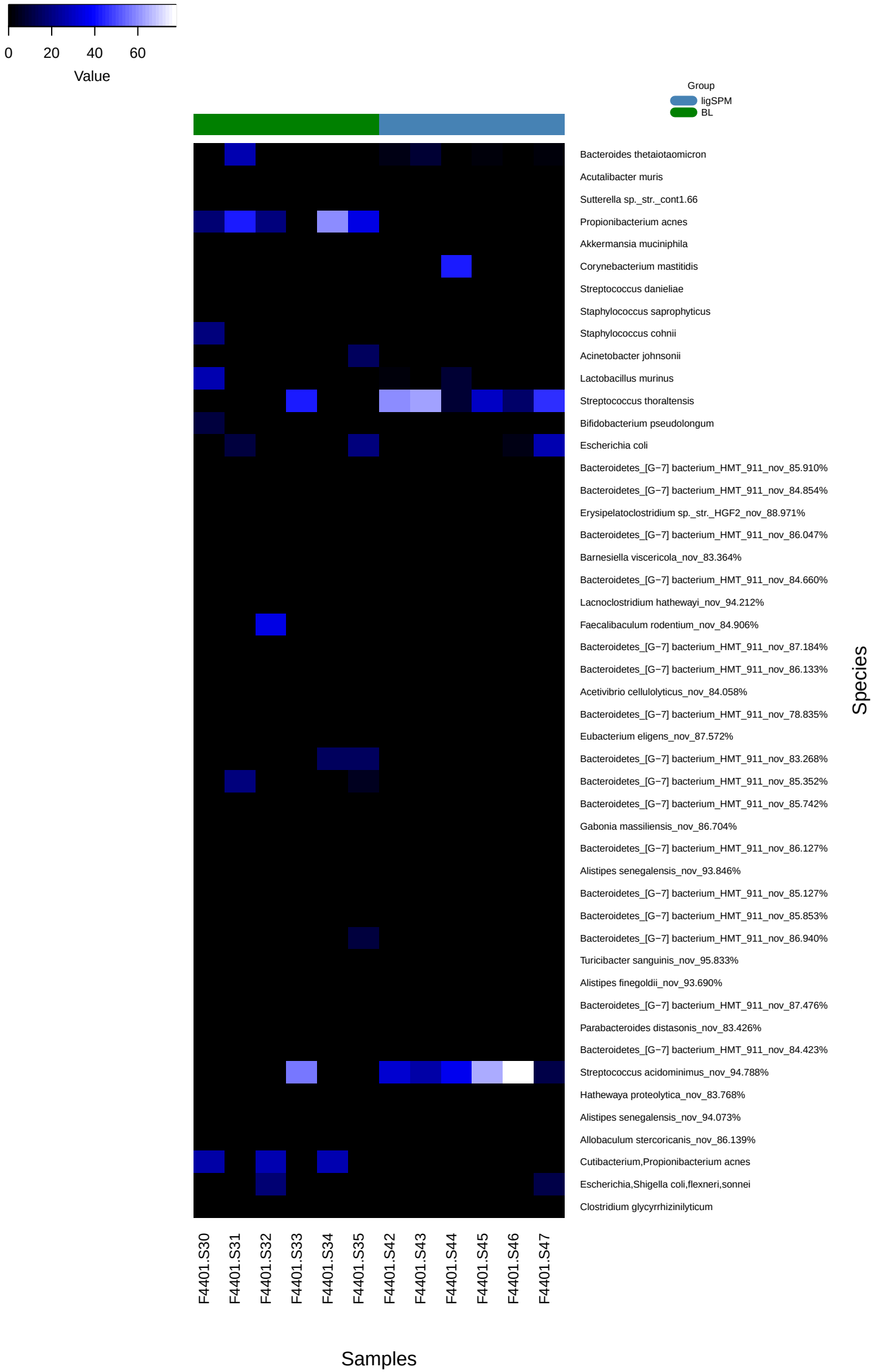
0
20
40
60
Value
Group
ligSPM
BL
Bacteroides thetaiotaomicron
Acutalibacter muris
Sutterella sp._str._cont1.66
Propionibacterium acnes
Akkermansia muciniphila
Corynebacterium mastitidis
Streptococcus danieliae
Staphylococcus saprophyticus
Staphylococcus cohnii
Acinetobacter johnsonii
Lactobacillus murinus
Streptococcus thoraltensis
Bifidobacterium pseudolongum
Escherichia coli
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.910%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.854%
Erysipelatoclostridium sp._str._HGF2_nov_88.971%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.047%
Barnesiella viscericola_nov_83.364%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.660%
Lacnoclostridium hathewayi_nov_94.212%
Faecalibaculum rodentium_nov_84.906%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_87.184%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.133%
Acetivibrio cellulolyticus_nov_84.058%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_78.835%
Eubacterium eligens_nov_87.572%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_83.268%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.352%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.742%
Gabonia massiliensis_nov_86.704%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.127%
Alistipes senegalensis_nov_93.846%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.127%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_85.853%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_86.940%
Turicibacter sanguinis_nov_95.833%
Alistipes finegoldii_nov_93.690%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_87.476%
Parabacteroides distasonis_nov_83.426%
Bacteroidetes_[G−7] bacterium_HMT_911_nov_84.423%
Streptococcus acidominimus_nov_94.788%
Hathewaya proteolytica_nov_83.768%
Alistipes senegalensis_nov_94.073%
Allobaculum stercoricanis_nov_86.139%
Cutibacterium,Propionibacterium acnes
Escherichia,Shigella coli,flexneri,sonnei
Clostridium glycyrrhizinilyticum
Species
F4401.S30
F4401.S31
F4401.S32
F4401.S33
F4401.S34
F4401.S35
F4401.S42
F4401.S43
F4401.S44
F4401.S45
F4401.S46
F4401.S47
Samples