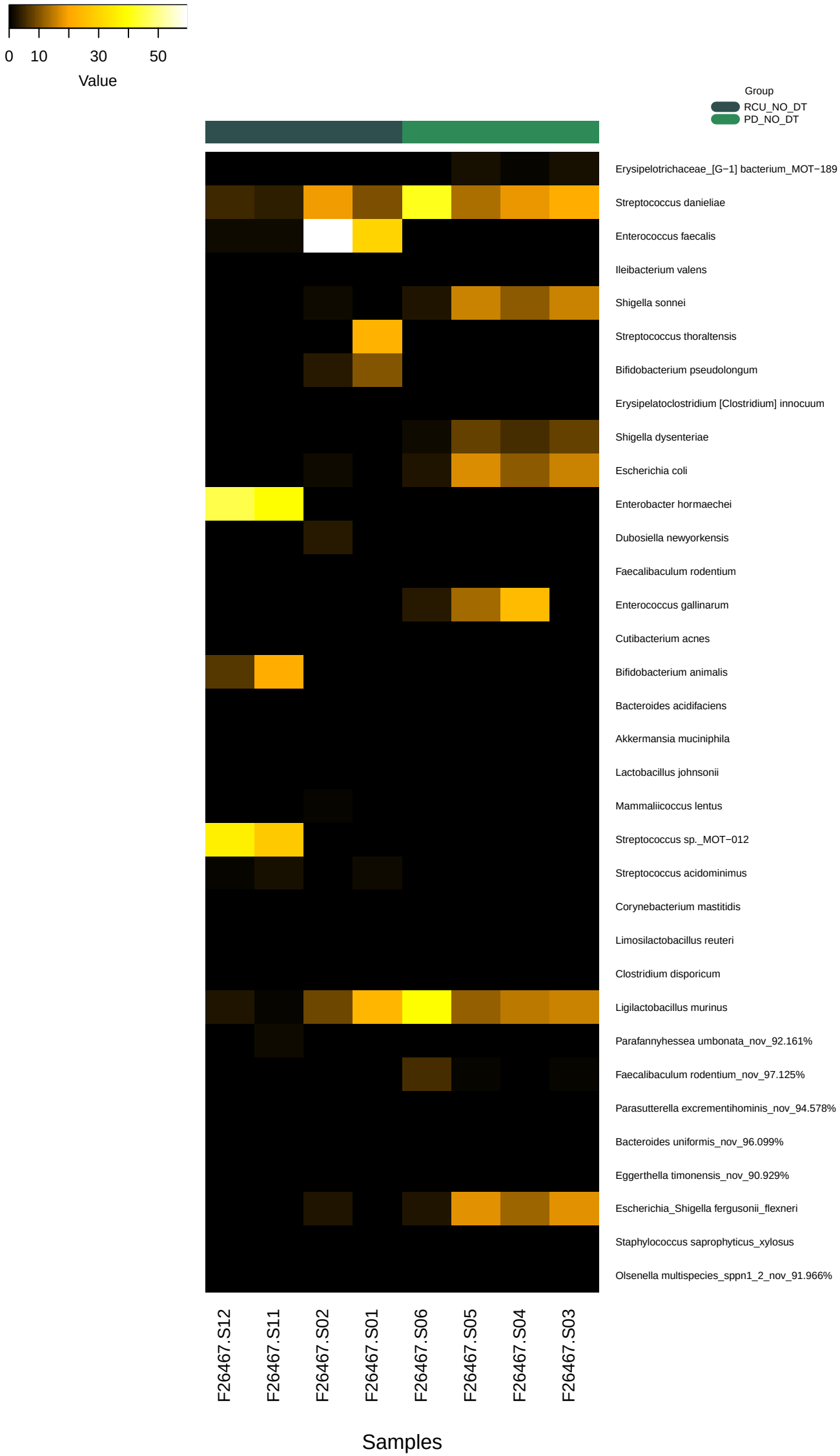
0
10
30
50
Value
Group
RCU_NO_DT
PD_NO_DT
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Streptococcus danieliae
Enterococcus faecalis
Ileibacterium valens
Shigella sonnei
Streptococcus thoraltensis
Bifidobacterium pseudolongum
Erysipelatoclostridium [Clostridium] innocuum
Shigella dysenteriae
Escherichia coli
Enterobacter hormaechei
Dubosiella newyorkensis
Faecalibaculum rodentium
Enterococcus gallinarum
Cutibacterium acnes
Bifidobacterium animalis
Bacteroides acidifaciens
Akkermansia muciniphila
Lactobacillus johnsonii
Mammaliicoccus lentus
Streptococcus sp._MOT−012
Streptococcus acidominimus
Corynebacterium mastitidis
Limosilactobacillus reuteri
Clostridium disporicum
Ligilactobacillus murinus
Parafannyhessea umbonata_nov_92.161%
Faecalibaculum rodentium_nov_97.125%
Parasutterella excrementihominis_nov_94.578%
Bacteroides uniformis_nov_96.099%
Eggerthella timonensis_nov_90.929%
Escherichia_Shigella fergusonii_flexneri
Staphylococcus saprophyticus_xylosus
Olsenella multispecies_sppn1_2_nov_91.966%
Species
F26467.S12
F26467.S11
F26467.S02
F26467.S01
F26467.S06
F26467.S05
F26467.S04
F26467.S03
Samples