] bacterium_MOT−159_nov_94.118%
SPN427 Lachnoclostridium [Clostridium] populeti_nov_96.145%
SPN4271 Mucispirillum schaedleri_nov_97.002%
SPN43 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.672%
SPN432 Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.842%
SPN4332 Acetatifactor muris_nov_92.777%
SPN4388 Eubacterium ventriosum_nov_96.388%
SPN4397 Anaerotaenia torta_nov_97.059%
SPN443 Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.506%
SPN4468 Eubacteriales_[G−4] bacterium_MOT−164_nov_97.743%
SPN450 Parasutterella excrementihominis_nov_94.004%
SPN452 Duncaniella freteri_nov_88.699%
SPN4531 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.929%
SPN4532 Butyricicoccus pullicaecorum_nov_94.407%
SPN460 Kineothrix alysoides_nov_93.213%
SPN4600 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_96.154%
SPN4612 Ruminococcus albus_nov_92.081%
SPN5458 Alistipes putredinis_nov_96.104%
SPN5466 Pseudoflavonifractor phocaeensis_nov_89.213%
SPN5474 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.043%
SPN5479 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_93.018%
SPN5483 Longibaculum muris_nov_92.094%
SPN5486 Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.775%
SPN5496 Millionella massiliensis_nov_92.641%
SPN550 Parabacteroides distasonis_nov_97.624%
SPN5502 Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.974%
SPN5506 Millionella massiliensis_nov_92.424%
SPN5512 Odoribacter splanchnicus_nov_93.290%
SPN5518 Mailhella massiliensis_nov_92.111%
SPN5520 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.693%
SPN5526 Eubacteriales_[G−4] bacterium_MOT−164_nov_95.485%
SPN5532 Prevotella sp._MOT−128_nov_91.540%
SPN5537 Ihubacter massiliensis_nov_96.840%
SPN5544 Mediterraneibacter [Ruminococcus] gnavus_nov_93.213%
SPN5556 Povalibacter uvarum_nov_90.618%
SPN5561 Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351_nov_95.506%
SPN5571 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.323%
SPN5579 Muribaculum intestinale_nov_87.957%
SPN5581 Azoarcus rhizosphaerae_nov_96.781%
SPN5588 Amedibacillus dolichus_nov_91.649%
SPN559 Prevotella sp._MOT−128_nov_92.191%
SPN56 Parasutterella excrementihominis_nov_93.576%
SPN566 Povalibacter uvarum_nov_91.702%
SPN571 Prevotellamassilia timonensis_nov_93.521%
SPN572 Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.641%
SPN582 Erysipelatoclostridium ramosum_nov_87.975%
SPN587 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_93.243%
SPN599 Collimonas fungivorans_nov_91.258%
SPN6 Parabacteroides distasonis_nov_96.963%
SPN608 Azoarcus olearius_nov_95.289%
SPN611 Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.625%
SPN612 Lawsonibacter asaccharolyticus_nov_97.303%
SPN62 Aminipila butyrica_nov_91.461%
SPN672 Alistipes putredinis_nov_96.753%
SPN69 Clostridiales_[F−1][G−1] bacterium_HMT_093_nov_89.910%
SPN739 Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.757%
SPN78 Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351_nov_96.833%
SPN794 Lachnospiraceae_[G−11] bacterium_MOT−178_nov_96.847%
SPN84 Clostridium oryzae_nov_88.764%
SPN90 Muribaculum intestinale_nov_92.208%
SPN928 Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.441%
SPN96 Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.483%
SPN987 Roseburia faecis_nov_97.072%
SPN990 Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.087%
SPP116 Clostridium subterminale_sulfidigenes_thiosulfatireducens
SPP117 Vibrio alginolyticus_galatheae_harveyi_hyugaensis_natrieg ...(5 species)
SPP124 Staphylococcus capitis_caprae_epidermidis
SPP131 Lactobacillus_Limosilactobacillus reuteri_reuteri_clade_938
SPP140 Blautia hansenii_hominis_marasmi
SPP147 Enterocloster bolteae_clostridioformis
SPP16 Bifidobacterium choerinum_pseudolongum
SPP18 Clostridium disporicum_saudiense
SPP19 Lachnospiraceae_[G−12] bacterium_MOT−179_bacterium_MOT−180
SPP26 Streptococcus acidominimus_sp._MOT−012
SPP36 Bacillus_Priestia aryabhattai_zanthoxyli
SPP39 Hungatella effluvii_hathewayi
SPP4 Paracoccus carotinifaciens_hibiscisoli_marcusii_nototheniae
SPP42 Arthrobacter_Pseudarthrobacter humicola_oryzae_oxydans_pascens_phenanthrenivorar
SPP43 Escherichia_Shigella coli_fergusonii_flexneri_sonnei
SPP45 Arthrobacter_Paeniglutamicibacter_Pseudarthrobacter cryotolerans_globiformis_psychro
SPP48 Bacteroides acidifaciens_xylanisolvens
SPP54 Bacteroides acidifaciens_acidofaciens
SPP58 Delftia lacustris_tsuruhatensis
SPP60 Staphylococcus argenteus_aureus_roterodami_simiae