
Species
SP16 Limosilactobacillus reuteri
SP17 Helicobacter ganmani
SP18 Oscillospiraceae_[G−7] bacterium_MOT−154
SP19 Clostridium disporicum
SP20 Ileibacterium valens
SP21 Rodentibacter pneumotropicus
SP22 Lachnospiraceae_[G−11] bacterium_MOT−177
SP23 Acutalibacter muris
SP24 Muribaculum intestinale
SP25 Peptococcaceae_[G−1] bacterium_MOT−146
SP26 Bacteroides caecimuris
SP27 Adlercreutzia mucosicola
SP28 Porphyromonas sp._MOT−131
SP29 Muribacter muris
SP3 Romboutsia ilealis
SP30 Turicimonas muris
SP31 Parabacteroides goldsteinii
SP32 Lactobacillus intestinalis
SP33 Brachyspira intermedia
SP34 Blautia hominis
SP35 Parabacteroides distasonis
SP36 Adlercreutzia caecimuris
SP37 Prevotella sp._MOT−128
SP38 Rodentibacter heylii
SP39 Mollicutes_[G−1] bacterium_MOT−186
SP4 Carnobacteriaceae_[G−1] bacterium_MOT−198
SP40 Parvibacter caecicola
SP41 Bifidobacterium pseudolongum
SP42 Phocaeicola sartorii
SP43 Corynebacterium mastitidis
SP45 Muribacter sp._MOT−142
SP46 Lactobacillus taiwanensis
SP5 Erysipelotrichaceae_[G−1] bacterium_MOT−189
SP50 Eubacteriales_[G−1] bacterium_MOT−159
SP52 Enterococcus gallinarum
SP54 Lactobacillus johnsonii
SP55 Muribaculaceae_[G−1] bacterium_MOT−129
SP56 Oscillospiraceae_[G−3] bacterium_MOT−150
SP57 Lachnospiraceae_[G−3] bacterium_MOT−168
SP58 Streptococcus acidominimus
SP6 Lachnospiraceae_[G−11] bacterium_MOT−178
SP63 Adlercreutzia muris
SP64 Erysipelatoclostridium [Clostridium] cocleatum
SP67 Ruminococcus gauvreauii
SP69 Escherichia fergusonii
SP7 Streptococcus sp._MOT−012
SP71 Escherichia marmotae
SP75 Akkermansia muciniphila
SP8 Lachnospiraceae_[G−14] bacterium_MOT−185
SP81 Alistipes timonensis
SP82 Alistipes sp._MOT−127
SP83 Eubacteriales_[G−1] bacterium_MOT−160
SP85 Lachnospiraceae_[G−2] bacterium_MOT−167
SP89 Mollicutes_[G−2] bacterium_MOT−188
SP9 Streptococcus danieliae
SP92 Weeksellaceae_[G_1] bacterium_MOT−126
SP93 Helicobacter typhlonius
SP94 Muribacter sp._MOT−143
SPN108 Lachnospiraceae_[G−11] bacterium_MOT−177_nov_95.287%
SPN109 Muribaculaceae_[G−1] bacterium_MOT−129_nov_93.228%
SPN11 Butyricicoccus pullicaecorum_nov_94.194%
SPN111 Enterocloster clostridioformis_nov_92.769%
SPN120 Lawsonibacter asaccharolyticus_nov_87.740%
SPN123 Lachnospiraceae_[G−11] bacterium_MOT−176_nov_95.797%
SPN131 Oscillospiraceae_[G−3] bacterium_MOT−150_nov_94.639%
SPN135 Prevotellamassilia timonensis_nov_89.413%
SPN136 Mailhella massiliensis_nov_91.242%
SPN14 Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.993%
SPN142 Gluceribacter canis_nov_92.397%
SPN148 Lachnospiraceae_[G−14] bacterium_MOT−184_nov_94.648%
SPN217 Eubacteriales_[G−4] bacterium_MOT−165_nov_93.034%
SPN218 Bacteroidetes_[G−3] bacterium_HMT_365_nov_87.666%
SPN219 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.463%
SPN22 Adlercreutzia caecimuris_nov_92.996%
SPN228 Oscillospiraceae_[G−3] bacterium_MOT−150_nov_95.045%
SPN230 Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.593%
SPN239 Rikenella microfusus_nov_93.227%
SPN242 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.542%
SPN250 Lacrimispora saccharolytica_nov_91.495%
SPN254 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.231%
SPN255 Ihubacter massiliensis_nov_96.039%
SPN261 Lachnospiraceae_[G−14] bacterium_MOT−182_nov_90.752%
SPN27 Duncaniella freteri_nov_92.271%
SPN271 Mailhella massiliensis_nov_91.995%
SPN278 Lachnospiraceae_[G−4] bacterium_MOT−169_nov_94.441%
SPN280 Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351_nov_96.439%
SPN281 Bacteroides uniformis_nov_96.825%
SPN282 Duncaniella freteri_nov_91.090%
SPN283 Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.535%
SPN284 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.136%
SPN285 Alistipes putredinis_nov_96.014%
SPN286 Oscillospiraceae_[G−6] bacterium_MOT−153_nov_92.971%
SPN287 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_96.335%
SPN288 Duncaniella freteri_nov_91.008%
SPN289 Eubacterium xylanophilum_nov_92.240%
SPN290 Lachnospiraceae_[G−11] bacterium_MOT−178_nov_95.238%
SPN291 Olsenella massiliensis_nov_94.126%
SPN292 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.027%
SPN293 Eubacterium xylanophilum_nov_91.588%
SPN294 Faecalicatena contorta_nov_92.382%
SPN295 Lachnospiraceae_[G−4] bacterium_MOT−169_nov_95.127%
SPN296 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_91.446%
SPN297 Faecalicatena contorta_nov_94.303%
SPN298 Muribaculum intestinale_nov_93.086%
SPN299 Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.945%
SPN300 Muribaculaceae_[G−1] bacterium_MOT−129_nov_93.694%
SPN301 Christensenella timonensis_nov_84.737%
SPN302 Lachnospiraceae_[G−12] bacterium_MOT−180_nov_95.782%
SPN303 Clostridium swellfunianum_nov_85.704%
SPN304 Lachnospiraceae_[G−13] bacterium_MOT−181_nov_89.500%
SPN305 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.211%
SPN306 Mediterraneibacter [Ruminococcus] gnavus_nov_94.433%
SPN307 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.126%
SPN308 Oscillibacter valericigenes_nov_95.386%
SPN309 Eubacteriales_[G−2] bacterium_MOT−162_nov_94.260%
SPN310 Oscillibacter valericigenes_nov_94.219%
SPN311 Clostridiales_[F−1][G−1] bacterium_HMT_093_nov_87.886%
SPN312 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.082%
SPN313 Odoribacter splanchnicus_nov_91.442%
SPN314 Lachnospiraceae_[G−8] bacterium_MOT−173_nov_88.988%
SPN315 Duncaniella freteri_nov_90.897%
SPN316 Duncaniella freteri_nov_90.761%
SPN317 Odoribacter splanchnicus_nov_93.862%
SPN318 Lachnospiraceae_[G−6] bacterium_MOT−171_nov_93.621%
SPN319 Lachnospiraceae_[G−3] bacterium_MOT−168_nov_96.845%
SPN32 Muribaculaceae_[G−1] bacterium_MOT−129_nov_90.642%
SPN320 Eubacterium coprostanoligenes_nov_94.861%
SPN321 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.884%
SPN322 Alistipes senegalensis_nov_95.604%
SPN323 Enterocloster bolteae_nov_93.210%
SPN324 Mucispirillum schaedleri_nov_96.136%
SPN325 Lachnospiraceae_[G−12] bacterium_MOT−180_nov_96.794%
SPN326 Faecalimonas umbilicata_nov_92.686%
SPN327 Duncaniella freteri_nov_92.339%
SPN328 Roseburia intestinalis_nov_88.569%
SPN329 Faecalicatena contorta_nov_92.181%
SPN330 Neglectibacter timonensis_nov_96.655%
SPN331 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.483%
SPN332 Oscillospiraceae_[G−3] bacterium_MOT−150_nov_94.353%
SPN333 Alistipes putredinis_nov_94.697%
SPN346 Clostridium merdae_nov_94.317%
SPN347 Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.775%
SPN348 Adlercreutzia equolifaciens_nov_92.350%
SPN349 Lacrimispora saccharolytica_nov_93.333%
SPN350 Kiloniella majae_nov_86.042%
SPN351 Paraeggerthella hongkongensis_nov_90.300%
SPN352 Lacrimispora amygdalina_nov_93.968%
SPN353 Adlercreutzia equolifaciens_nov_91.562%
SPN354 Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.565%
SPN355 Eubacterium oxidoreducens_nov_92.539%
SPN356 Hungatella xylanolytica_nov_94.045%
SPN357 Christensenella hongkongensis_nov_86.023%
SPN358 Eubacterium xylanophilum_nov_92.177%
SPN359 Muribaculum intestinale_nov_90.610%
SPN360 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_89.145%
SPN361 Oscillospiraceae_[G−3] bacterium_MOT−150_nov_90.391%
SPN362 Anaerotignum lactatifermentans_nov_96.362%
SPN363 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_89.933%
SPN364 Lachnospiraceae_[G−5] bacterium_MOT−170_nov_97.663%
SPN365 Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.132%
SPN366 Mordavella massiliensis_nov_92.730%
SPN367 Eubacteriales_[G−4] bacterium_MOT−164_nov_97.785%
SPN368 Caproiciproducens galactitolivorans_nov_91.253%
SPN369 Eubacterium xylanophilum_nov_91.287%
SPN370 Lachnospiraceae_[G−14] bacterium_MOT−184_nov_96.502%
SPN371 Eisenbergiella massiliensis_nov_90.373%
SPN372 Lawsonibacter asaccharolyticus_nov_94.722%
SPN373 Butyribacter intestini_nov_90.370%
SPN374 Lacrimispora amygdalina_nov_91.078%
SPN375 Kiloniella laminariae_nov_86.152%
SPN376 Alistipes shahii_nov_97.378%
SPN377 Alistipes senegalensis_nov_95.658%
SPN378 Acutalibacter muris_nov_97.598%
SPN379 Odoribacter splanchnicus_nov_94.199%
SPN38 Duncaniella freteri_nov_93.493%
SPN380 Longibaculum muris_nov_94.769%
SPN381 Faecalicatena contorta_nov_95.339%
SPN382 Lachnospiraceae_[G−11] bacterium_MOT−178_nov_93.568%
SPN383 Lachnospiraceae_[G−11] bacterium_MOT−177_nov_94.726%
SPN384 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_93.805%
SPN385 Flavonifractor plautii_nov_93.823%
SPN386 Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.244%
SPN387 Lachnospiraceae_[G−11] bacterium_MOT−177_nov_96.783%
SPN388 Lacrimispora celerecrescens_nov_93.664%
SPN389 Duncaniella freteri_nov_89.222%
SPN390 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_89.153%
SPN391 Oscillospiraceae_[G−2] bacterium_MOT−149_nov_94.574%
SPN392 Anaerotaenia torta_nov_88.866%
SPN393 Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.149%
SPN394 Lachnospiraceae_[G−2] bacterium_MOT−167_nov_93.033%
SPN395 Lachnospiraceae_[G−14] bacterium_MOT−183_nov_96.540%
SPN396 Oscillospiraceae_[G−2] bacterium_MOT−149_nov_97.730%
SPN397 Mollicutes_[G−2] bacterium_MOT−188_nov_87.231%
SPN398 Eubacterium xylanophilum_nov_92.376%
SPN399 Mollicutes_[G−2] bacterium_MOT−187_nov_97.281%
SPN400 Eubacteriales_[G−1] bacterium_MOT−160_nov_97.924%
SPN401 Prevotella sp._MOT−128_nov_91.581%
SPN402 Lachnospiraceae_[G−10] bacterium_MOT−175_nov_89.349%
SPN411 Marvinbryantia formatexigens_nov_91.655%
SPN43 Desulfovibrio desulfuricans_nov_92.438%
SPN50 Defluviitalea saccharophila_nov_82.903%
SPN54 Alistipes putredinis_nov_97.032%
SPN62 Parabacteroides merdae_nov_95.530%
SPN65 Oscillospiraceae_[G−1] bacterium_MOT−147_nov_97.735%
SPN74 Mediterraneibacter [Ruminococcus] gnavus_nov_94.983%
SPN76 Adlercreutzia caecimuris_nov_91.655%
SPN86 Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.736%
SPN87 Culturomica massiliensis_nov_91.758%
SPN98 Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.360%
SPN99 Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.252%