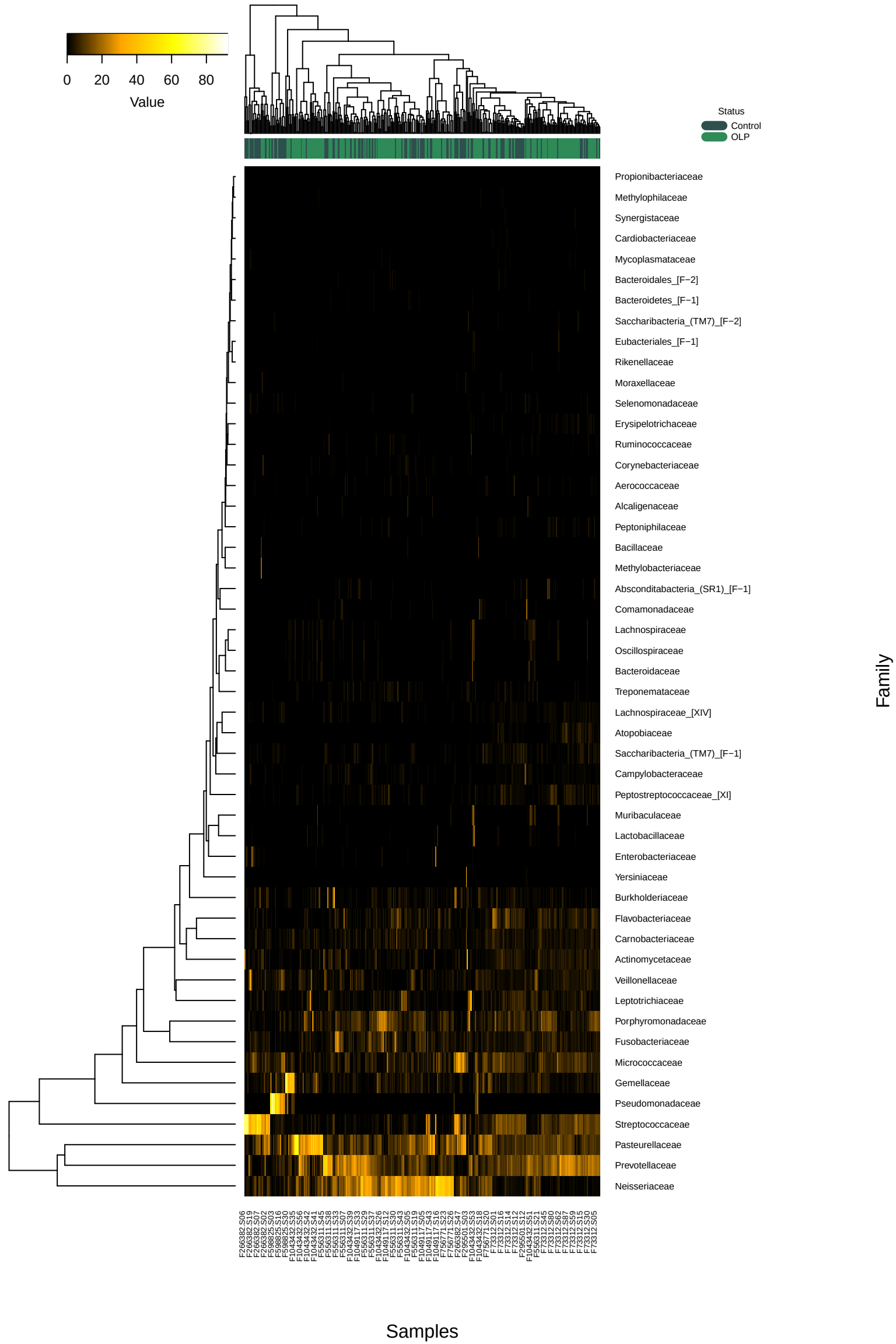
0
20
40
60
80
Value
Status
Control
OLP
Propionibacteriaceae
Methylophilaceae
Synergistaceae
Cardiobacteriaceae
Mycoplasmataceae
Bacteroidales_[F−2]
Bacteroidetes_[F−1]
Saccharibacteria_(TM7)_[F−2]
Eubacteriales_[F−1]
Rikenellaceae
Moraxellaceae
Selenomonadaceae
Erysipelotrichaceae
Ruminococcaceae
Corynebacteriaceae
Aerococcaceae
Alcaligenaceae
Peptoniphilaceae
Bacillaceae
Methylobacteriaceae
Absconditabacteria_(SR1)_[F−1]
Comamonadaceae
Lachnospiraceae
Oscillospiraceae
Bacteroidaceae
Treponemataceae
Lachnospiraceae_[XIV]
Atopobiaceae
Saccharibacteria_(TM7)_[F−1]
Campylobacteraceae
Peptostreptococcaceae_[XI]
Muribaculaceae
Lactobacillaceae
Enterobacteriaceae
Yersiniaceae
Burkholderiaceae
Flavobacteriaceae
Carnobacteriaceae
Actinomycetaceae
Veillonellaceae
Leptotrichiaceae
Porphyromonadaceae
Fusobacteriaceae
Micrococcaceae
Gemellaceae
Pseudomonadaceae
Streptococcaceae
Pasteurellaceae
Prevotellaceae
Neisseriaceae
Family
F266382.S06
F266382.S19
F266382.S07
F266382.S02
F598825.S03
F598825.S16
F598825.S10
F1043432.S35
F1043432.S56
F1043432.S42
F1043432.S41
F556311.S45
F556311.S38
F556311.S03
F556311.S07
F1043432.S39
F1049117.S33
F556311.S29
F556311.S37
F1043432.S26
F1049117.S12
F556311.S30
F556311.S43
F1043432.S05
F556311.S19
F1049117.S05
F1049117.S43
F1049117.S16
F756771.S23
F756771.S26
F266382.S47
F266382.S53
F1043432.S18
F756771.S20
F73312.S01
F73312.S16
F73312.S14
F73312.S12
F266382.S55
F1043432.S51
F556311.S21
F73312.S45
F73312.S80
F73312.S62
F73312.S87
F73312.S99
F73312.S15
F73312.S10
F73312.S05
Samples