0
20
40
60
80
Value
Group
OVX
OVX_E2
Enterococcus faecalis
Ligilactobacillus murinus
Bifidobacterium pseudolongum
Staphylococcus saprophyticus_xylosus
Phocaeicola vulgatus
Bacteroides uniformis
Parasutterella excrementihominis_nov_94.578%
Turicimonas muris
Alistipes senegalensis_nov_96.920%
Olsenella multispecies_sppn1_2_nov_91.966%
Parafannyhessea umbonata_nov_92.161%
Faecalicatena fissicatena_nov_95.407%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.125%
Anaeromassilibacillus senegalensis_nov_92.489%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.097%
Alistipes putredinis_nov_92.213%
Erysipelatoclostridium [Clostridium] cocleatum
Alistipes shahii_nov_93.827%
Adlercreutzia equolifaciens
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_93.737%
Faecalicatena fissicatena_nov_93.528%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_95.000%
Clostridium disporicum
Adlercreutzia caecimuris_nov_93.290%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.351%
Alistipes putredinis_nov_92.813%
Lawsonia intracellularis_nov_89.336%
Faecalicatena multispecies_sppn3_2_nov_93.737%
Bacteroides caecimuris
Faecalicoccus acidiformans_nov_89.600%
Parvibacter caecicola_nov_97.609%
Turicibacter sanguinis_nov_95.923%
Duncaniella freteri_nov_87.375%
Mammaliicoccus sciuri
Staphylococcus saprophyticus
Parabacteroides goldsteinii
Lactobacillus johnsonii
Gluceribacter canis_nov_93.501%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Lachnospiraceae_[G−3] bacterium_MOT−168
Faecalicatena multispecies_sppn2_2_nov_92.067%
Lachnospiraceae_[G−2] bacterium_HMT_096_nov_91.632%
Mammaliicoccus lentus
Acinetobacter septicus
Ihubacter massiliensis_nov_94.572%
Alistipes sp._MOT−127
Serratia marcescens_nematodiphila
Dubosiella newyorkensis
Serratia marcescens
Faecalibaculum rodentium
Species
F29493.S07
F29493.S08
F29493.S12
F29493.S06
F29493.S11
F29493.S09
F29493.S10
F29493.S05
Samples