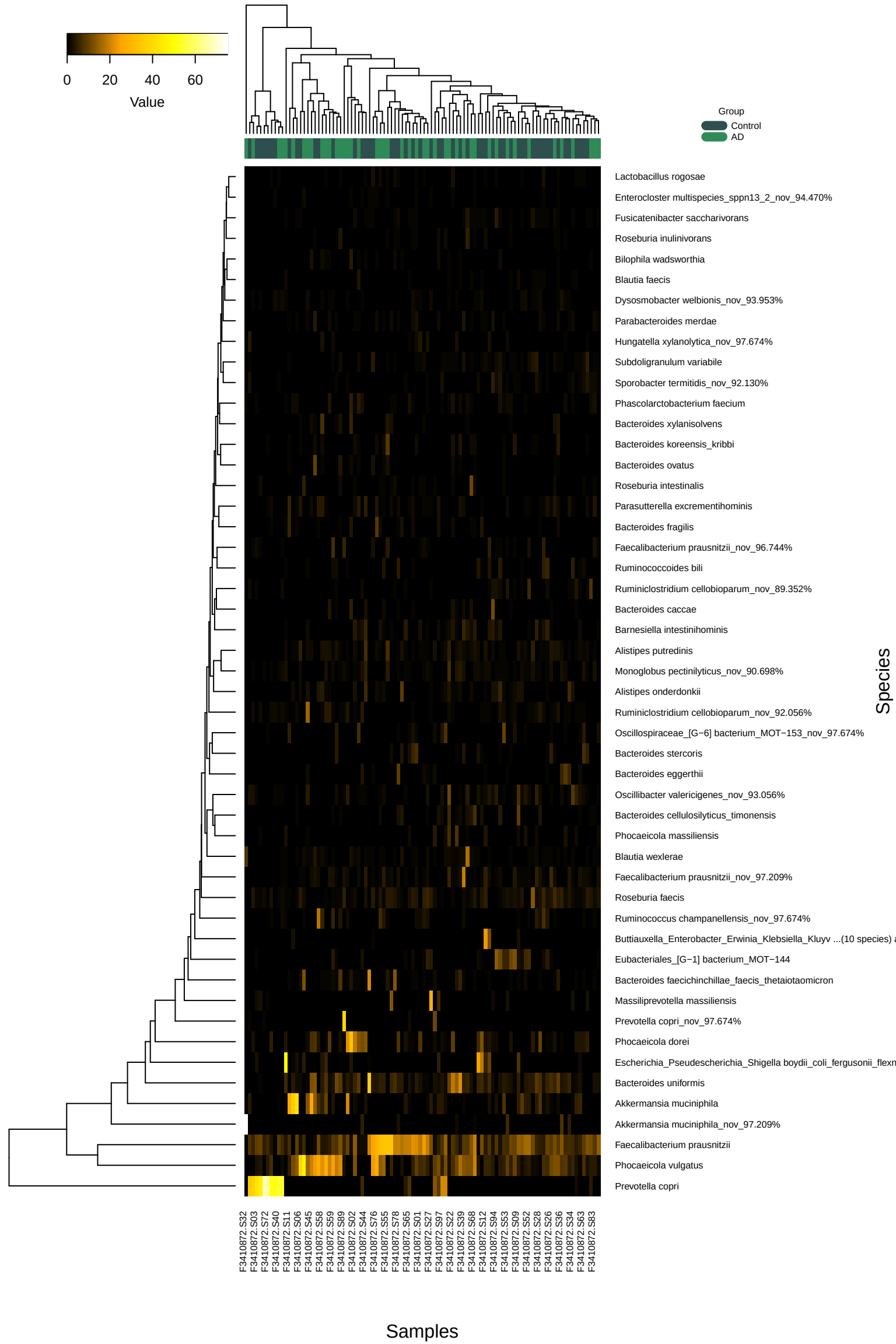
0
20
40
60
Value
Group
Control
AD
Lactobacillus rogosae
Enterocloster multispecies_sppn13_2_nov_94.470%
Fusicatenibacter saccharivorans
Roseburia inulinivorans
Bilophila wadsworthia
Blautia faecis
Dysosmobacter welbionis_nov_93.953%
Parabacteroides merdae
Hungatella xylanolytica_nov_97.674%
Subdoligranulum variabile
Sporobacter termitidis_nov_92.130%
Phascolarctobacterium faecium
Bacteroides xylanisolvens
Bacteroides koreensis_kribbi
Bacteroides ovatus
Roseburia intestinalis
Parasutterella excrementihominis
Bacteroides fragilis
Faecalibacterium prausnitzii_nov_96.744%
Ruminococcoides bili
Ruminiclostridium cellobioparum_nov_89.352%
Bacteroides caccae
Barnesiella intestinihominis
Alistipes putredinis
Monoglobus pectinilyticus_nov_90.698%
Alistipes onderdonkii
Ruminiclostridium cellobioparum_nov_92.056%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.674%
Bacteroides stercoris
Bacteroides eggerthii
Oscillibacter valericigenes_nov_93.056%
Bacteroides cellulosilyticus_timonensis
Phocaeicola massiliensis
Blautia wexlerae
Faecalibacterium prausnitzii_nov_97.209%
Roseburia faecis
Ruminococcus champanellensis_nov_97.674%
Buttiauxella_Enterobacter_Erwinia_Klebsiella_Kluyv ...(10 species) a
Eubacteriales_[G−1] bacterium_MOT−144
Bacteroides faecichinchillae_faecis_thetaiotaomicron
Massiliprevotella massiliensis
Prevotella copri_nov_97.674%
Phocaeicola dorei
Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexn
Bacteroides uniformis
Akkermansia muciniphila
Akkermansia muciniphila_nov_97.209%
Faecalibacterium prausnitzii
Phocaeicola vulgatus
Prevotella copri
Species
F3410872.S32
F3410872.S03
F3410872.S72
F3410872.S40
F3410872.S11
F3410872.S06
F3410872.S45
F3410872.S58
F3410872.S59
F3410872.S89
F3410872.S02
F3410872.S44
F3410872.S76
F3410872.S55
F3410872.S78
F3410872.S65
F3410872.S01
F3410872.S27
F3410872.S97
F3410872.S22
F3410872.S39
F3410872.S68
F3410872.S12
F3410872.S94
F3410872.S53
F3410872.S09
F3410872.S52
F3410872.S28
F3410872.S26
F3410872.S36
F3410872.S34
F3410872.S63
F3410872.S83
Samples