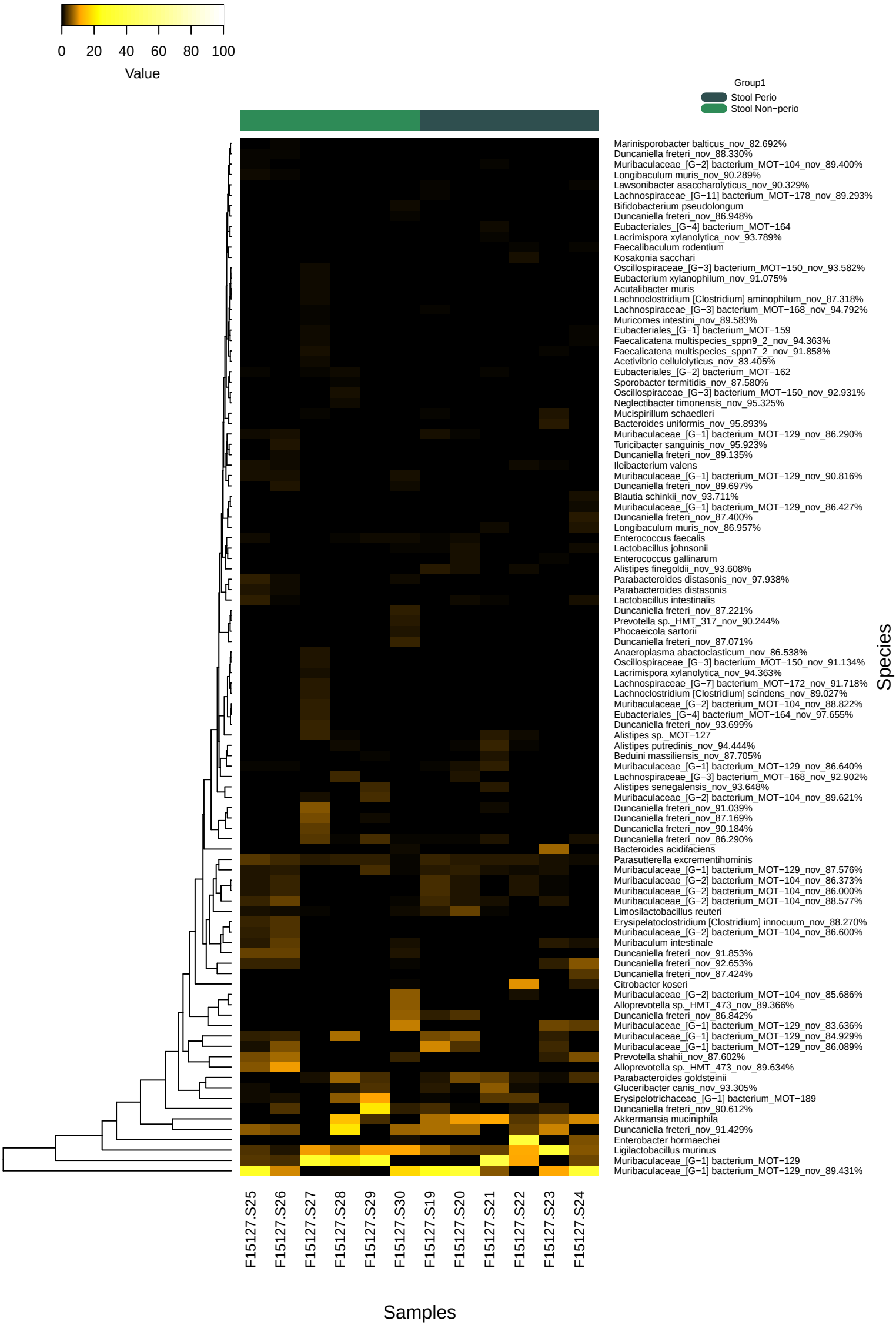
0
20
40
60
80
100
Value
Group1
Stool Perio
Stool Non−perio
Marinisporobacter balticus_nov_82.692%
Duncaniella freteri_nov_88.330%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.400%
Longibaculum muris_nov_90.289%
Lawsonibacter asaccharolyticus_nov_90.329%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_89.293%
Bifidobacterium pseudolongum
Duncaniella freteri_nov_86.948%
Eubacteriales_[G−4] bacterium_MOT−164
Lacrimispora xylanolytica_nov_93.789%
Faecalibaculum rodentium
Kosakonia sacchari
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.582%
Eubacterium xylanophilum_nov_91.075%
Acutalibacter muris
Lachnoclostridium [Clostridium] aminophilum_nov_87.318%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.792%
Muricomes intestini_nov_89.583%
Eubacteriales_[G−1] bacterium_MOT−159
Faecalicatena multispecies_sppn9_2_nov_94.363%
Faecalicatena multispecies_sppn7_2_nov_91.858%
Acetivibrio cellulolyticus_nov_83.405%
Eubacteriales_[G−2] bacterium_MOT−162
Sporobacter termitidis_nov_87.580%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_92.931%
Neglectibacter timonensis_nov_95.325%
Mucispirillum schaedleri
Bacteroides uniformis_nov_95.893%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.290%
Turicibacter sanguinis_nov_95.923%
Duncaniella freteri_nov_89.135%
Ileibacterium valens
Muribaculaceae_[G−1] bacterium_MOT−129_nov_90.816%
Duncaniella freteri_nov_89.697%
Blautia schinkii_nov_93.711%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.427%
Duncaniella freteri_nov_87.400%
Longibaculum muris_nov_86.957%
Enterococcus faecalis
Lactobacillus johnsonii
Enterococcus gallinarum
Alistipes finegoldii_nov_93.608%
Parabacteroides distasonis_nov_97.938%
Parabacteroides distasonis
Lactobacillus intestinalis
Duncaniella freteri_nov_87.221%
Prevotella sp._HMT_317_nov_90.244%
Phocaeicola sartorii
Duncaniella freteri_nov_87.071%
Anaeroplasma abactoclasticum_nov_86.538%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_91.134%
Lacrimispora xylanolytica_nov_94.363%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.718%
Lachnoclostridium [Clostridium] scindens_nov_89.027%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.822%
Eubacteriales_[G−4] bacterium_MOT−164_nov_97.655%
Duncaniella freteri_nov_93.699%
Alistipes sp._MOT−127
Alistipes putredinis_nov_94.444%
Beduini massiliensis_nov_87.705%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.640%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_92.902%
Alistipes senegalensis_nov_93.648%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.621%
Duncaniella freteri_nov_91.039%
Duncaniella freteri_nov_87.169%
Duncaniella freteri_nov_90.184%
Duncaniella freteri_nov_86.290%
Bacteroides acidifaciens
Parasutterella excrementihominis
Muribaculaceae_[G−1] bacterium_MOT−129_nov_87.576%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.373%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.000%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.577%
Limosilactobacillus reuteri
Erysipelatoclostridium [Clostridium] innocuum_nov_88.270%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.600%
Muribaculum intestinale
Duncaniella freteri_nov_91.853%
Duncaniella freteri_nov_92.653%
Duncaniella freteri_nov_87.424%
Citrobacter koseri
Muribaculaceae_[G−2] bacterium_MOT−104_nov_85.686%
Alloprevotella sp._HMT_473_nov_89.366%
Duncaniella freteri_nov_86.842%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_83.636%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_84.929%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.089%
Prevotella shahii_nov_87.602%
Alloprevotella sp._HMT_473_nov_89.634%
Parabacteroides goldsteinii
Gluceribacter canis_nov_93.305%
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Duncaniella freteri_nov_90.612%
Akkermansia muciniphila
Duncaniella freteri_nov_91.429%
Enterobacter hormaechei
Ligilactobacillus murinus
Muribaculaceae_[G−1] bacterium_MOT−129
Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.431%
Species
F15127.S25
F15127.S26
F15127.S27
F15127.S28
F15127.S29
F15127.S30
F15127.S19
F15127.S20
F15127.S21
F15127.S22
F15127.S23
F15127.S24
Samples