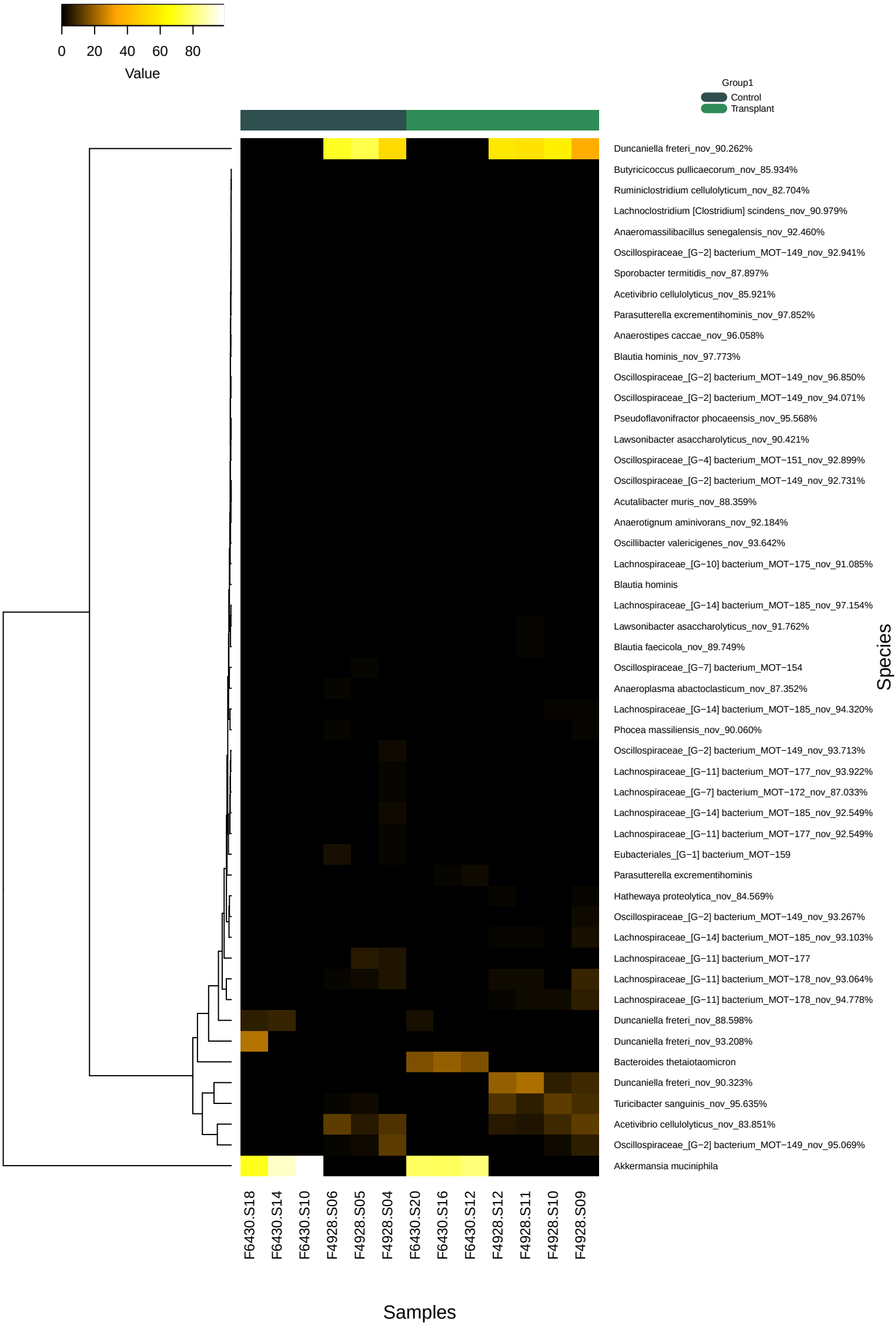
0
20
40
60
80
Value
Group1
Control
Transplant
Duncaniella freteri_nov_90.262%
Butyricicoccus pullicaecorum_nov_85.934%
Ruminiclostridium cellulolyticum_nov_82.704%
Lachnoclostridium [Clostridium] scindens_nov_90.979%
Anaeromassilibacillus senegalensis_nov_92.460%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_92.941%
Sporobacter termitidis_nov_87.897%
Acetivibrio cellulolyticus_nov_85.921%
Parasutterella excrementihominis_nov_97.852%
Anaerostipes caccae_nov_96.058%
Blautia hominis_nov_97.773%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.850%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_94.071%
Pseudoflavonifractor phocaeensis_nov_95.568%
Lawsonibacter asaccharolyticus_nov_90.421%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_92.899%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_92.731%
Acutalibacter muris_nov_88.359%
Anaerotignum aminivorans_nov_92.184%
Oscillibacter valericigenes_nov_93.642%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_91.085%
Blautia hominis
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_97.154%
Lawsonibacter asaccharolyticus_nov_91.762%
Blautia faecicola_nov_89.749%
Oscillospiraceae_[G−7] bacterium_MOT−154
Anaeroplasma abactoclasticum_nov_87.352%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_94.320%
Phocea massiliensis_nov_90.060%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.713%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_93.922%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_87.033%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.549%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_92.549%
Eubacteriales_[G−1] bacterium_MOT−159
Parasutterella excrementihominis
Hathewaya proteolytica_nov_84.569%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.267%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_93.103%
Lachnospiraceae_[G−11] bacterium_MOT−177
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_93.064%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_94.778%
Duncaniella freteri_nov_88.598%
Duncaniella freteri_nov_93.208%
Bacteroides thetaiotaomicron
Duncaniella freteri_nov_90.323%
Turicibacter sanguinis_nov_95.635%
Acetivibrio cellulolyticus_nov_83.851%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.069%
Akkermansia muciniphila
Species
F6430.S18
F6430.S14
F6430.S10
F4928.S06
F4928.S05
F4928.S04
F6430.S20
F6430.S16
F6430.S12
F4928.S12
F4928.S11
F4928.S10
F4928.S09
Samples