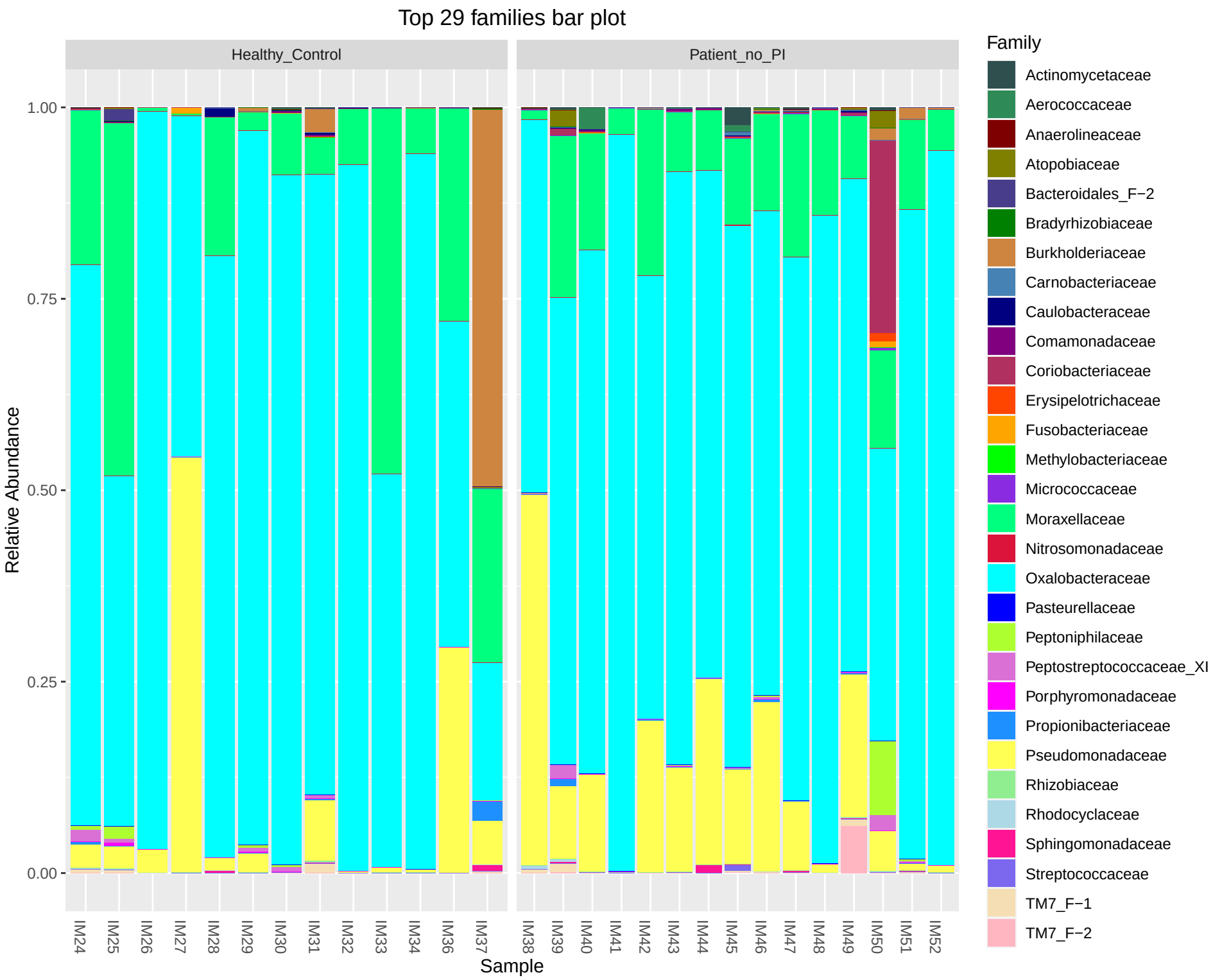
Top 29 families bar plot
Healthy_Control
Patient_no_PI
Relative Abundance
1.00
0.75
0.50
0.25
0.00
IM24
IM25
IM26
IM27
IM28
IM29
IM30
IM31
IM32
IM33
IM34
IM36
IM37
IM38
IM39
IM40
IM41
IM42
IM43
IM44
IM45
IM46
IM47
IM48
IM49
IM50
IM51
IM52
Sample
Family
Actinomycetaceae
Aerococcaceae
Anaerolineaceae
Atopobiaceae
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