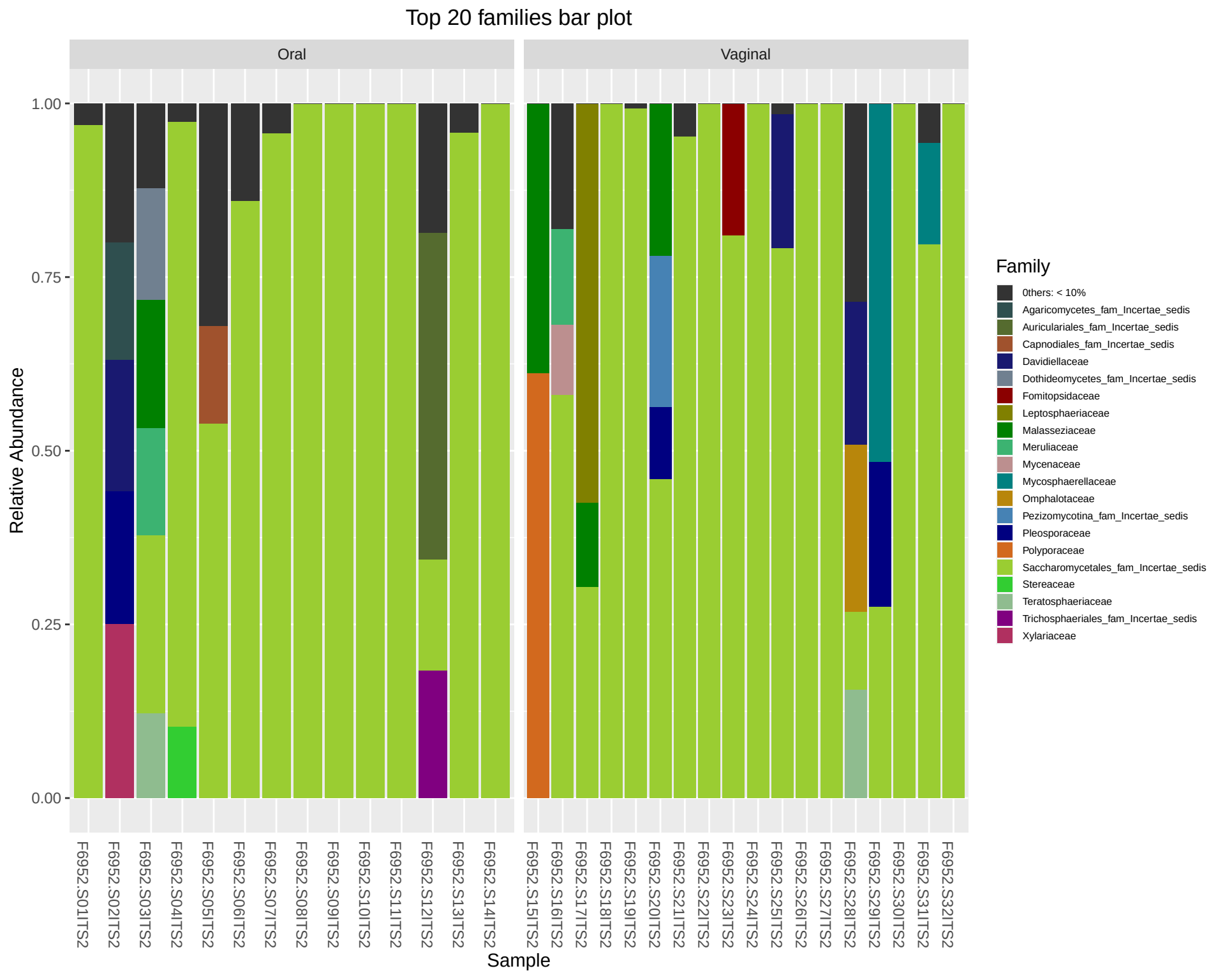
Top 20 families bar plot
Oral
Vaginal
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F6952.S01ITS2
F6952.S02ITS2
F6952.S03ITS2
F6952.S04ITS2
F6952.S05ITS2
F6952.S06ITS2
F6952.S07ITS2
F6952.S08ITS2
F6952.S09ITS2
F6952.S10ITS2
F6952.S11ITS2
F6952.S12ITS2
F6952.S13ITS2
F6952.S14ITS2
F6952.S15ITS2
F6952.S16ITS2
F6952.S17ITS2
F6952.S18ITS2
F6952.S19ITS2
F6952.S20ITS2
F6952.S21ITS2
F6952.S22ITS2
F6952.S23ITS2
F6952.S24ITS2
F6952.S25ITS2
F6952.S26ITS2
F6952.S27ITS2
F6952.S28ITS2
F6952.S29ITS2
F6952.S30ITS2
F6952.S31ITS2
F6952.S32ITS2
Sample
Family
0thers: < 10%
Agaricomycetes_fam_Incertae_sedis
Auriculariales_fam_Incertae_sedis
Capnodiales_fam_Incertae_sedis
Davidiellaceae
Dothideomycetes_fam_Incertae_sedis
Fomitopsidaceae
Leptosphaeriaceae
Malasseziaceae
Meruliaceae
Mycenaceae
Mycosphaerellaceae
Omphalotaceae
Pezizomycotina_fam_Incertae_sedis
Pleosporaceae
Polyporaceae
Saccharomycetales_fam_Incertae_sedis
Stereaceae
Teratosphaeriaceae
Trichosphaeriales_fam_Incertae_sedis
Xylariaceae