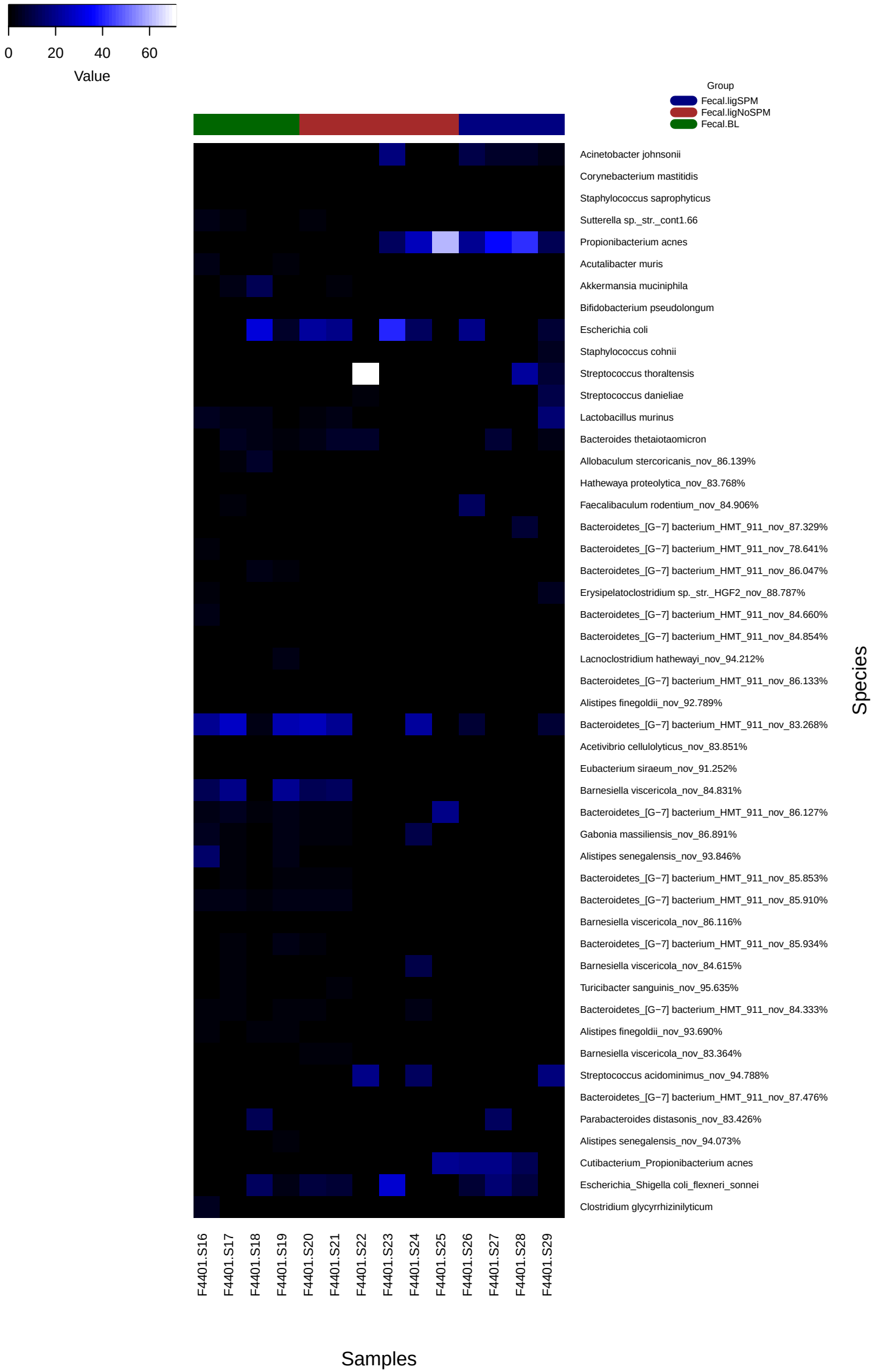
0
20
40
60
Value
Group
Fecal.ligSPM
Fecal.ligNoSPM
Fecal.BL
Acinetobacter johnsonii
Corynebacterium mastitidis
Staphylococcus saprophyticus
Sutterella sp._str._cont1.66
Propionibacterium acnes
Acutalibacter muris
Akkermansia muciniphila
Bifidobacterium pseudolongum
Escherichia coli
Staphylococcus cohnii
Streptococcus thoraltensis
Streptococcus danieliae
Lactobacillus murinus
Bacteroides thetaiotaomicron
Allobaculum stercoricanis_nov_86.139%
Hathewaya proteolytica_nov_83.768%
Faecalibaculum rodentium_nov_84.906%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_87.329%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_78.641%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.047%
Erysipelatoclostridium sp._str._HGF2_nov_88.787%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.660%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.854%
Lacnoclostridium hathewayi_nov_94.212%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.133%
Alistipes finegoldii_nov_92.789%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_83.268%
Acetivibrio cellulolyticus_nov_83.851%
Eubacterium siraeum_nov_91.252%
Barnesiella viscericola_nov_84.831%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.127%
Gabonia massiliensis_nov_86.891%
Alistipes senegalensis_nov_93.846%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.853%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.910%
Barnesiella viscericola_nov_86.116%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.934%
Barnesiella viscericola_nov_84.615%
Turicibacter sanguinis_nov_95.635%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.333%
Alistipes finegoldii_nov_93.690%
Barnesiella viscericola_nov_83.364%
Streptococcus acidominimus_nov_94.788%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_87.476%
Parabacteroides distasonis_nov_83.426%
Alistipes senegalensis_nov_94.073%
Cutibacterium_Propionibacterium acnes
Escherichia_Shigella coli_flexneri_sonnei
Clostridium glycyrrhizinilyticum
Species
F4401.S16
F4401.S17
F4401.S18
F4401.S19
F4401.S20
F4401.S21
F4401.S22
F4401.S23
F4401.S24
F4401.S25
F4401.S26
F4401.S27
F4401.S28
F4401.S29
Samples