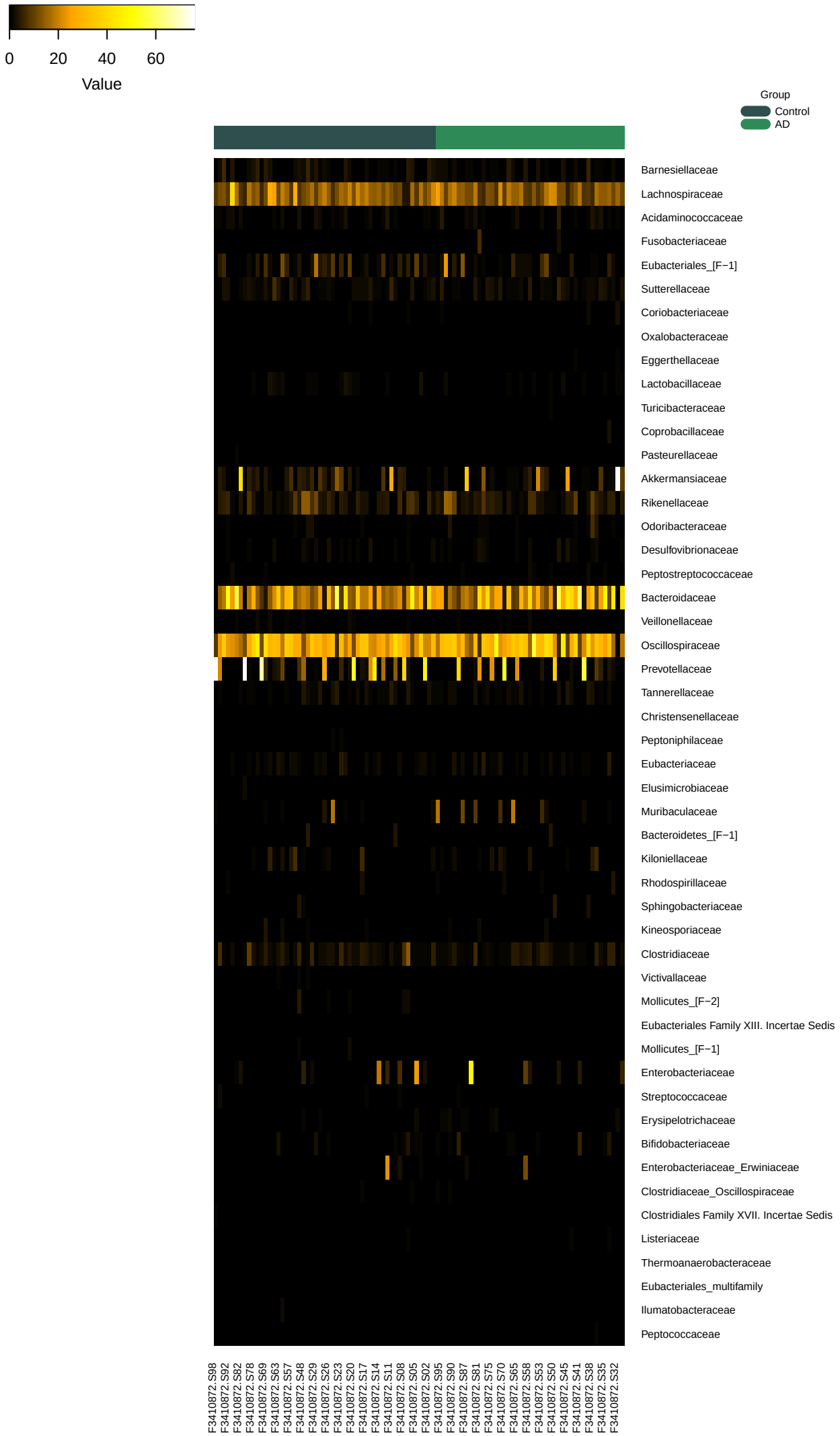
0
20
40
60
Value
Group
Control
AD
Barnesiellaceae
Lachnospiraceae
Acidaminococcaceae
Fusobacteriaceae
Eubacteriales_[F−1]
Sutterellaceae
Coriobacteriaceae
Oxalobacteraceae
Eggerthellaceae
Lactobacillaceae
Turicibacteraceae
Coprobacillaceae
Pasteurellaceae
Akkermansiaceae
Rikenellaceae
Odoribacteraceae
Desulfovibrionaceae
Peptostreptococcaceae
Bacteroidaceae
Veillonellaceae
Oscillospiraceae
Prevotellaceae
Tannerellaceae
Christensenellaceae
Peptoniphilaceae
Eubacteriaceae
Elusimicrobiaceae
Muribaculaceae
Bacteroidetes_[F−1]
Kiloniellaceae
Rhodospirillaceae
Sphingobacteriaceae
Kineosporiaceae
Clostridiaceae
Victivallaceae
Mollicutes_[F−2]
Eubacteriales Family XIII. Incertae Sedis
Mollicutes_[F−1]
Enterobacteriaceae
Streptococcaceae
Erysipelotrichaceae
Bifidobacteriaceae
Enterobacteriaceae_Erwiniaceae
Clostridiaceae_Oscillospiraceae
Clostridiales Family XVII. Incertae Sedis
Listeriaceae
Thermoanaerobacteraceae
Eubacteriales_multifamily
Ilumatobacteraceae
Peptococcaceae
Family
F3410872.S98
F3410872.S92
F3410872.S82
F3410872.S78
F3410872.S69
F3410872.S63
F3410872.S57
F3410872.S48
F3410872.S29
F3410872.S26
F3410872.S23
F3410872.S20
F3410872.S17
F3410872.S14
F3410872.S11
F3410872.S08
F3410872.S05
F3410872.S02
F3410872.S95
F3410872.S90
F3410872.S87
F3410872.S81
F3410872.S75
F3410872.S70
F3410872.S65
F3410872.S58
F3410872.S53
F3410872.S50
F3410872.S45
F3410872.S41
F3410872.S38
F3410872.S35
F3410872.S32
Samples