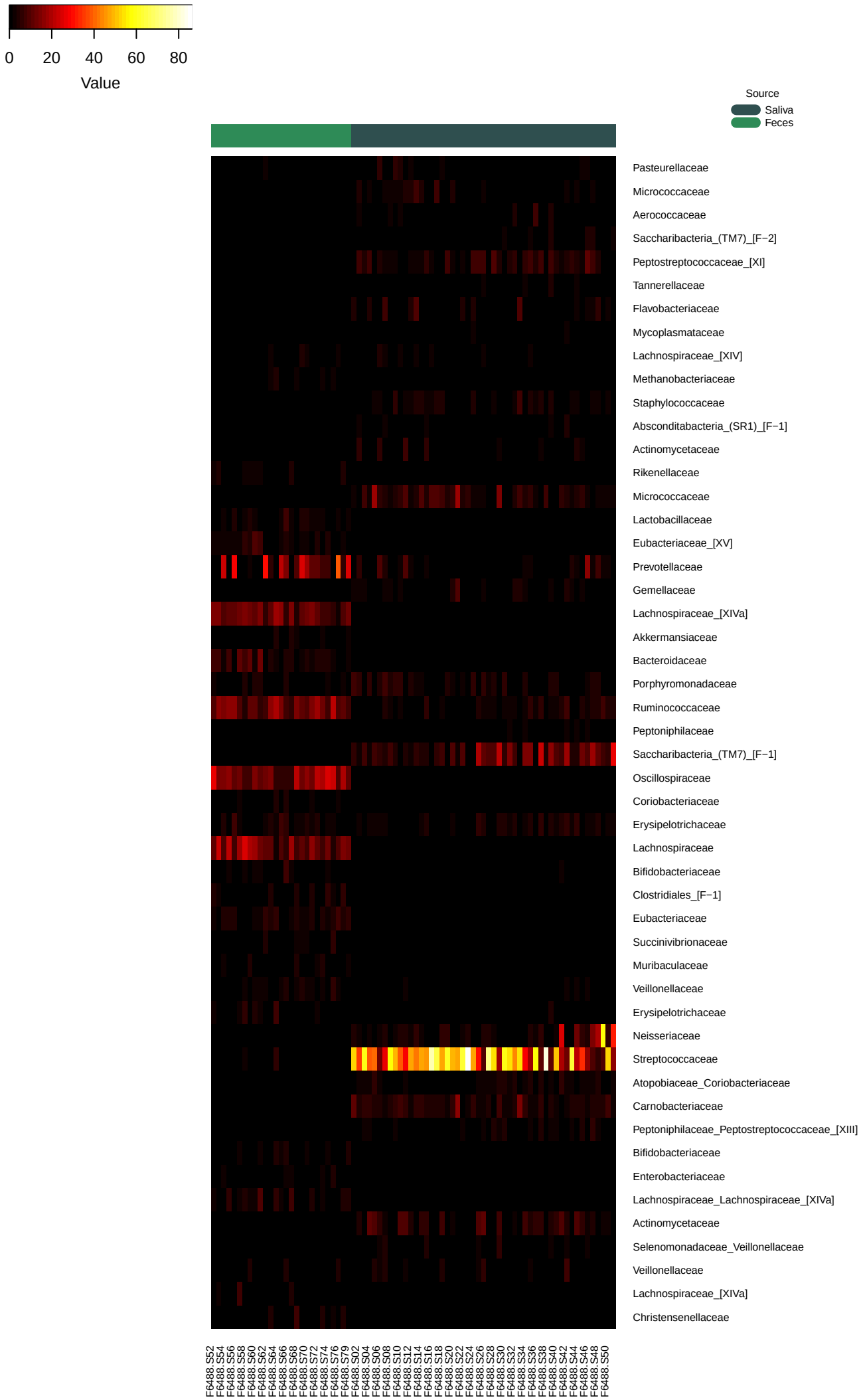
0
20
40
60
80
Value
Source
Saliva
Feces
Pasteurellaceae
Micrococcaceae
Aerococcaceae
Saccharibacteria_(TM7)_[F−2]
Peptostreptococcaceae_[XI]
Tannerellaceae
Flavobacteriaceae
Mycoplasmataceae
Lachnospiraceae_[XIV]
Methanobacteriaceae
Staphylococcaceae
Absconditabacteria_(SR1)_[F−1]
Actinomycetaceae
Rikenellaceae
Micrococcaceae
Lactobacillaceae
Eubacteriaceae_[XV]
Prevotellaceae
Gemellaceae
Lachnospiraceae_[XIVa]
Akkermansiaceae
Bacteroidaceae
Porphyromonadaceae
Ruminococcaceae
Peptoniphilaceae
Saccharibacteria_(TM7)_[F−1]
Oscillospiraceae
Coriobacteriaceae
Erysipelotrichaceae
Lachnospiraceae
Bifidobacteriaceae
Clostridiales_[F−1]
Eubacteriaceae
Succinivibrionaceae
Muribaculaceae
Veillonellaceae
Erysipelotrichaceae
Neisseriaceae
Streptococcaceae
Atopobiaceae_Coriobacteriaceae
Carnobacteriaceae
Peptoniphilaceae_Peptostreptococcaceae_[XIII]
Bifidobacteriaceae
Enterobacteriaceae
Lachnospiraceae_Lachnospiraceae_[XIVa]
Actinomycetaceae
Selenomonadaceae_Veillonellaceae
Veillonellaceae
Lachnospiraceae_[XIVa]
Christensenellaceae
Family
F6488.S52
F6488.S54
F6488.S56
F6488.S58
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F6488.S02
F6488.S04
F6488.S06
F6488.S08
F6488.S10
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F6488.S38
F6488.S40
F6488.S42
F6488.S44
F6488.S46
F6488.S48
F6488.S50
Samples