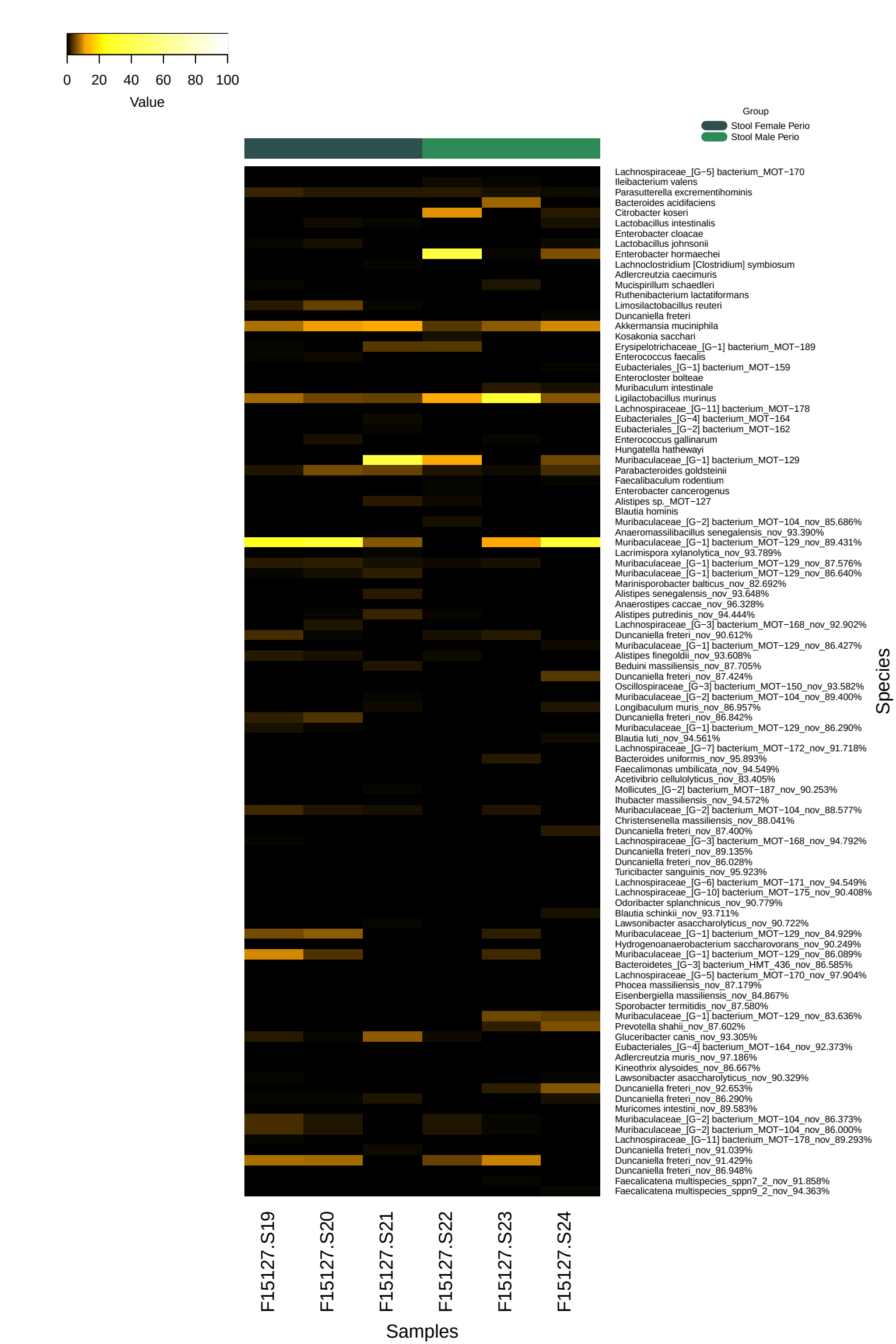
0
20
40
60
80
100
Value
Group
Stool Female Perio
Stool Male Perio
Lachnospiraceae_[G−5] bacterium_MOT−170
Ileibacterium valens
Parasutterella excrementihominis
Bacteroides acidifaciens
Citrobacter koseri
Lactobacillus intestinalis
Enterobacter cloacae
Lactobacillus johnsonii
Enterobacter hormaechei
Lachnoclostridium [Clostridium] symbiosum
Adlercreutzia caecimuris
Mucispirillum schaedleri
Ruthenibacterium lactatiformans
Limosilactobacillus reuteri
Duncaniella freteri
Akkermansia muciniphila
Kosakonia sacchari
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Enterococcus faecalis
Eubacteriales_[G−1] bacterium_MOT−159
Enterocloster bolteae
Muribaculum intestinale
Ligilactobacillus murinus
Lachnospiraceae_[G−11] bacterium_MOT−178
Eubacteriales_[G−4] bacterium_MOT−164
Eubacteriales_[G−2] bacterium_MOT−162
Enterococcus gallinarum
Hungatella hathewayi
Muribaculaceae_[G−1] bacterium_MOT−129
Parabacteroides goldsteinii
Faecalibaculum rodentium
Enterobacter cancerogenus
Alistipes sp._MOT−127
Blautia hominis
Muribaculaceae_[G−2] bacterium_MOT−104_nov_85.686%
Anaeromassilibacillus senegalensis_nov_93.390%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.431%
Lacrimispora xylanolytica_nov_93.789%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_87.576%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.640%
Marinisporobacter balticus_nov_82.692%
Alistipes senegalensis_nov_93.648%
Anaerostipes caccae_nov_96.328%
Alistipes putredinis_nov_94.444%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_92.902%
Duncaniella freteri_nov_90.612%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.427%
Alistipes finegoldii_nov_93.608%
Beduini massiliensis_nov_87.705%
Duncaniella freteri_nov_87.424%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_93.582%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.400%
Longibaculum muris_nov_86.957%
Duncaniella freteri_nov_86.842%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.290%
Blautia luti_nov_94.561%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.718%
Bacteroides uniformis_nov_95.893%
Faecalimonas umbilicata_nov_94.549%
Acetivibrio cellulolyticus_nov_83.405%
Mollicutes_[G−2] bacterium_MOT−187_nov_90.253%
Ihubacter massiliensis_nov_94.572%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.577%
Christensenella massiliensis_nov_88.041%
Duncaniella freteri_nov_87.400%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.792%
Duncaniella freteri_nov_89.135%
Duncaniella freteri_nov_86.028%
Turicibacter sanguinis_nov_95.923%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.549%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_90.408%
Odoribacter splanchnicus_nov_90.779%
Blautia schinkii_nov_93.711%
Lawsonibacter asaccharolyticus_nov_90.722%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_84.929%
Hydrogenoanaerobacterium saccharovorans_nov_90.249%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.089%
Bacteroidetes_[G−3] bacterium_HMT_436_nov_86.585%
Lachnospiraceae_[G−5] bacterium_MOT−170_nov_97.904%
Phocea massiliensis_nov_87.179%
Eisenbergiella massiliensis_nov_84.867%
Sporobacter termitidis_nov_87.580%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_83.636%
Prevotella shahii_nov_87.602%
Gluceribacter canis_nov_93.305%
Eubacteriales_[G−4] bacterium_MOT−164_nov_92.373%
Adlercreutzia muris_nov_97.186%
Kineothrix alysoides_nov_86.667%
Lawsonibacter asaccharolyticus_nov_90.329%
Duncaniella freteri_nov_92.653%
Duncaniella freteri_nov_86.290%
Muricomes intestini_nov_89.583%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.373%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.000%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_89.293%
Duncaniella freteri_nov_91.039%
Duncaniella freteri_nov_91.429%
Duncaniella freteri_nov_86.948%
Faecalicatena multispecies_sppn7_2_nov_91.858%
Faecalicatena multispecies_sppn9_2_nov_94.363%
Species
F15127.S19
F15127.S20
F15127.S21
F15127.S22
F15127.S23
F15127.S24
Samples