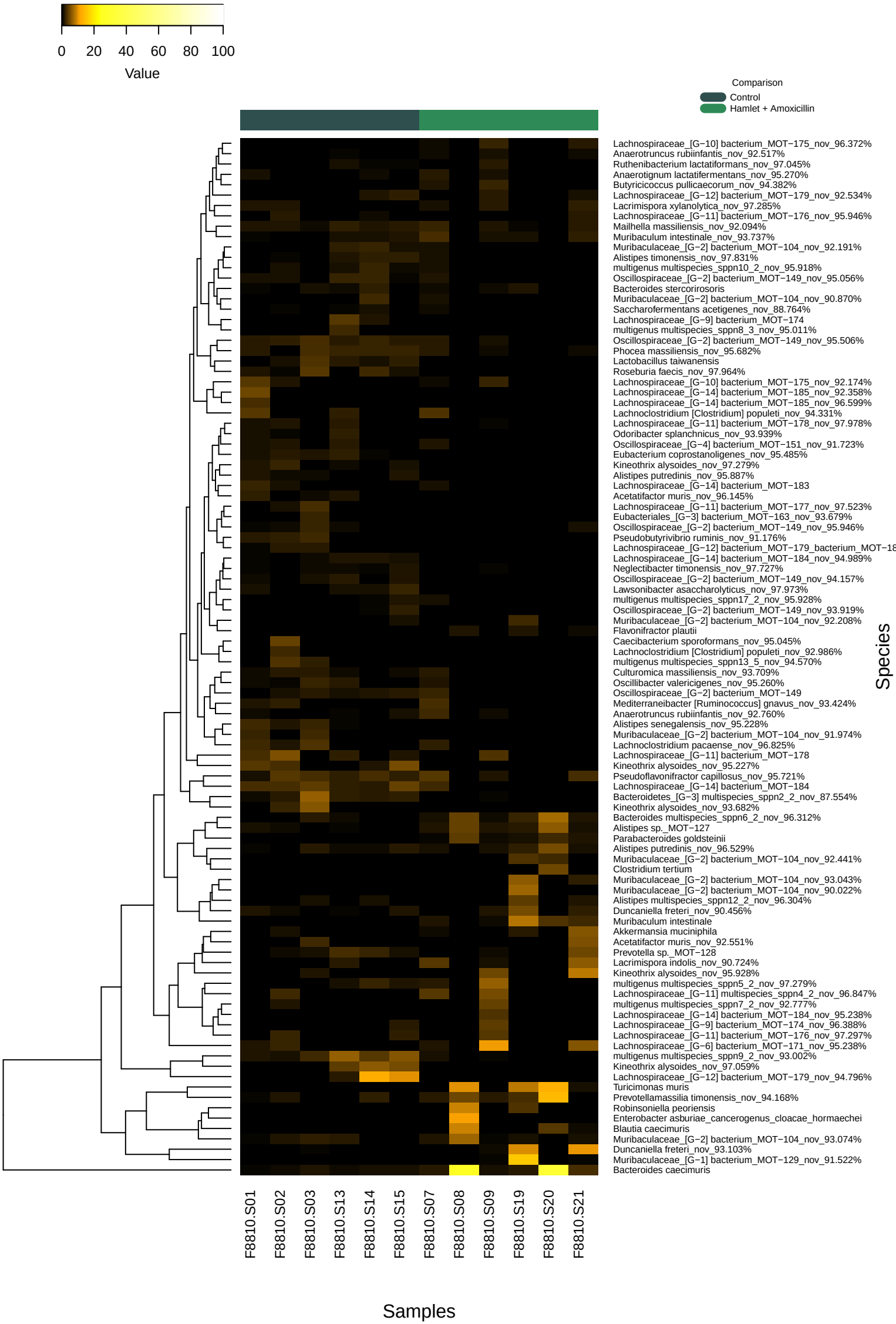
0
20
40
60
80
100
Value
Comparison
Control
Hamlet + Amoxicillin
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_96.372%
Anaerotruncus rubiinfantis_nov_92.517%
Ruthenibacterium lactatiformans_nov_97.045%
Anaerotignum lactatifermentans_nov_95.270%
Butyricicoccus pullicaecorum_nov_94.382%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_92.534%
Lacrimispora xylanolytica_nov_97.285%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_95.946%
Mailhella massiliensis_nov_92.094%
Muribaculum intestinale_nov_93.737%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.191%
Alistipes timonensis_nov_97.831%
multigenus multispecies_sppn10_2_nov_95.918%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.056%
Bacteroides stercorirosoris
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.870%
Saccharofermentans acetigenes_nov_88.764%
Lachnospiraceae_[G−9] bacterium_MOT−174
multigenus multispecies_sppn8_3_nov_95.011%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.506%
Phocea massiliensis_nov_95.682%
Lactobacillus taiwanensis
Roseburia faecis_nov_97.964%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_92.174%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.358%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_96.599%
Lachnoclostridium [Clostridium] populeti_nov_94.331%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_97.978%
Odoribacter splanchnicus_nov_93.939%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_91.723%
Eubacterium coprostanoligenes_nov_95.485%
Kineothrix alysoides_nov_97.279%
Alistipes putredinis_nov_95.887%
Lachnospiraceae_[G−14] bacterium_MOT−183
Acetatifactor muris_nov_96.145%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_97.523%
Eubacteriales_[G−3] bacterium_MOT−163_nov_93.679%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.946%
Pseudobutyrivibrio ruminis_nov_91.176%
Lachnospiraceae_[G−12] bacterium_MOT−179_bacterium_MOT−18
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_94.989%
Neglectibacter timonensis_nov_97.727%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_94.157%
Lawsonibacter asaccharolyticus_nov_97.973%
multigenus multispecies_sppn17_2_nov_95.928%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.919%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.208%
Flavonifractor plautii
Caecibacterium sporoformans_nov_95.045%
Lachnoclostridium [Clostridium] populeti_nov_92.986%
multigenus multispecies_sppn13_5_nov_94.570%
Culturomica massiliensis_nov_93.709%
Oscillibacter valericigenes_nov_95.260%
Oscillospiraceae_[G−2] bacterium_MOT−149
Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
Anaerotruncus rubiinfantis_nov_92.760%
Alistipes senegalensis_nov_95.228%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.974%
Lachnoclostridium pacaense_nov_96.825%
Lachnospiraceae_[G−11] bacterium_MOT−178
Kineothrix alysoides_nov_95.227%
Pseudoflavonifractor capillosus_nov_95.721%
Lachnospiraceae_[G−14] bacterium_MOT−184
Bacteroidetes_[G−3] multispecies_sppn2_2_nov_87.554%
Kineothrix alysoides_nov_93.682%
Bacteroides multispecies_sppn6_2_nov_96.312%
Alistipes sp._MOT−127
Parabacteroides goldsteinii
Alistipes putredinis_nov_96.529%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.441%
Clostridium tertium
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.043%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.022%
Alistipes multispecies_sppn12_2_nov_96.304%
Duncaniella freteri_nov_90.456%
Muribaculum intestinale
Akkermansia muciniphila
Acetatifactor muris_nov_92.551%
Prevotella sp._MOT−128
Lacrimispora indolis_nov_90.724%
Kineothrix alysoides_nov_95.928%
multigenus multispecies_sppn5_2_nov_97.279%
Lachnospiraceae_[G−11] multispecies_sppn4_2_nov_96.847%
multigenus multispecies_sppn7_2_nov_92.777%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_95.238%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.388%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_97.297%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_95.238%
multigenus multispecies_sppn9_2_nov_93.002%
Kineothrix alysoides_nov_97.059%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_94.796%
Turicimonas muris
Prevotellamassilia timonensis_nov_94.168%
Robinsoniella peoriensis
Enterobacter asburiae_cancerogenus_cloacae_hormaechei
Blautia caecimuris
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.S01
F8810.S02
F8810.S03
F8810.S13
F8810.S14
F8810.S15
F8810.S07
F8810.S08
F8810.S09
F8810.S19
F8810.S20
F8810.S21
Samples