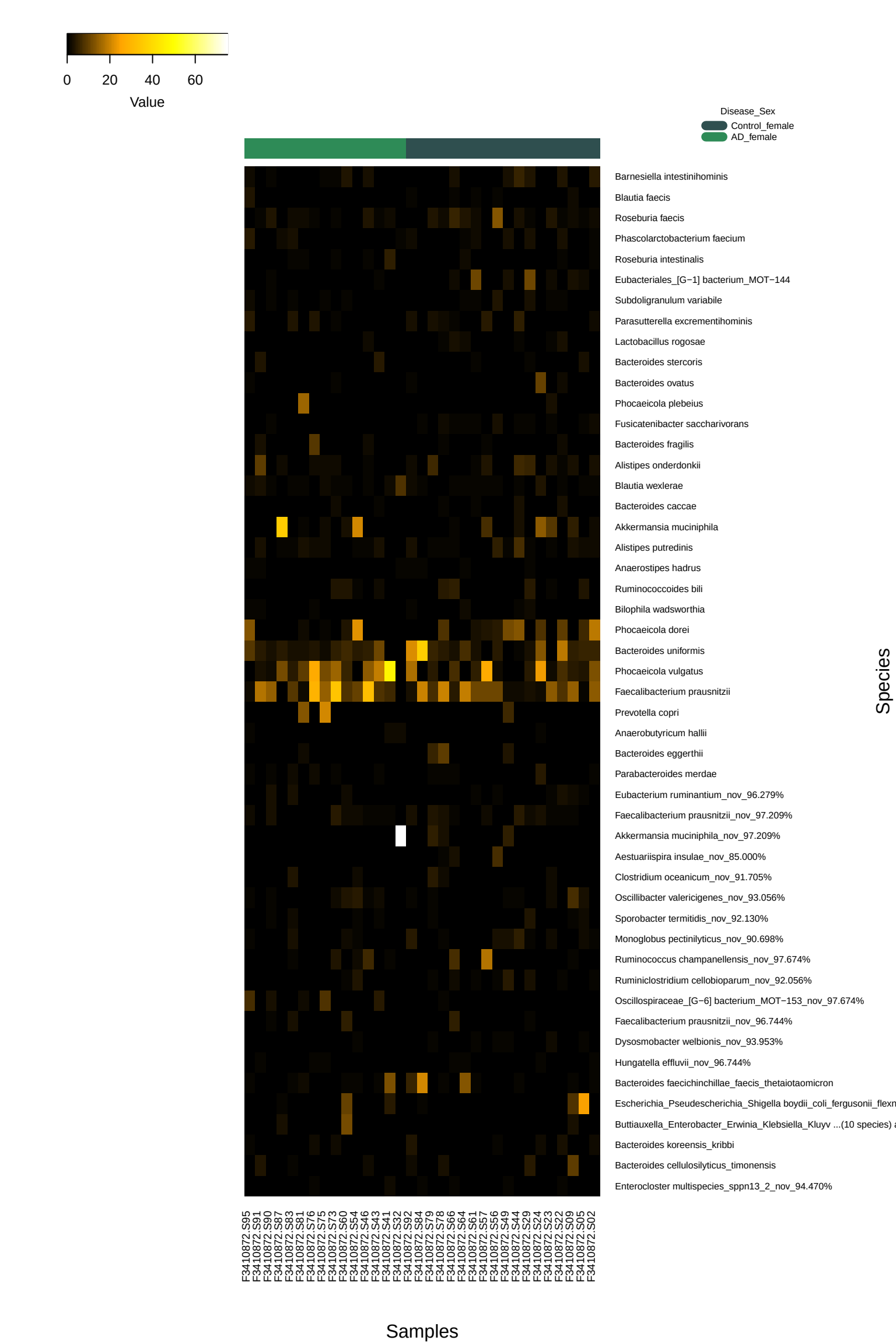
0
20
40
60
Value
Disease_Sex
Control_female
AD_female
Barnesiella intestinihominis
Blautia faecis
Roseburia faecis
Phascolarctobacterium faecium
Roseburia intestinalis
Eubacteriales_[G−1] bacterium_MOT−144
Subdoligranulum variabile
Parasutterella excrementihominis
Lactobacillus rogosae
Bacteroides stercoris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