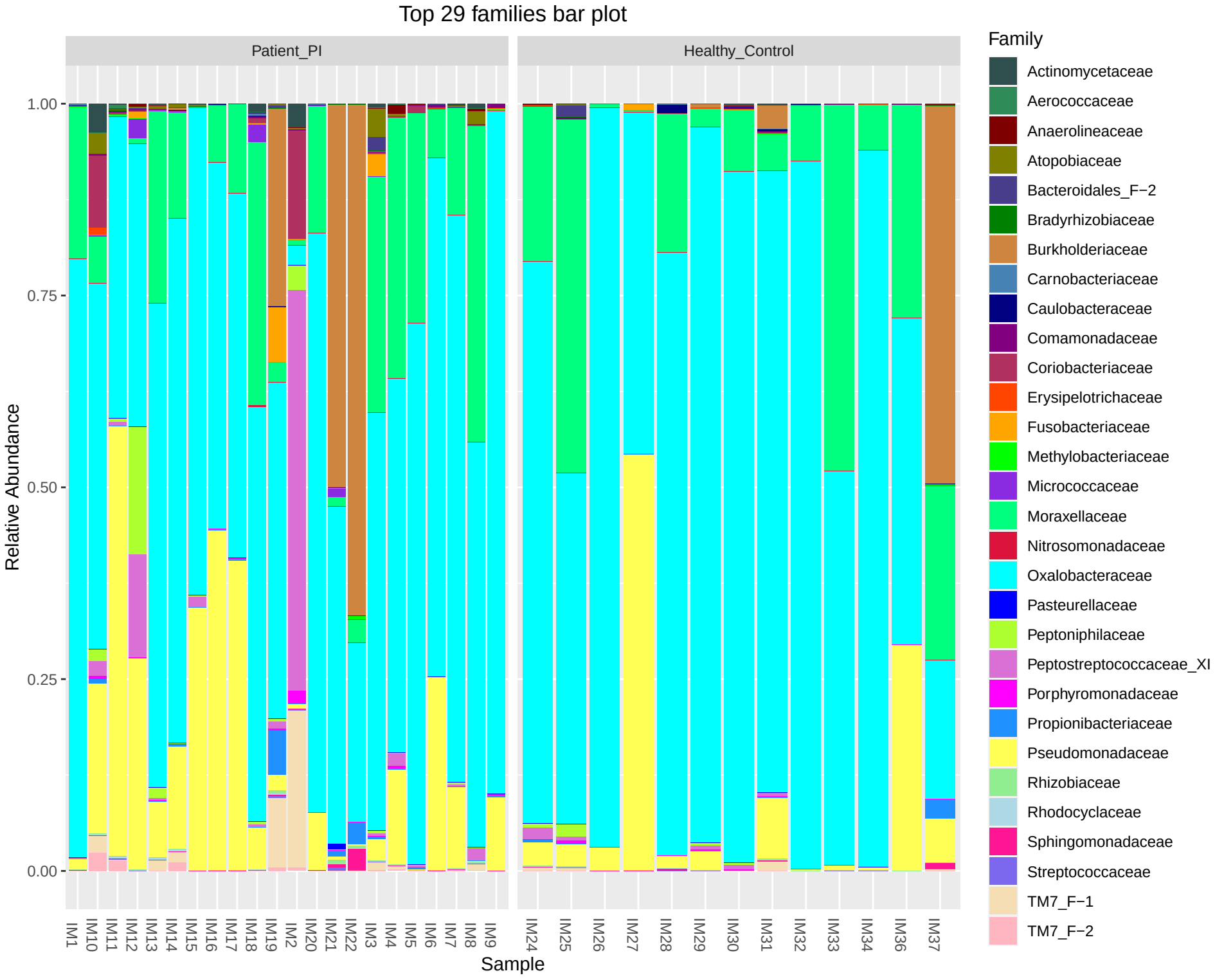
Top 29 families bar plot
Patient_PI
Healthy_Control
Relative Abundance
1.00
0.75
0.50
0.25
0.00
IM1
IM10
IM11
IM12
IM13
IM14
IM15
IM16
IM17
IM18
IM19
IM2
IM20
IM21
IM22
IM3
IM4
IM5
IM6
IM7
IM8
IM9
IM24
IM25
IM26
IM27
IM28
IM29
IM30
IM31
IM32
IM33
IM34
IM36
IM37
Sample
Family
Actinomycetaceae
Aerococcaceae
Anaerolineaceae
Atopobiaceae
Bacteroidales_F−2
Bradyrhizobiaceae
Burkholderiaceae
Carnobacteriaceae
Caulobacteraceae
Comamonadaceae
Coriobacteriaceae
Erysipelotrichaceae
Fusobacteriaceae
Methylobacteriaceae
Micrococcaceae
Moraxellaceae
Nitrosomonadaceae
Oxalobacteraceae
Pasteurellaceae
Peptoniphilaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Propionibacteriaceae
Pseudomonadaceae
Rhizobiaceae
Rhodocyclaceae
Sphingomonadaceae
Streptococcaceae
TM7_F−1
TM7_F−2