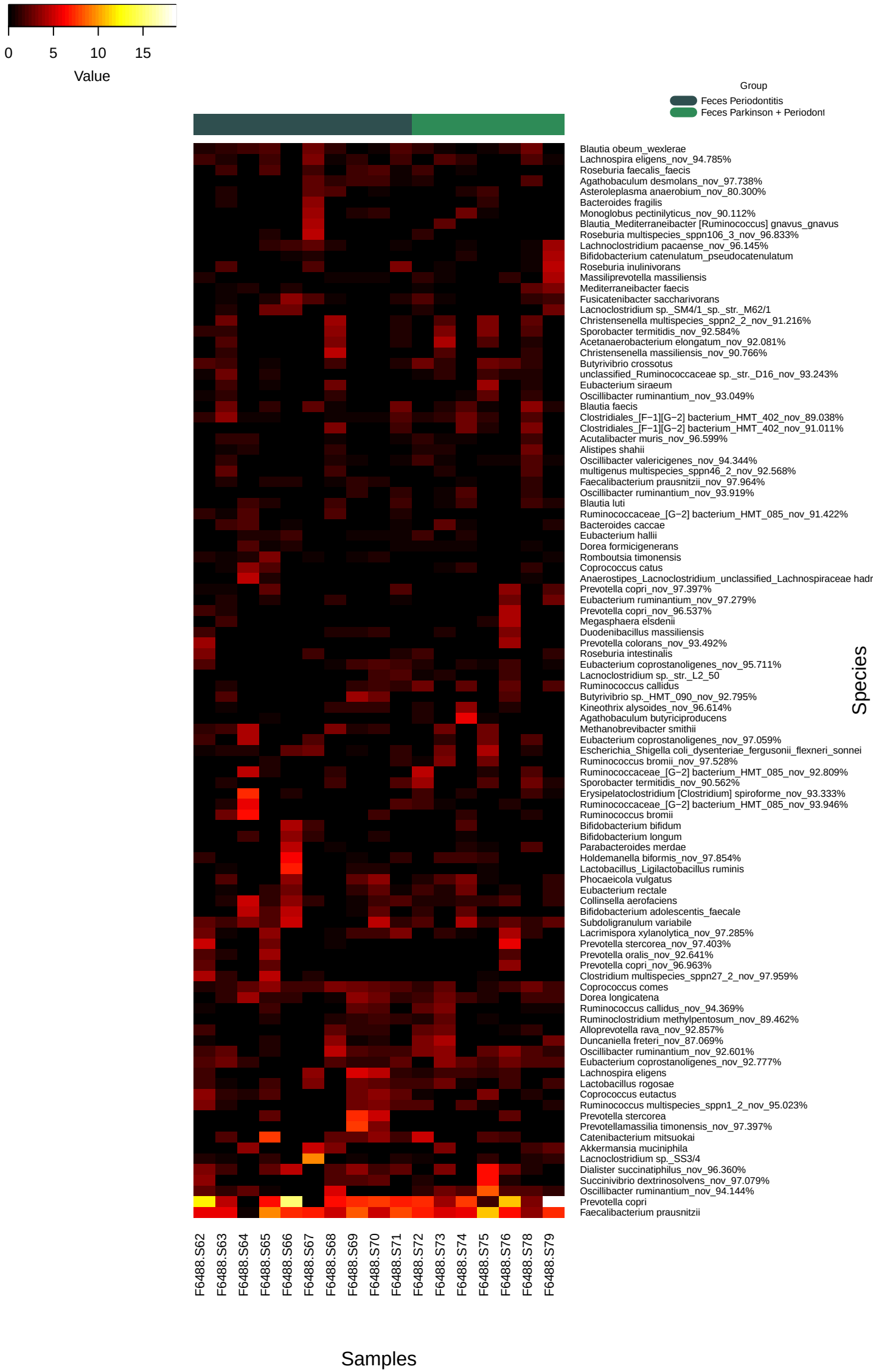
0
5
10
15
Value
Group
Feces Periodontitis
Feces Parkinson + Periodont
Blautia obeum_wexlerae
Lachnospira eligens_nov_94.785%
Roseburia faecalis_faecis
Agathobaculum desmolans_nov_97.738%
Asteroleplasma anaerobium_nov_80.300%
Bacteroides fragilis
Monoglobus pectinilyticus_nov_90.112%
Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
Roseburia multispecies_sppn106_3_nov_96.833%
Lachnoclostridium pacaense_nov_96.145%
Bifidobacterium catenulatum_pseudocatenulatum
Roseburia inulinivorans
Massiliprevotella massiliensis
Mediterraneibacter faecis
Fusicatenibacter saccharivorans
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Christensenella multispecies_sppn2_2_nov_91.216%
Sporobacter termitidis_nov_92.584%
Acetanaerobacterium elongatum_nov_92.081%
Christensenella massiliensis_nov_90.766%
Butyrivibrio crossotus
unclassified_Ruminococcaceae sp._str._D16_nov_93.243%
Eubacterium siraeum
Oscillibacter ruminantium_nov_93.049%
Blautia faecis
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_91.011%
Acutalibacter muris_nov_96.599%
Alistipes shahii
Oscillibacter valericigenes_nov_94.344%
multigenus multispecies_sppn46_2_nov_92.568%
Faecalibacterium prausnitzii_nov_97.964%
Oscillibacter ruminantium_nov_93.919%
Blautia luti
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_91.422%
Bacteroides caccae
Eubacterium hallii
Dorea formicigenerans
Romboutsia timonensis
Coprococcus catus
Anaerostipes_Lacnoclostridium_unclassified_Lachnospiraceae hadr
Prevotella copri_nov_97.397%
Eubacterium ruminantium_nov_97.279%
Prevotella copri_nov_96.537%
Megasphaera elsdenii
Duodenibacillus massiliensis
Prevotella colorans_nov_93.492%
Roseburia intestinalis
Eubacterium coprostanoligenes_nov_95.711%
Lacnoclostridium sp._str._L2_50
Ruminococcus callidus
Butyrivibrio sp._HMT_090_nov_92.795%
Kineothrix alysoides_nov_96.614%
Agathobaculum butyriciproducens
Methanobrevibacter smithii
Eubacterium coprostanoligenes_nov_97.059%
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Ruminococcus bromii_nov_97.528%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_92.809%
Sporobacter termitidis_nov_90.562%
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.946%
Ruminococcus bromii
Bifidobacterium bifidum
Bifidobacterium longum
Parabacteroides merdae
Holdemanella biformis_nov_97.854%
Lactobacillus_Ligilactobacillus ruminis
Phocaeicola vulgatus
Eubacterium rectale
Collinsella aerofaciens
Bifidobacterium adolescentis_faecale
Subdoligranulum variabile
Lacrimispora xylanolytica_nov_97.285%
Prevotella stercorea_nov_97.403%
Prevotella oralis_nov_92.641%
Prevotella copri_nov_96.963%
Clostridium multispecies_sppn27_2_nov_97.959%
Coprococcus comes
Dorea longicatena
Ruminococcus callidus_nov_94.369%
Ruminoclostridium methylpentosum_nov_89.462%
Alloprevotella rava_nov_92.857%
Duncaniella freteri_nov_87.069%
Oscillibacter ruminantium_nov_92.601%
Eubacterium coprostanoligenes_nov_92.777%
Lachnospira eligens
Lactobacillus rogosae
Coprococcus eutactus
Ruminococcus multispecies_sppn1_2_nov_95.023%
Prevotella stercorea
Prevotellamassilia timonensis_nov_97.397%
Catenibacterium mitsuokai
Akkermansia muciniphila
Lacnoclostridium sp._SS3/4
Dialister succinatiphilus_nov_96.360%
Succinivibrio dextrinosolvens_nov_97.079%
Oscillibacter ruminantium_nov_94.144%
Prevotella copri
Faecalibacterium prausnitzii
Species
F6488.S62
F6488.S63
F6488.S64
F6488.S65
F6488.S66
F6488.S67
F6488.S68
F6488.S69
F6488.S70
F6488.S71
F6488.S72
F6488.S73
F6488.S74
F6488.S75
F6488.S76
F6488.S78
F6488.S79
Samples