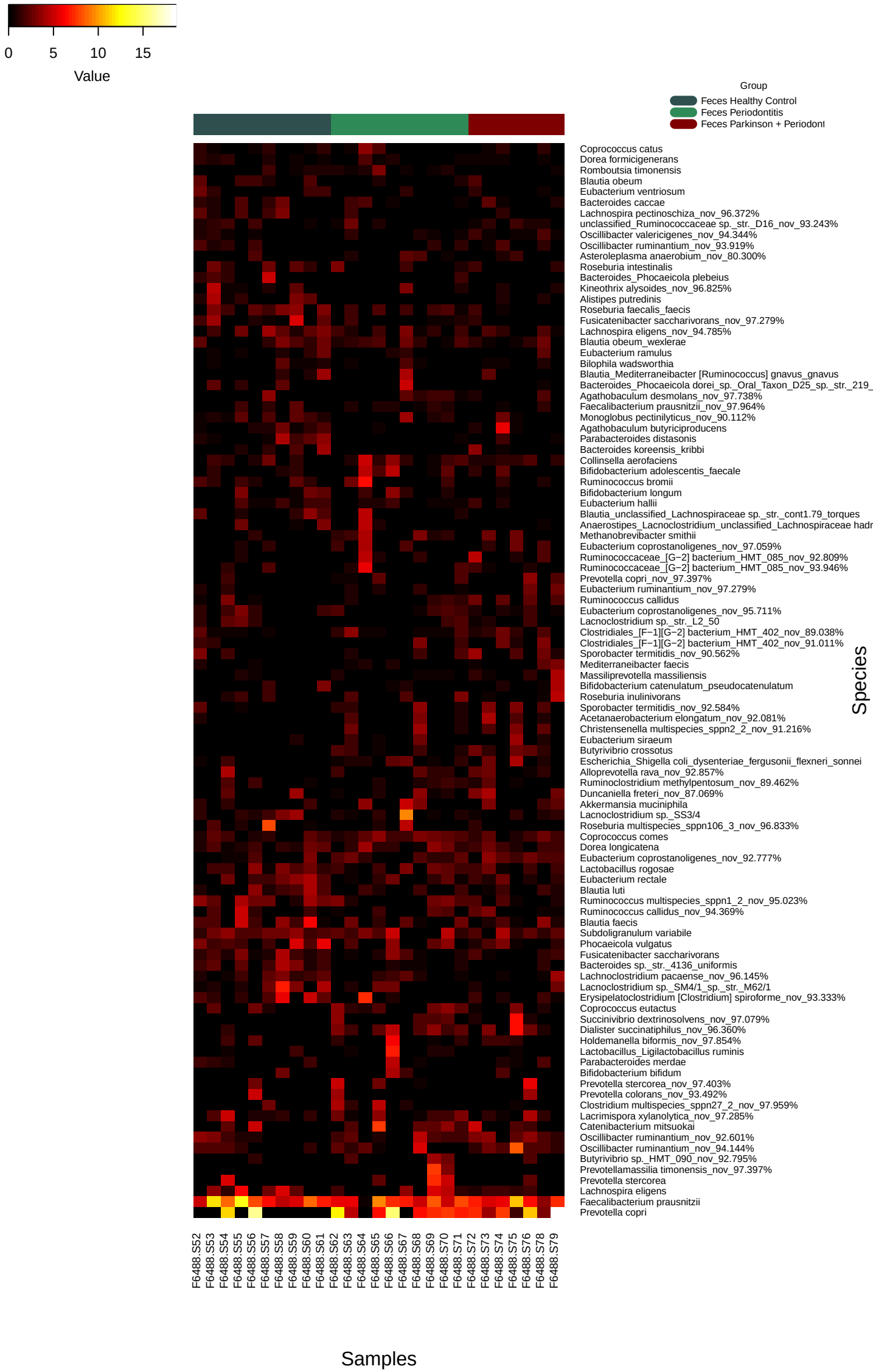
0
5
10
15
Value
Group
Feces Healthy Control
Feces Periodontitis
Feces Parkinson + Periodont
Coprococcus catus
Dorea formicigenerans
Romboutsia timonensis
Blautia obeum
Eubacterium ventriosum
Bacteroides caccae
Lachnospira pectinoschiza_nov_96.372%
unclassified_Ruminococcaceae sp._str._D16_nov_93.243%
Oscillibacter valericigenes_nov_94.344%
Oscillibacter ruminantium_nov_93.919%
Asteroleplasma anaerobium_nov_80.300%
Roseburia intestinalis
Bacteroides_Phocaeicola plebeius
Kineothrix alysoides_nov_96.825%
Alistipes putredinis
Roseburia faecalis_faecis
Fusicatenibacter saccharivorans_nov_97.279%
Lachnospira eligens_nov_94.785%
Blautia obeum_wexlerae
Eubacterium ramulus
Bilophila wadsworthia
Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
Bacteroides_Phocaeicola dorei_sp._Oral_Taxon_D25_sp._str._219_
Agathobaculum desmolans_nov_97.738%
Faecalibacterium prausnitzii_nov_97.964%
Monoglobus pectinilyticus_nov_90.112%
Agathobaculum butyriciproducens
Parabacteroides distasonis
Bacteroides koreensis_kribbi
Collinsella aerofaciens
Bifidobacterium adolescentis_faecale
Ruminococcus bromii
Bifidobacterium longum
Eubacterium hallii
Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torques
Anaerostipes_Lacnoclostridium_unclassified_Lachnospiraceae hadr
Methanobrevibacter smithii
Eubacterium coprostanoligenes_nov_97.059%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_92.809%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.946%
Prevotella copri_nov_97.397%
Eubacterium ruminantium_nov_97.279%
Ruminococcus callidus
Eubacterium coprostanoligenes_nov_95.711%
Lacnoclostridium sp._str._L2_50
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_91.011%
Sporobacter termitidis_nov_90.562%
Mediterraneibacter faecis
Massiliprevotella massiliensis
Bifidobacterium catenulatum_pseudocatenulatum
Roseburia inulinivorans
Sporobacter termitidis_nov_92.584%
Acetanaerobacterium elongatum_nov_92.081%
Christensenella multispecies_sppn2_2_nov_91.216%
Eubacterium siraeum
Butyrivibrio crossotus
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Alloprevotella rava_nov_92.857%
Ruminoclostridium methylpentosum_nov_89.462%
Duncaniella freteri_nov_87.069%
Akkermansia muciniphila
Lacnoclostridium sp._SS3/4
Roseburia multispecies_sppn106_3_nov_96.833%
Coprococcus comes
Dorea longicatena
Eubacterium coprostanoligenes_nov_92.777%
Lactobacillus rogosae
Eubacterium rectale
Blautia luti
Ruminococcus multispecies_sppn1_2_nov_95.023%
Ruminococcus callidus_nov_94.369%
Blautia faecis
Subdoligranulum variabile
Phocaeicola vulgatus
Fusicatenibacter saccharivorans
Bacteroides sp._str._4136_uniformis
Lachnoclostridium pacaense_nov_96.145%
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Coprococcus eutactus
Succinivibrio dextrinosolvens_nov_97.079%
Dialister succinatiphilus_nov_96.360%
Holdemanella biformis_nov_97.854%
Lactobacillus_Ligilactobacillus ruminis
Parabacteroides merdae
Bifidobacterium bifidum
Prevotella stercorea_nov_97.403%
Prevotella colorans_nov_93.492%
Clostridium multispecies_sppn27_2_nov_97.959%
Lacrimispora xylanolytica_nov_97.285%
Catenibacterium mitsuokai
Oscillibacter ruminantium_nov_92.601%
Oscillibacter ruminantium_nov_94.144%
Butyrivibrio sp._HMT_090_nov_92.795%
Prevotellamassilia timonensis_nov_97.397%
Prevotella stercorea
Lachnospira eligens
Faecalibacterium prausnitzii
Prevotella copri
Species
F6488.S52
F6488.S53
F6488.S54
F6488.S55
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F6488.S72
F6488.S73
F6488.S74
F6488.S75
F6488.S76
F6488.S78
F6488.S79
Samples