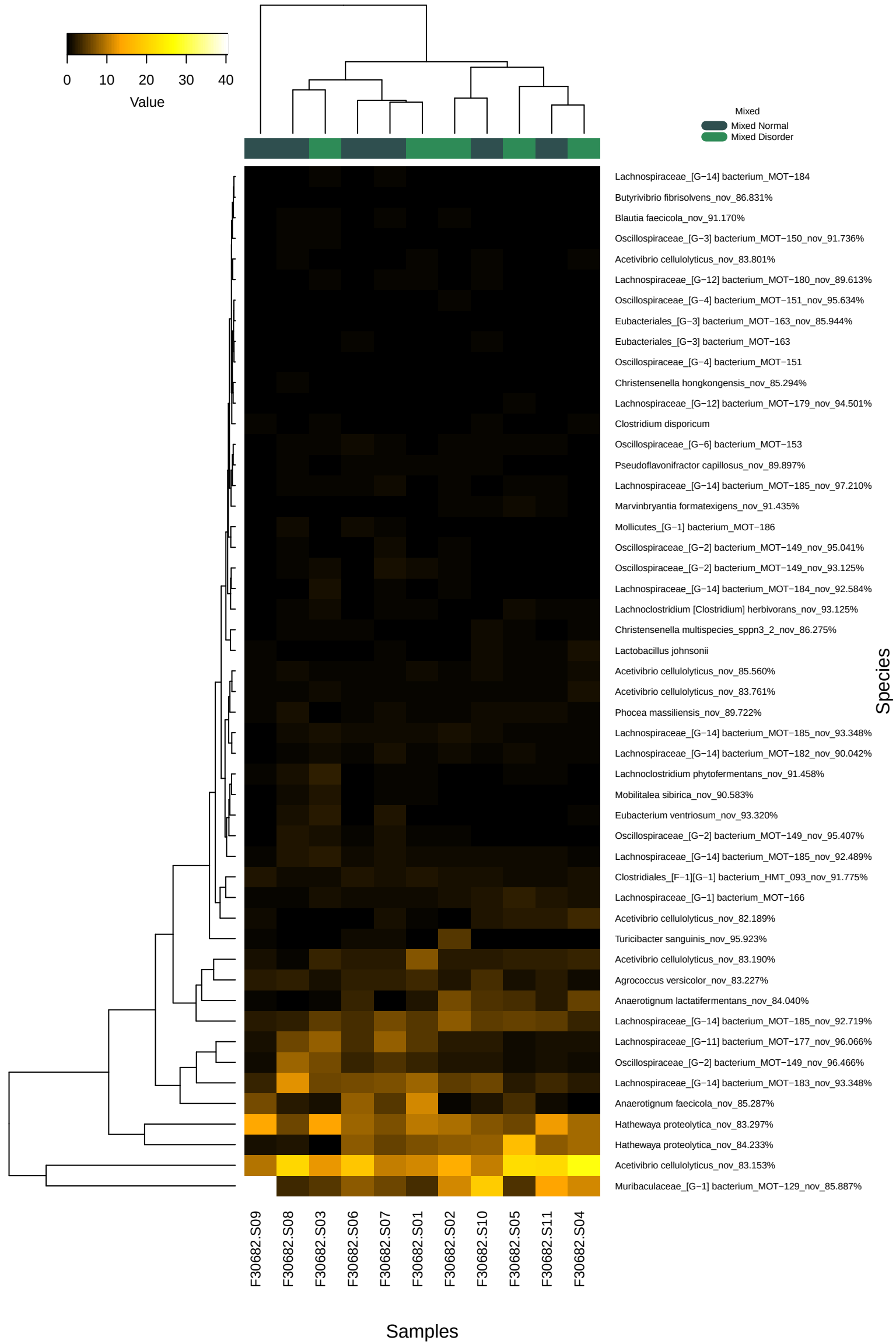
0
10
20
30
40
Value
Mixed
Mixed Normal
Mixed Disorder
Lachnospiraceae_[G−14] bacterium_MOT−184
Butyrivibrio fibrisolvens_nov_86.831%
Blautia faecicola_nov_91.170%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_91.736%
Acetivibrio cellulolyticus_nov_83.801%
Lachnospiraceae_[G−12] bacterium_MOT−180_nov_89.613%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_95.634%
Eubacteriales_[G−3] bacterium_MOT−163_nov_85.944%
Eubacteriales_[G−3] bacterium_MOT−163
Oscillospiraceae_[G−4] bacterium_MOT−151
Christensenella hongkongensis_nov_85.294%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_94.501%
Clostridium disporicum
Oscillospiraceae_[G−6] bacterium_MOT−153
Pseudoflavonifractor capillosus_nov_89.897%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_97.210%
Marvinbryantia formatexigens_nov_91.435%
Mollicutes_[G−1] bacterium_MOT−186
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.041%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.125%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_92.584%
Lachnoclostridium [Clostridium] herbivorans_nov_93.125%
Christensenella multispecies_sppn3_2_nov_86.275%
Lactobacillus johnsonii
Acetivibrio cellulolyticus_nov_85.560%
Acetivibrio cellulolyticus_nov_83.761%
Phocea massiliensis_nov_89.722%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_93.348%
Lachnospiraceae_[G−14] bacterium_MOT−182_nov_90.042%
Lachnoclostridium phytofermentans_nov_91.458%
Mobilitalea sibirica_nov_90.583%
Eubacterium ventriosum_nov_93.320%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.407%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.489%
Clostridiales_[F−1][G−1] bacterium_HMT_093_nov_91.775%
Lachnospiraceae_[G−1] bacterium_MOT−166
Acetivibrio cellulolyticus_nov_82.189%
Turicibacter sanguinis_nov_95.923%
Acetivibrio cellulolyticus_nov_83.190%
Agrococcus versicolor_nov_83.227%
Anaerotignum lactatifermentans_nov_84.040%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.719%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_96.066%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.466%
Lachnospiraceae_[G−14] bacterium_MOT−183_nov_93.348%
Anaerotignum faecicola_nov_85.287%
Hathewaya proteolytica_nov_83.297%
Hathewaya proteolytica_nov_84.233%
Acetivibrio cellulolyticus_nov_83.153%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Species
F30682.S09
F30682.S08
F30682.S03
F30682.S06
F30682.S07
F30682.S01
F30682.S02
F30682.S10
F30682.S05
F30682.S11
F30682.S04
Samples