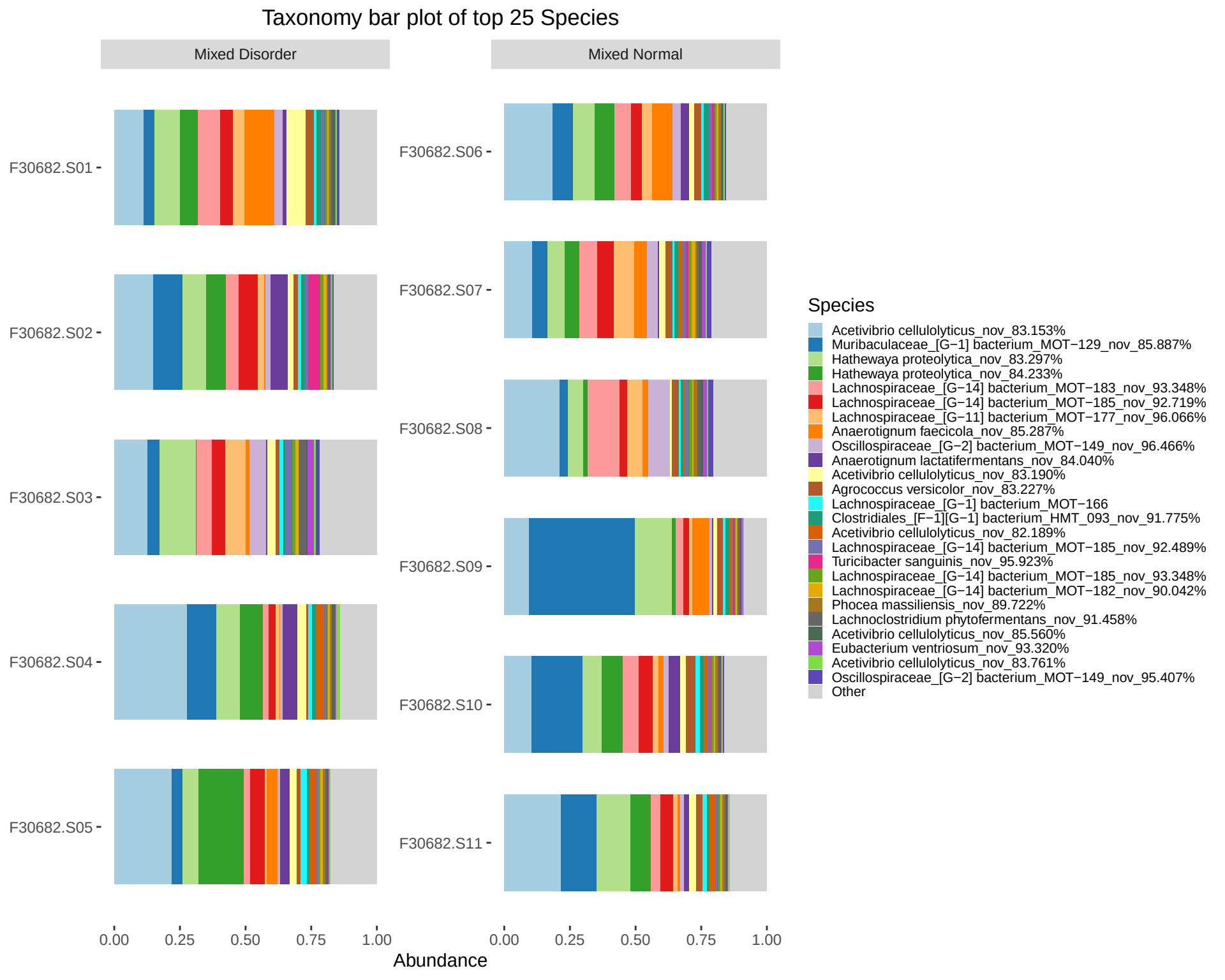
Taxonomy bar plot of top 25 Species
Mixed Disorder
Mixed Normal
F30682.S01
F30682.S02
F30682.S03
F30682.S04
F30682.S05
F30682.S06
F30682.S07
F30682.S08
F30682.S09
F30682.S10
F30682.S11
0.00
0.25
0.50
0.75
1.00
Abundance
Species
Acetivibrio cellulolyticus_nov_83.153%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Hathewaya proteolytica_nov_83.297%
Hathewaya proteolytica_nov_84.233%
Lachnospiraceae_[G−14] bacterium_MOT−183_nov_93.348%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.719%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_96.066%
Anaerotignum faecicola_nov_85.287%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.466%
Anaerotignum lactatifermentans_nov_84.040%
Acetivibrio cellulolyticus_nov_83.190%
Agrococcus versicolor_nov_83.227%
Lachnospiraceae_[G−1] bacterium_MOT−166
Clostridiales_[F−1][G−1] bacterium_HMT_093_nov_91.775%
Acetivibrio cellulolyticus_nov_82.189%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.489%
Turicibacter sanguinis_nov_95.923%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_93.348%
Lachnospiraceae_[G−14] bacterium_MOT−182_nov_90.042%
Phocea massiliensis_nov_89.722%
Lachnoclostridium phytofermentans_nov_91.458%
Acetivibrio cellulolyticus_nov_85.560%
Eubacterium ventriosum_nov_93.320%
Acetivibrio cellulolyticus_nov_83.761%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.407%
Other