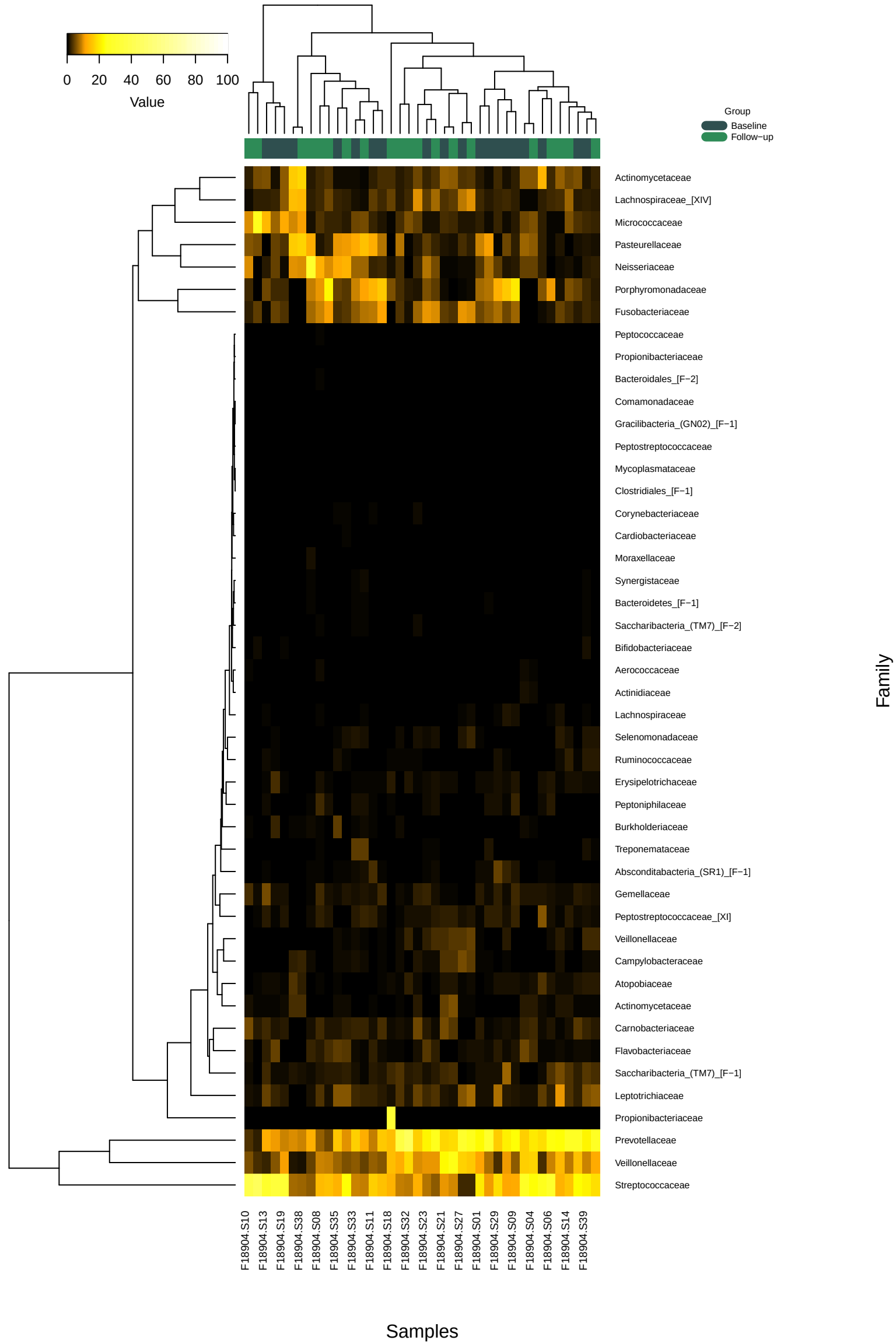
0
20
40
60
80
100
Value
Group
Baseline
Follow−up
Actinomycetaceae
Lachnospiraceae_[XIV]
Micrococcaceae
Pasteurellaceae
Neisseriaceae
Porphyromonadaceae
Fusobacteriaceae
Peptococcaceae
Propionibacteriaceae
Bacteroidales_[F−2]
Comamonadaceae
Gracilibacteria_(GN02)_[F−1]
Peptostreptococcaceae
Mycoplasmataceae
Clostridiales_[F−1]
Corynebacteriaceae
Cardiobacteriaceae
Moraxellaceae
Synergistaceae
Bacteroidetes_[F−1]
Saccharibacteria_(TM7)_[F−2]
Bifidobacteriaceae
Aerococcaceae
Actinidiaceae
Lachnospiraceae
Selenomonadaceae
Ruminococcaceae
Erysipelotrichaceae
Peptoniphilaceae
Burkholderiaceae
Treponemataceae
Absconditabacteria_(SR1)_[F−1]
Gemellaceae
Peptostreptococcaceae_[XI]
Veillonellaceae
Campylobacteraceae
Atopobiaceae
Actinomycetaceae
Carnobacteriaceae
Flavobacteriaceae
Saccharibacteria_(TM7)_[F−1]
Leptotrichiaceae
Propionibacteriaceae
Prevotellaceae
Veillonellaceae
Streptococcaceae
F18904.S10
F18904.S13
F18904.S19
F18904.S38
F18904.S08
F18904.S35
F18904.S33
F18904.S11
F18904.S18
F18904.S32
F18904.S23
F18904.S21
F18904.S27
F18904.S01
F18904.S29
F18904.S09
F18904.S04
F18904.S06
F18904.S14
F18904.S39
Samples
Family