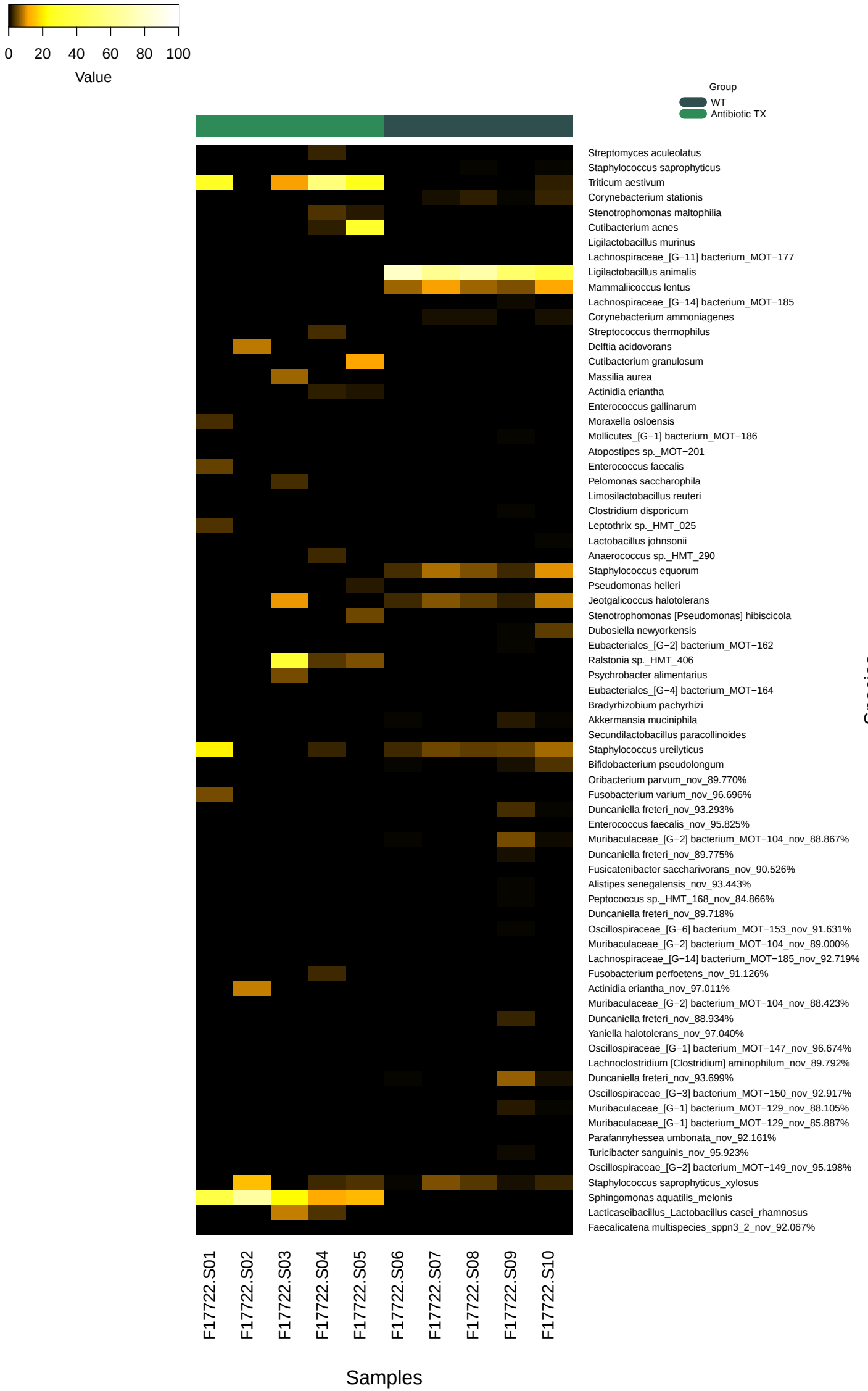
0
20
40
60
80
100
Value
Group
WT
Antibiotic TX
Streptomyces aculeolatus
Staphylococcus saprophyticus
Triticum aestivum
Corynebacterium stationis
Stenotrophomonas maltophilia
Cutibacterium acnes
Ligilactobacillus murinus
Lachnospiraceae_[G−11] bacterium_MOT−177
Ligilactobacillus animalis
Mammaliicoccus lentus
Lachnospiraceae_[G−14] bacterium_MOT−185
Corynebacterium ammoniagenes
Streptococcus thermophilus
Delftia acidovorans
Cutibacterium granulosum
Massilia aurea
Actinidia eriantha
Enterococcus gallinarum
Moraxella osloensis
Mollicutes_[G−1] bacterium_MOT−186
Atopostipes sp._MOT−201
Enterococcus faecalis
Pelomonas saccharophila
Limosilactobacillus reuteri
Clostridium disporicum
Leptothrix sp._HMT_025
Lactobacillus johnsonii
Anaerococcus sp._HMT_290
Staphylococcus equorum
Pseudomonas helleri
Jeotgalicoccus halotolerans
Stenotrophomonas [Pseudomonas] hibiscicola
Dubosiella newyorkensis
Eubacteriales_[G−2] bacterium_MOT−162
Ralstonia sp._HMT_406
Psychrobacter alimentarius
Eubacteriales_[G−4] bacterium_MOT−164
Bradyrhizobium pachyrhizi
Akkermansia muciniphila
Secundilactobacillus paracollinoides
Staphylococcus ureilyticus
Bifidobacterium pseudolongum
Oribacterium parvum_nov_89.770%
Fusobacterium varium_nov_96.696%
Duncaniella freteri_nov_93.293%
Enterococcus faecalis_nov_95.825%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.867%
Duncaniella freteri_nov_89.775%
Fusicatenibacter saccharivorans_nov_90.526%
Alistipes senegalensis_nov_93.443%
Peptococcus sp._HMT_168_nov_84.866%
Duncaniella freteri_nov_89.718%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_91.631%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.000%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.719%
Fusobacterium perfoetens_nov_91.126%
Actinidia eriantha_nov_97.011%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.423%
Duncaniella freteri_nov_88.934%
Yaniella halotolerans_nov_97.040%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_96.674%
Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
Duncaniella freteri_nov_93.699%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_92.917%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.105%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Parafannyhessea umbonata_nov_92.161%
Turicibacter sanguinis_nov_95.923%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.198%
Staphylococcus saprophyticus_xylosus
Sphingomonas aquatilis_melonis
Lacticaseibacillus_Lactobacillus casei_rhamnosus
Faecalicatena multispecies_sppn3_2_nov_92.067%
Species
F17722.S01
F17722.S02
F17722.S03
F17722.S04
F17722.S05
F17722.S06
F17722.S07
F17722.S08
F17722.S09
F17722.S10
Samples