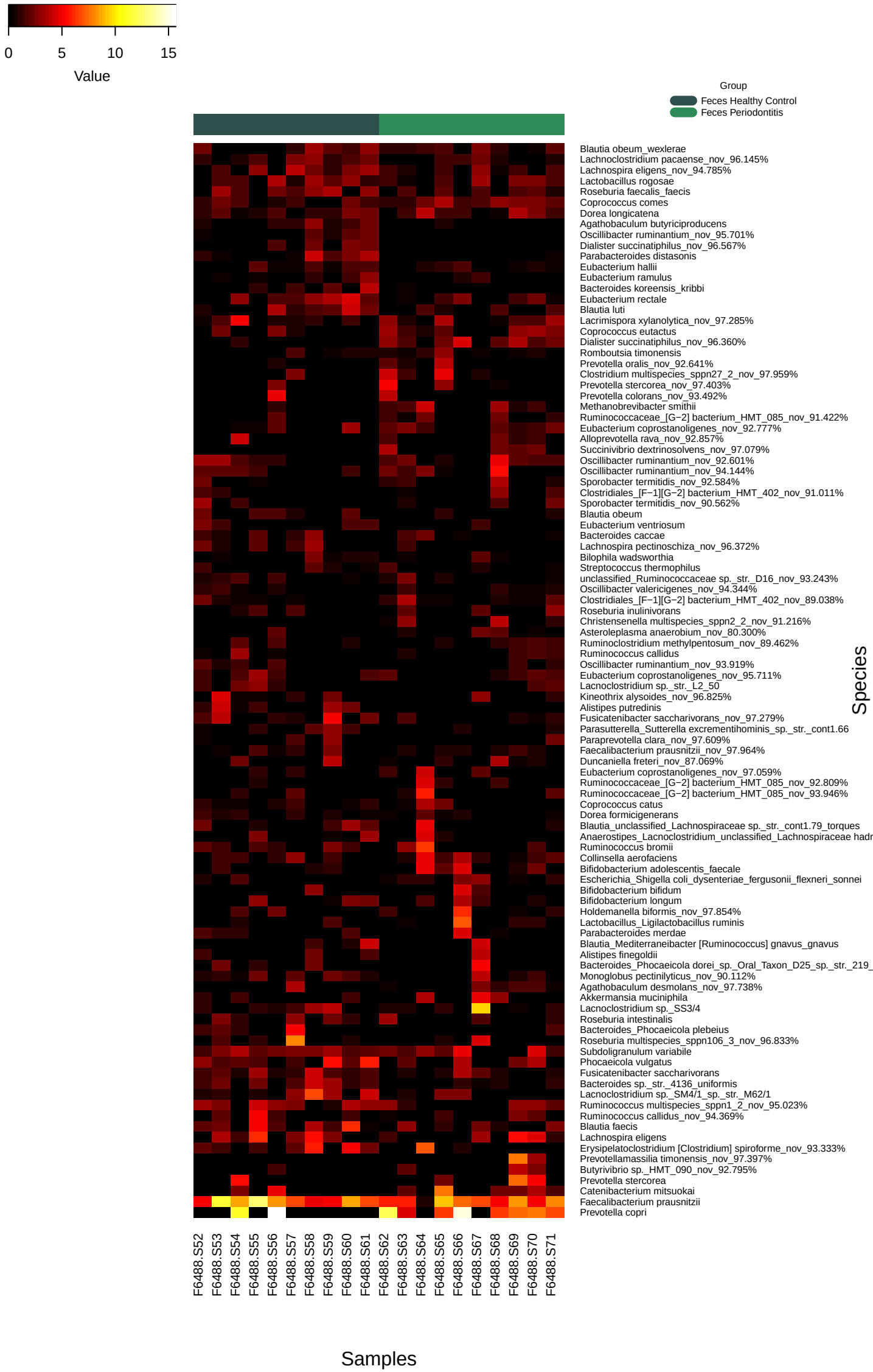
0
5
10
15
Value
Group
Feces Healthy Control
Feces Periodontitis
Blautia obeum_wexlerae
Lachnoclostridium pacaense_nov_96.145%
Lachnospira eligens_nov_94.785%
Lactobacillus rogosae
Roseburia faecalis_faecis
Coprococcus comes
Dorea longicatena
Agathobaculum butyriciproducens
Oscillibacter ruminantium_nov_95.701%
Dialister succinatiphilus_nov_96.567%
Parabacteroides distasonis
Eubacterium hallii
Eubacterium ramulus
Bacteroides koreensis_kribbi
Eubacterium rectale
Blautia luti
Lacrimispora xylanolytica_nov_97.285%
Coprococcus eutactus
Dialister succinatiphilus_nov_96.360%
Romboutsia timonensis
Prevotella oralis_nov_92.641%
Clostridium multispecies_sppn27_2_nov_97.959%
Prevotella stercorea_nov_97.403%
Prevotella colorans_nov_93.492%
Methanobrevibacter smithii
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_91.422%
Eubacterium coprostanoligenes_nov_92.777%
Alloprevotella rava_nov_92.857%
Succinivibrio dextrinosolvens_nov_97.079%
Oscillibacter ruminantium_nov_92.601%
Oscillibacter ruminantium_nov_94.144%
Sporobacter termitidis_nov_92.584%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_91.011%
Sporobacter termitidis_nov_90.562%
Blautia obeum
Eubacterium ventriosum
Bacteroides caccae
Lachnospira pectinoschiza_nov_96.372%
Bilophila wadsworthia
Streptococcus thermophilus
unclassified_Ruminococcaceae sp._str._D16_nov_93.243%
Oscillibacter valericigenes_nov_94.344%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Roseburia inulinivorans
Christensenella multispecies_sppn2_2_nov_91.216%
Asteroleplasma anaerobium_nov_80.300%
Ruminoclostridium methylpentosum_nov_89.462%
Ruminococcus callidus
Oscillibacter ruminantium_nov_93.919%
Eubacterium coprostanoligenes_nov_95.711%
Lacnoclostridium sp._str._L2_50
Kineothrix alysoides_nov_96.825%
Alistipes putredinis
Fusicatenibacter saccharivorans_nov_97.279%
Parasutterella_Sutterella excrementihominis_sp._str._cont1.66
Paraprevotella clara_nov_97.609%
Faecalibacterium prausnitzii_nov_97.964%
Duncaniella freteri_nov_87.069%
Eubacterium coprostanoligenes_nov_97.059%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_92.809%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.946%
Coprococcus catus
Dorea formicigenerans
Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torques
Anaerostipes_Lacnoclostridium_unclassified_Lachnospiraceae hadr
Ruminococcus bromii
Collinsella aerofaciens
Bifidobacterium adolescentis_faecale
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Bifidobacterium bifidum
Bifidobacterium longum
Holdemanella biformis_nov_97.854%
Lactobacillus_Ligilactobacillus ruminis
Parabacteroides merdae
Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
Alistipes finegoldii
Bacteroides_Phocaeicola dorei_sp._Oral_Taxon_D25_sp._str._219_
Monoglobus pectinilyticus_nov_90.112%
Agathobaculum desmolans_nov_97.738%
Akkermansia muciniphila
Lacnoclostridium sp._SS3/4
Roseburia intestinalis
Bacteroides_Phocaeicola plebeius
Roseburia multispecies_sppn106_3_nov_96.833%
Subdoligranulum variabile
Phocaeicola vulgatus
Fusicatenibacter saccharivorans
Bacteroides sp._str._4136_uniformis
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Ruminococcus multispecies_sppn1_2_nov_95.023%
Ruminococcus callidus_nov_94.369%
Blautia faecis
Lachnospira eligens
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Prevotellamassilia timonensis_nov_97.397%
Butyrivibrio sp._HMT_090_nov_92.795%
Prevotella stercorea
Catenibacterium mitsuokai
Faecalibacterium prausnitzii
Prevotella copri
Species
F6488.S52
F6488.S53
F6488.S54
F6488.S55
F6488.S56
F6488.S57
F6488.S58
F6488.S59
F6488.S60
F6488.S61
F6488.S62
F6488.S63
F6488.S64
F6488.S65
F6488.S66
F6488.S67
F6488.S68
F6488.S69
F6488.S70
F6488.S71
Samples