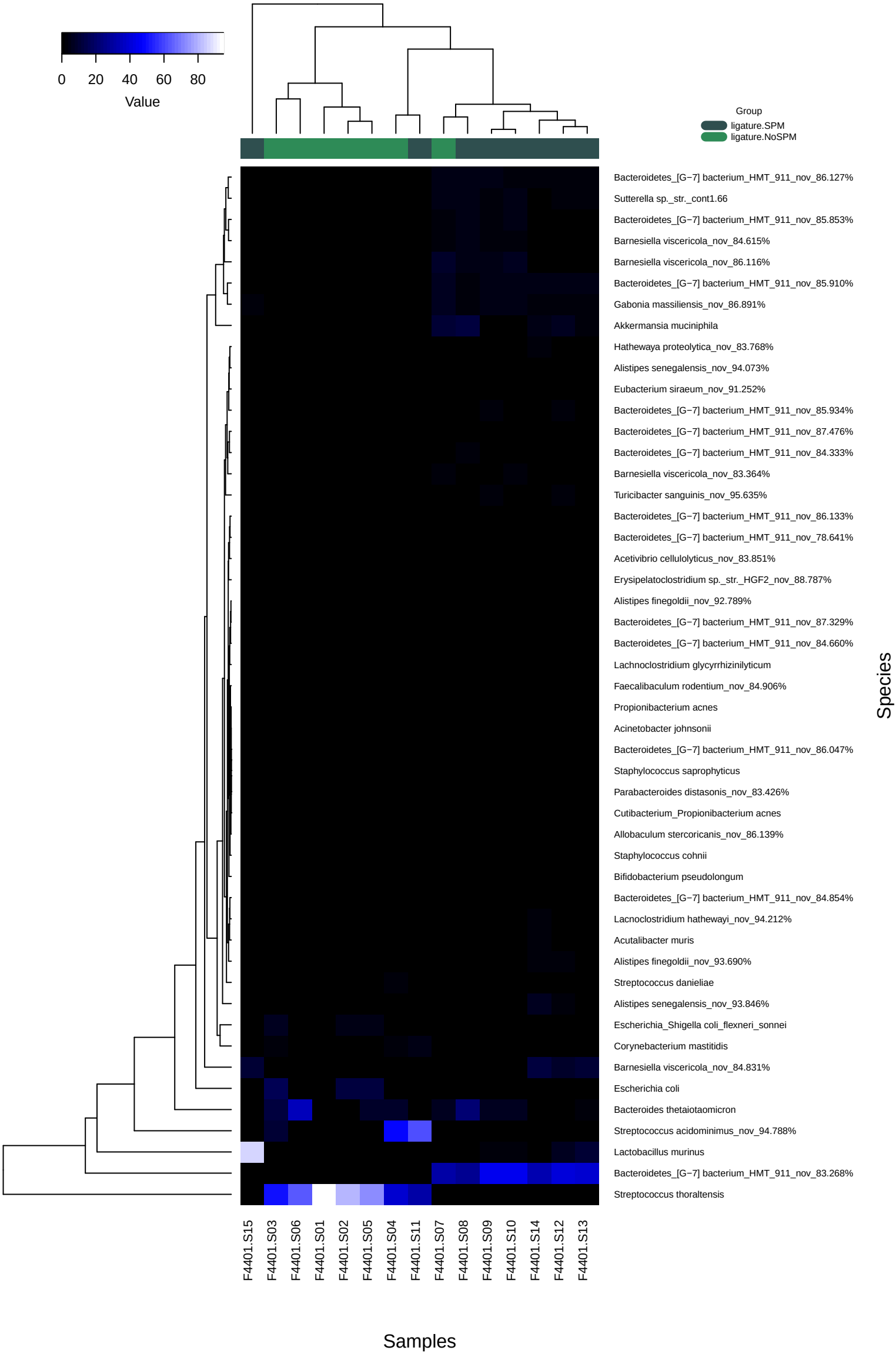
0
20
40
60
80
Value
Group
ligature.SPM
ligature.NoSPM
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.127%
Sutterella sp._str._cont1.66
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.853%
Barnesiella viscericola_nov_84.615%
Barnesiella viscericola_nov_86.116%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.910%
Gabonia massiliensis_nov_86.891%
Akkermansia muciniphila
Hathewaya proteolytica_nov_83.768%
Alistipes senegalensis_nov_94.073%
Eubacterium siraeum_nov_91.252%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.934%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_87.476%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.333%
Barnesiella viscericola_nov_83.364%
Turicibacter sanguinis_nov_95.635%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.133%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_78.641%
Acetivibrio cellulolyticus_nov_83.851%
Erysipelatoclostridium sp._str._HGF2_nov_88.787%
Alistipes finegoldii_nov_92.789%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_87.329%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.660%
Lachnoclostridium glycyrrhizinilyticum
Faecalibaculum rodentium_nov_84.906%
Propionibacterium acnes
Acinetobacter johnsonii
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.047%
Staphylococcus saprophyticus
Parabacteroides distasonis_nov_83.426%
Cutibacterium_Propionibacterium acnes
Allobaculum stercoricanis_nov_86.139%
Staphylococcus cohnii
Bifidobacterium pseudolongum
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.854%
Lacnoclostridium hathewayi_nov_94.212%
Acutalibacter muris
Alistipes finegoldii_nov_93.690%
Streptococcus danieliae
Alistipes senegalensis_nov_93.846%
Escherichia_Shigella coli_flexneri_sonnei
Corynebacterium mastitidis
Barnesiella viscericola_nov_84.831%
Escherichia coli
Bacteroides thetaiotaomicron
Streptococcus acidominimus_nov_94.788%
Lactobacillus murinus
Bacteroidetes_[G−7] bacterium_HMT_911_nov_83.268%
Streptococcus thoraltensis
Species
F4401.S15
F4401.S03
F4401.S06
F4401.S01
F4401.S02
F4401.S05
F4401.S04
F4401.S11
F4401.S07
F4401.S08
F4401.S09
F4401.S10
F4401.S14
F4401.S12
F4401.S13
Samples