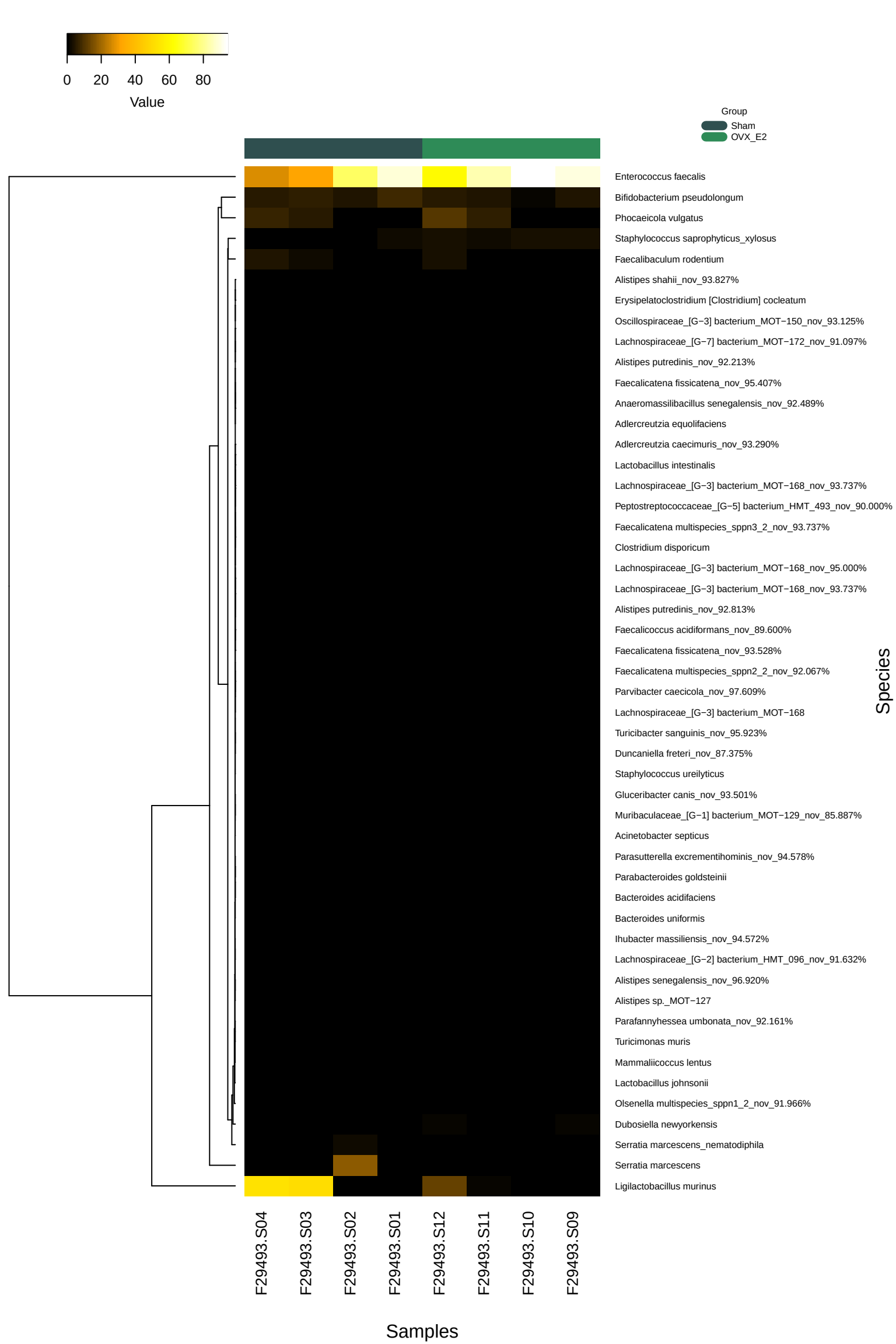
0
20
40
60
80
Value
Group
Sham
OVX_E2
Enterococcus faecalis
Bifidobacterium pseudolongum
Phocaeicola vulgatus
Staphylococcus saprophyticus_xylosus
Faecalibaculum rodentium
Alistipes shahii_nov_93.827%
Erysipelatoclostridium [Clostridium] cocleatum
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.125%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.097%
Alistipes putredinis_nov_92.213%
Faecalicatena fissicatena_nov_95.407%
Anaeromassilibacillus senegalensis_nov_92.489%
Adlercreutzia equolifaciens
Adlercreutzia caecimuris_nov_93.290%
Lactobacillus intestinalis
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_93.737%
Peptostreptococcaceae_[G−5] bacterium_HMT_493_nov_90.000%
Faecalicatena multispecies_sppn3_2_nov_93.737%
Clostridium disporicum
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_95.000%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_93.737%
Alistipes putredinis_nov_92.813%
Faecalicoccus acidiformans_nov_89.600%
Faecalicatena fissicatena_nov_93.528%
Faecalicatena multispecies_sppn2_2_nov_92.067%
Parvibacter caecicola_nov_97.609%
Lachnospiraceae_[G−3] bacterium_MOT−168
Turicibacter sanguinis_nov_95.923%
Duncaniella freteri_nov_87.375%
Staphylococcus ureilyticus
Gluceribacter canis_nov_93.501%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Acinetobacter septicus
Parasutterella excrementihominis_nov_94.578%
Parabacteroides goldsteinii
Bacteroides acidifaciens
Bacteroides uniformis
Ihubacter massiliensis_nov_94.572%
Lachnospiraceae_[G−2] bacterium_HMT_096_nov_91.632%
Alistipes senegalensis_nov_96.920%
Alistipes sp._MOT−127
Parafannyhessea umbonata_nov_92.161%
Turicimonas muris
Mammaliicoccus lentus
Lactobacillus johnsonii
Olsenella multispecies_sppn1_2_nov_91.966%
Dubosiella newyorkensis
Serratia marcescens_nematodiphila
Serratia marcescens
Ligilactobacillus murinus
Species
F29493.S04
F29493.S03
F29493.S02
F29493.S01
F29493.S12
F29493.S11
F29493.S10
F29493.S09
Samples