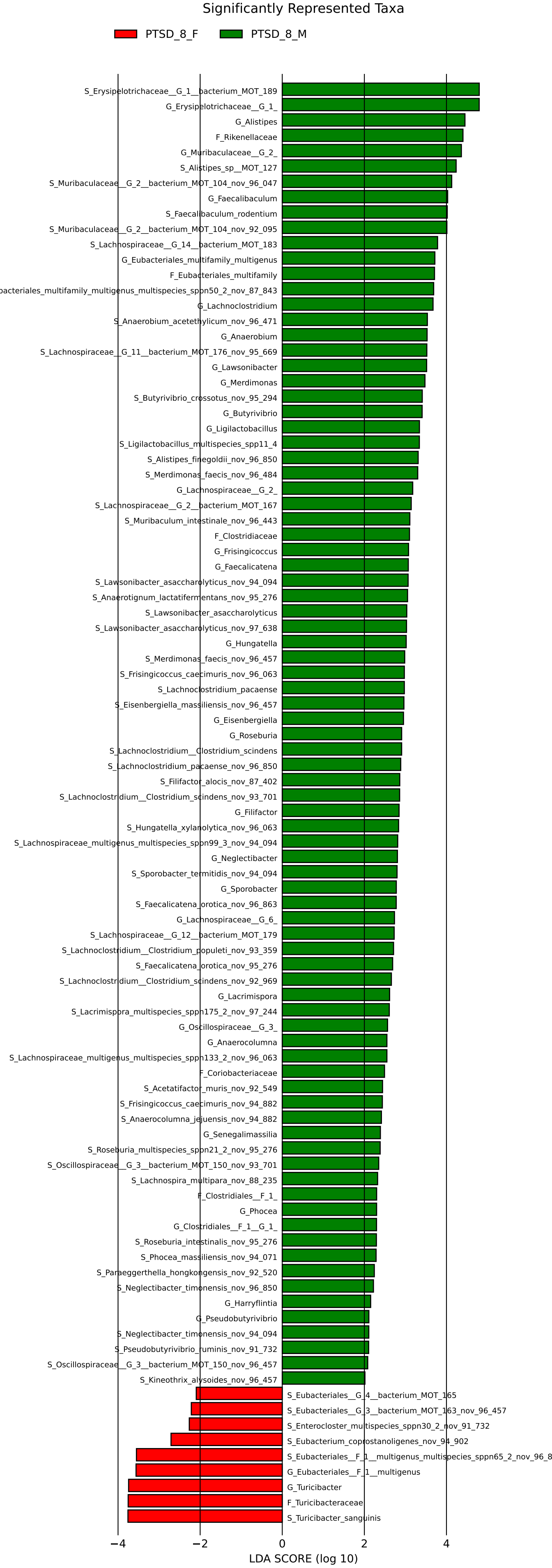
Significantly Represented Taxa
PTSD_8_F
PTSD_8_M
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
G_Erysipelotrichaceae__G_1_
G_Alistipes
F_Rikenellaceae
G_Muribaculaceae__G_2_
S_Alistipes_sp__MOT_127
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_96_047
G_Faecalibaculum
S_Faecalibaculum_rodentium
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_92_095
S_Lachnospiraceae__G_14__bacterium_MOT_183
G_Eubacteriales_multifamily_multigenus
F_Eubacteriales_multifamily
bacteriales_multifamily_multigenus_multispecies_spon50_2_nov_87_843
G_Lachnoclostridium
S_Anaerobium_acetethylicum_nov_96_471
G_Anaerobium
S_Lachnospiraceae__G_11__bacterium_MOT_176_nov_95_669
G_Lawsonibacter
G_Merdimonas
S_Butyrivibrio_crossotus_nov_95_294
G_Butyrivibrio
G_Ligilactobacillus
S_Ligilactobacillus_multispecies_spp11_4
S_Alistipes_finegoldii_nov_96_850
S_Merdimonas_faecis_nov_96_484
G_Lachnospiraceae__G_2_
S_Lachnospiraceae__G_2__bacterium_MOT_167
S_Muribaculum_intestinale_nov_96_443
F_Clostridiaceae
G_Frisingicoccus
G_Faecalicatena
S_Lawsonibacter_asaccharolyticus_nov_94_094
S_Anaerotignum_lactatifermentans_nov_95_276
S_Lawsonibacter_asaccharolyticus
S_Lawsonibacter_asaccharolyticus_nov_97_638
G_Hungatella
S_Merdimonas_faecis_nov_96_457
S_Frisingicoccus_caecimuris_nov_96_063
S_Lachnoclostridium_pacaense
S_Eisenbergiella_massiliensis_nov_96_457
G_Eisenbergiella
G_Roseburia
S_Lachnoclostridium__Clostridium_scindens
S_Lachnoclostridium_pacaense_nov_96_850
S_Filifactor_alocis_nov_87_402
S_Lachnoclostridium__Clostridium_scindens_nov_93_701
G_Filifactor
S_Hungatella_xylanolytica_nov_96_063
S_Lachnospiraceae_multigenus_multispecies_spon99_3_nov_94_094
G_Neglectibacter
S_Sporobacter_termitidis_nov_94_094
G_Sporobacter
S_Faecalicatena_orotica_nov_96_863
G_Lachnospiraceae__G_6_
S_Lachnospiraceae__G_12__bacterium_MOT_179
S_Lachnoclostridium__Clostridium_populeti_nov_93_359
S_Faecalicatena_orotica_nov_95_276
S_Lachnoclostridium__Clostridium_scindens_nov_92_969
G_Lacrimispora
S_Lacrimispora_multispecies_sppn175_2_nov_97_244
G_Oscillospiraceae__G_3_
G_Anaerocolumna
S_Lachnospiraceae_multigenus_multispecies_sppn133_2_nov_96_063
F_Coriobacteriaceae
S_Acetatifactor_muris_nov_92_549
S_Frisingicoccus_caecimuris_nov_94_882
S_Anaerocolumna_jejuensis_nov_94_882
G_Senegalimassilia
S_Roseburia_multispecies_spon21_2_nov_95_276
S_Oscillospiraceae__G_3__bacterium_MOT_150_nov_93_701
S_Lachnospira_multipara_nov_88_235
F_Clostridiales__F_1_
G_Phocea
G_Clostridiales__F_1__G_1_
S_Roseburia_intestinalis_nov_95_276
S_Phocea_massiliensis_nov_94_071
S_Paraeggerthella_hongkongensis_nov_92_520
S_Neglectibacter_timonensis_nov_96_850
G_Harryflintia
G_Pseudobutyrivibrio
S_Neglectibacter_timonensis_nov_94_094
S_Pseudobutyrivibrio_ruminis_nov_91_732
S_Oscillospiraceae__G_3__bacterium_MOT_150_nov_96_457
S_Kineothrix_alysoides_nov_96_457
S_Eubacteriales__G_4__bacterium_MOT_165
S_Eubacteriales__G_3__bacterium_MOT_163_nov_96_457
S_Enterocloster_multispecies_sppn30_2_nov_91_732
S_Eubacterium_coprostanoligenes_nov_94_902
S_Eubacteriales__F_1__multigenus_multispecies_sppn65_2_nov_96_8
G_Eubacteriales__F_1__multigenus
G_Turicibacter
F_Turicibacteraceae
S_Turicibacter_sanguinis
−4
−2
0
2
4
LDA SCORE (log 10)