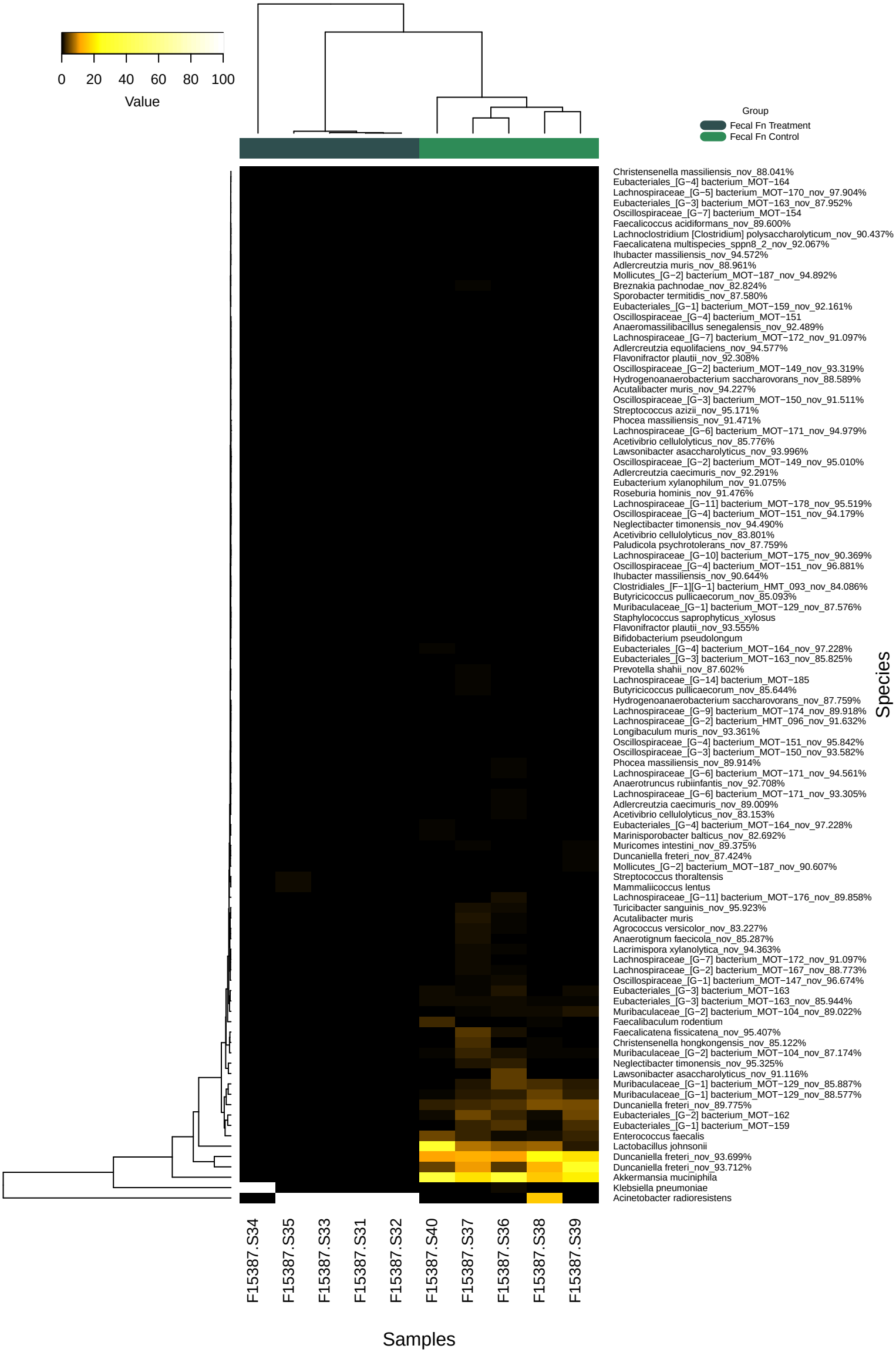
0
20
40
60
80
100
Value
Group
Fecal Fn Treatment
Fecal Fn Control
Christensenella massiliensis_nov_88.041%
Eubacteriales_[G−4] bacterium_MOT−164
Lachnospiraceae_[G−5] bacterium_MOT−170_nov_97.904%
Eubacteriales_[G−3] bacterium_MOT−163_nov_87.952%
Oscillospiraceae_[G−7] bacterium_MOT−154
Faecalicoccus acidiformans_nov_89.600%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.437%
Faecalicatena multispecies_sppn8_2_nov_92.067%
Ihubacter massiliensis_nov_94.572%
Adlercreutzia muris_nov_88.961%
Mollicutes_[G−2] bacterium_MOT−187_nov_94.892%
Breznakia pachnodae_nov_82.824%
Sporobacter termitidis_nov_87.580%
Eubacteriales_[G−1] bacterium_MOT−159_nov_92.161%
Oscillospiraceae_[G−4] bacterium_MOT−151
Anaeromassilibacillus senegalensis_nov_92.489%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.097%
Adlercreutzia equolifaciens_nov_94.577%
Flavonifractor plautii_nov_92.308%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.319%
Hydrogenoanaerobacterium saccharovorans_nov_88.589%
Acutalibacter muris_nov_94.227%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_91.511%
Streptococcus azizii_nov_95.171%
Phocea massiliensis_nov_91.471%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.979%
Acetivibrio cellulolyticus_nov_85.776%
Lawsonibacter asaccharolyticus_nov_93.996%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.010%
Adlercreutzia caecimuris_nov_92.291%
Eubacterium xylanophilum_nov_91.075%
Roseburia hominis_nov_91.476%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_95.519%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_94.179%
Neglectibacter timonensis_nov_94.490%
Acetivibrio cellulolyticus_nov_83.801%
Paludicola psychrotolerans_nov_87.759%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_90.369%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_96.881%
Ihubacter massiliensis_nov_90.644%
Clostridiales_[F−1][G−1] bacterium_HMT_093_nov_84.086%
Butyricicoccus pullicaecorum_nov_85.093%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_87.576%
Staphylococcus saprophyticus_xylosus
Flavonifractor plautii_nov_93.555%
Bifidobacterium pseudolongum
Eubacteriales_[G−4] bacterium_MOT−164_nov_97.228%
Eubacteriales_[G−3] bacterium_MOT−163_nov_85.825%
Prevotella shahii_nov_87.602%
Lachnospiraceae_[G−14] bacterium_MOT−185
Butyricicoccus pullicaecorum_nov_85.644%
Hydrogenoanaerobacterium saccharovorans_nov_87.759%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_89.918%
Lachnospiraceae_[G−2] bacterium_HMT_096_nov_91.632%
Longibaculum muris_nov_93.361%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_95.842%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.582%
Phocea massiliensis_nov_89.914%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.561%
Anaerotruncus rubiinfantis_nov_92.708%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_93.305%
Adlercreutzia caecimuris_nov_89.009%
Acetivibrio cellulolyticus_nov_83.153%
Eubacteriales_[G−4] bacterium_MOT−164_nov_97.228%
Marinisporobacter balticus_nov_82.692%
Muricomes intestini_nov_89.375%
Duncaniella freteri_nov_87.424%
Mollicutes_[G−2] bacterium_MOT−187_nov_90.607%
Streptococcus thoraltensis
Mammaliicoccus lentus
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_89.858%
Turicibacter sanguinis_nov_95.923%
Acutalibacter muris
Agrococcus versicolor_nov_83.227%
Anaerotignum faecicola_nov_85.287%
Lacrimispora xylanolytica_nov_94.363%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.097%
Lachnospiraceae_[G−2] bacterium_MOT−167_nov_88.773%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_96.674%
Eubacteriales_[G−3] bacterium_MOT−163
Eubacteriales_[G−3] bacterium_MOT−163_nov_85.944%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.022%
Faecalibaculum rodentium
Faecalicatena fissicatena_nov_95.407%
Christensenella hongkongensis_nov_85.122%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_87.174%
Neglectibacter timonensis_nov_95.325%
Lawsonibacter asaccharolyticus_nov_91.116%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.577%
Duncaniella freteri_nov_89.775%
Eubacteriales_[G−2] bacterium_MOT−162
Eubacteriales_[G−1] bacterium_MOT−159
Enterococcus faecalis
Lactobacillus johnsonii
Duncaniella freteri_nov_93.699%
Duncaniella freteri_nov_93.712%
Akkermansia muciniphila
Klebsiella pneumoniae
Acinetobacter radioresistens
Species
F15387.S34
F15387.S35
F15387.S33
F15387.S31
F15387.S32
F15387.S40
F15387.S37
F15387.S36
F15387.S38
F15387.S39
Samples