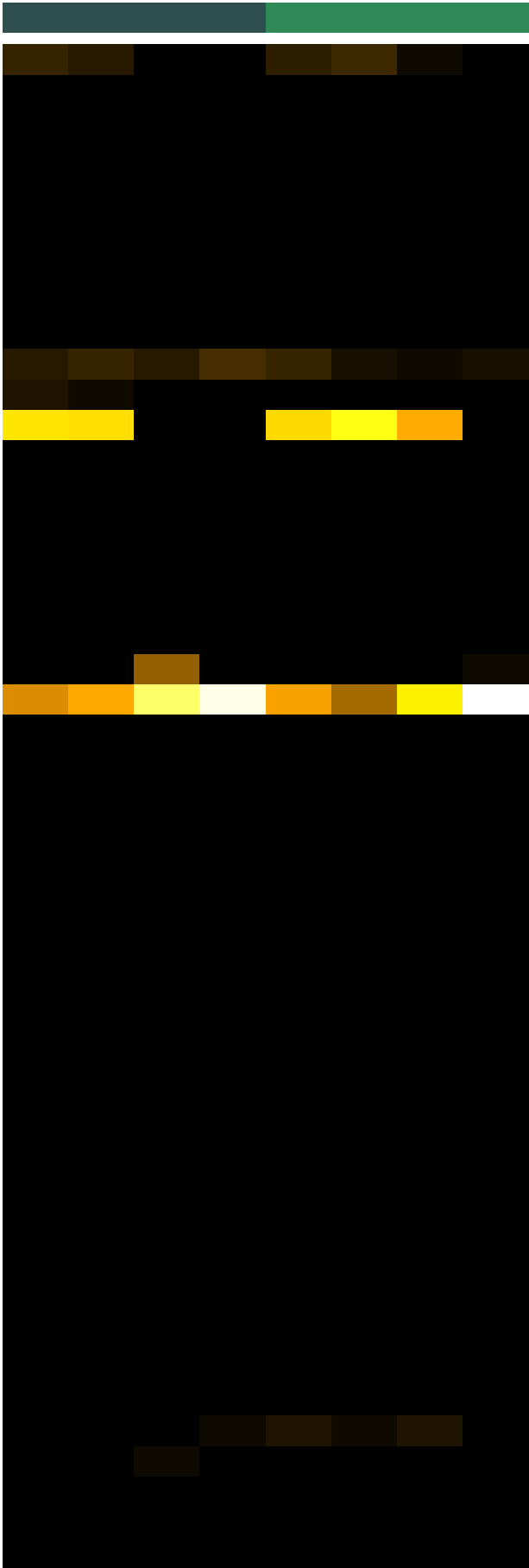


Group

Sham

OVX



Phocaeicola vulgatus
Lachnospiraceae_[G-3] bacterium_MOT-168
Bacteroides caecimuris
Staphylococcus saprophyticus
Turicimonas muris
Alistipes sp._MOT-127
Mammaliicoccus sciuri
Bacteroides acidifaciens
Adlercreutzia equolifaciens
Bacteroides uniformis
Bifidobacterium pseudolongum
Faecalibaculum rodentium
Ligilactobacillus murinus
Lactobacillus intestinalis
Parabacteroides goldsteinii
Lactobacillus johnsonii
Romboutsia ilealis
Dubosiella newyorkensis
Clostridium disporicum
Staphylococcus ureilyticus
Serratia marcescens
Enterococcus faecalis
Mammaliicoccus lentus
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.000%
Alistipes putredinis_nov_92.213%
Alistipes shahii_nov_93.827%
Faecalicoccus acidiformans_nov_89.600%
Parafannyhessea umbonata_nov_92.161%
Parvibacter caecicola_nov_97.609%
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_93.737%
Adlercreutzia caecimuris_nov_93.290%
Parasutterella excrementihominis_nov_94.578%
Peptostreptococcaceae_[G-5] bacterium_HMT_493_nov_90.000%
Alistipes senegalensis_nov_96.920%
Adlercreutzia caecimuris_nov_92.291%
Ihubacter massiliensis_nov_94.572%
Lachnospiraceae_[G-2] bacterium_HMT_096_nov_91.632%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
Turicibacter sanguinis_nov_95.923%
Duncaniella freteri_nov_87.375%
Faecalicatena fissicatena_nov_95.407%
Glucerbacter canis_nov_93.501%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.097%
Anaeromassilibacillus senegalensis_nov_92.489%
Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.125%
Staphylococcus saprophyticus_xylosus
Serratia marcescens_nematodiphila
Olsenella multispecies_sppn1_2_nov_91.966%
Faecalicatena multispecies_sppn2_2_nov_92.067%
Faecalicatena multispecies_sppn3_2_nov_93.737%

Species

F29493.S04

F29493.S03

F29493.S02

F29493.S01

F29493.S08

F29493.S07

F29493.S06

F29493.S05

Samples