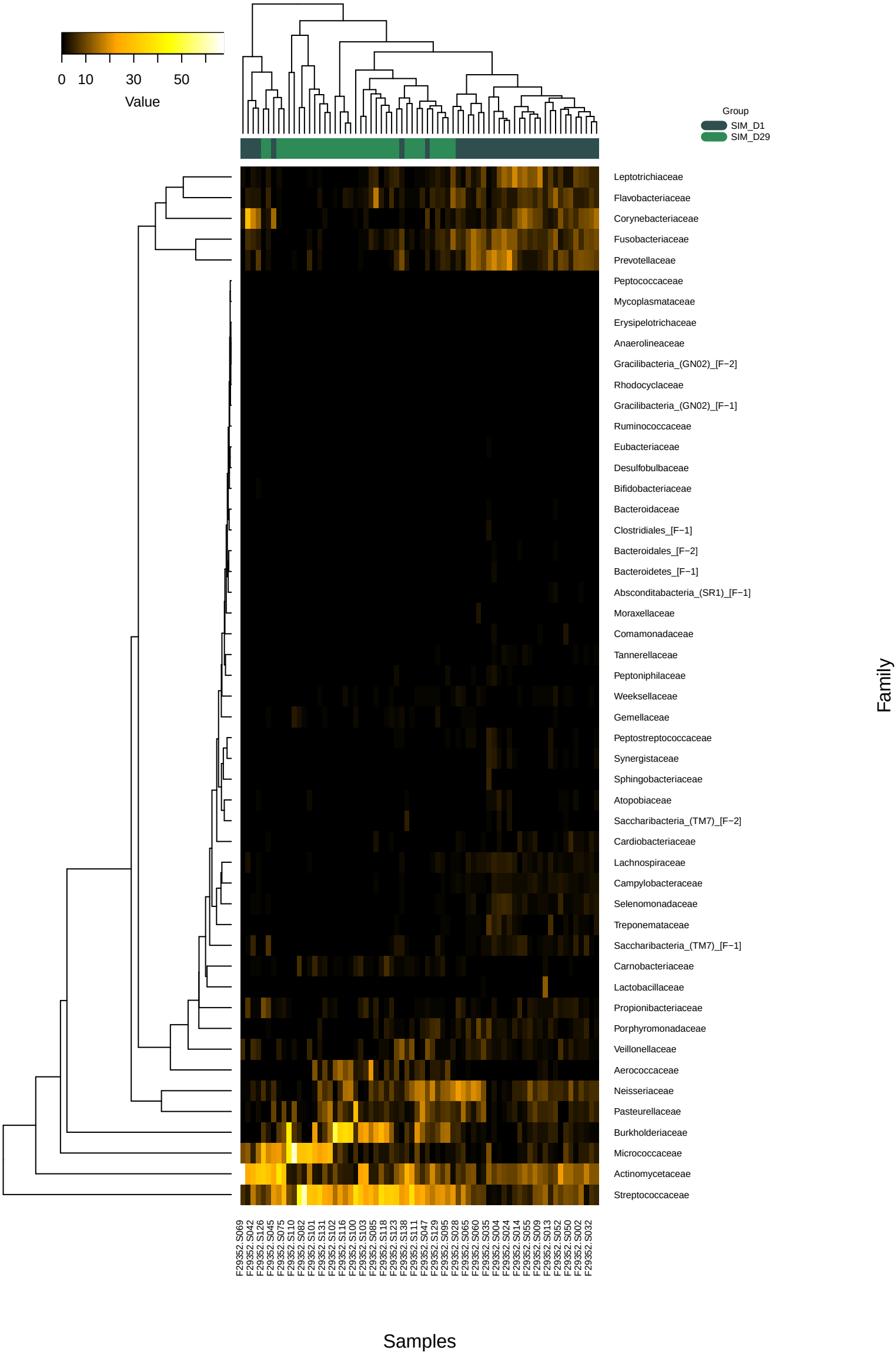
0
10
30
50
Value
Group
SIM_D1
SIM_D29
Leptotrichiaceae
Flavobacteriaceae
Corynebacteriaceae
Fusobacteriaceae
Prevotellaceae
Peptococcaceae
Mycoplasmataceae
Erysipelotrichaceae
Anaerolineaceae
Gracilibacteria_(GN02)_[F−2]
Rhodocyclaceae
Gracilibacteria_(GN02)_[F−1]
Ruminococcaceae
Eubacteriaceae
Desulfobulbaceae
Bifidobacteriaceae
Bacteroidaceae
Clostridiales_[F−1]
Bacteroidales_[F−2]
Bacteroidetes_[F−1]
Absconditabacteria_(SR1)_[F−1]
Moraxellaceae
Comamonadaceae
Tannerellaceae
Peptoniphilaceae
Weeksellaceae
Gemellaceae
Peptostreptococcaceae
Synergistaceae
Sphingobacteriaceae
Atopobiaceae
Saccharibacteria_(TM7)_[F−2]
Cardiobacteriaceae
Lachnospiraceae
Campylobacteraceae
Selenomonadaceae
Treponemataceae
Saccharibacteria_(TM7)_[F−1]
Carnobacteriaceae
Lactobacillaceae
Propionibacteriaceae
Porphyromonadaceae
Veillonellaceae
Aerococcaceae
Neisseriaceae
Pasteurellaceae
Burkholderiaceae
Micrococcaceae
Actinomycetaceae
Streptococcaceae
Family
F29352.S069
F29352.S042
F29352.S126
F29352.S045
F29352.S075
F29352.S110
F29352.S082
F29352.S101
F29352.S131
F29352.S102
F29352.S116
F29352.S100
F29352.S103
F29352.S085
F29352.S118
F29352.S123
F29352.S138
F29352.S111
F29352.S047
F29352.S129
F29352.S095
F29352.S028
F29352.S065
F29352.S060
F29352.S035
F29352.S004
F29352.S024
F29352.S014
F29352.S055
F29352.S009
F29352.S013
F29352.S052
F29352.S050
F29352.S002
F29352.S032
Samples