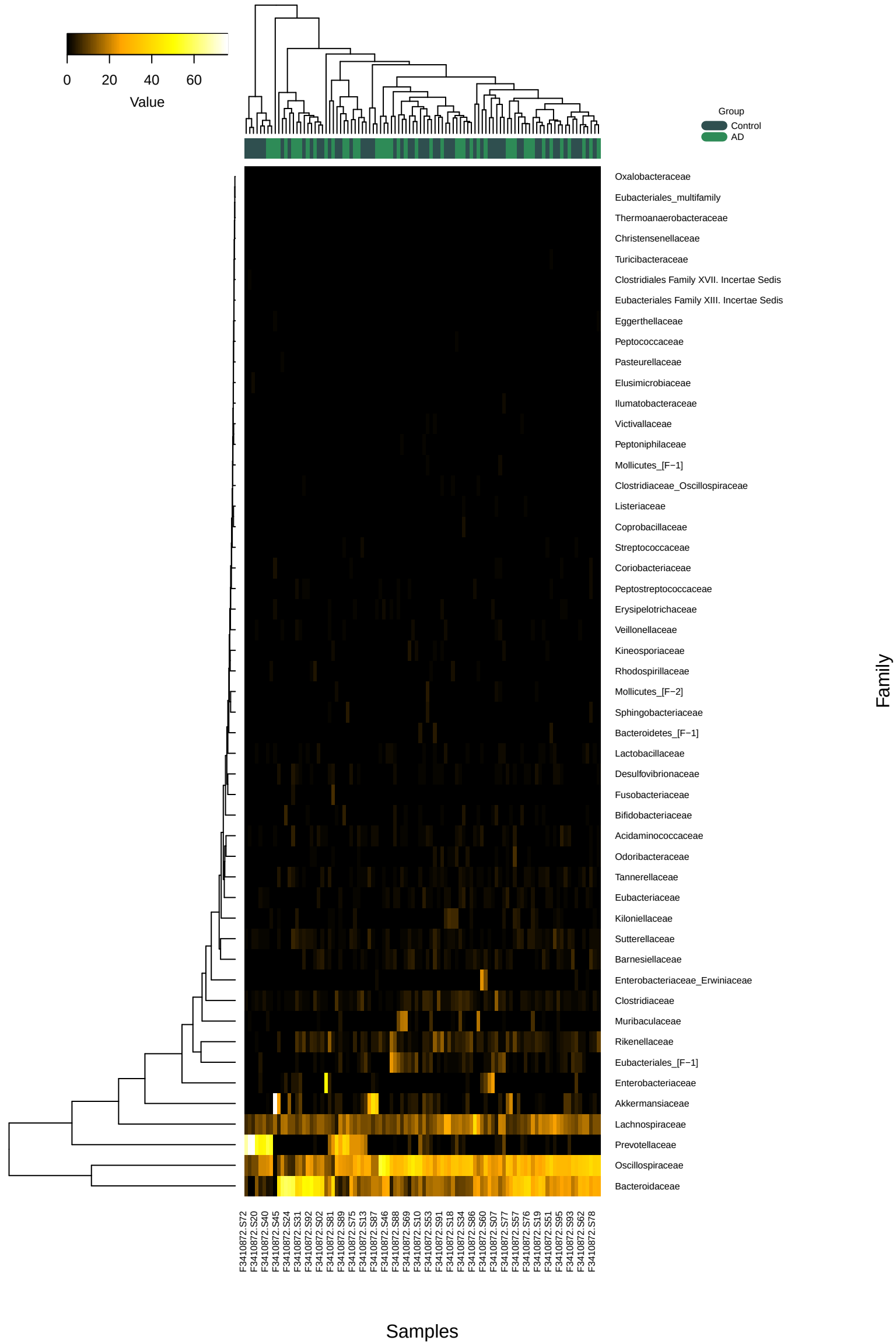
0
20
40
60
Value
Group
Control
AD
Oxalobacteraceae
Eubacteriales_multifamily
Thermoanaerobacteraceae
Christensenellaceae
Turicibacteraceae
Clostridiales Family XVII. Incertae Sedis
Eubacteriales Family XIII. Incertae Sedis
Eggerthellaceae
Peptococcaceae
Pasteurellaceae
Elusimicrobiaceae
Ilumatobacteraceae
Victivallaceae
Peptoniphilaceae
Mollicutes_[F−1]
Clostridiaceae_Oscillospiraceae
Listeriaceae
Coprobacillaceae
Streptococcaceae
Coriobacteriaceae
Peptostreptococcaceae
Erysipelotrichaceae
Veillonellaceae
Kineosporiaceae
Rhodospirillaceae
Mollicutes_[F−2]
Sphingobacteriaceae
Bacteroidetes_[F−1]
Lactobacillaceae
Desulfovibrionaceae
Fusobacteriaceae
Bifidobacteriaceae
Acidaminococcaceae
Odoribacteraceae
Tannerellaceae
Eubacteriaceae
Kiloniellaceae
Sutterellaceae
Barnesiellaceae
Enterobacteriaceae_Erwiniaceae
Clostridiaceae
Muribaculaceae
Rikenellaceae
Eubacteriales_[F−1]
Enterobacteriaceae
Akkermansiaceae
Lachnospiraceae
Prevotellaceae
Oscillospiraceae
Bacteroidaceae
Family
F3410872.S72
F3410872.S20
F3410872.S40
F3410872.S45
F3410872.S24
F3410872.S31
F3410872.S92
F3410872.S02
F3410872.S81
F3410872.S89
F3410872.S75
F3410872.S13
F3410872.S87
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F3410872.S91
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F3410872.S60
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F3410872.S77
F3410872.S57
F3410872.S76
F3410872.S19
F3410872.S51
F3410872.S95
F3410872.S93
F3410872.S62
F3410872.S78
Samples