

Species

- SP1 Lactobacillus taiwanensis
- SP11 Pantoea agglomerans
- SP12 Turicimonas muris
- SP2 Pantoea plantarum
- SP21 Bacteroides stercorisoris
- SP27 Streptococcus thermophilus
- SP28 Streptococcus mutans
- SP34 Staphylococcus sciuri
- SP5 Akkermansia muciniphila
- SP8 Muribaculum intestinale
- SPN10 Absiella tortuosum_nov_91.863%
- SPN11 Muribaculum intestinale_nov_92.657%
- SPN12 Eubacterium coprostanoligenes_nov_95.485%
- SPN13 Muribaculum intestinale_nov_90.909%
- SPN14 Anaerostipes butyraticus_nov_97.052%
- SPN15 Muribaculum intestinale_nov_89.827%
- SPN17 Alistipes timonensis_nov_97.609%
- SPN18 Bacteroidetes_[G-7] bacterium_HMT_911_nov_87.284%
- SPN19 Muribaculum intestinale_nov_87.312%
- SPN2 Muribaculum intestinale_nov_91.974%
- SPN20 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_92.325%
- SPN21 Lachnospirillum asparagiforme_nov_96.364%
- SPN22 Neglecta timonensis_nov_97.506%
- SPN23 Mailhella massiliensis_nov_91.863%
- SPN25 Prevotellamassilia timonensis_nov_93.900%
- SPN26 Curvibacter gracilis_nov_97.645%
- SPN27 Muribaculum intestinale_nov_90.870%
- SPN28 Phocaea massiliensis_nov_95.000%
- SPN29 Muribaculum intestinale_nov_90.476%
- SPN3 Muribaculum intestinale_nov_91.775%
- SPN30 Firmicutes_[G] sp._Oral_Taxon_A55_nov_91.221%
- SPN31 Erysipelatoclostridium ramosum_nov_88.186%
- SPN32 Ruminococcaceae_[G-2] bacterium_HMT_085_nov_91.648%
- SPN39 Ruminococcaceae_[G-1] bacterium_HMT_075_nov_90.293%
- SPN4 Blautia gnavus_nov_93.197%
- SPN5 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_97.079%
- SPN59 Ruminococcaceae_[G-1] bacterium_HMT_075_nov_90.724%
- SPN6 Muribaculum intestinale_nov_92.424%
- SPN69 Lachnospirillum xylanolyticum_nov_97.279%
- SPN7 Muribaculum intestinale_nov_90.239%
- SPN72 Longibaculum muris_nov_93.790%
- SPN8 Muribaculum intestinale_nov_87.097%
- SPN9 Muribaculum intestinale_nov_92.641%
- SPP1 Alcaligenes aquatilis,faecalis
- SPP2 Corynebacterium amycolatum,sp._Oral_Taxon_B71
- SPP3 Cutibacterium,Propionibacterium acnes,sp._str._LG
- SPP4 Lactobacillus animalis,apodemi,murinus
- SPP7 Lactobacillus reuteri,reuteri_clade_938
- SPPN1 Lachnospirillum multispecies_sppn1_2_nov_96.154%
- SPPN10 Roseburia multispecies_sppn10_2_nov_96.145%
- SPPN11 Lachnospirillum multispecies_sppn11_2_nov_96.364%
- SPPN12 multigenus multispecies_sppn12_3_nov_92.609%
- SPPN2 Lachnospirillum multispecies_sppn2_2_nov_94.091%
- SPPN3 Ruminoclostridium thermocellum
- SPPN4 Roseburia multispecies_sppn4_2_nov_97.964%
- SPPN5 Lachnospirillum multispecies_sppn5_2_nov_96.825%
- SPPN6 Lachnospirillum multispecies_sppn6_2_nov_95.465%
- SPPN7 multigenus multispecies_sppn7_3_nov_97.052%
- SPPN8 Leifsonia multispecies_sppn8_4_nov_97.303%
- SPPN9 Lachnospirillum multispecies_sppn9_2_nov_95.909%