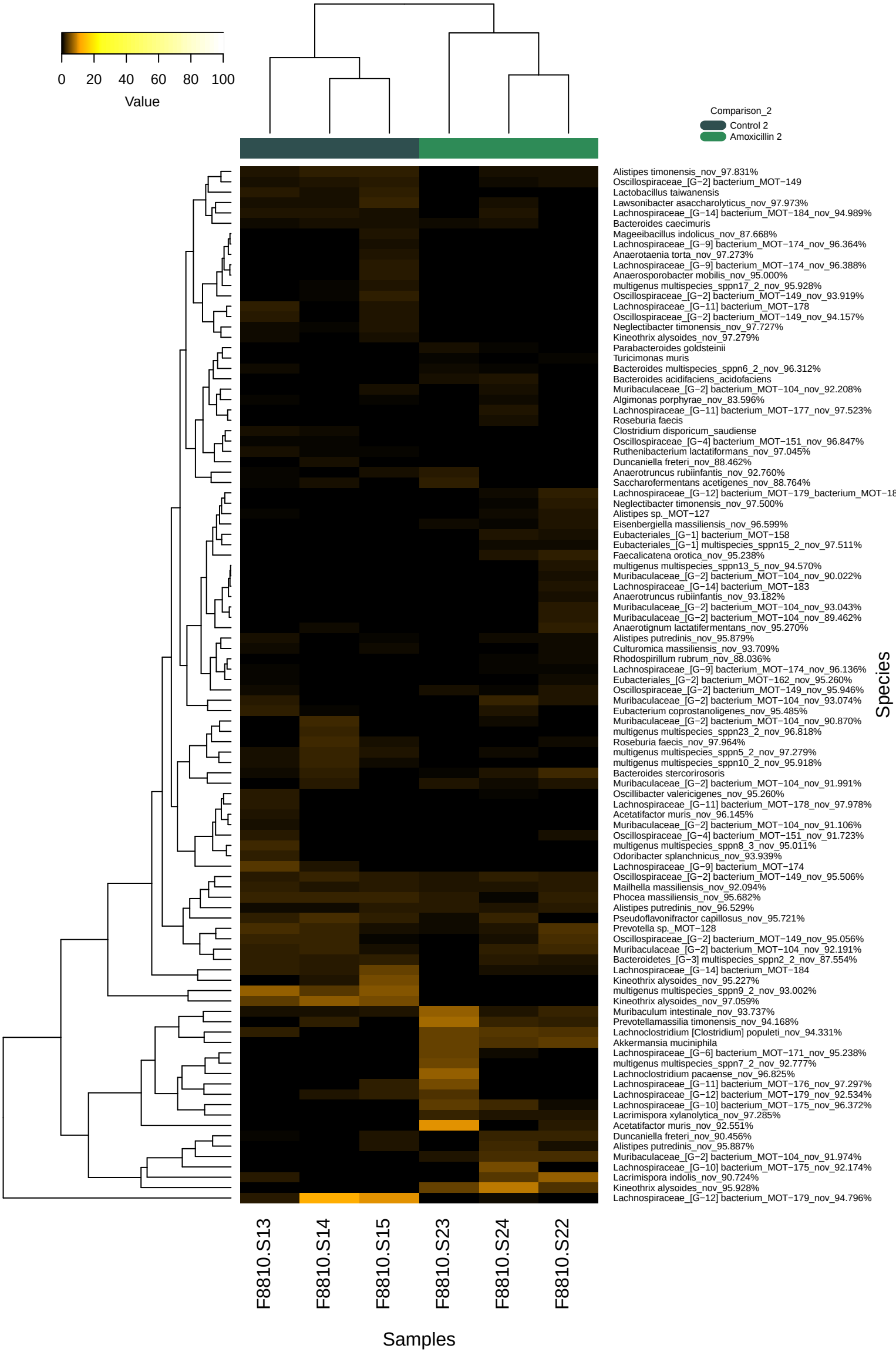
0
20
40
60
80
100
Value
Comparison_2
Control 2
Amoxicillin 2
Alistipes timonensis_nov_97.831%
Oscillospiraceae_[G−2] bacterium_MOT−149
Lactobacillus taiwanensis
Lawsonibacter asaccharolyticus_nov_97.973%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_94.989%
Bacteroides caecimuris
Mageeibacillus indolicus_nov_87.668%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.364%
Anaerotaenia torta_nov_97.273%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.388%
Anaerosporobacter mobilis_nov_95.000%
multigenus multispecies_sppn17_2_nov_95.928%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.919%
Lachnospiraceae_[G−11] bacterium_MOT−178
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_94.157%
Neglectibacter timonensis_nov_97.727%
Kineothrix alysoides_nov_97.279%
Parabacteroides goldsteinii
Turicimonas muris
Bacteroides multispecies_sppn6_2_nov_96.312%
Bacteroides acidifaciens_acidofaciens
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.208%
Algimonas porphyrae_nov_83.596%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_97.523%
Roseburia faecis
Clostridium disporicum_saudiense
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_96.847%
Ruthenibacterium lactatiformans_nov_97.045%
Duncaniella freteri_nov_88.462%
Anaerotruncus rubiinfantis_nov_92.760%
Saccharofermentans acetigenes_nov_88.764%
Lachnospiraceae_[G−12] bacterium_MOT−179_bacterium_MOT−18
Neglectibacter timonensis_nov_97.500%
Alistipes sp._MOT−127
Eisenbergiella massiliensis_nov_96.599%
Eubacteriales_[G−1] bacterium_MOT−158
Eubacteriales_[G−1] multispecies_sppn15_2_nov_97.511%
Faecalicatena orotica_nov_95.238%
multigenus multispecies_sppn13_5_nov_94.570%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.022%
Lachnospiraceae_[G−14] bacterium_MOT−183
Anaerotruncus rubiinfantis_nov_93.182%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.043%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.462%
Anaerotignum lactatifermentans_nov_95.270%
Alistipes putredinis_nov_95.879%
Culturomica massiliensis_nov_93.709%
Rhodospirillum rubrum_nov_88.036%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.136%
Eubacteriales_[G−2] bacterium_MOT−162_nov_95.260%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.946%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.074%
Eubacterium coprostanoligenes_nov_95.485%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.870%
multigenus multispecies_sppn23_2_nov_96.818%
Roseburia faecis_nov_97.964%
multigenus multispecies_sppn5_2_nov_97.279%
multigenus multispecies_sppn10_2_nov_95.918%
Bacteroides stercorirosoris
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.991%
Oscillibacter valericigenes_nov_95.260%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_97.978%
Acetatifactor muris_nov_96.145%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.106%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_91.723%
multigenus multispecies_sppn8_3_nov_95.011%
Odoribacter splanchnicus_nov_93.939%
Lachnospiraceae_[G−9] bacterium_MOT−174
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.506%
Mailhella massiliensis_nov_92.094%
Phocea massiliensis_nov_95.682%
Alistipes putredinis_nov_96.529%
Pseudoflavonifractor capillosus_nov_95.721%
Prevotella sp._MOT−128
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.056%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.191%
Bacteroidetes_[G−3] multispecies_sppn2_2_nov_87.554%
Lachnospiraceae_[G−14] bacterium_MOT−184
Kineothrix alysoides_nov_95.227%
multigenus multispecies_sppn9_2_nov_93.002%
Kineothrix alysoides_nov_97.059%
Muribaculum intestinale_nov_93.737%
Prevotellamassilia timonensis_nov_94.168%
Lachnoclostridium [Clostridium] populeti_nov_94.331%
Akkermansia muciniphila
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_95.238%
multigenus multispecies_sppn7_2_nov_92.777%
Lachnoclostridium pacaense_nov_96.825%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_97.297%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_92.534%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_96.372%
Lacrimispora xylanolytica_nov_97.285%
Acetatifactor muris_nov_92.551%
Duncaniella freteri_nov_90.456%
Alistipes putredinis_nov_95.887%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.974%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_92.174%
Lacrimispora indolis_nov_90.724%
Kineothrix alysoides_nov_95.928%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_94.796%
Species
F8810.S13
F8810.S14
F8810.S15
F8810.S23
F8810.S24
F8810.S22
Samples