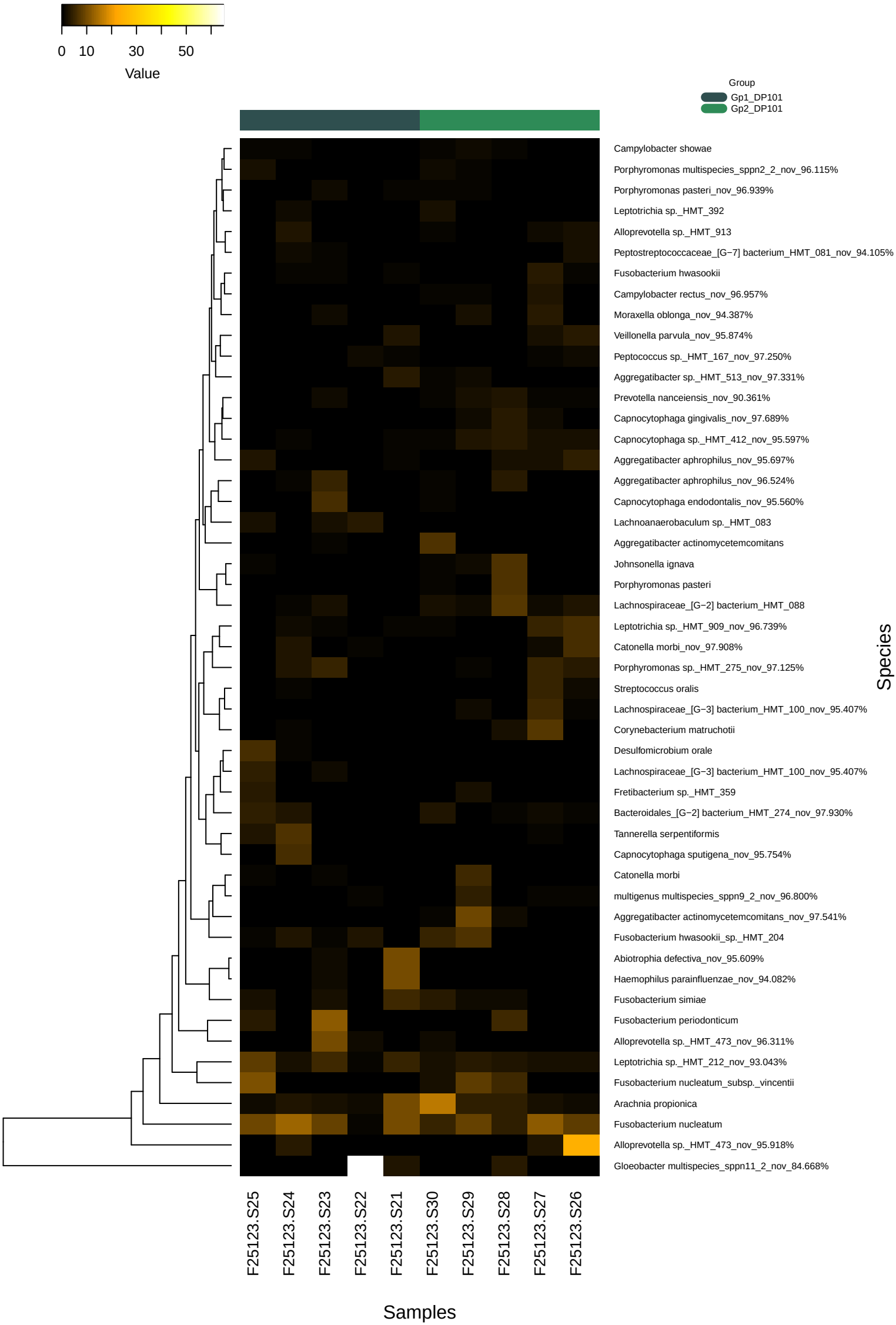
0
10
30
50
Value
Group
Gp1_DP101
Gp2_DP101
Campylobacter showae
Porphyromonas multispecies_sppn2_2_nov_96.115%
Porphyromonas pasteri_nov_96.939%
Leptotrichia sp._HMT_392
Alloprevotella sp._HMT_913
Peptostreptococcaceae_[G−7] bacterium_HMT_081_nov_94.105%
Fusobacterium hwasookii
Campylobacter rectus_nov_96.957%
Moraxella oblonga_nov_94.387%
Veillonella parvula_nov_95.874%
Peptococcus sp._HMT_167_nov_97.250%
Aggregatibacter sp._HMT_513_nov_97.331%
Prevotella nanceiensis_nov_90.361%
Capnocytophaga gingivalis_nov_97.689%
Capnocytophaga sp._HMT_412_nov_95.597%
Aggregatibacter aphrophilus_nov_95.697%
Aggregatibacter aphrophilus_nov_96.524%
Capnocytophaga endodontalis_nov_95.560%
Lachnoanaerobaculum sp._HMT_083
Aggregatibacter actinomycetemcomitans
Johnsonella ignava
Porphyromonas pasteri
Lachnospiraceae_[G−2] bacterium_HMT_088
Leptotrichia sp._HMT_909_nov_96.739%
Catonella morbi_nov_97.908%
Porphyromonas sp._HMT_275_nov_97.125%
Streptococcus oralis
Lachnospiraceae_[G−3] bacterium_HMT_100_nov_95.407%
Corynebacterium matruchotii
Desulfomicrobium orale
Lachnospiraceae_[G−3] bacterium_HMT_100_nov_95.407%
Fretibacterium sp._HMT_359
Bacteroidales_[G−2] bacterium_HMT_274_nov_97.930%
Tannerella serpentiformis
Capnocytophaga sputigena_nov_95.754%
Catonella morbi
multigenus multispecies_sppn9_2_nov_96.800%
Aggregatibacter actinomycetemcomitans_nov_97.541%
Fusobacterium hwasookii_sp._HMT_204
Abiotrophia defectiva_nov_95.609%
Haemophilus parainfluenzae_nov_94.082%
Fusobacterium simiae
Fusobacterium periodonticum
Alloprevotella sp._HMT_473_nov_96.311%
Leptotrichia sp._HMT_212_nov_93.043%
Fusobacterium nucleatum_subsp._vincentii
Arachnia propionica
Fusobacterium nucleatum
Alloprevotella sp._HMT_473_nov_95.918%
Gloeobacter multispecies_sppn11_2_nov_84.668%
Species
F25123.S25
F25123.S24
F25123.S23
F25123.S22
F25123.S21
F25123.S30
F25123.S29
F25123.S28
F25123.S27
F25123.S26
Samples