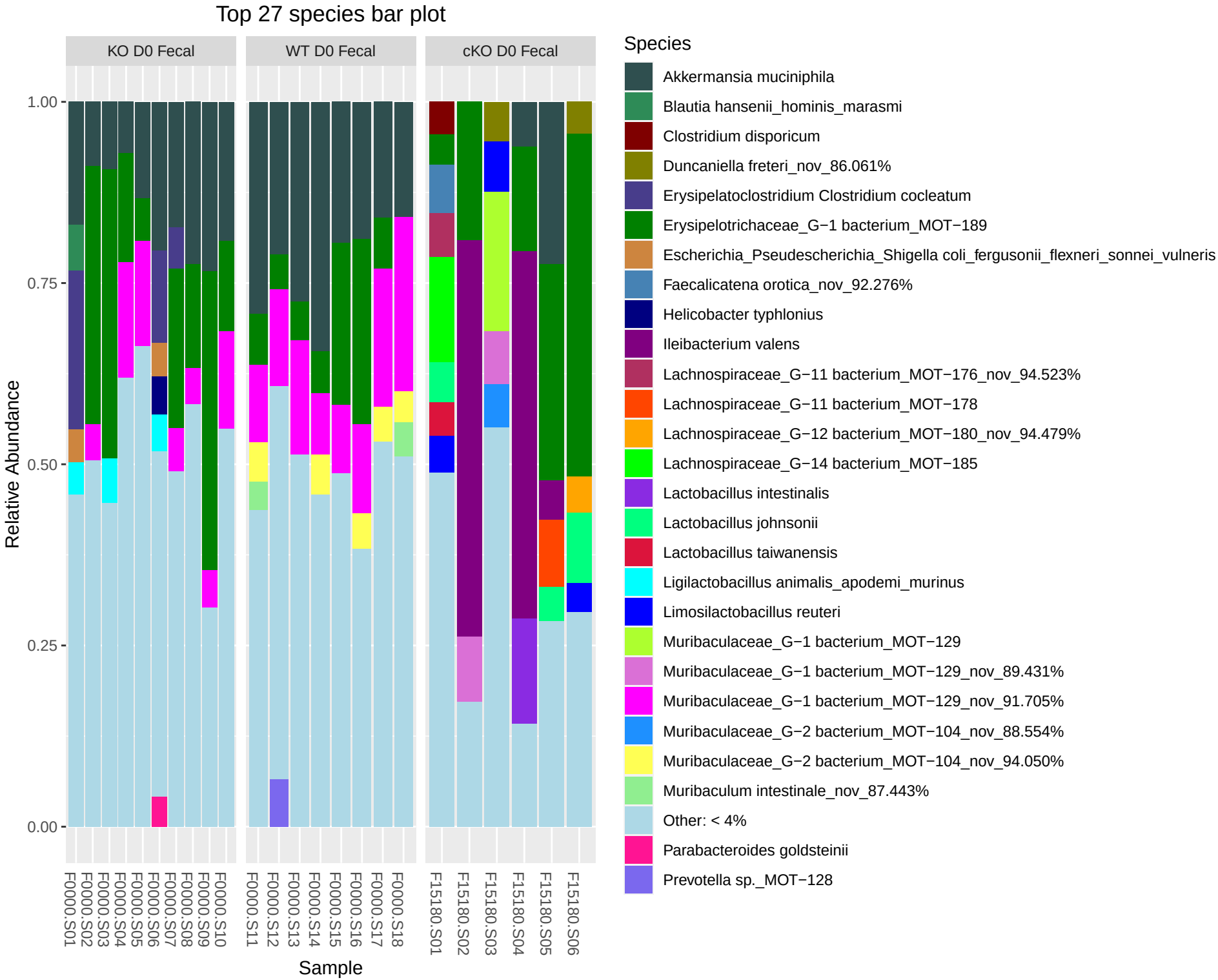
Top 27 species bar plot
KO D0 Fecal
WT D0 Fecal
cKO D0 Fecal
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F0000.S01
F0000.S02
F0000.S03
F0000.S04
F0000.S05
F0000.S06
F0000.S07
F0000.S08
F0000.S09
F0000.S10
F0000.S11
F0000.S12
F0000.S13
F0000.S14
F0000.S15
F0000.S16
F0000.S17
F0000.S18
F15180.S01
F15180.S02
F15180.S03
F15180.S04
F15180.S05
F15180.S06
Sample
Species
Akkermansia muciniphila
Blautia hansenii_hominis_marasmi
Clostridium disporicum
Duncaniella freteri_nov_86.061%
Erysipelatoclostridium Clostridium cocleatum
Erysipelotrichaceae_G−1 bacterium_MOT−189
Escherichia_Pseudescherichia_Shigella coli_fergusonii_flexneri_sonnei_vulneris
Faecalicatena orotica_nov_92.276%
Helicobacter typhlonius
Ileibacterium valens
Lachnospiraceae_G−11 bacterium_MOT−176_nov_94.523%
Lachnospiraceae_G−11 bacterium_MOT−178
Lachnospiraceae_G−12 bacterium_MOT−180_nov_94.479%
Lachnospiraceae_G−14 bacterium_MOT−185
Lactobacillus intestinalis
Lactobacillus johnsonii
Lactobacillus taiwanensis
Ligilactobacillus animalis_apodemi_murinus
Limosilactobacillus reuteri
Muribaculaceae_G−1 bacterium_MOT−129
Muribaculaceae_G−1 bacterium_MOT−129_nov_89.431%
Muribaculaceae_G−1 bacterium_MOT−129_nov_91.705%
Muribaculaceae_G−2 bacterium_MOT−104_nov_88.554%
Muribaculaceae_G−2 bacterium_MOT−104_nov_94.050%
Muribaculum intestinale_nov_87.443%
Other: < 4%
Parabacteroides goldsteinii
Prevotella sp._MOT−128