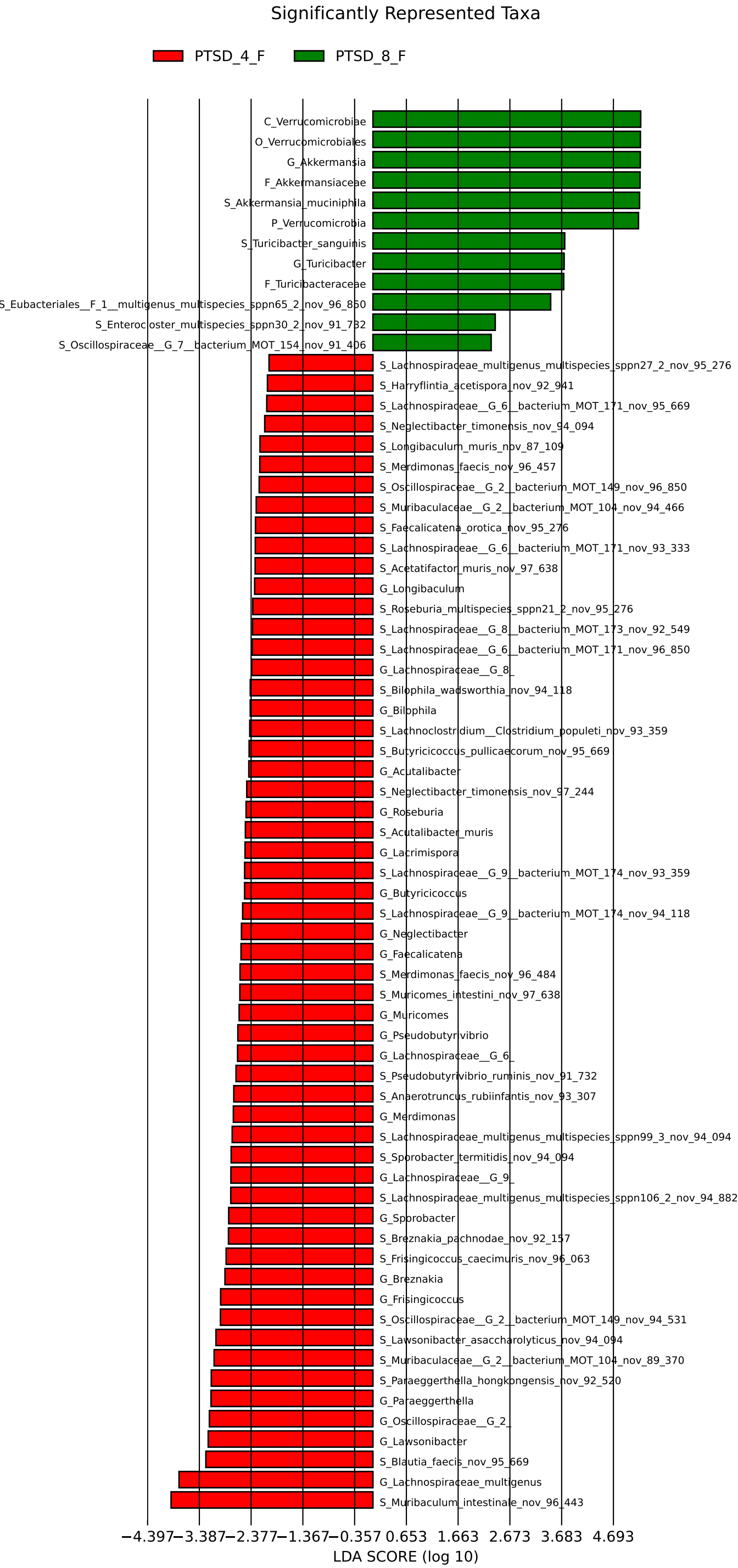
Significantly Represented Taxa
PTSD_4_F
PTSD_8_F
C_Verrucomicrobiae
O_Verrucomicrobiales
G_Akkermansia
F_Akkermansiaceae
S_Akkermansia_muciniphila
P_Verrucomicrobia
S_Turicibacter_sanguinis
G_Turicibacter
F_Turicibacteraceae
S_Eubacteriales__F_1__multigenus_multispecies_sppn65_2_nov_96_850
S_Enterocloster_multispecies_sppn30_2_nov_91_732
S_Oscillospiraceae__G_7__bacterium_MOT_154_nov_91_406
S_Lachnospiraceae_multigenus_multispecies_sppn27_2_nov_95_276
S_Harryflintia_acetispora_nov_92_941
S_Lachnospiraceae__G_6__bacterium_MOT_171_nov_95_669
S_Neglectibacter_timonensis_nov_94_094
S_Longibaculum_muris_nov_87_109
S_Merdimonas_faecis_nov_96_457
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_96_850
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_94_466
S_Faecalicatena_orotica_nov_95_276
S_Lachnospiraceae__G_6__bacterium_MOT_171_nov_93_333
S_Acetatifactor_muris_nov_97_638
G_Longibaculum
S_Roseburia_multispecies_sppn21_2_nov_95_276
S_Lachnospiraceae__G_8__bacterium_MOT_173_nov_92_549
S_Lachnospiraceae__G_6__bacterium_MOT_171_nov_96_850
G_Lachnospiraceae__G_8_
S_Bilophila_wadsworthia_nov_94_118
G_Bilophila
S_Lachnoclostridium__Clostridium_populeti_nov_93_359
S_Butyricicoccus_pullicaecorum_nov_95_669
G_Acutalibacter
S_Neglectibacter_timonensis_nov_97_244
G_Roseburia
S_Acutalibacter_muris
G_Lacrimispora
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_93_359
G_Butyricicoccus
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_94_118
G_Neglectibacter
G_Faecalicatena
S_Merdimonas_faecis_nov_96_484
S_Muricomes_intestini_nov_97_638
G_Muricomes
G_Pseudobutyrivibrio
G_Lachnospiraceae__G_6_
S_Pseudobutyrivibrio_ruminis_nov_91_732
S_Anaerotruncus_rubiinfantis_nov_93_307
G_Merdimonas
S_Lachnospiraceae_multigenus_multispecies_sppn99_3_nov_94_094
S_Sporobacter_termitidis_nov_94_094
G_Lachnospiraceae__G_9_
S_Lachnospiraceae_multigenus_multispecies_sppn106_2_nov_94_882
G_Sporobacter
S_Breznakia_pachnodae_nov_92_157
S_Frisingicoccus_caecimuris_nov_96_063
G_Breznakia
G_Frisingicoccus
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_94_531
S_Lawsonibacter_asaccharolyticus_nov_94_094
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_89_370
S_Paraeggerthella_hongkongensis_nov_92_520
G_Paraeggerthella
G_Oscillospiraceae__G_2_
G_Lawsonibacter
S_Blautia_faecis_nov_95_669
G_Lachnospiraceae_multigenus
S_Muribaculum_intestinale_nov_96_443
−4.397
−3.387
−2.377
−1.367
−0.357
0.653
1.663
2.673
3.683
4.693
LDA SCORE (log 10)