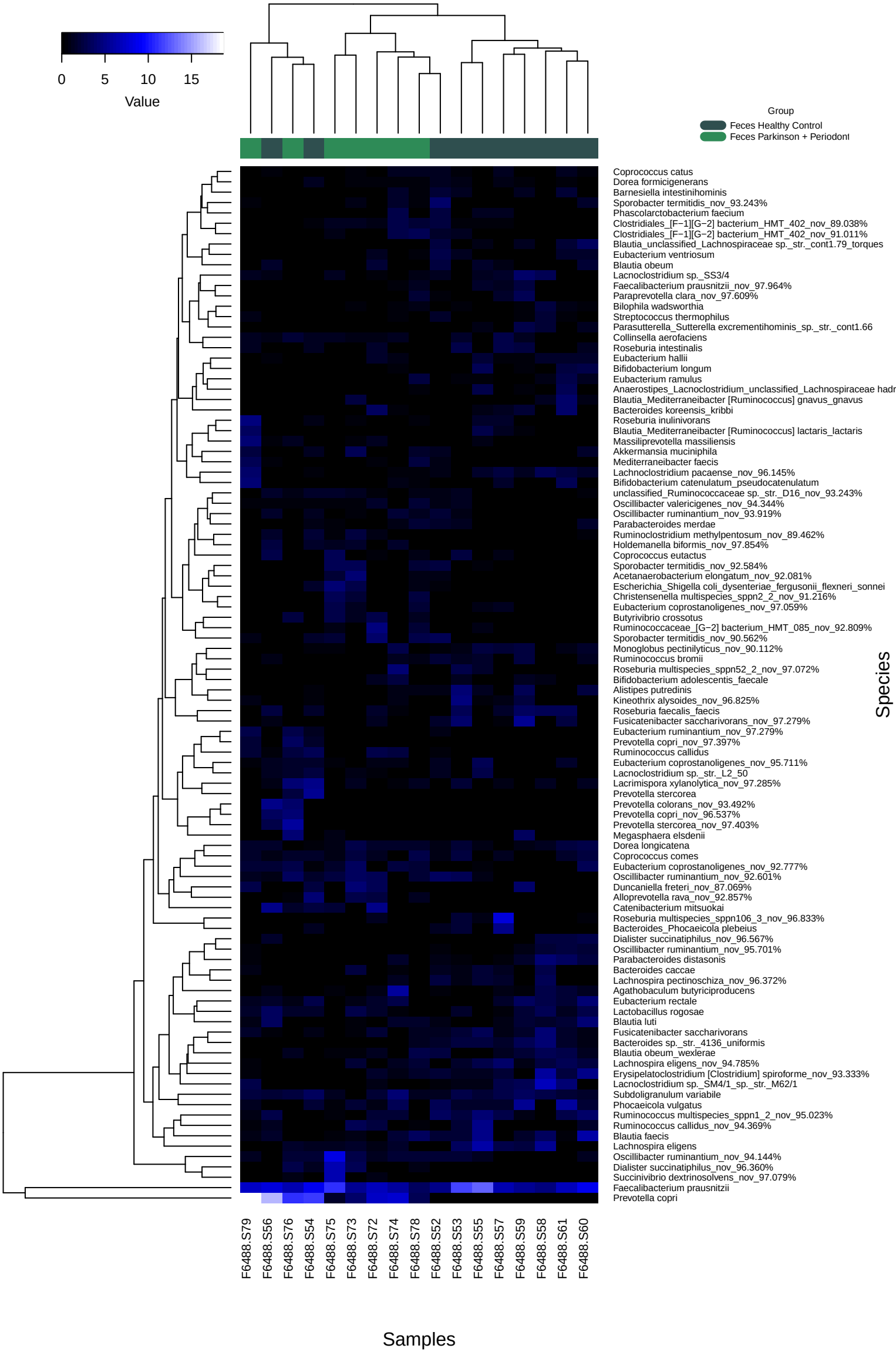
0
5
10
15
Value
Group
Feces Healthy Control
Feces Parkinson + Periodont
Coprococcus catus
Dorea formicigenerans
Barnesiella intestinihominis
Sporobacter termitidis_nov_93.243%
Phascolarctobacterium faecium
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_91.011%
Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torques
Eubacterium ventriosum
Blautia obeum
Lacnoclostridium sp._SS3/4
Faecalibacterium prausnitzii_nov_97.964%
Paraprevotella clara_nov_97.609%
Bilophila wadsworthia
Streptococcus thermophilus
Parasutterella_Sutterella excrementihominis_sp._str._cont1.66
Collinsella aerofaciens
Roseburia intestinalis
Eubacterium hallii
Bifidobacterium longum
Eubacterium ramulus
Anaerostipes_Lacnoclostridium_unclassified_Lachnospiraceae hadr
Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
Bacteroides koreensis_kribbi
Roseburia inulinivorans
Blautia_Mediterraneibacter [Ruminococcus] lactaris_lactaris
Massiliprevotella massiliensis
Akkermansia muciniphila
Mediterraneibacter faecis
Lachnoclostridium pacaense_nov_96.145%
Bifidobacterium catenulatum_pseudocatenulatum
unclassified_Ruminococcaceae sp._str._D16_nov_93.243%
Oscillibacter valericigenes_nov_94.344%
Oscillibacter ruminantium_nov_93.919%
Parabacteroides merdae
Ruminoclostridium methylpentosum_nov_89.462%
Holdemanella biformis_nov_97.854%
Coprococcus eutactus
Sporobacter termitidis_nov_92.584%
Acetanaerobacterium elongatum_nov_92.081%
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Christensenella multispecies_sppn2_2_nov_91.216%
Eubacterium coprostanoligenes_nov_97.059%
Butyrivibrio crossotus
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_92.809%
Sporobacter termitidis_nov_90.562%
Monoglobus pectinilyticus_nov_90.112%
Ruminococcus bromii
Roseburia multispecies_sppn52_2_nov_97.072%
Bifidobacterium adolescentis_faecale
Alistipes putredinis
Kineothrix alysoides_nov_96.825%
Roseburia faecalis_faecis
Fusicatenibacter saccharivorans_nov_97.279%
Eubacterium ruminantium_nov_97.279%
Prevotella copri_nov_97.397%
Ruminococcus callidus
Eubacterium coprostanoligenes_nov_95.711%
Lacnoclostridium sp._str._L2_50
Lacrimispora xylanolytica_nov_97.285%
Prevotella stercorea
Prevotella colorans_nov_93.492%
Prevotella copri_nov_96.537%
Prevotella stercorea_nov_97.403%
Megasphaera elsdenii
Dorea longicatena
Coprococcus comes
Eubacterium coprostanoligenes_nov_92.777%
Oscillibacter ruminantium_nov_92.601%
Duncaniella freteri_nov_87.069%
Alloprevotella rava_nov_92.857%
Catenibacterium mitsuokai
Roseburia multispecies_sppn106_3_nov_96.833%
Bacteroides_Phocaeicola plebeius
Dialister succinatiphilus_nov_96.567%
Oscillibacter ruminantium_nov_95.701%
Parabacteroides distasonis
Bacteroides caccae
Lachnospira pectinoschiza_nov_96.372%
Agathobaculum butyriciproducens
Eubacterium rectale
Lactobacillus rogosae
Blautia luti
Fusicatenibacter saccharivorans
Bacteroides sp._str._4136_uniformis
Blautia obeum_wexlerae
Lachnospira eligens_nov_94.785%
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Subdoligranulum variabile
Phocaeicola vulgatus
Ruminococcus multispecies_sppn1_2_nov_95.023%
Ruminococcus callidus_nov_94.369%
Blautia faecis
Lachnospira eligens
Oscillibacter ruminantium_nov_94.144%
Dialister succinatiphilus_nov_96.360%
Succinivibrio dextrinosolvens_nov_97.079%
Faecalibacterium prausnitzii
Prevotella copri
Species
F6488.S79
F6488.S56
F6488.S76
F6488.S54
F6488.S75
F6488.S73
F6488.S72
F6488.S74
F6488.S78
F6488.S52
F6488.S53
F6488.S55
F6488.S57
F6488.S59
F6488.S58
F6488.S61
F6488.S60
Samples