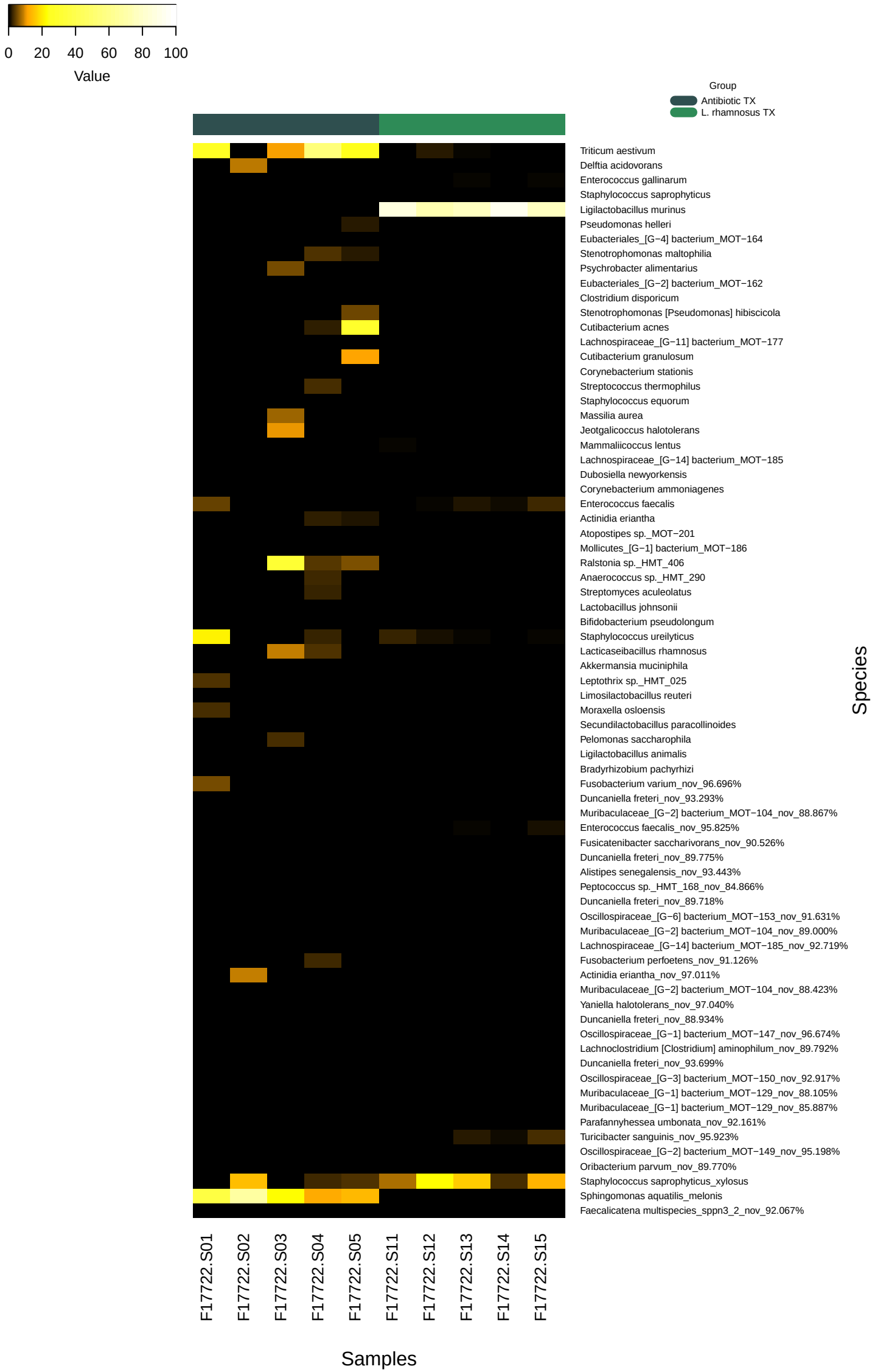
0
20
40
60
80
100
Value
Group
Antibiotic TX
L. rhamnosus TX
Triticum aestivum
Delftia acidovorans
Enterococcus gallinarum
Staphylococcus saprophyticus
Ligilactobacillus murinus
Pseudomonas helleri
Eubacteriales_[G−4] bacterium_MOT−164
Stenotrophomonas maltophilia
Psychrobacter alimentarius
Eubacteriales_[G−2] bacterium_MOT−162
Clostridium disporicum
Stenotrophomonas [Pseudomonas] hibiscicola
Cutibacterium acnes
Lachnospiraceae_[G−11] bacterium_MOT−177
Cutibacterium granulosum
Corynebacterium stationis
Streptococcus thermophilus
Staphylococcus equorum
Massilia aurea
Jeotgalicoccus halotolerans
Mammaliicoccus lentus
Lachnospiraceae_[G−14] bacterium_MOT−185
Dubosiella newyorkensis
Corynebacterium ammoniagenes
Enterococcus faecalis
Actinidia eriantha
Atopostipes sp._MOT−201
Mollicutes_[G−1] bacterium_MOT−186
Ralstonia sp._HMT_406
Anaerococcus sp._HMT_290
Streptomyces aculeolatus
Lactobacillus johnsonii
Bifidobacterium pseudolongum
Staphylococcus ureilyticus
Lacticaseibacillus rhamnosus
Akkermansia muciniphila
Leptothrix sp._HMT_025
Limosilactobacillus reuteri
Moraxella osloensis
Secundilactobacillus paracollinoides
Pelomonas saccharophila
Ligilactobacillus animalis
Bradyrhizobium pachyrhizi
Fusobacterium varium_nov_96.696%
Duncaniella freteri_nov_93.293%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.867%
Enterococcus faecalis_nov_95.825%
Fusicatenibacter saccharivorans_nov_90.526%
Duncaniella freteri_nov_89.775%
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%
Yaniella halotolerans_nov_97.040%
Duncaniella freteri_nov_88.934%
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%
Oribacterium parvum_nov_89.770%
Staphylococcus saprophyticus_xylosus
Sphingomonas aquatilis_melonis
Faecalicatena multispecies_sppn3_2_nov_92.067%
Species
F17722.S01
F17722.S02
F17722.S03
F17722.S04
F17722.S05
F17722.S11
F17722.S12
F17722.S13
F17722.S14
F17722.S15
Samples