0
20
40
60
80
Value
Status
Control
OLP
Unaffected
Unknown
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
F266382.S14
F266382.S18
F266382.S07
F295501.S07
F266382.S51
F266382.S24
F104432.S43
F266382.S45
F266382.S46
F556311.S20
F104432.S57
F598825.S09
F598825.S11
F598825.S04
F1049117.S31
F1049117.S20
F266382.S40
F266382.S28
F266382.S23
F1049117.S34
F104432.S31
F556311.S40
F556311.S51
F556311.S14
F756771.S08
F756771.S17
F556311.S24
F556771.S33
F556771.S34
F556311.S58
F104432.S22
F556311.S42
F104432.S07
F104432.S42
F1049117.S01
F598825.S26
F266382.S46
F73312.S64
F73312.S67
F73312.S28
F104432.S15
F73312.S03
F73312.S75
F73312.S73
F73312.S74
F73312.S15
F73312.S33
F73312.S34
Samples