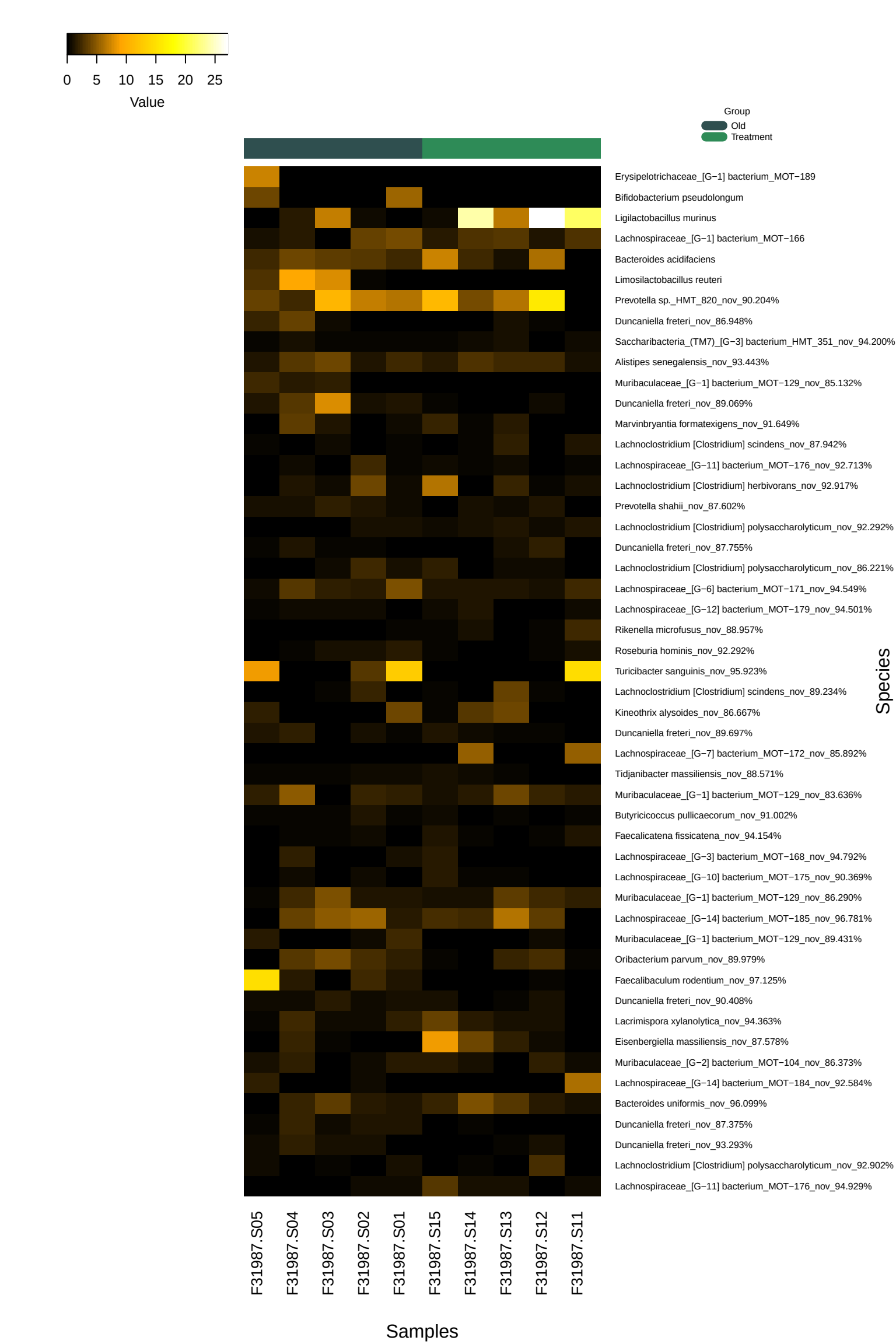

0
5
10
15
20
25
Value
Group
Old
Treatment
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Bifidobacterium pseudolongum
Ligilactobacillus murinus
Lachnospiraceae_[G−1] bacterium_MOT−166
Bacteroides acidifaciens
Limosilactobacillus reuteri
Prevotella sp._HMT_820_nov_90.204%
Duncaniella freteri_nov_86.948%
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351_nov_94.200%
Alistipes senegalensis_nov_93.443%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.132%
Duncaniella freteri_nov_89.069%
Marvinbryantia formatexigens_nov_91.649%
Lachnoclostridium [Clostridium] scindens_nov_87.942%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_92.713%
Lachnoclostridium [Clostridium] herbivorans_nov_92.917%
Prevotella shahii_nov_87.602%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.292%
Duncaniella freteri_nov_87.755%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_86.221%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.549%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_94.501%
Rikenella microfusus_nov_88.957%
Roseburia hominis_nov_92.292%
Turicibacter sanguinis_nov_95.923%
Lachnoclostridium [Clostridium] scindens_nov_89.234%
Kineothrix alysoides_nov_86.667%
Duncaniella freteri_nov_89.697%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_85.892%
Tidjanibacter massiliensis_nov_88.571%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_83.636%
Butyricicoccus pullicaecorum_nov_91.002%
Faecalicatena fissicatena_nov_94.154%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.792%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_90.369%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.290%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_96.781%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.431%
Oribacterium parvum_nov_89.979%
Faecalibaculum rodentium_nov_97.125%
Duncaniella freteri_nov_90.408%
Lacrimispora xylanolytica_nov_94.363%
Eisenbergiella massiliensis_nov_87.578%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.373%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_92.584%
Bacteroides uniformis_nov_96.099%
Duncaniella freteri_nov_87.375%
Duncaniella freteri_nov_93.293%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.902%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_94.929%
Species
F31987.S05
F31987.S04
F31987.S03
F31987.S02
F31987.S01
F31987.S15
F31987.S14
F31987.S13
F31987.S12
F31987.S11
Samples