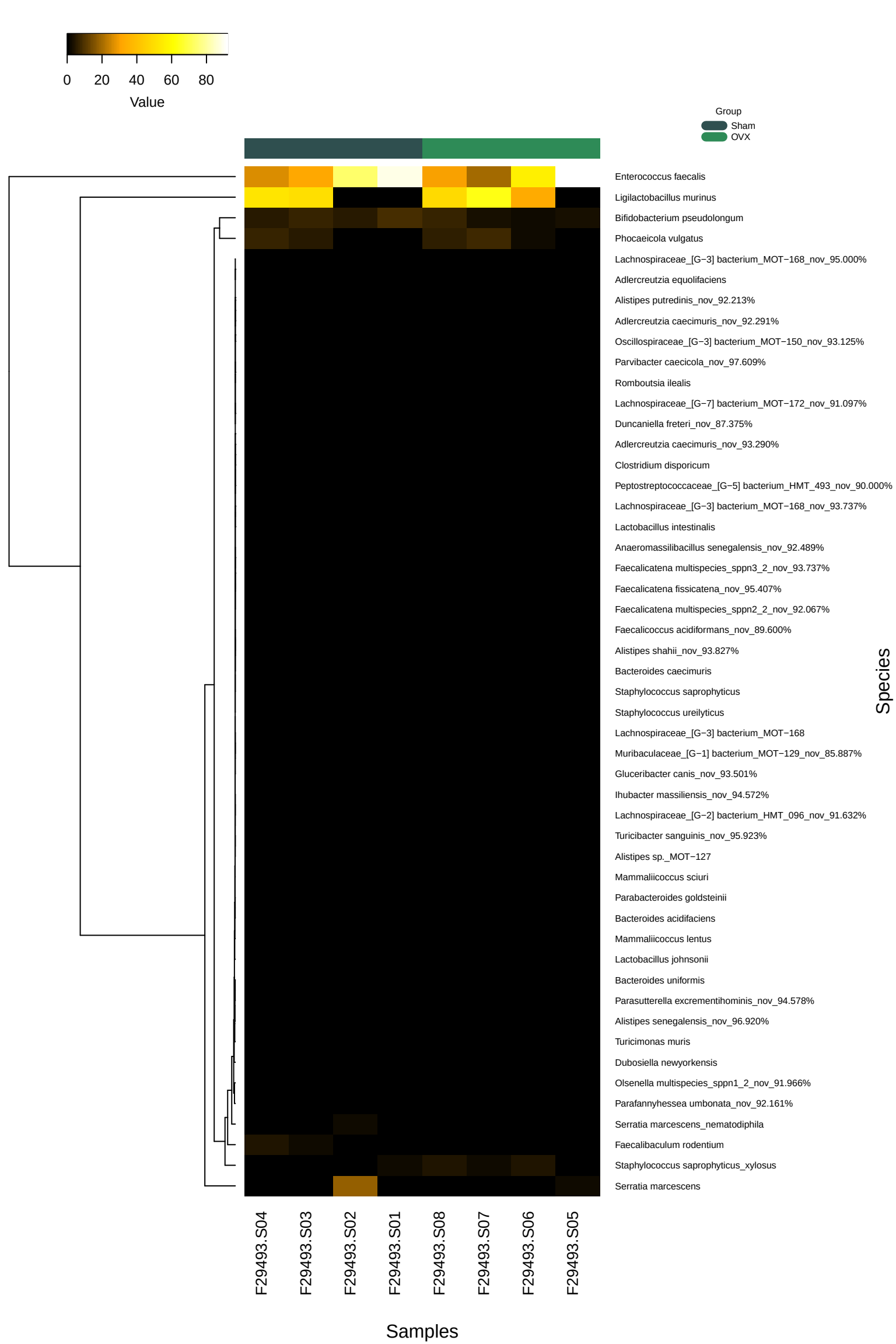
0
20
40
60
80
Value
Group
Sham
OVX
Enterococcus faecalis
Ligilactobacillus murinus
Bifidobacterium pseudolongum
Phocaeicola vulgatus
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_95.000%
Adlercreutzia equolifaciens
Alistipes putredinis_nov_92.213%
Adlercreutzia caecimuris_nov_92.291%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.125%
Parvibacter caecicola_nov_97.609%
Romboutsia ilealis
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.097%
Duncaniella freteri_nov_87.375%
Adlercreutzia caecimuris_nov_93.290%
Clostridium disporicum
Peptostreptococcaceae_[G−5] bacterium_HMT_493_nov_90.000%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_93.737%
Lactobacillus intestinalis
Anaeromassilibacillus senegalensis_nov_92.489%
Faecalicatena multispecies_sppn3_2_nov_93.737%
Faecalicatena fissicatena_nov_95.407%
Faecalicatena multispecies_sppn2_2_nov_92.067%
Faecalicoccus acidiformans_nov_89.600%
Alistipes shahii_nov_93.827%
Bacteroides caecimuris
Staphylococcus saprophyticus
Staphylococcus ureilyticus
Lachnospiraceae_[G−3] bacterium_MOT−168
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Gluceribacter canis_nov_93.501%
Ihubacter massiliensis_nov_94.572%
Lachnospiraceae_[G−2] bacterium_HMT_096_nov_91.632%
Turicibacter sanguinis_nov_95.923%
Alistipes sp._MOT−127
Mammaliicoccus sciuri
Parabacteroides goldsteinii
Bacteroides acidifaciens
Mammaliicoccus lentus
Lactobacillus johnsonii
Bacteroides uniformis
Parasutterella excrementihominis_nov_94.578%
Alistipes senegalensis_nov_96.920%
Turicimonas muris
Dubosiella newyorkensis
Olsenella multispecies_sppn1_2_nov_91.966%
Parafannyhessea umbonata_nov_92.161%
Serratia marcescens_nematodiphila
Faecalibaculum rodentium
Staphylococcus saprophyticus_xylosus
Serratia marcescens
Species
F29493.S04
F29493.S03
F29493.S02
F29493.S01
F29493.S08
F29493.S07
F29493.S06
F29493.S05
Samples