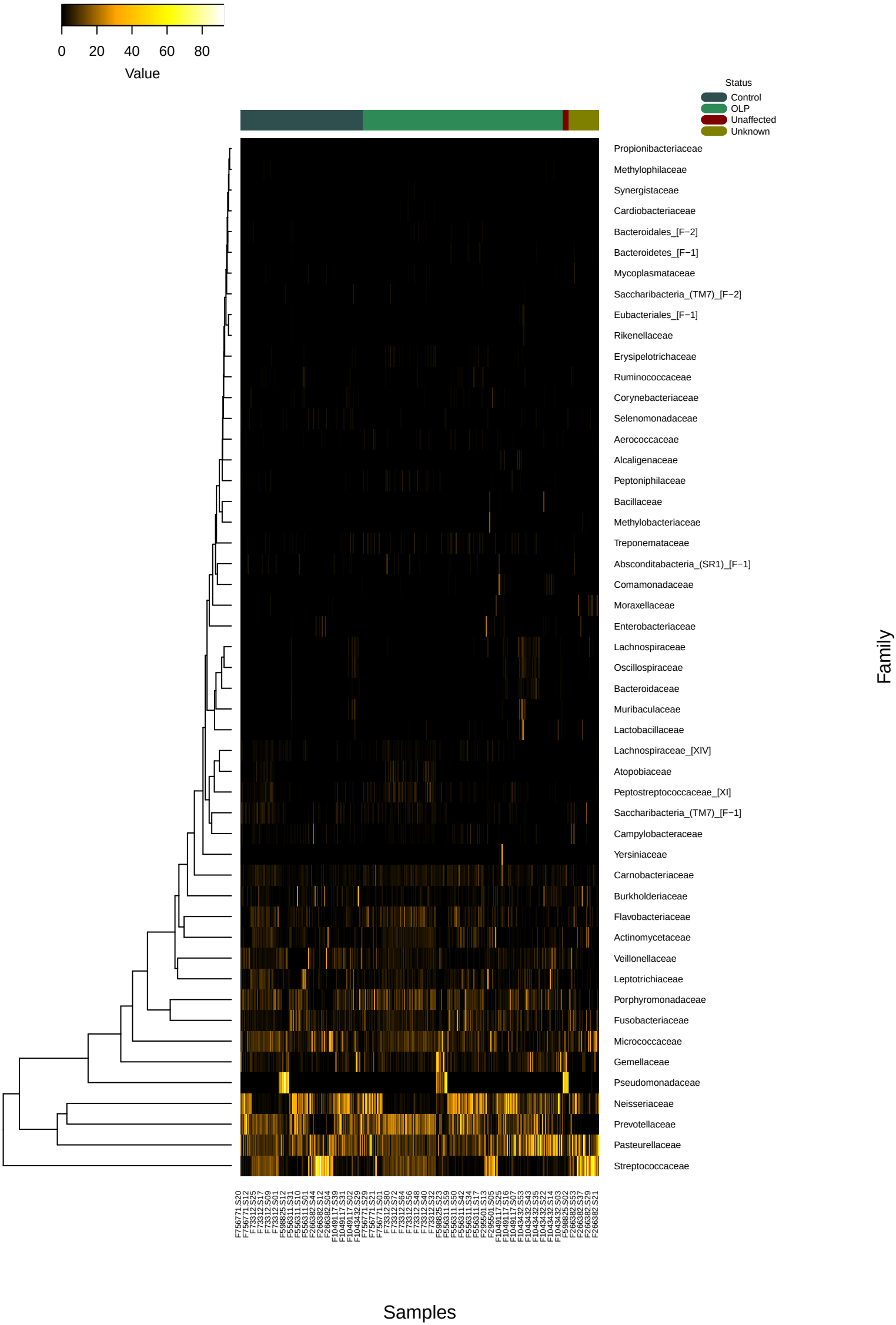
0
20
40
60
80
Value
Status
Control
OLP
Unaffected
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Propionibacteriaceae
Methylophilaceae
Synergistaceae
Cardiobacteriaceae
Bacteroidales_[F−2]
Bacteroidetes_[F−1]
Mycoplasmataceae
Saccharibacteria_(TM7)_[F−2]
Eubacteriales_[F−1]
Rikenellaceae
Erysipelotrichaceae
Ruminococcaceae
Corynebacteriaceae
Selenomonadaceae
Aerococcaceae
Alcaligenaceae
Peptoniphilaceae
Bacillaceae
Methylobacteriaceae
Treponemataceae
Absconditabacteria_(SR1)_[F−1]
Comamonadaceae
Moraxellaceae
Enterobacteriaceae
Lachnospiraceae
Oscillospiraceae
Bacteroidaceae
Muribaculaceae
Lactobacillaceae
Lachnospiraceae_[XIV]
Atopobiaceae
Peptostreptococcaceae_[XI]
Saccharibacteria_(TM7)_[F−1]
Campylobacteraceae
Yersiniaceae
Carnobacteriaceae
Burkholderiaceae
Flavobacteriaceae
Actinomycetaceae
Veillonellaceae
Leptotrichiaceae
Porphyromonadaceae
Fusobacteriaceae
Micrococcaceae
Gemellaceae
Pseudomonadaceae
Neisseriaceae
Prevotellaceae
Pasteurellaceae
Streptococcaceae
Family
F56771.S20
F56771.S12
F73312.S25
F73312.S17
F73312.S09
F73312.S01
F59825.S12
F556311.S11
F556311.S10
F556311.S01
F266382.S44
F266382.S12
F266382.S04
F1049117.S39
F1049117.S31
F1049117.S02
F1043432.S29
F756771.S29
F756771.S21
F756771.S01
F73312.S80
F73312.S72
F73312.S64
F73312.S56
F73312.S48
F73312.S40
F73312.S32
F59825.S23
F556311.S59
F556311.S50
F556311.S42
F556311.S34
F556311.S17
F295501.S13
F295501.S05
F1049117.S25
F1049117.S16
F1049117.S07
F1043432.S43
F1043432.S35
F1043432.S22
F1043432.S14
F1043432.S03
F59825.S02
F266382.S53
F266382.S37
F266382.S29
F266382.S21
Samples