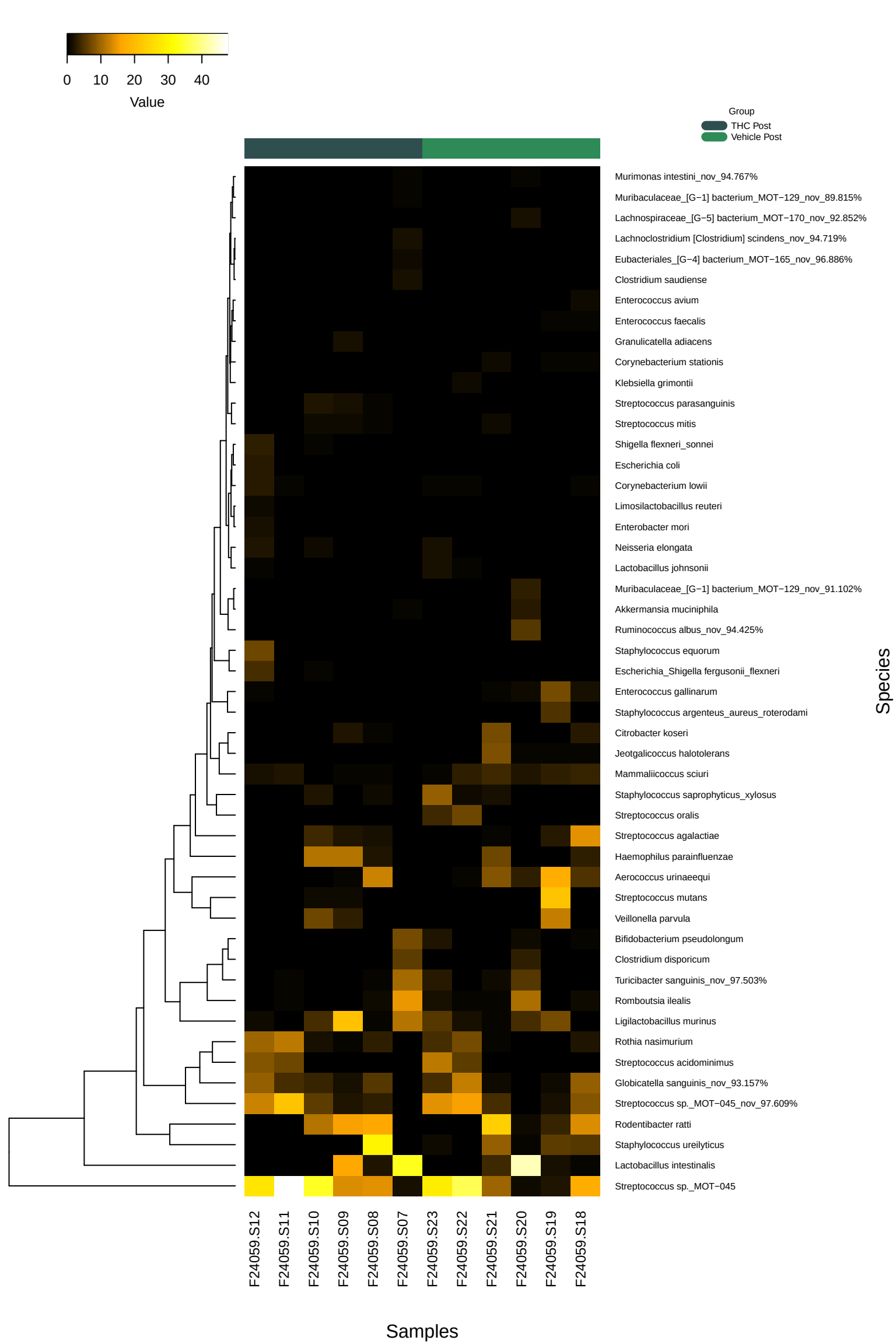
0
10
20
30
40
Value
Group
THC Post
Vehicle Post
Murimonas intestini_nov_94.767%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.815%
Lachnospiraceae_[G−5] bacterium_MOT−170_nov_92.852%
Lachnoclostridium [Clostridium] scindens_nov_94.719%
Eubacteriales_[G−4] bacterium_MOT−165_nov_96.886%
Clostridium saudiense
Enterococcus avium
Enterococcus faecalis
Granulicatella adiacens
Corynebacterium stationis
Klebsiella grimontii
Streptococcus parasanguinis
Streptococcus mitis
Shigella flexneri_sonnei
Escherichia coli
Corynebacterium lowii
Limosilactobacillus reuteri
Enterobacter mori
Neisseria elongata
Lactobacillus johnsonii
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.102%
Akkermansia muciniphila
Ruminococcus albus_nov_94.425%
Staphylococcus equorum
Escherichia_Shigella fergusonii_flexneri
Enterococcus gallinarum
Staphylococcus argenteus_aureus_roterodami
Citrobacter koseri
Jeotgalicoccus halotolerans
Mammaliicoccus sciuri
Staphylococcus saprophyticus_xylosus
Streptococcus oralis
Streptococcus agalactiae
Haemophilus parainfluenzae
Aerococcus urinaeequi
Streptococcus mutans
Veillonella parvula
Bifidobacterium pseudolongum
Clostridium disporicum
Turicibacter sanguinis_nov_97.503%
Romboutsia ilealis
Ligilactobacillus murinus
Rothia nasimurium
Streptococcus acidominimus
Globicatella sanguinis_nov_93.157%
Streptococcus sp._MOT−045_nov_97.609%
Rodentibacter ratti
Staphylococcus ureilyticus
Lactobacillus intestinalis
Streptococcus sp._MOT−045
Species
F24059.S12
F24059.S11
F24059.S10
F24059.S09
F24059.S08
F24059.S07
F24059.S23
F24059.S22
F24059.S21
F24059.S20
F24059.S19
F24059.S18
Samples