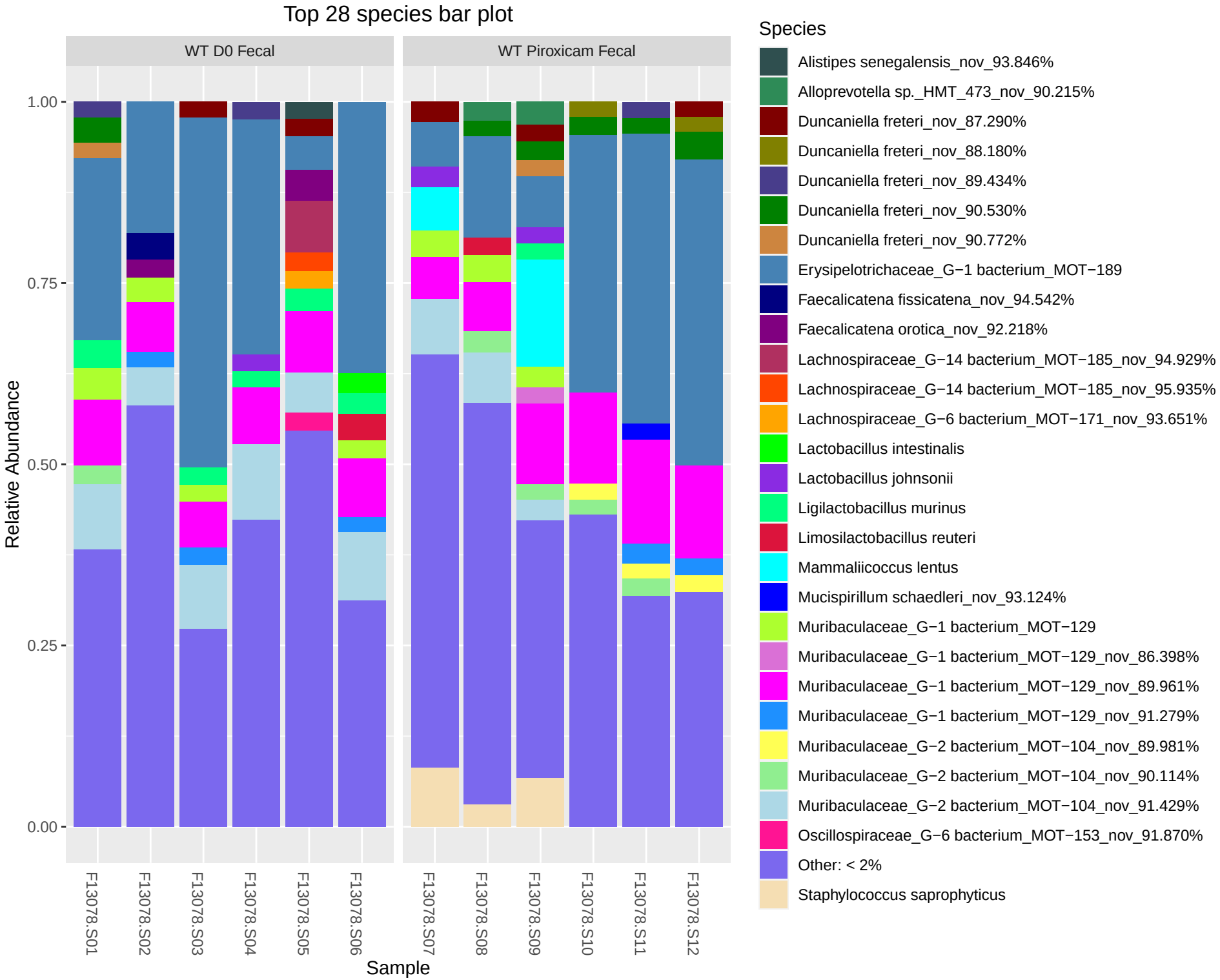
Top 28 species bar plot
WT D0 Fecal
WT Piroxicam Fecal
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F13078.S01
F13078.S02
F13078.S03
F13078.S04
F13078.S05
F13078.S06
F13078.S07
F13078.S08
F13078.S09
F13078.S10
F13078.S11
F13078.S12
Sample
Species
Alistipes senegalensis_nov_93.846%
Alloprevotella sp._HMT_473_nov_90.215%
Duncaniella freteri_nov_87.290%
Duncaniella freteri_nov_88.180%
Duncaniella freteri_nov_89.434%
Duncaniella freteri_nov_90.530%
Duncaniella freteri_nov_90.772%
Erysipelotrichaceae_G−1 bacterium_MOT−189
Faecalicatena fissicatena_nov_94.542%
Faecalicatena orotica_nov_92.218%
Lachnospiraceae_G−14 bacterium_MOT−185_nov_94.929%
Lachnospiraceae_G−14 bacterium_MOT−185_nov_95.935%
Lachnospiraceae_G−6 bacterium_MOT−171_nov_93.651%
Lactobacillus intestinalis
Lactobacillus johnsonii
Ligilactobacillus murinus
Limosilactobacillus reuteri
Mammaliicoccus lentus
Mucispirillum schaedleri_nov_93.124%
Muribaculaceae_G−1 bacterium_MOT−129
Muribaculaceae_G−1 bacterium_MOT−129_nov_86.398%
Muribaculaceae_G−1 bacterium_MOT−129_nov_89.961%
Muribaculaceae_G−1 bacterium_MOT−129_nov_91.279%
Muribaculaceae_G−2 bacterium_MOT−104_nov_89.981%
Muribaculaceae_G−2 bacterium_MOT−104_nov_90.114%
Muribaculaceae_G−2 bacterium_MOT−104_nov_91.429%
Oscillospiraceae_G−6 bacterium_MOT−153_nov_91.870%
Other: < 2%
Staphylococcus saprophyticus