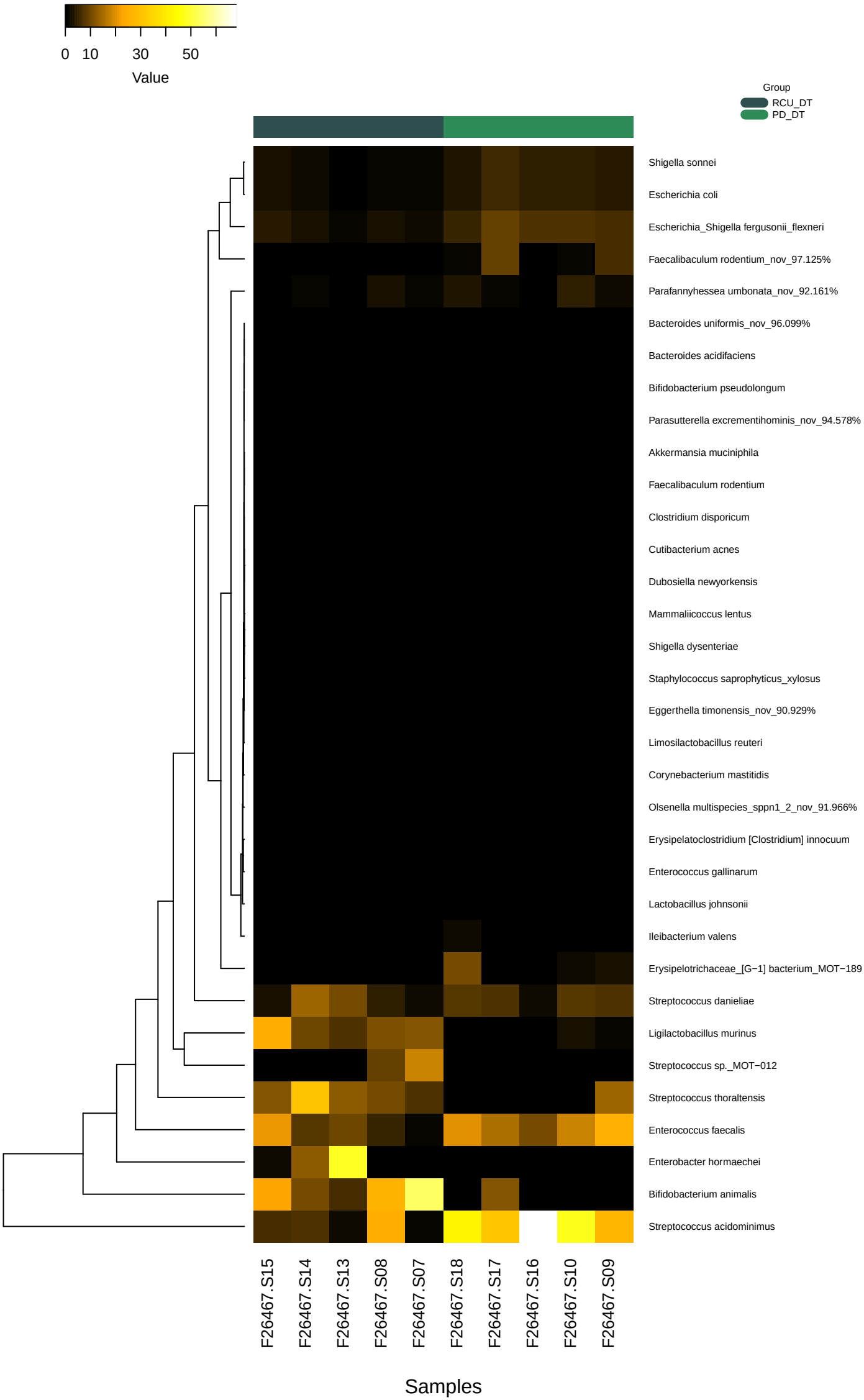
0
10
30
50
Value
Group
RCU_DT
PD_DT
Shigella sonnei
Escherichia coli
Escherichia_Shigella fergusonii_flexneri
Faecalibaculum rodentium_nov_97.125%
Parafannyhessea umbonata_nov_92.161%
Bacteroides uniformis_nov_96.099%
Bacteroides acidifaciens
Bifidobacterium pseudolongum
Parasutterella excrementihominis_nov_94.578%
Akkermansia muciniphila
Faecalibaculum rodentium
Clostridium disporicum
Cutibacterium acnes
Dubosiella newyorkensis
Mammaliicoccus lentus
Shigella dysenteriae
Staphylococcus saprophyticus_xylosus
Eggerthella timonensis_nov_90.929%
Limosilactobacillus reuteri
Corynebacterium mastitidis
Olsenella multispecies_sppn1_2_nov_91.966%
Erysipelatoclostridium [Clostridium] innocuum
Enterococcus gallinarum
Lactobacillus johnsonii
Ileibacterium valens
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Streptococcus danieliae
Ligilactobacillus murinus
Streptococcus sp._MOT−012
Streptococcus thoraltensis
Enterococcus faecalis
Enterobacter hormaechei
Bifidobacterium animalis
Streptococcus acidominimus
Species
F26467.S15
F26467.S14
F26467.S13
F26467.S08
F26467.S07
F26467.S18
F26467.S17
F26467.S16
F26467.S10
F26467.S09
Samples