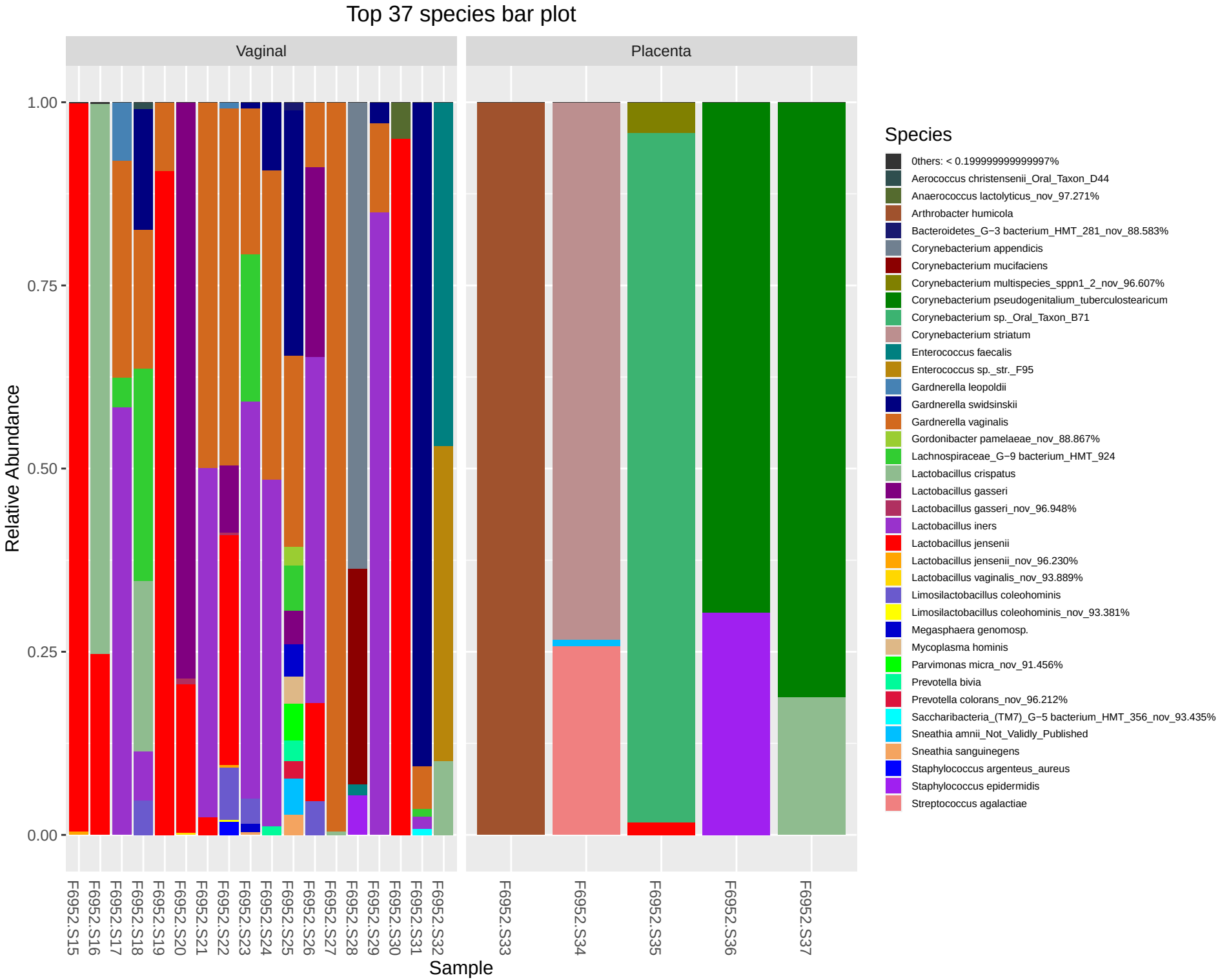
Top 37 species bar plot
Vaginal
Placenta
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F6952.S15
F6952.S16
F6952.S17
F6952.S18
F6952.S19
F6952.S20
F6952.S21
F6952.S22
F6952.S23
F6952.S24
F6952.S25
F6952.S26
F6952.S27
F6952.S28
F6952.S29
F6952.S30
F6952.S31
F6952.S32
F6952.S33
F6952.S34
F6952.S35
F6952.S36
F6952.S37
Sample
Species
0thers: < 0.199999999999997%
Aerococcus christensenii_Oral_Taxon_D44
Anaerococcus lactolyticus_nov_97.271%
Arthrobacter humicola
Bacteroidetes_G−3 bacterium_HMT_281_nov_88.583%
Corynebacterium appendicis
Corynebacterium mucifaciens
Corynebacterium multispecies_sppn1_2_nov_96.607%
Corynebacterium pseudogenitalium_tuberculostearicum
Corynebacterium sp._Oral_Taxon_B71
Corynebacterium striatum
Enterococcus faecalis
Enterococcus sp._str._F95
Gardnerella leopoldii
Gardnerella swidsinskii
Gardnerella vaginalis
Gordonibacter pamelaeae_nov_88.867%
Lachnospiraceae_G−9 bacterium_HMT_924
Lactobacillus crispatus
Lactobacillus gasseri
Lactobacillus gasseri_nov_96.948%
Lactobacillus iners
Lactobacillus jensenii
Lactobacillus jensenii_nov_96.230%
Lactobacillus vaginalis_nov_93.889%
Limosilactobacillus coleohominis
Limosilactobacillus coleohominis_nov_93.381%
Megasphaera genomosp.
Mycoplasma hominis
Parvimonas micra_nov_91.456%
Prevotella bivia
Prevotella colorans_nov_96.212%
Saccharibacteria_(TM7)_G−5 bacterium_HMT_356_nov_93.435%
Sneathia amnii_Not_Validly_Published
Sneathia sanguinegens
Staphylococcus argenteus_aureus
Staphylococcus epidermidis
Streptococcus agalactiae