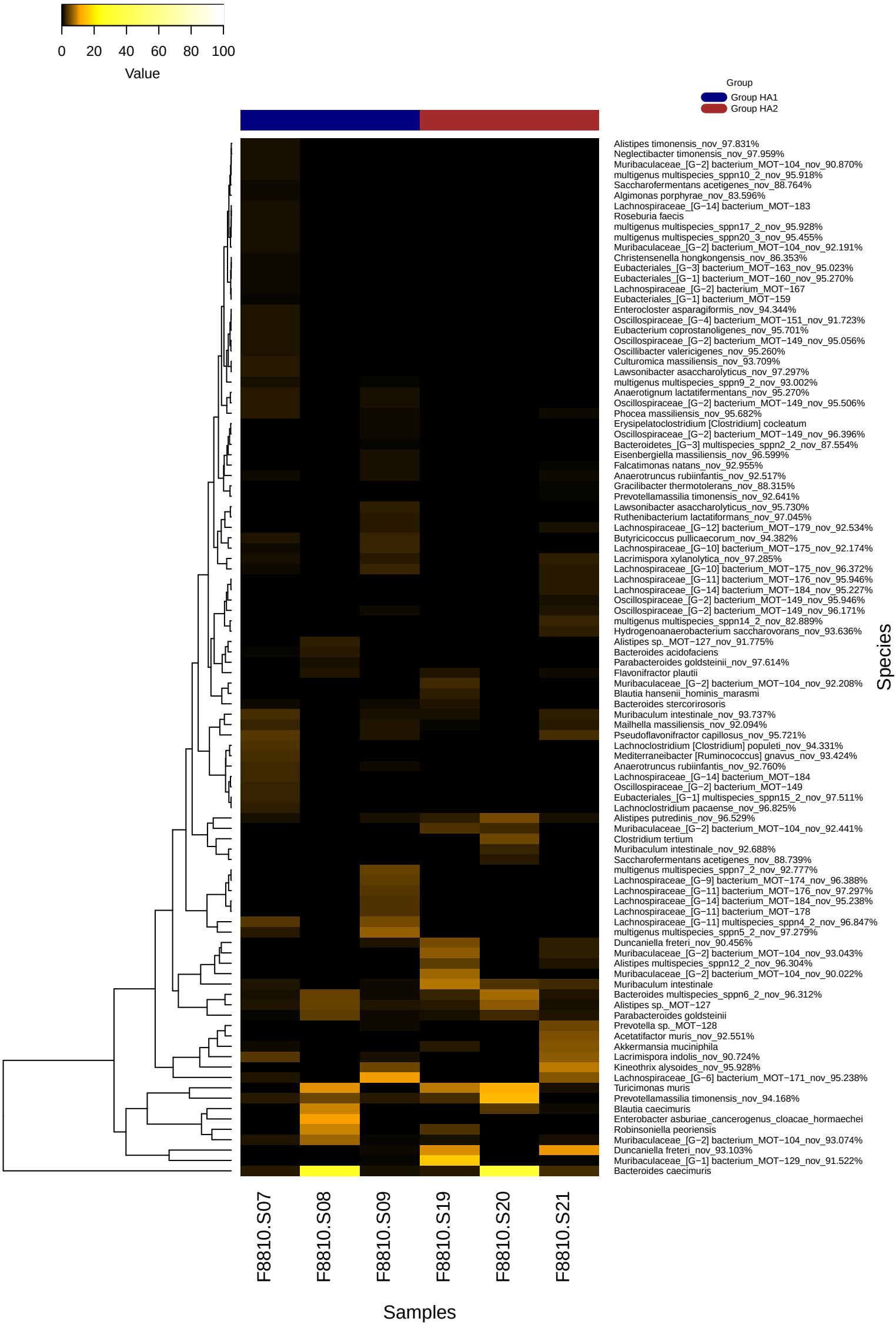
0
20
40
60
80
100
Value
Group
Group HA1
Group HA2
Alistipes timonensis_nov_97.831%
Neglectibacter timonensis_nov_97.959%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.870%
multigenus multispecies_sppn10_2_nov_95.918%
Saccharofermentans acetigenes_nov_88.764%
Algimonas porphyrae_nov_83.596%
Lachnospiraceae_[G−14] bacterium_MOT−183
Roseburia faecis
multigenus multispecies_sppn17_2_nov_95.928%
multigenus multispecies_sppn20_3_nov_95.455%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.191%
Christensenella hongkongensis_nov_86.353%
Eubacteriales_[G−3] bacterium_MOT−163_nov_95.023%
Eubacteriales_[G−1] bacterium_MOT−160_nov_95.270%
Lachnospiraceae_[G−2] bacterium_MOT−167
Eubacteriales_[G−1] bacterium_MOT−159
Enterocloster asparagiformis_nov_94.344%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_91.723%
Eubacterium coprostanoligenes_nov_95.701%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.056%
Oscillibacter valericigenes_nov_95.260%
Culturomica massiliensis_nov_93.709%
Lawsonibacter asaccharolyticus_nov_97.297%
multigenus multispecies_sppn9_2_nov_93.002%
Anaerotignum lactatifermentans_nov_95.270%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.506%
Phocea massiliensis_nov_95.682%
Erysipelatoclostridium [Clostridium] cocleatum
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.396%
Bacteroidetes_[G−3] multispecies_sppn2_2_nov_87.554%
Eisenbergiella massiliensis_nov_96.599%
Falcatimonas natans_nov_92.955%
Anaerotruncus rubiinfantis_nov_92.517%
Gracilibacter thermotolerans_nov_88.315%
Prevotellamassilia timonensis_nov_92.641%
Lawsonibacter asaccharolyticus_nov_95.730%
Ruthenibacterium lactatiformans_nov_97.045%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_92.534%
Butyricicoccus pullicaecorum_nov_94.382%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_92.174%
Lacrimispora xylanolytica_nov_97.285%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_96.372%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_95.946%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_95.227%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.946%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.171%
multigenus multispecies_sppn14_2_nov_82.889%
Hydrogenoanaerobacterium saccharovorans_nov_93.636%
Alistipes sp._MOT−127_nov_91.775%
Bacteroides acidofaciens
Parabacteroides goldsteinii_nov_97.614%
Flavonifractor plautii
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.208%
Blautia hansenii_hominis_marasmi
Bacteroides stercorirosoris
Muribaculum intestinale_nov_93.737%
Mailhella massiliensis_nov_92.094%
Pseudoflavonifractor capillosus_nov_95.721%
Lachnoclostridium [Clostridium] populeti_nov_94.331%
Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
Anaerotruncus rubiinfantis_nov_92.760%
Lachnospiraceae_[G−14] bacterium_MOT−184
Oscillospiraceae_[G−2] bacterium_MOT−149
Eubacteriales_[G−1] multispecies_sppn15_2_nov_97.511%
Lachnoclostridium pacaense_nov_96.825%
Alistipes putredinis_nov_96.529%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.441%
Clostridium tertium
Muribaculum intestinale_nov_92.688%
Saccharofermentans acetigenes_nov_88.739%
multigenus multispecies_sppn7_2_nov_92.777%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.388%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_97.297%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_95.238%
Lachnospiraceae_[G−11] bacterium_MOT−178
Lachnospiraceae_[G−11] multispecies_sppn4_2_nov_96.847%
multigenus multispecies_sppn5_2_nov_97.279%
Duncaniella freteri_nov_90.456%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.043%
Alistipes multispecies_sppn12_2_nov_96.304%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.022%
Muribaculum intestinale
Bacteroides multispecies_sppn6_2_nov_96.312%
Alistipes sp._MOT−127
Parabacteroides goldsteinii
Prevotella sp._MOT−128
Acetatifactor muris_nov_92.551%
Akkermansia muciniphila
Lacrimispora indolis_nov_90.724%
Kineothrix alysoides_nov_95.928%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_95.238%
Turicimonas muris
Prevotellamassilia timonensis_nov_94.168%
Blautia caecimuris
Enterobacter asburiae_cancerogenus_cloacae_hormaechei
Robinsoniella peoriensis
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.074%
Duncaniella freteri_nov_93.103%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.522%
Bacteroides caecimuris
Species
F8810.S07
F8810.S08
F8810.S09
F8810.S19
F8810.S20
F8810.S21
Samples