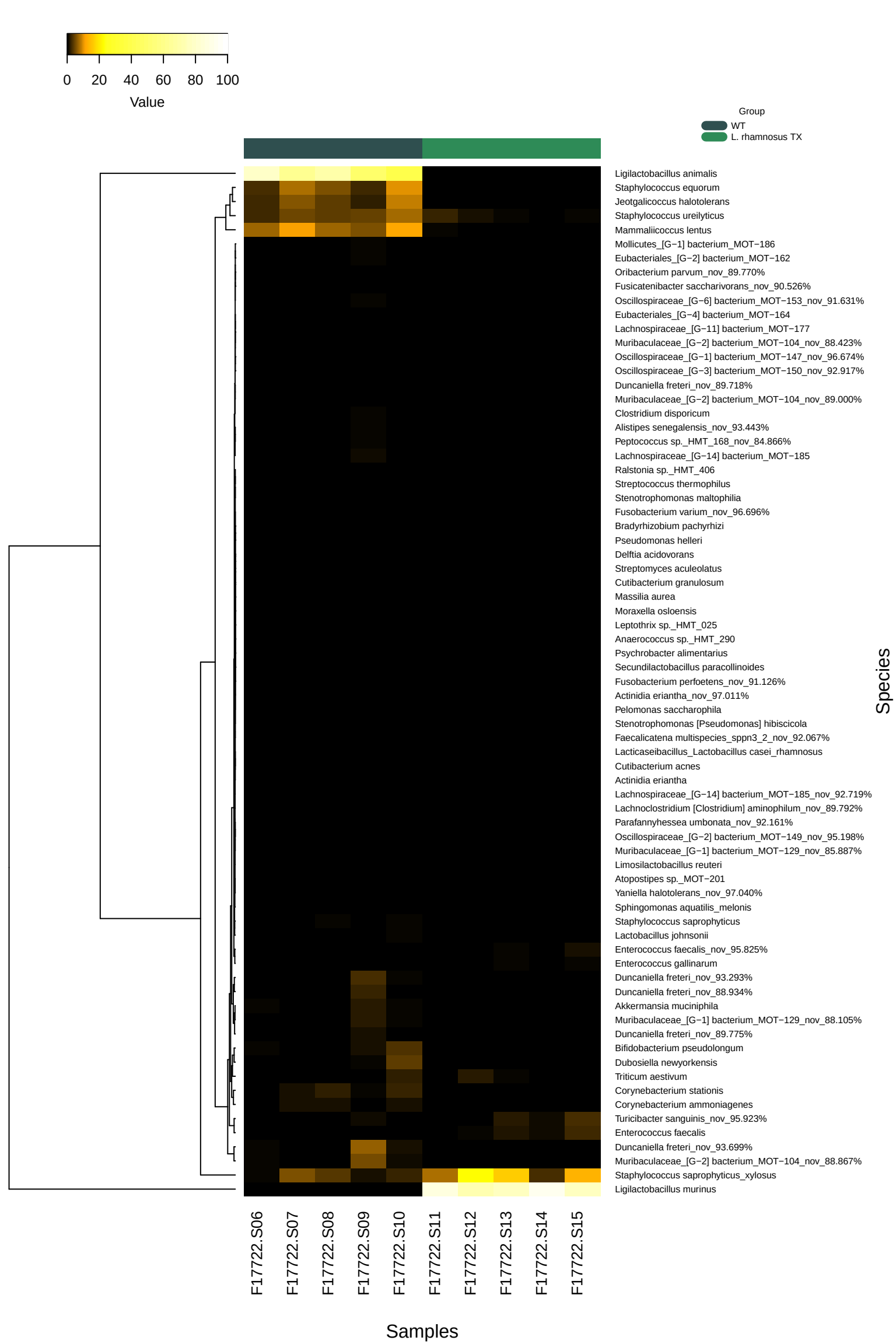
0
20
40
60
80
100
Value
Group
WT
L. rhamnosus TX
Ligilactobacillus animalis
Staphylococcus equorum
Jeotgalicoccus halotolerans
Staphylococcus ureilyticus
Mammaliicoccus lentus
Mollicutes_[G−1] bacterium_MOT−186
Eubacteriales_[G−2] bacterium_MOT−162
Oribacterium parvum_nov_89.770%
Fusicatenibacter saccharivorans_nov_90.526%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_91.631%
Eubacteriales_[G−4] bacterium_MOT−164
Lachnospiraceae_[G−11] bacterium_MOT−177
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.423%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_96.674%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_92.917%
Duncaniella freteri_nov_89.718%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.000%
Clostridium disporicum
Alistipes senegalensis_nov_93.443%
Peptococcus sp._HMT_168_nov_84.866%
Lachnospiraceae_[G−14] bacterium_MOT−185
Ralstonia sp._HMT_406
Streptococcus thermophilus
Stenotrophomonas maltophilia
Fusobacterium varium_nov_96.696%
Bradyrhizobium pachyrhizi
Pseudomonas helleri
Delftia acidovorans
Streptomyces aculeolatus
Cutibacterium granulosum
Massilia aurea
Moraxella osloensis
Leptothrix sp._HMT_025
Anaerococcus sp._HMT_290
Psychrobacter alimentarius
Secundilactobacillus paracollinoides
Fusobacterium perfoetens_nov_91.126%
Actinidia eriantha_nov_97.011%
Pelomonas saccharophila
Stenotrophomonas [Pseudomonas] hibiscicola
Faecalicatena multispecies_sppn3_2_nov_92.067%
Lacticaseibacillus_Lactobacillus casei_rhamnosus
Cutibacterium acnes
Actinidia eriantha
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.719%
Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
Parafannyhessea umbonata_nov_92.161%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.198%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Limosilactobacillus reuteri
Atopostipes sp._MOT−201
Yaniella halotolerans_nov_97.040%
Sphingomonas aquatilis_melonis
Staphylococcus saprophyticus
Lactobacillus johnsonii
Enterococcus faecalis_nov_95.825%
Enterococcus gallinarum
Duncaniella freteri_nov_93.293%
Duncaniella freteri_nov_88.934%
Akkermansia muciniphila
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.105%
Duncaniella freteri_nov_89.775%
Bifidobacterium pseudolongum
Dubosiella newyorkensis
Triticum aestivum
Corynebacterium stationis
Corynebacterium ammoniagenes
Turicibacter sanguinis_nov_95.923%
Enterococcus faecalis
Duncaniella freteri_nov_93.699%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.867%
Staphylococcus saprophyticus_xylosus
Ligilactobacillus murinus
Species
F17722.S06
F17722.S07
F17722.S08
F17722.S09
F17722.S10
F17722.S11
F17722.S12
F17722.S13
F17722.S14
F17722.S15
Samples