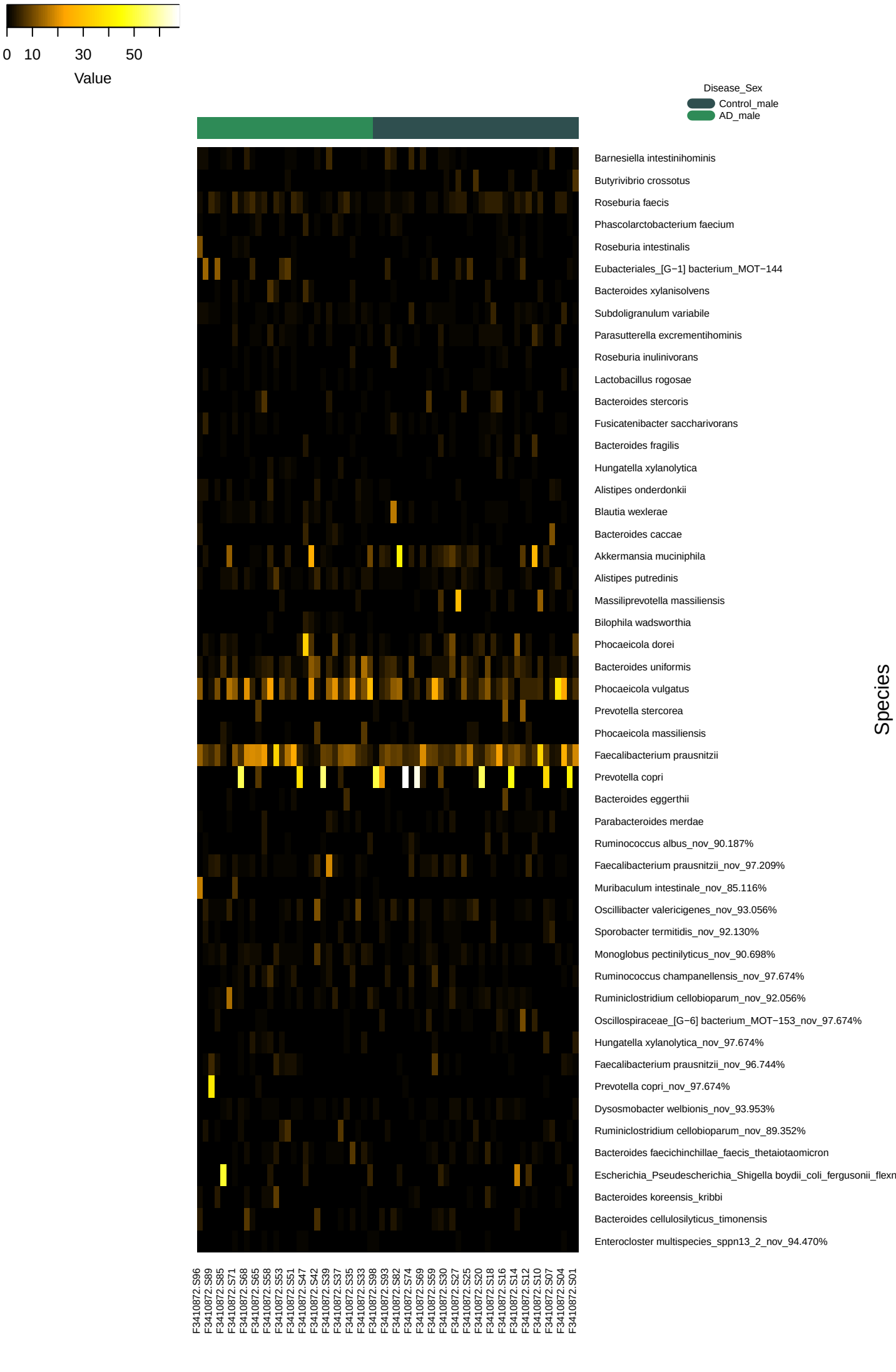
0 10 30 50
Value
Disease_Sex
Control_male
AD_male
Barnesiella intestinihominis
Butyrivibrio crossotus
Roseburia faecis
Phascolarctobacterium faecium
Roseburia intestinalis
Eubacteriales_[G−1] bacterium_MOT−144
Bacteroides xylanisolvens
Subdoligranulum variabile
Parasutterella excrementihominis
Roseburia inulinivorans
Lactobacillus rogosae
Bacteroides stercoris
Fusicatenibacter saccharivorans
Bacteroides fragilis
Hungatella xylanolytica
Alistipes onderdonkii
Blautia wexlerae
Bacteroides caccae
Akkermansia muciniphila
Alistipes putredinis
Massiliprevotella massiliensis
Bilophila wadsworthia
Phocaeicola dorei
Bacteroides uniformis
Phocaeicola vulgatus
Prevotella stercorea
Phocaeicola massiliensis
Faecalibacterium prausnitzii
Prevotella copri
Bacteroides eggerthii
Parabacteroides merdae
Ruminococcus albus_nov_90.187%
Faecalibacterium prausnitzii_nov_97.209%
Muribaculum intestinale_nov_85.116%
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%
Hungatella xylanolytica_nov_97.674%
Faecalibacterium prausnitzii_nov_96.744%
Prevotella copri_nov_97.674%
Dysosmobacter welbionis_nov_93.953%
Ruminiclostridium cellobioparum_nov_89.352%
Bacteroides faecichinchillae_faecis_thetaiotaomicron
Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexn
Bacteroides koreensis_kribbi
Bacteroides cellulosilyticus_timonensis
Enterocloster multispecies_sppn13_2_nov_94.470%
Species
F3410872.S96
F3410872.S89
F3410872.S85
F3410872.S71
F3410872.S68
F3410872.S65
F3410872.S58
F3410872.S53
F3410872.S51
F3410872.S47
F3410872.S42
F3410872.S39
F3410872.S37
F3410872.S35
F3410872.S33
F3410872.S98
F3410872.S93
F3410872.S82
F3410872.S74
F3410872.S69
F3410872.S59
F3410872.S30
F3410872.S27
F3410872.S25
F3410872.S20
F3410872.S18
F3410872.S16
F3410872.S14
F3410872.S12
F3410872.S10
F3410872.S07
F3410872.S04
F3410872.S01
Samples