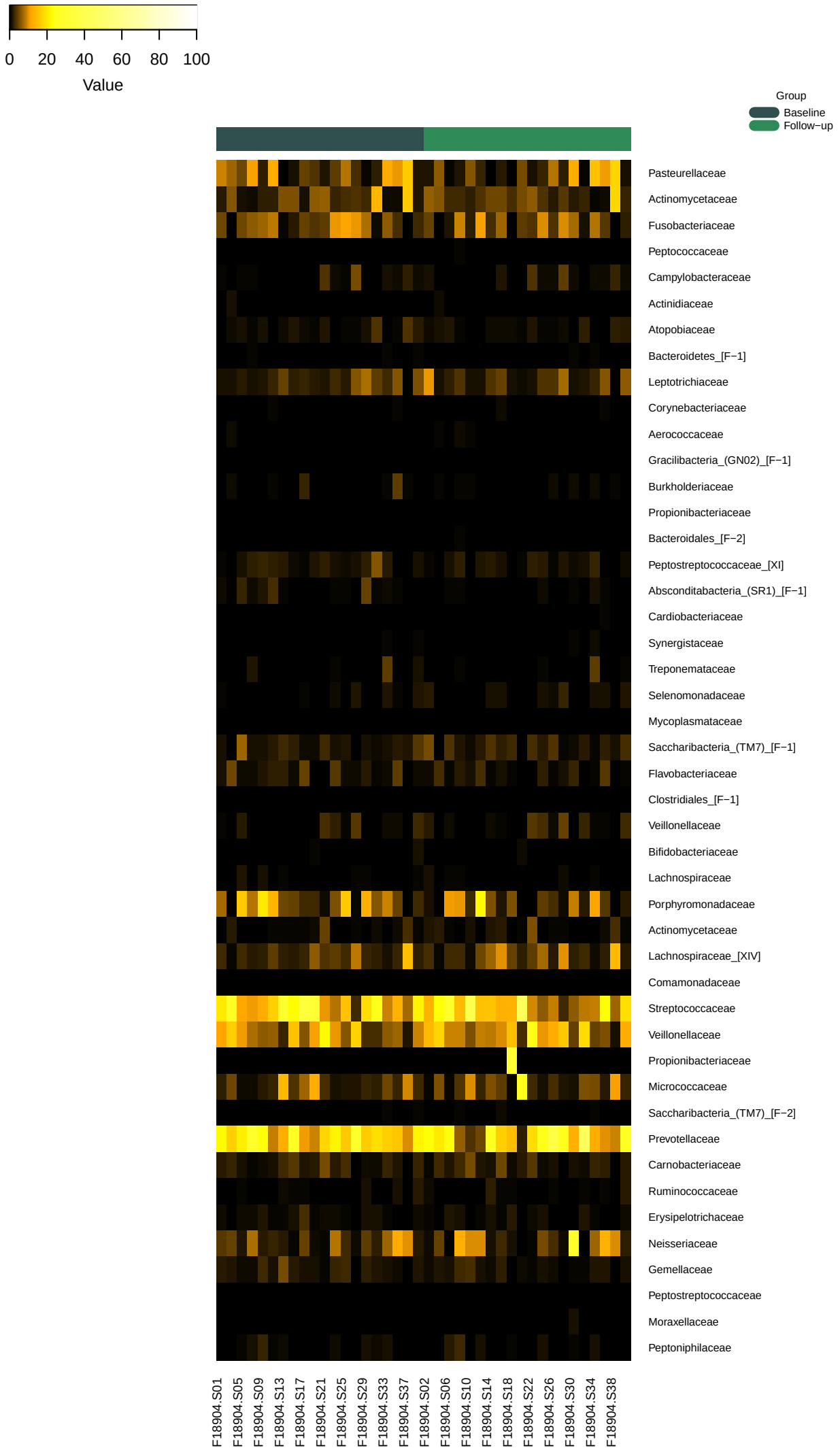
0
20
40
60
80
100
Value
Group
Baseline
Follow−up
Pasteurellaceae
Actinomycetaceae
Fusobacteriaceae
Peptococcaceae
Campylobacteraceae
Actinidiaceae
Atopobiaceae
Bacteroidetes_[F−1]
Leptotrichiaceae
Corynebacteriaceae
Aerococcaceae
Gracilibacteria_(GN02)_[F−1]
Burkholderiaceae
Propionibacteriaceae
Bacteroidales_[F−2]
Peptostreptococcaceae_[XI]
Absconditabacteria_(SR1)_[F−1]
Cardiobacteriaceae
Synergistaceae
Treponemataceae
Selenomonadaceae
Mycoplasmataceae
Saccharibacteria_(TM7)_[F−1]
Flavobacteriaceae
Clostridiales_[F−1]
Veillonellaceae
Bifidobacteriaceae
Lachnospiraceae
Porphyromonadaceae
Actinomycetaceae
Lachnospiraceae_[XIV]
Comamonadaceae
Streptococcaceae
Veillonellaceae
Propionibacteriaceae
Micrococcaceae
Saccharibacteria_(TM7)_[F−2]
Prevotellaceae
Carnobacteriaceae
Ruminococcaceae
Erysipelotrichaceae
Neisseriaceae
Gemellaceae
Peptostreptococcaceae
Moraxellaceae
Peptoniphilaceae
Family
F18904.S01
F18904.S05
F18904.S09
F18904.S13
F18904.S17
F18904.S21
F18904.S25
F18904.S29
F18904.S33
F18904.S37
F18904.S02
F18904.S06
F18904.S10
F18904.S14
F18904.S18
F18904.S22
F18904.S26
F18904.S30
F18904.S34
F18904.S38
Samples