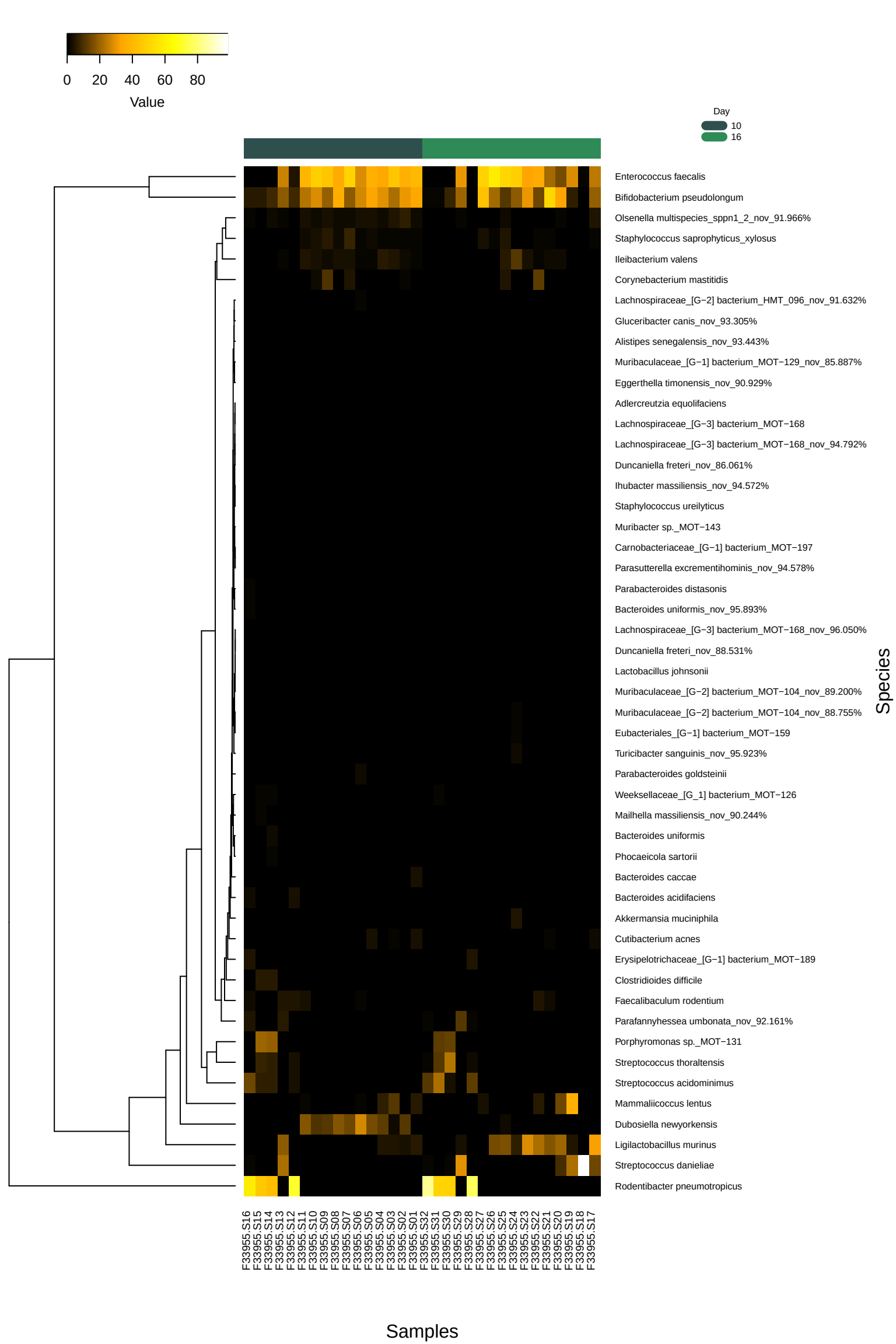

0
20
40
60
80
Value
Day
10
16
Enterococcus faecalis
Bifidobacterium pseudolongum
Olsenella multispecies_sppn1_2_nov_91.966%
Staphylococcus saprophyticus_xylosus
Ileibacterium valens
Corynebacterium mastitidis
Lachnospiraceae_[G−2] bacterium_HMT_096_nov_91.632%
Gluceribacter canis_nov_93.305%
Alistipes senegalensis_nov_93.443%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Eggerthella timonensis_nov_90.929%
Adlercreutzia equolifaciens
Lachnospiraceae_[G−3] bacterium_MOT−168
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.792%
Duncaniella freteri_nov_86.061%
Ihubacter massiliensis_nov_94.572%
Staphylococcus ureilyticus
Muribacter sp._MOT−143
Carnobacteriaceae_[G−1] bacterium_MOT−197
Parasutterella excrementihominis_nov_94.578%
Parabacteroides distasonis
Bacteroides uniformis_nov_95.893%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_96.050%
Duncaniella freteri_nov_88.531%
Lactobacillus johnsonii
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.200%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.755%
Eubacteriales_[G−1] bacterium_MOT−159
Turicibacter sanguinis_nov_95.923%
Parabacteroides goldsteinii
Weeksellaceae_[G_1] bacterium_MOT−126
Mailhella massiliensis_nov_90.244%
Bacteroides uniformis
Phocaeicola sartorii
Bacteroides caccae
Bacteroides acidifaciens
Akkermansia muciniphila
Cutibacterium acnes
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Clostridioides difficile
Faecalibaculum rodentium
Parafannyhessea umbonata_nov_92.161%
Porphyromonas sp._MOT−131
Streptococcus thoraltensis
Streptococcus acidominimus
Mammaliicoccus lentus
Dubosiella newyorkensis
Ligilactobacillus murinus
Streptococcus danieliae
Rodentibacter pneumotropicus
Species
F3955.S16
F3955.S15
F3955.S14
F3955.S13
F3955.S12
F3955.S11
F3955.S10
F3955.S09
F3955.S08
F3955.S07
F3955.S06
F3955.S05
F3955.S04
F3955.S03
F3955.S02
F3955.S01
F3955.S32
F3955.S31
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F3955.S25
F3955.S24
F3955.S23
F3955.S22
F3955.S21
F3955.S20
F3955.S19
F3955.S18
F3955.S17
Samples