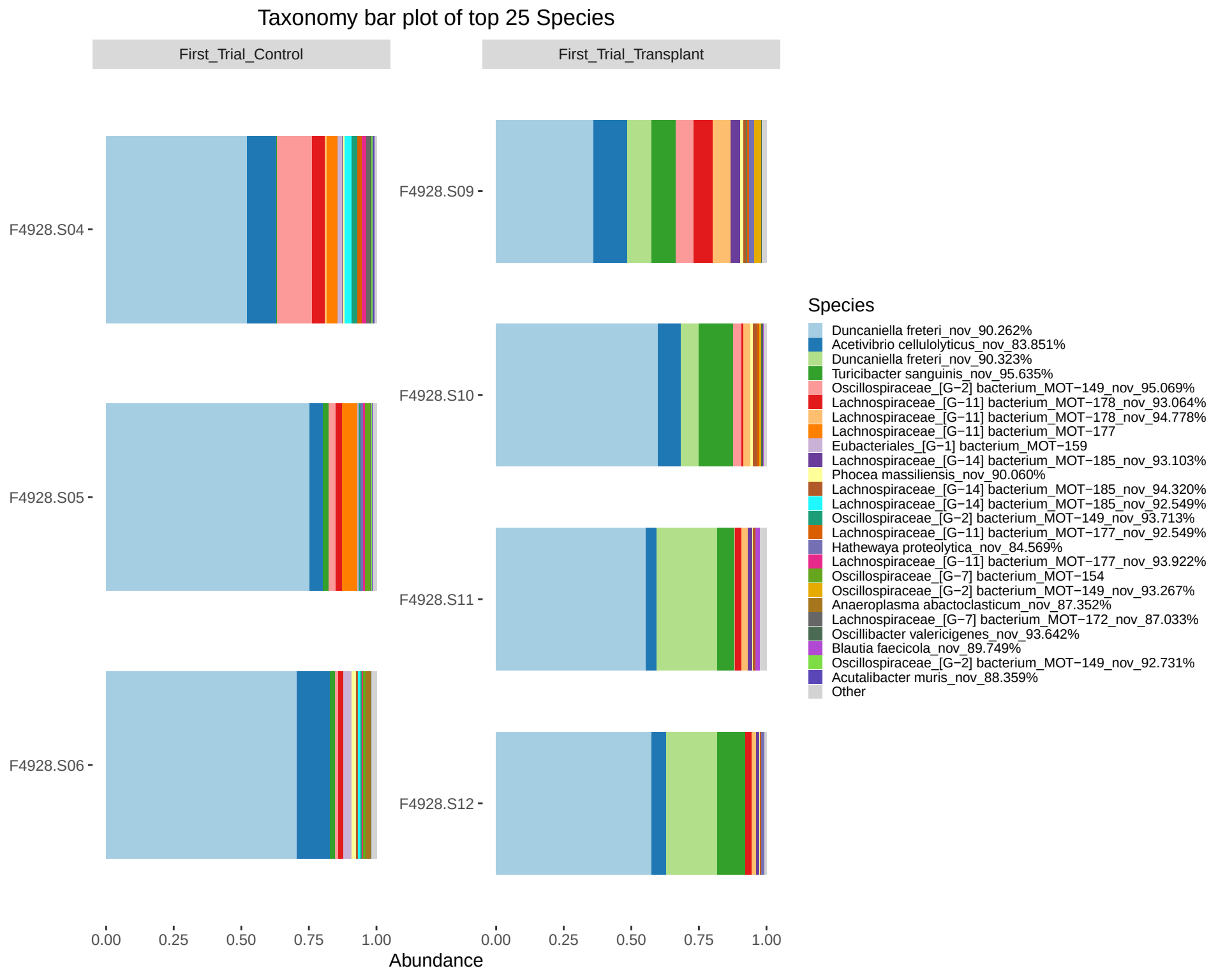
Taxonomy bar plot of top 25 Species
First_Trial_Control
First_Trial_Transplant
F4928.S04
F4928.S05
F4928.S06
F4928.S09
F4928.S10
F4928.S11
F4928.S12
0.00
0.25
0.50
0.75
1.00
Abundance
Species
Duncaniella freteri_nov_90.262%
Acetivibrio cellulolyticus_nov_83.851%
Duncaniella freteri_nov_90.323%
Turicibacter sanguinis_nov_95.635%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.069%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_93.064%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_94.778%
Lachnospiraceae_[G−11] bacterium_MOT−177
Eubacteriales_[G−1] bacterium_MOT−159
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_93.103%
Phocea massiliensis_nov_90.060%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_94.320%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.549%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.713%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_92.549%
Hathewaya proteolytica_nov_84.569%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_93.922%
Oscillospiraceae_[G−7] bacterium_MOT−154
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.267%
Anaeroplasma abactoclasticum_nov_87.352%
Lachnospiraceae_[G−7] bacterium_MOT−172_nov_87.033%
Oscillibacter valericigenes_nov_93.642%
Blautia faecicola_nov_89.749%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_92.731%
Acutalibacter muris_nov_88.359%
Other