0 10 30 50
Value
Disease_Sex
Control_male
AD_male
Lactobacillus rogosae
Enterocloster multispecies_sppn13_2_nov_94.470%
Hungatella xylanolytica
Dysosmobacter welbionis_nov_93.953%
Fusicatenibacter saccharivorans
Roseburia inulinivorans
Parabacteroides merdae
Phascolarctobacterium faecium
Bilophila wadsworthia
Hungatella xylanolytica_nov_97.674%
Sporobacter termitidis_nov_92.130%
Alistipes onderdonkii
Subdoligranulum variabile
Ruminococcus albus_nov_90.187%
Parasutterella excrementihominis
Bacteroides fragilis
Bacteroides faecichinchillae_faecis_thetaiotaomicron
Bacteroides xylanisolvens
Bacteroides koreensis_kribbi
Ruminococcus champanellensis_nov_97.674%
Faecalibacterium prausnitzii_nov_96.744%
Bacteroides eggerthii
Ruminiclostridium cellobioparum_nov_89.352%
Butyrivibrio crossotus
Alistipes putredinis
Monoglobus pectinilyticus_nov_90.698%
Phocaeicola massiliensis
Barnesiella intestinihominis
Bacteroides cellulosilyticus_timonensis
Bacteroides caccae
Bacteroides stercoris
Ruminiclostridium cellobioparum_nov_92.056%
Oscillibacter valericigenes_nov_93.056%
Blautia wexlerae
Roseburia faecis
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.674%
Prevotella stercorea
Roseburia intestinalis
Muribaculum intestinale_nov_85.116%
Faecalibacterium prausnitzii_nov_97.209%
Eubacteriales_[G−1] bacterium_MOT−144
Massiliprevotella massiliensis
Prevotella copri_nov_97.674%
Bacteroides uniformis
Phocaeicola dorei
Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexn
Akkermansia muciniphila
Faecalibacterium prausnitzii
Phocaeicola vulgatus
Prevotella copri
Species
F3410872.S08
F3410872.S03
F3410872.S80
F3410872.S20
F3410872.S70
F3410872.S85
F3410872.S11
F3410872.S06
F3410872.S31
F3410872.S59
F3410872.S89
F3410872.S14
F3410872.S68
F3410872.S55
F3410872.S69
F3410872.S01
F3410872.S67
F3410872.S62
F3410872.S77
F3410872.S86
F3410872.S42
F3410872.S48
F3410872.S13
F3410872.S88
F3410872.S52
F3410872.S28
F3410872.S19
F3410872.S71
F3410872.S34
F3410872.S21
F3410872.S53
F3410872.S36
F3410872.S18
Samples