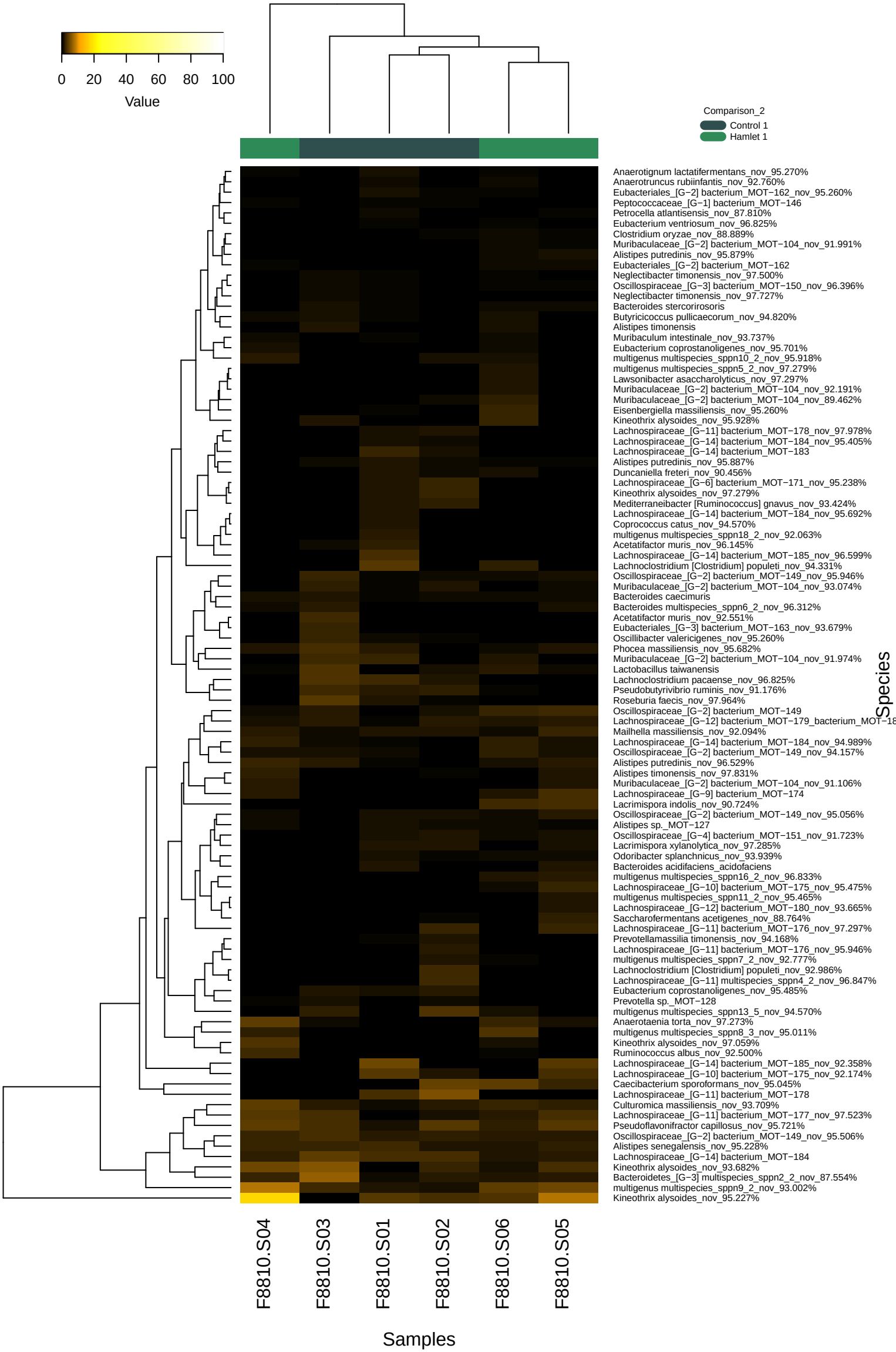
0
20
40
60
80
100
Value
Comparison_2
Control 1
Hamlet 1
Anaerotignum lactatifermentans_nov_95.270%
Anaerotruncus rubiinfantis_nov_92.760%
Eubacteriales_[G−2] bacterium_MOT−162_nov_95.260%
Peptococcaceae_[G−1] bacterium_MOT−146
Petrocella atlantisensis_nov_87.810%
Eubacterium ventriosum_nov_96.825%
Clostridium oryzae_nov_88.889%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.991%
Alistipes putredinis_nov_95.879%
Eubacteriales_[G−2] bacterium_MOT−162
Neglectibacter timonensis_nov_97.500%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_96.396%
Neglectibacter timonensis_nov_97.727%
Bacteroides stercorirosoris
Butyricicoccus pullicaecorum_nov_94.820%
Alistipes timonensis
Muribaculum intestinale_nov_93.737%
Eubacterium coprostanoligenes_nov_95.701%
multigenus multispecies_sppn10_2_nov_95.918%
multigenus multispecies_sppn5_2_nov_97.279%
Lawsonibacter asaccharolyticus_nov_97.297%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.191%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.462%
Eisenbergiella massiliensis_nov_95.260%
Kineothrix alysoides_nov_95.928%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_97.978%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_95.405%
Lachnospiraceae_[G−14] bacterium_MOT−183
Alistipes putredinis_nov_95.887%
Duncaniella freteri_nov_90.456%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_95.238%
Kineothrix alysoides_nov_97.279%
Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_95.692%
Coprococcus catus_nov_94.570%
multigenus multispecies_sppn18_2_nov_92.063%
Acetatifactor muris_nov_96.145%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_96.599%
Lachnoclostridium [Clostridium] populeti_nov_94.331%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.946%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.074%
Bacteroides caecimuris
Bacteroides multispecies_sppn6_2_nov_96.312%
Acetatifactor muris_nov_92.551%
Eubacteriales_[G−3] bacterium_MOT−163_nov_93.679%
Oscillibacter valericigenes_nov_95.260%
Phocea massiliensis_nov_95.682%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.974%
Lactobacillus taiwanensis
Lachnoclostridium pacaense_nov_96.825%
Pseudobutyrivibrio ruminis_nov_91.176%
Roseburia faecis_nov_97.964%
Oscillospiraceae_[G−2] bacterium_MOT−149
Lachnospiraceae_[G−12] bacterium_MOT−179_bacterium_MOT−18
Mailhella massiliensis_nov_92.094%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_94.989%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_94.157%
Alistipes putredinis_nov_96.529%
Alistipes timonensis_nov_97.831%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.106%
Lachnospiraceae_[G−9] bacterium_MOT−174
Lacrimispora indolis_nov_90.724%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.056%
Alistipes sp._MOT−127
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_91.723%
Lacrimispora xylanolytica_nov_97.285%
Odoribacter splanchnicus_nov_93.939%
Bacteroides acidifaciens_acidofaciens
multigenus multispecies_sppn16_2_nov_96.833%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_95.475%
multigenus multispecies_sppn11_2_nov_95.465%
Lachnospiraceae_[G−12] bacterium_MOT−180_nov_93.665%
Saccharofermentans acetigenes_nov_88.764%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_97.297%
Prevotellamassilia timonensis_nov_94.168%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_95.946%
multigenus multispecies_sppn7_2_nov_92.777%
Lachnoclostridium [Clostridium] populeti_nov_92.986%
Lachnospiraceae_[G−11] multispecies_sppn4_2_nov_96.847%
Eubacterium coprostanoligenes_nov_95.485%
Prevotella sp._MOT−128
multigenus multispecies_sppn13_5_nov_94.570%
Anaerotaenia torta_nov_97.273%
multigenus multispecies_sppn8_3_nov_95.011%
Kineothrix alysoides_nov_97.059%
Ruminococcus albus_nov_92.500%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.358%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_92.174%
Caecibacterium sporoformans_nov_95.045%
Lachnospiraceae_[G−11] bacterium_MOT−178
Culturomica massiliensis_nov_93.709%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_97.523%
Pseudoflavonifractor capillosus_nov_95.721%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.506%
Alistipes senegalensis_nov_95.228%
Lachnospiraceae_[G−14] bacterium_MOT−184
Kineothrix alysoides_nov_93.682%
Bacteroidetes_[G−3] multispecies_sppn2_2_nov_87.554%
multigenus multispecies_sppn9_2_nov_93.002%
Kineothrix alysoides_nov_95.227%
Species
F8810.S04
F8810.S03
F8810.S01
F8810.S02
F8810.S06
F8810.S05
Samples