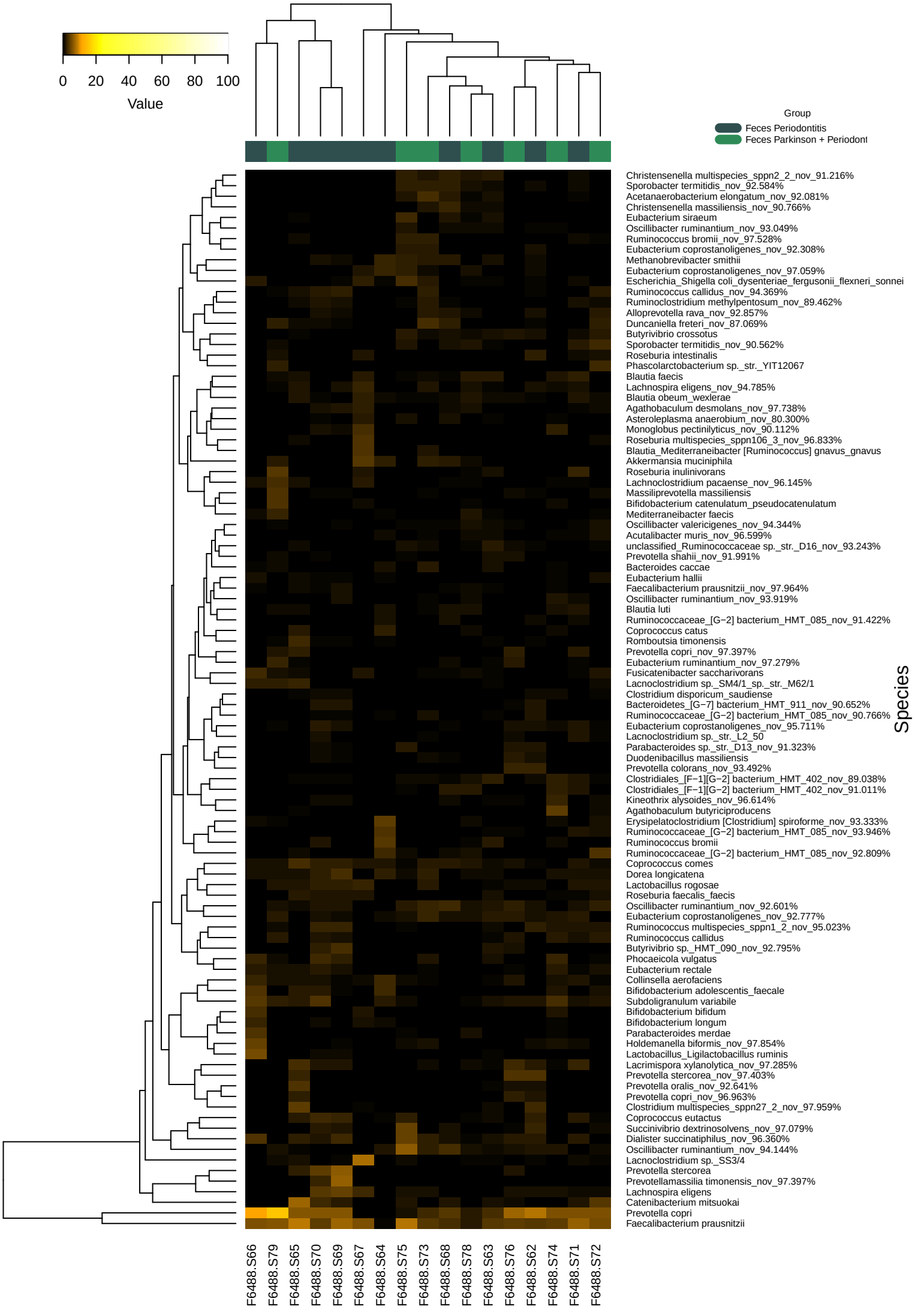
0
20
40
60
80
100
Value
Group
Feces Periodontitis
Feces Parkinson + Periodont
Christensenella multispecies_sppn2_2_nov_91.216%
Sporobacter termitidis_nov_92.584%
Acetanaerobacterium elongatum_nov_92.081%
Christensenella massiliensis_nov_90.766%
Eubacterium siraeum
Oscillibacter ruminantium_nov_93.049%
Ruminococcus bromii_nov_97.528%
Eubacterium coprostanoligenes_nov_92.308%
Methanobrevibacter smithii
Eubacterium coprostanoligenes_nov_97.059%
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Ruminococcus callidus_nov_94.369%
Ruminoclostridium methylpentosum_nov_89.462%
Alloprevotella rava_nov_92.857%
Duncaniella freteri_nov_87.069%
Butyrivibrio crossotus
Sporobacter termitidis_nov_90.562%
Roseburia intestinalis
Phascolarctobacterium sp._str._YIT12067
Blautia faecis
Lachnospira eligens_nov_94.785%
Blautia obeum_wexlerae
Agathobaculum desmolans_nov_97.738%
Asteroleplasma anaerobium_nov_80.300%
Monoglobus pectinilyticus_nov_90.112%
Roseburia multispecies_sppn106_3_nov_96.833%
Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
Akkermansia muciniphila
Roseburia inulinivorans
Lachnoclostridium pacaense_nov_96.145%
Massiliprevotella massiliensis
Bifidobacterium catenulatum_pseudocatenulatum
Mediterraneibacter faecis
Oscillibacter valericigenes_nov_94.344%
Acutalibacter muris_nov_96.599%
unclassified_Ruminococcaceae sp._str._D16_nov_93.243%
Prevotella shahii_nov_91.991%
Bacteroides caccae
Eubacterium hallii
Faecalibacterium prausnitzii_nov_97.964%
Oscillibacter ruminantium_nov_93.919%
Blautia luti
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_91.422%
Coprococcus catus
Romboutsia timonensis
Prevotella copri_nov_97.397%
Eubacterium ruminantium_nov_97.279%
Fusicatenibacter saccharivorans
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Clostridium disporicum_saudiense
Bacteroidetes_[G−7] bacterium_HMT_911_nov_90.652%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_90.766%
Eubacterium coprostanoligenes_nov_95.711%
Lacnoclostridium sp._str._L2_50
Parabacteroides sp._str._D13_nov_91.323%
Duodenibacillus massiliensis
Prevotella colorans_nov_93.492%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_91.011%
Kineothrix alysoides_nov_96.614%
Agathobaculum butyriciproducens
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.946%
Ruminococcus bromii
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_92.809%
Coprococcus comes
Dorea longicatena
Lactobacillus rogosae
Roseburia faecalis_faecis
Oscillibacter ruminantium_nov_92.601%
Eubacterium coprostanoligenes_nov_92.777%
Ruminococcus multispecies_sppn1_2_nov_95.023%
Ruminococcus callidus
Butyrivibrio sp._HMT_090_nov_92.795%
Phocaeicola vulgatus
Eubacterium rectale
Collinsella aerofaciens
Bifidobacterium adolescentis_faecale
Subdoligranulum variabile
Bifidobacterium bifidum
Bifidobacterium longum
Parabacteroides merdae
Holdemanella biformis_nov_97.854%
Lactobacillus_Ligilactobacillus ruminis
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 eutactus
Succinivibrio dextrinosolvens_nov_97.079%
Dialister succinatiphilus_nov_96.360%
Oscillibacter ruminantium_nov_94.144%
Lacnoclostridium sp._SS3/4
Prevotella stercorea
Prevotellamassilia timonensis_nov_97.397%
Lachnospira eligens
Catenibacterium mitsuokai
Prevotella copri
Faecalibacterium prausnitzii
Species
F6488.S66
F6488.S79
F6488.S65
F6488.S70
F6488.S69
F6488.S67
F6488.S64
F6488.S75
F6488.S73
F6488.S68
F6488.S78
F6488.S63
F6488.S76
F6488.S62
F6488.S74
F6488.S71
F6488.S72
Samples