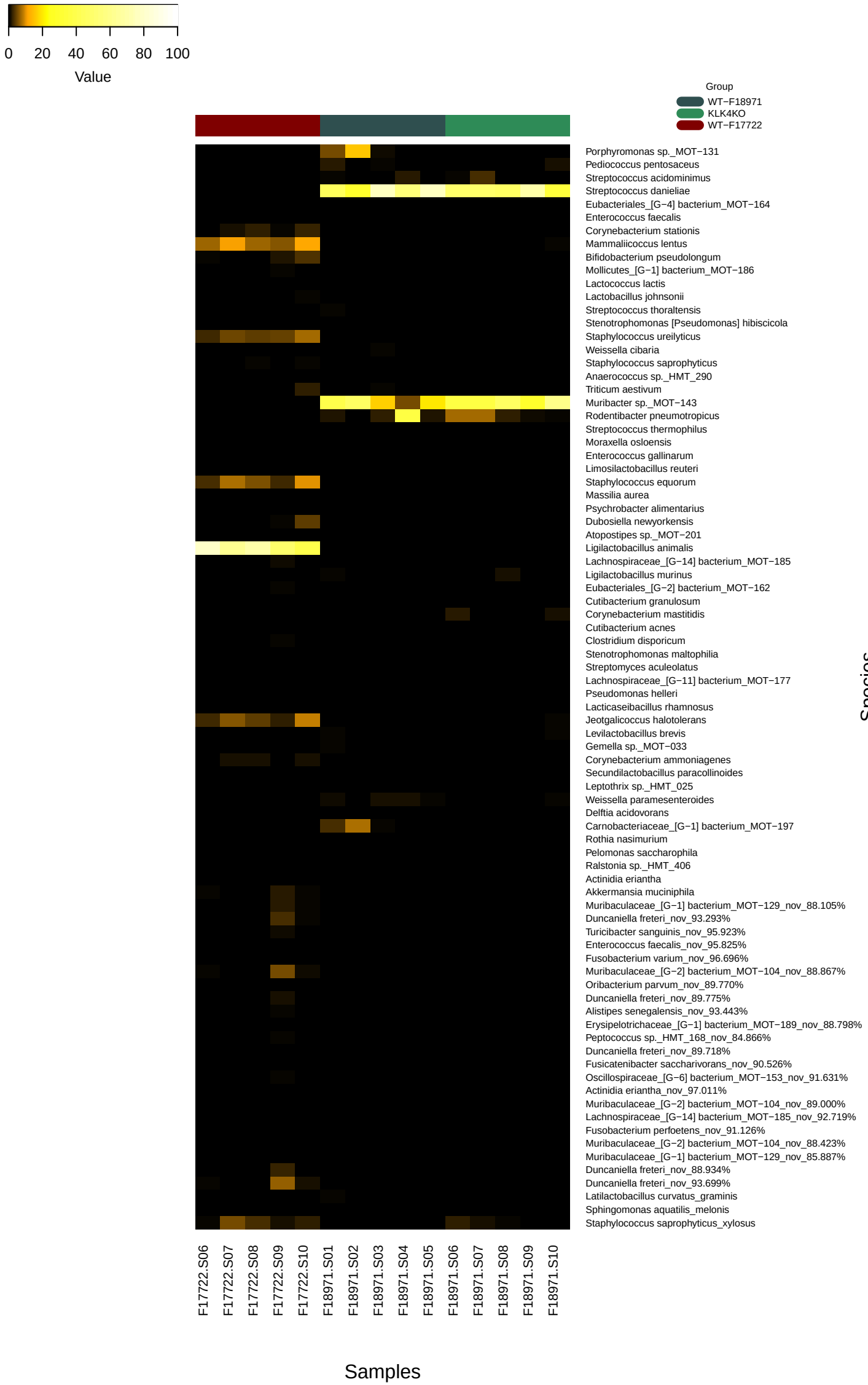
0
20
40
60
80
100
Value
Group
WT−F18971
KLK4KO
WT−F17722
Porphyromonas sp._MOT−131
Pediococcus pentosaceus
Streptococcus acidominimus
Streptococcus danieliae
Eubacteriales_[G−4] bacterium_MOT−164
Enterococcus faecalis
Corynebacterium stationis
Mammaliicoccus lentus
Bifidobacterium pseudolongum
Mollicutes_[G−1] bacterium_MOT−186
Lactococcus lactis
Lactobacillus johnsonii
Streptococcus thoraltensis
Stenotrophomonas [Pseudomonas] hibiscicola
Staphylococcus ureilyticus
Weissella cibaria
Staphylococcus saprophyticus
Anaerococcus sp._HMT_290
Triticum aestivum
Muribacter sp._MOT−143
Rodentibacter pneumotropicus
Streptococcus thermophilus
Moraxella osloensis
Enterococcus gallinarum
Limosilactobacillus reuteri
Staphylococcus equorum
Massilia aurea
Psychrobacter alimentarius
Dubosiella newyorkensis
Atopostipes sp._MOT−201
Ligilactobacillus animalis
Lachnospiraceae_[G−14] bacterium_MOT−185
Ligilactobacillus murinus
Eubacteriales_[G−2] bacterium_MOT−162
Cutibacterium granulosum
Corynebacterium mastitidis
Cutibacterium acnes
Clostridium disporicum
Stenotrophomonas maltophilia
Streptomyces aculeolatus
Lachnospiraceae_[G−11] bacterium_MOT−177
Pseudomonas helleri
Lacticaseibacillus rhamnosus
Jeotgalicoccus halotolerans
Levilactobacillus brevis
Gemella sp._MOT−033
Corynebacterium ammoniagenes
Secundilactobacillus paracollinoides
Leptothrix sp._HMT_025
Weissella paramesenteroides
Delftia acidovorans
Carnobacteriaceae_[G−1] bacterium_MOT−197
Rothia nasimurium
Pelomonas saccharophila
Ralstonia sp._HMT_406
Actinidia eriantha
Akkermansia muciniphila
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.105%
Duncaniella freteri_nov_93.293%
Turicibacter sanguinis_nov_95.923%
Enterococcus faecalis_nov_95.825%
Fusobacterium varium_nov_96.696%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.867%
Oribacterium parvum_nov_89.770%
Duncaniella freteri_nov_89.775%
Alistipes senegalensis_nov_93.443%
Erysipelotrichaceae_[G−1] bacterium_MOT−189_nov_88.798%
Peptococcus sp._HMT_168_nov_84.866%
Duncaniella freteri_nov_89.718%
Fusicatenibacter saccharivorans_nov_90.526%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_91.631%
Actinidia eriantha_nov_97.011%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.000%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.719%
Fusobacterium perfoetens_nov_91.126%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.423%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Duncaniella freteri_nov_88.934%
Duncaniella freteri_nov_93.699%
Latilactobacillus curvatus_graminis
Sphingomonas aquatilis_melonis
Staphylococcus saprophyticus_xylosus
Species
F17722.S06
F17722.S07
F17722.S08
F17722.S09
F17722.S10
F18971.S01
F18971.S02
F18971.S03
F18971.S04
F18971.S05
F18971.S06
F18971.S07
F18971.S08
F18971.S09
F18971.S10
Samples