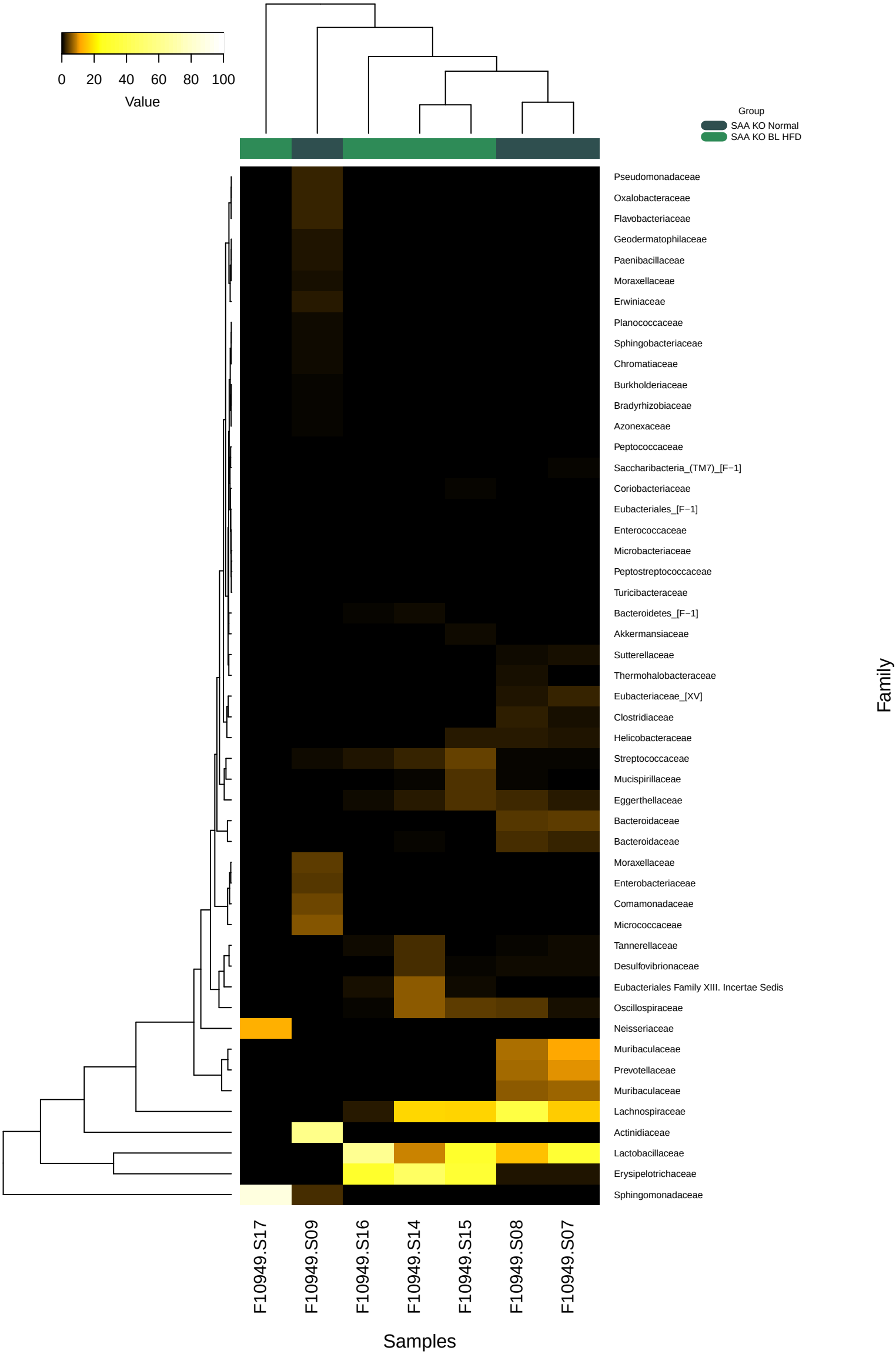
0
20
40
60
80
100
Value
Group
SAA KO Normal
SAA KO BL HFD
Pseudomonadaceae
Oxalobacteraceae
Flavobacteriaceae
Geodermatophilaceae
Paenibacillaceae
Moraxellaceae
Erwiniaceae
Planococcaceae
Sphingobacteriaceae
Chromatiaceae
Burkholderiaceae
Bradyrhizobiaceae
Azonexaceae
Peptococcaceae
Saccharibacteria_(TM7)_[F−1]
Coriobacteriaceae
Eubacteriales_[F−1]
Enterococcaceae
Microbacteriaceae
Peptostreptococcaceae
Turicibacteraceae
Bacteroidetes_[F−1]
Akkermansiaceae
Sutterellaceae
Thermohalobacteraceae
Eubacteriaceae_[XV]
Clostridiaceae
Helicobacteraceae
Streptococcaceae
Mucispirillaceae
Eggerthellaceae
Bacteroidaceae
Bacteroidaceae
Moraxellaceae
Enterobacteriaceae
Comamonadaceae
Micrococcaceae
Tannerellaceae
Desulfovibrionaceae
Eubacteriales Family XIII. Incertae Sedis
Oscillospiraceae
Neisseriaceae
Muribaculaceae
Prevotellaceae
Muribaculaceae
Lachnospiraceae
Actinidiaceae
Lactobacillaceae
Erysipelotrichaceae
Sphingomonadaceae
Family
F10949.S17
F10949.S09
F10949.S16
F10949.S14
F10949.S15
F10949.S08
F10949.S07
Samples