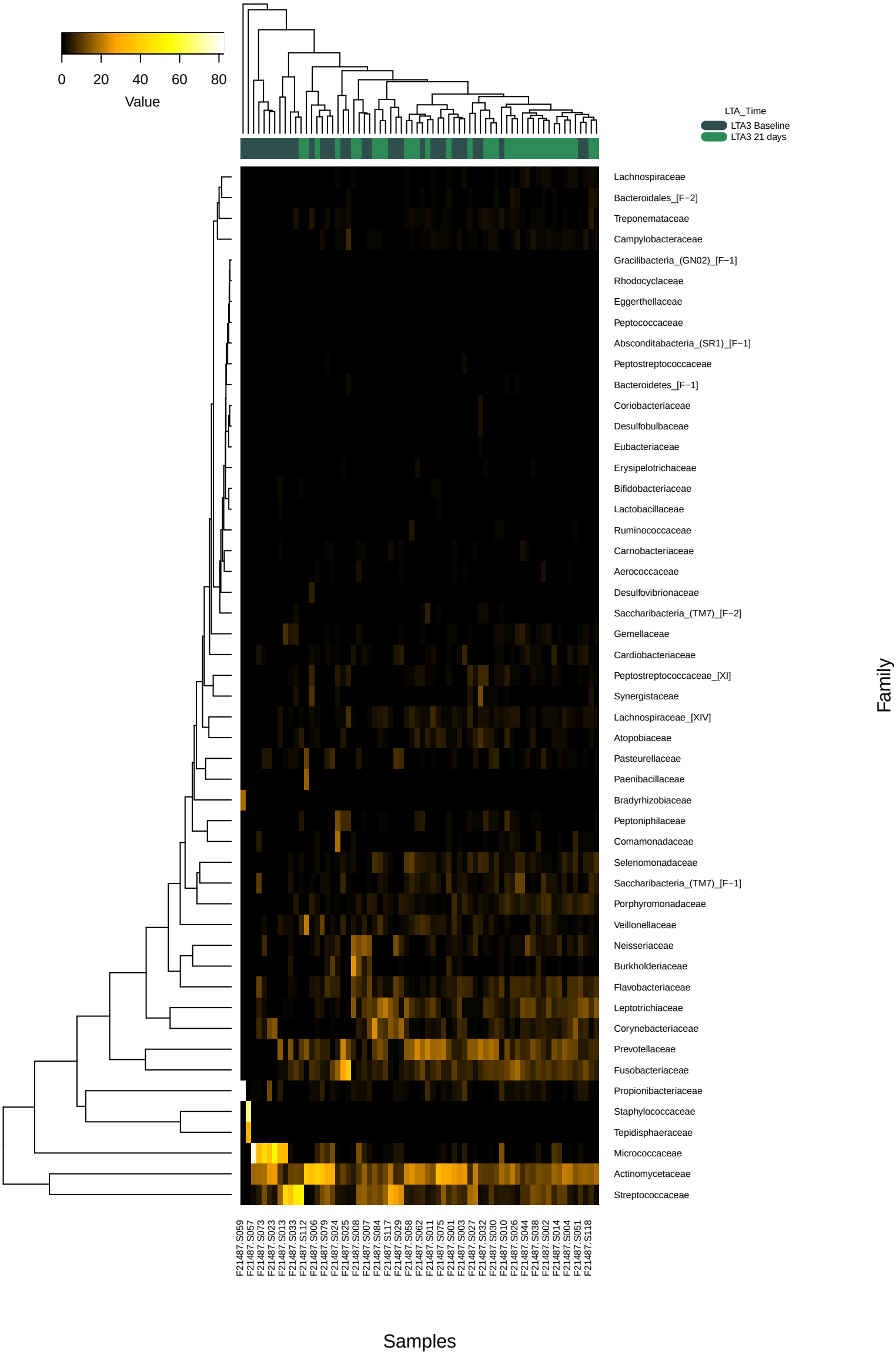
0
20
40
60
80
Value
LTA_Time
LTA3 Baseline
LTA3 21 days
Lachnospiraceae
Bacteroidales_[F−2]
Treponemataceae
Campylobacteraceae
Gracilibacteria_(GN02)_[F−1]
Rhodocyclaceae
Eggerthellaceae
Peptococcaceae
Absconditabacteria_(SR1)_[F−1]
Peptostreptococcaceae
Bacteroidetes_[F−1]
Coriobacteriaceae
Desulfobulbaceae
Eubacteriaceae
Erysipelotrichaceae
Bifidobacteriaceae
Lactobacillaceae
Ruminococcaceae
Carnobacteriaceae
Aerococcaceae
Desulfovibrionaceae
Saccharibacteria_(TM7)_[F−2]
Gemellaceae
Cardiobacteriaceae
Peptostreptococcaceae_[XI]
Synergistaceae
Lachnospiraceae_[XIV]
Atopobiaceae
Pasteurellaceae
Paenibacillaceae
Bradyrhizobiaceae
Peptoniphilaceae
Comamonadaceae
Selenomonadaceae
Saccharibacteria_(TM7)_[F−1]
Porphyromonadaceae
Veillonellaceae
Neisseriaceae
Burkholderiaceae
Flavobacteriaceae
Leptotrichiaceae
Corynebacteriaceae
Prevotellaceae
Fusobacteriaceae
Propionibacteriaceae
Staphylococcaceae
Tepidisphaeraceae
Micrococcaceae
Actinomycetaceae
Streptococcaceae
Family
F21487.S059
F21487.S057
F21487.S073
F21487.S023
F21487.S013
F21487.S033
F21487.S112
F21487.S006
F21487.S079
F21487.S024
F21487.S025
F21487.S008
F21487.S007
F21487.S084
F21487.S117
F21487.S029
F21487.S058
F21487.S062
F21487.S011
F21487.S075
F21487.S001
F21487.S003
F21487.S027
F21487.S032
F21487.S030
F21487.S010
F21487.S026
F21487.S044
F21487.S038
F21487.S002
F21487.S014
F21487.S004
F21487.S051
F21487.S118
Samples