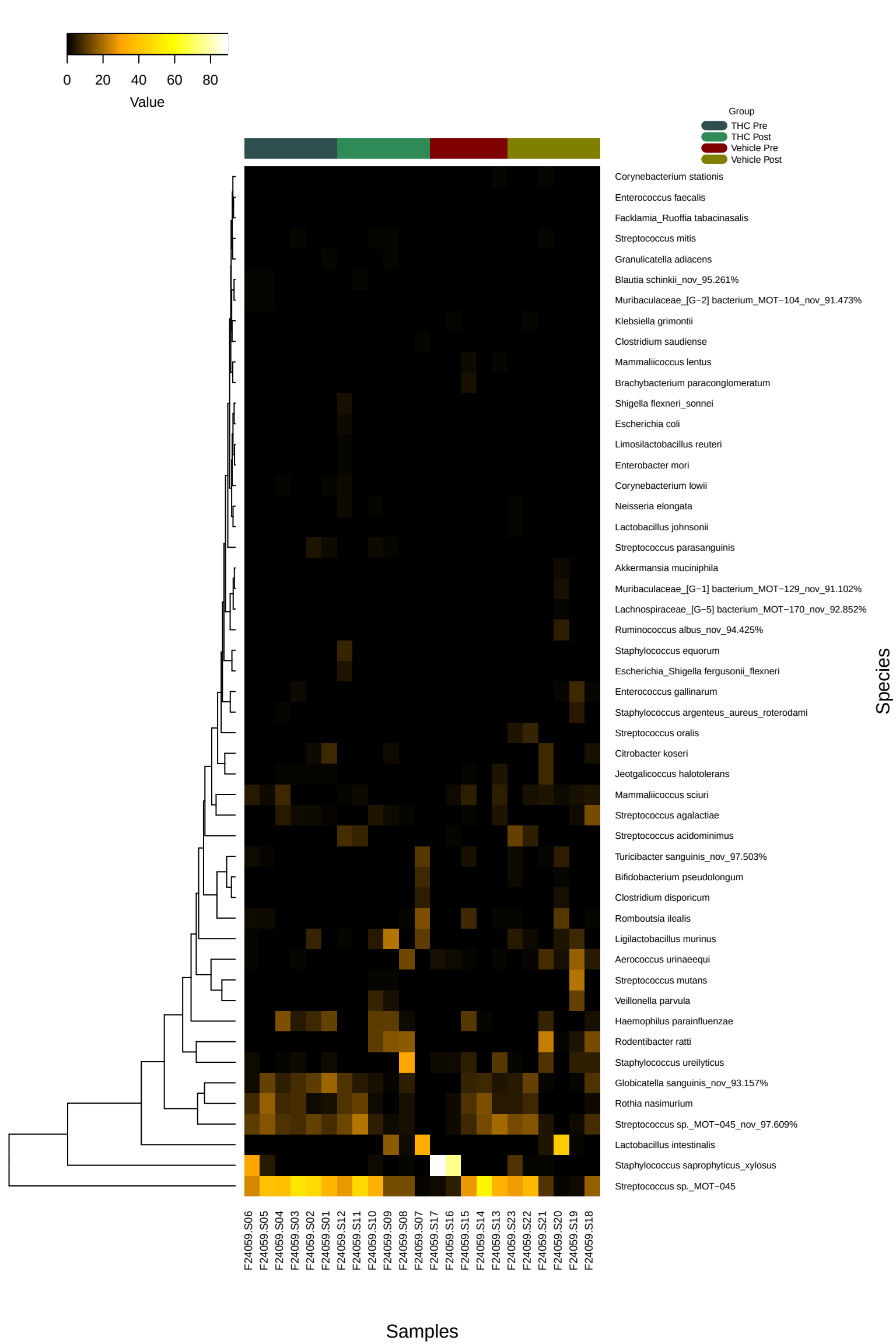
0
20
40
60
80
Value
Group
THC Pre
THC Post
Vehicle Pre
Vehicle Post
Corynebacterium stationis
Enterococcus faecalis
Facklamia_Ruoffia tabacinasalis
Streptococcus mitis
Granulicatella adiacens
Blautia schinkii_nov_95.261%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.473%
Klebsiella grimontii
Clostridium saudiense
Mammaliicoccus lentus
Brachybacterium paraconglomeratum
Shigella flexneri_sonnei
Escherichia coli
Limosilactobacillus reuteri
Enterobacter mori
Corynebacterium lowii
Neisseria elongata
Lactobacillus johnsonii
Streptococcus parasanguinis
Akkermansia muciniphila
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.102%
Lachnospiraceae_[G−5] bacterium_MOT−170_nov_92.852%
Ruminococcus albus_nov_94.425%
Staphylococcus equorum
Escherichia_Shigella fergusonii_flexneri
Enterococcus gallinarum
Staphylococcus argenteus_aureus_roterodami
Streptococcus oralis
Citrobacter koseri
Jeotgalicoccus halotolerans
Mammaliicoccus sciuri
Streptococcus agalactiae
Streptococcus acidominimus
Turicibacter sanguinis_nov_97.503%
Bifidobacterium pseudolongum
Clostridium disporicum
Romboutsia ilealis
Ligilactobacillus murinus
Aerococcus urinaeequi
Streptococcus mutans
Veillonella parvula
Haemophilus parainfluenzae
Rodentibacter ratti
Staphylococcus ureilyticus
Globicatella sanguinis_nov_93.157%
Rothia nasimurium
Streptococcus sp._MOT−045_nov_97.609%
Lactobacillus intestinalis
Staphylococcus saprophyticus_xylosus
Streptococcus sp._MOT−045
Species
F24059.S06
F24059.S05
F24059.S04
F24059.S03
F24059.S02
F24059.S01
F24059.S12
F24059.S11
F24059.S10
F24059.S09
F24059.S08
F24059.S07
F24059.S17
F24059.S16
F24059.S15
F24059.S14
F24059.S13
F24059.S23
F24059.S22
F24059.S21
F24059.S20
F24059.S19
F24059.S18
Samples