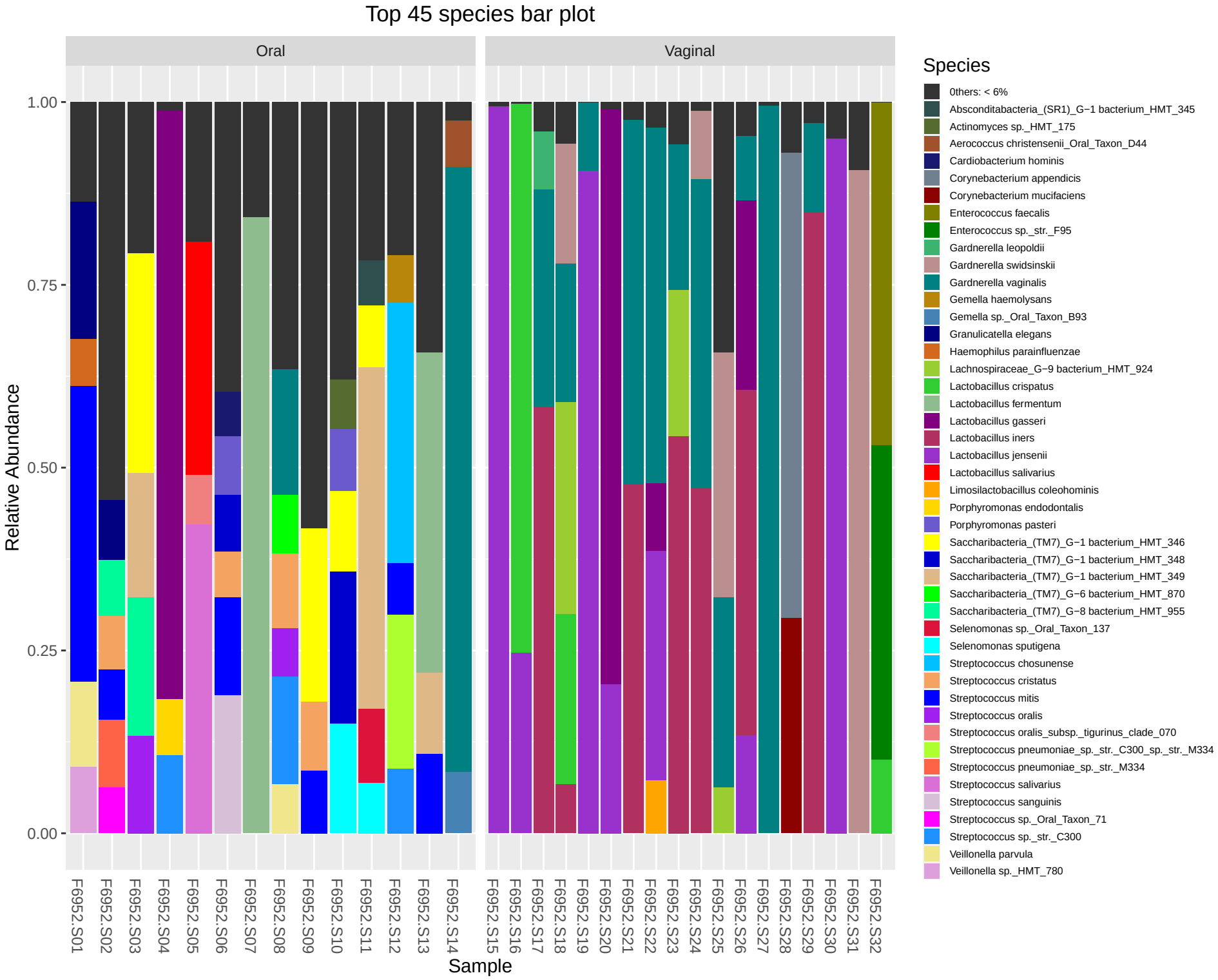
Top 45 species bar plot
Oral
Vaginal
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F6952.S01
F6952.S02
F6952.S03
F6952.S04
F6952.S05
F6952.S06
F6952.S07
F6952.S08
F6952.S09
F6952.S10
F6952.S11
F6952.S12
F6952.S13
F6952.S14
F6952.S15
F6952.S16
F6952.S17
F6952.S18
F6952.S19
F6952.S20
F6952.S21
F6952.S22
F6952.S23
F6952.S24
F6952.S25
F6952.S26
F6952.S27
F6952.S28
F6952.S29
F6952.S30
F6952.S31
F6952.S32
Sample
Species
0thers: < 6%
Absconditabacteria_(SR1)_G−1 bacterium_HMT_345
Actinomyces sp._HMT_175
Aerococcus christensenii_Oral_Taxon_D44
Cardiobacterium hominis
Corynebacterium appendicis
Corynebacterium mucifaciens
Enterococcus faecalis
Enterococcus sp._str._F95
Gardnerella leopoldii
Gardnerella swidsinskii
Gardnerella vaginalis
Gemella haemolysans
Gemella sp._Oral_Taxon_B93
Granulicatella elegans
Haemophilus parainfluenzae
Lachnospiraceae_G−9 bacterium_HMT_924
Lactobacillus crispatus
Lactobacillus fermentum
Lactobacillus gasseri
Lactobacillus iners
Lactobacillus jensenii
Lactobacillus salivarius
Limosilactobacillus coleohominis
Porphyromonas endodontalis
Porphyromonas pasteri
Saccharibacteria_(TM7)_G−1 bacterium_HMT_346
Saccharibacteria_(TM7)_G−1 bacterium_HMT_348
Saccharibacteria_(TM7)_G−1 bacterium_HMT_349
Saccharibacteria_(TM7)_G−6 bacterium_HMT_870
Saccharibacteria_(TM7)_G−8 bacterium_HMT_955
Selenomonas sp._Oral_Taxon_137
Selenomonas sputigena
Streptococcus chosunense
Streptococcus cristatus
Streptococcus mitis
Streptococcus oralis
Streptococcus oralis_subsp._tigurinus_clade_070
Streptococcus pneumoniae_sp._str._C300_sp._str._M334
Streptococcus pneumoniae_sp._str._M334
Streptococcus salivarius
Streptococcus sanguinis
Streptococcus sp._Oral_Taxon_71
Streptococcus sp._str._C300
Veillonella parvula
Veillonella sp._HMT_780