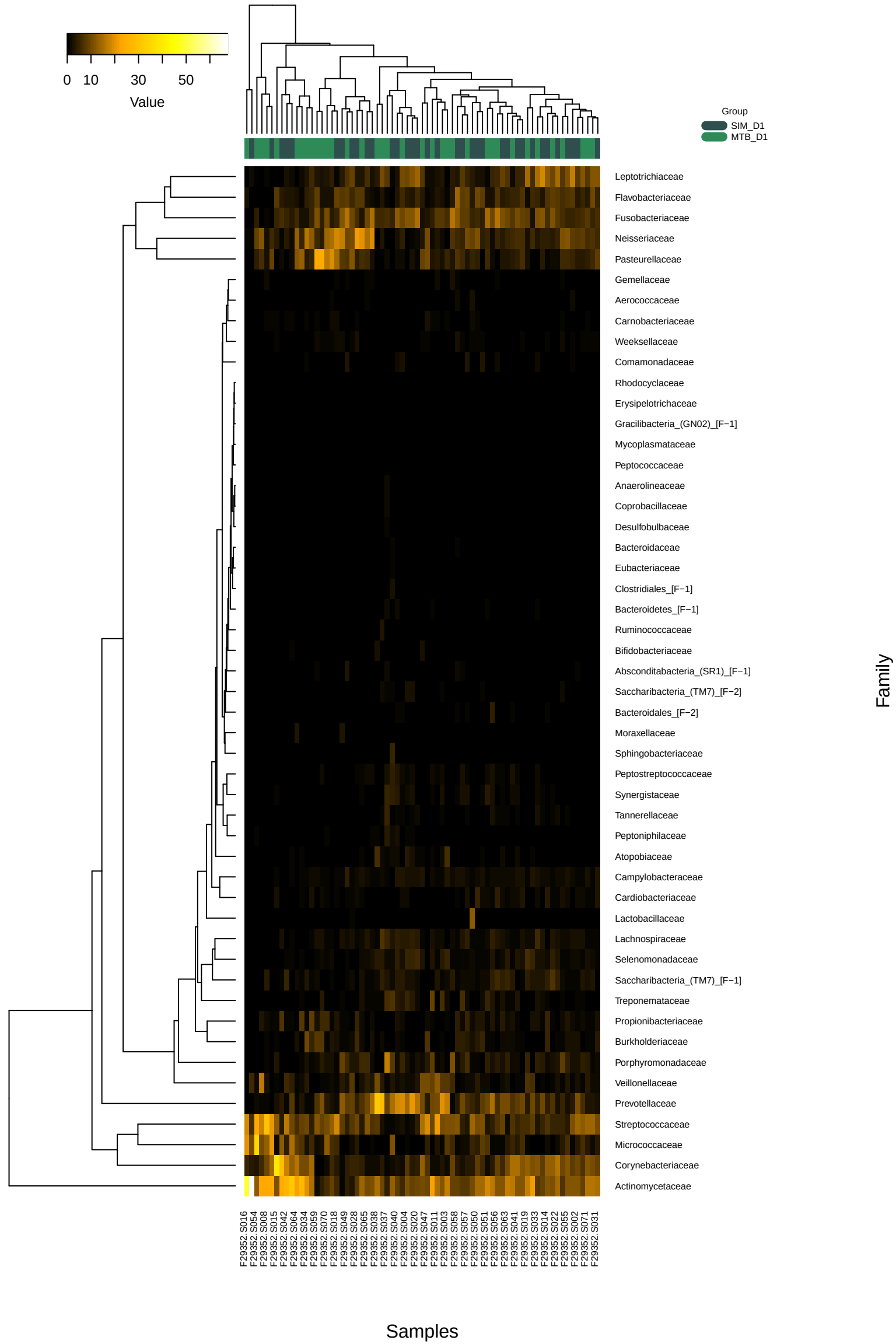
0 10 30 50
Value
Group
SIM_D1
MTB_D1
Leptotrichiaceae
Flavobacteriaceae
Fusobacteriaceae
Neisseriaceae
Pasteurellaceae
Gemellaceae
Aerococcaceae
Carnobacteriaceae
Weeksellaceae
Comamonadaceae
Rhodocyclaceae
Erysipelotrichaceae
Gracilibacteria_(GN02)_[F−1]
Mycoplasmataceae
Peptococcaceae
Anaerolineaceae
Coprobacillaceae
Desulfobulbaceae
Bacteroidaceae
Eubacteriaceae
Clostridiales_[F−1]
Bacteroidetes_[F−1]
Ruminococcaceae
Bifidobacteriaceae
Absconditabacteria_(SR1)_[F−1]
Saccharibacteria_(TM7)_[F−2]
Bacteroidales_[F−2]
Moraxellaceae
Sphingobacteriaceae
Peptostreptococcaceae
Synergistaceae
Tannerellaceae
Peptoniphilaceae
Atopobiaceae
Campylobacteraceae
Cardiobacteriaceae
Lactobacillaceae
Lachnospiraceae
Selenomonadaceae
Saccharibacteria_(TM7)_[F−1]
Treponemataceae
Propionibacteriaceae
Burkholderiaceae
Porphyromonadaceae
Veillonellaceae
Prevotellaceae
Streptococcaceae
Micrococcaceae
Corynebacteriaceae
Actinomycetaceae
F29352.S016
F29352.S054
F29352.S008
F29352.S015
F29352.S042
F29352.S064
F29352.S034
F29352.S059
F29352.S070
F29352.S018
F29352.S049
F29352.S028
F29352.S065
F29352.S038
F29352.S037
F29352.S040
F29352.S004
F29352.S020
F29352.S047
F29352.S011
F29352.S003
F29352.S058
F29352.S057
F29352.S050
F29352.S051
F29352.S056
F29352.S063
F29352.S041
F29352.S019
F29352.S033
F29352.S014
F29352.S022
F29352.S055
F29352.S002
F29352.S071
F29352.S031
Samples
Family