



Group
Stool Female Non-perio
Stool Male Non-perio



Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.623%
Alistipes finegoldii_nov_93.608%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.400%
Duncaniella freteri_nov_88.978%
Duncaniella freteri_nov_88.330%
Marinisporobacter balticus_nov_82.692%
Clostridium disporicum
Eubacteriales_[G-1] bacterium_MOT-158
Longibaculum muris_nov_86.957%
Beduini massiliensis_nov_87.705%
Mucispirillum schaedleri
Flavonifractor plautii_nov_92.308%
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.792%
Caproiciproducens galactitolivorans_nov_83.789%
Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_82.255%
Muricomes intestini_nov_89.583%
Eubacteriales_[G-4] bacterium_MOT-164
Bacteroides acidifaciens
Duncaniella freteri_nov_86.948%
Bifidobacterium pseudolongum
Eubacteriales_[G-4] bacterium_MOT-164_nov_97.655%
Duncaniella freteri_nov_89.069%
Enterococcus casseliflavus
Neglectibacter timonensis_nov_95.325%
Alistipes putredinis_nov_94.444%
Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.931%
Eubacteriales_[G-2] bacterium_MOT-162
Sporobacter termitidis_nov_87.580%
Hatthewaya proteolytica_nov_83.514%
Enterococcus faecalis
Acutalibacter muris
Lachnoclostridium [Clostridium] aminophilum_nov_87.318%
Eubacteriales_[G-1] bacterium_MOT-159
Acetivibrio cellulolyticus_nov_83.405%
Duncaniella freteri_nov_92.915%
Faecalicatena multispecies_sppn9_2_nov_94.363%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.279%
Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.582%
Eubacterium xylanophilum_nov_91.075%
Faecalicatena multispecies_sppn7_2_nov_91.858%
Lacrimispora xylanolytica_nov_94.363%
Limosilactobacillus reuteri
Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.816%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.290%
Ileibacterium valens
Longibaculum muris_nov_90.289%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.640%
Turicibacter sanguinis_nov_95.923%
Duncaniella freteri_nov_89.135%
Duncaniella freteri_nov_89.697%
Duncaniella freteri_nov_87.221%
Prevotella sp._HMT_317_nov_90.244%
Phocaeicola sartorii
Enterobacter hormaechei
Duncaniella freteri_nov_87.071%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.822%
Eubacteriales_[G-4] bacterium_MOT-164_nov_97.655%
Duncaniella freteri_nov_93.699%
Alistipes sp._MOT-127
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.718%
Lachnoclostridium [Clostridium] scindens_nov_89.027%
Anaeroplasma abactoclasticum_nov_86.538%
Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.134%
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_92.902%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.373%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.000%
Duncaniella freteri_nov_92.653%
Lactobacillus intestinalis
Parabacteroides distasonis
Parabacteroides distasonis_nov_97.938%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.621%
Alistipes senegalensis_nov_93.648%
Gluciribacter canis_nov_93.305%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.576%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.577%
Muribaculum intestinale
Erysipelatoclostridium [Clostridium] innocuum_nov_88.270%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.600%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.089%
Parasutterella excrementihominis
Duncaniella freteri_nov_87.169%
Duncaniella freteri_nov_91.039%
Duncaniella freteri_nov_90.184%
Duncaniella freteri_nov_86.290%
Alloprevotella sp._HMT_473_nov_89.366%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_85.686%
Duncaniella freteri_nov_86.842%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_83.636%
Prevotella shahii_nov_87.602%
Duncaniella freteri_nov_91.853%
Alloprevotella sp._HMT_473_nov_89.634%
Duncaniella freteri_nov_90.612%
Erysipelotrichaceae_[G-1] bacterium_MOT-189
Ligilactobacillus murinus
Muribaculaceae_[G-1] bacterium_MOT-129_nov_84.929%
Parabacteroides goldsteinii
Akkermansia muciniphila
Duncaniella freteri_nov_91.429%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
Muribaculaceae_[G-1] bacterium_MOT-129

F15127.S28
F15127.S27
F15127.S29
F15127.S30
F15127.S25
F15127.S26

Samples

Species