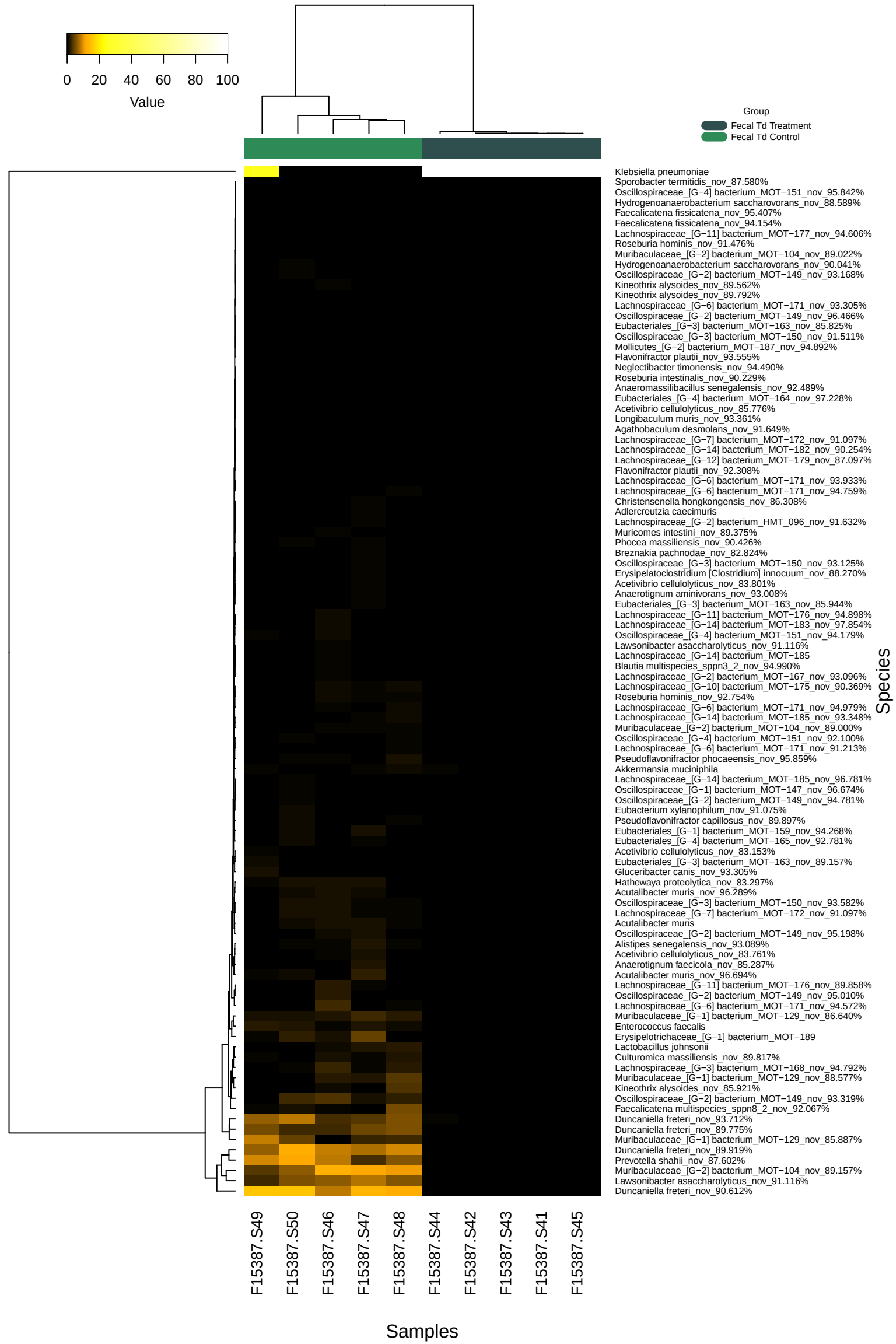
0 20 40 60 80 100
Value
Group
Fecal Td Treatment
Fecal Td Control
Klebsiella pneumoniae
Sporobacter termitidis_nov_87.580%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_95.842%
Hydrogenoanaerobacterium saccharovorans_nov_88.589%
Faecalicatena fissicatena_nov_95.407%
Faecalicatena fissicatena_nov_94.154%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_94.606%
Roseburia hominis_nov_91.476%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.022%
Hydrogenoanaerobacterium saccharovorans_nov_90.041%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.168%
Kineothrix alysoides_nov_89.562%
Kineothrix alysoides_nov_89.792%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_93.305%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.466%
Eubacteriales_[G−3] bacterium_MOT−163_nov_85.825%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_91.511%
Mollicutes_[G−2] bacterium_MOT−187_nov_94.892%
Flavonifractor plautii_nov_93.555%
Neglectibacter timonensis_nov_94.490%
Roseburia intestinalis_nov_90.229%
Anaeromassilibacillus senegalensis_nov_92.489%
Eubacteriales_[G−4] bacterium_MOT−164_nov_97.228%
Acetivibrio cellulolyticus_nov_85.776%
Longibaculum muris_nov_93.361%
Agathobaculum desmolans_nov_91.649%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.097%
Lachnospiraceae_[G−14] bacterium_MOT−182_nov_90.254%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_87.097%
Flavonifractor plautii_nov_92.308%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_93.933%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.759%
Christensenella hongkongensis_nov_86.308%
Adlercreutzia caecimuris
Lachnospiraceae_[G−2] bacterium_HMT_096_nov_91.632%
Muricomes intestini_nov_89.375%
Phocea massiliensis_nov_90.426%
Breznakia pachnodae_nov_82.824%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.125%
Erysipelatoclostridium [Clostridium] innocuum_nov_88.270%
Acetivibrio cellulolyticus_nov_83.801%
Anaerotignum aminivorans_nov_93.008%
Eubacteriales_[G−3] bacterium_MOT−163_nov_85.944%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_94.898%
Lachnospiraceae_[G−14] bacterium_MOT−183_nov_97.854%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_94.179%
Lawsonibacter asaccharolyticus_nov_91.116%
Lachnospiraceae_[G−14] bacterium_MOT−185
Blautia multispecies_sppn3_2_nov_94.990%
Lachnospiraceae_[G−2] bacterium_MOT−167_nov_93.096%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_90.369%
Roseburia hominis_nov_92.754%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.979%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_93.348%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.000%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_92.100%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_91.213%
Pseudoflavonifractor phocaeensis_nov_95.859%
Akkermansia muciniphila
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_96.781%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_96.674%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_94.781%
Eubacterium xylanophilum_nov_91.075%
Pseudoflavonifractor capillosus_nov_89.897%
Eubacteriales_[G−1] bacterium_MOT−159_nov_94.268%
Eubacteriales_[G−4] bacterium_MOT−165_nov_92.781%
Acetivibrio cellulolyticus_nov_83.153%
Eubacteriales_[G−3] bacterium_MOT−163_nov_89.157%
Gluceribacter canis_nov_93.305%
Hathewaya proteolytica_nov_83.297%
Acutalibacter muris_nov_96.289%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.582%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.097%
Acutalibacter muris
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.198%
Alistipes senegalensis_nov_93.089%
Acetivibrio cellulolyticus_nov_83.761%
Anaerotignum faecicola_nov_85.287%
Acutalibacter muris_nov_96.694%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_89.858%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.010%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.572%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.640%
Enterococcus faecalis
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Lactobacillus johnsonii
Culturomica massiliensis_nov_89.817%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.792%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.577%
Kineothrix alysoides_nov_85.921%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.319%
Faecalicatena multispecies_sppn8_2_nov_92.067%
Duncaniella freteri_nov_93.712%
Duncaniella freteri_nov_89.775%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Duncaniella freteri_nov_89.919%
Prevotella shahii_nov_87.602%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.157%
Lawsonibacter asaccharolyticus_nov_91.116%
Duncaniella freteri_nov_90.612%
Species
F15387.S49
F15387.S50
F15387.S46
F15387.S47
F15387.S48
F15387.S44
F15387.S42
F15387.S43
F15387.S41
F15387.S45
Samples