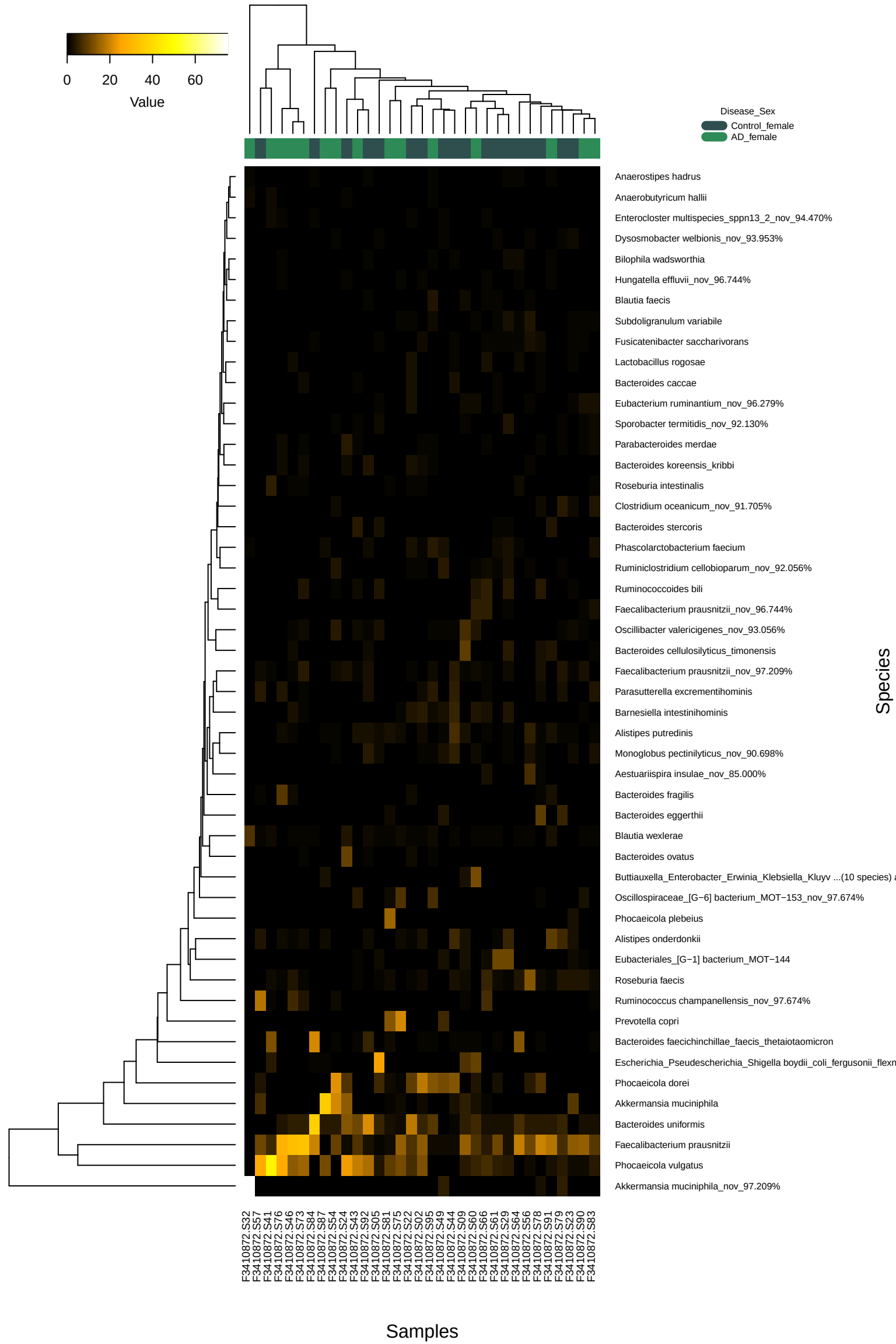
0
20
40
60
Value
Disease_Sex
Control_female
AD_female
Anaerostipes hadrus
Anaerobutyricum hallii
Enterocloster multispecies_sppn13_2_nov_94.470%
Dysosmobacter welbionis_nov_93.953%
Bilophila wadsworthia
Hungatella effluvii_nov_96.744%
Blautia faecis
Subdoligranulum variabile
Fusicatenibacter saccharivorans
Lactobacillus rogosae
Bacteroides caccae
Eubacterium ruminantium_nov_96.279%
Sporobacter termitidis_nov_92.130%
Parabacteroides merdae
Bacteroides koreensis_kribbi
Roseburia intestinalis
Clostridium oceanicum_nov_91.705%
Bacteroides stercoris
Phascolarctobacterium faecium
Ruminiclostridium cellobioparum_nov_92.056%
Ruminococcoides bili
Faecalibacterium prausnitzii_nov_96.744%
Oscillibacter valericigenes_nov_93.056%
Bacteroides cellulosilyticus_timonensis
Faecalibacterium prausnitzii_nov_97.209%
Parasutterella excrementihominis
Barnesiella intestinihominis
Alistipes putredinis
Monoglobus pectinilyticus_nov_90.698%
Aestuariispira insulae_nov_85.000%
Bacteroides fragilis
Bacteroides eggerthii
Blautia wexlerae
Bacteroides ovatus
Buttiauxella_Enterobacter_Erwinia_Klebsiella_Kluyv ...(10 species) a
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.674%
Phocaeicola plebeius
Alistipes onderdonkii
Eubacteriales_[G−1] bacterium_MOT−144
Roseburia faecis
Ruminococcus champanellensis_nov_97.674%
Prevotella copri
Bacteroides faecichinchillae_faecis_thetaiotaomicron
Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexn
Phocaeicola dorei
Akkermansia muciniphila
Bacteroides uniformis
Faecalibacterium prausnitzii
Phocaeicola vulgatus
Akkermansia muciniphila_nov_97.209%
Species
F3410872.S32
F3410872.S57
F3410872.S41
F3410872.S76
F3410872.S46
F3410872.S73
F3410872.S84
F3410872.S87
F3410872.S54
F3410872.S24
F3410872.S43
F3410872.S92
F3410872.S05
F3410872.S81
F3410872.S75
F3410872.S22
F3410872.S02
F3410872.S95
F3410872.S49
F3410872.S44
F3410872.S09
F3410872.S60
F3410872.S66
F3410872.S61
F3410872.S29
F3410872.S64
F3410872.S56
F3410872.S78
F3410872.S91
F3410872.S79
F3410872.S23
F3410872.S90
F3410872.S83
Samples