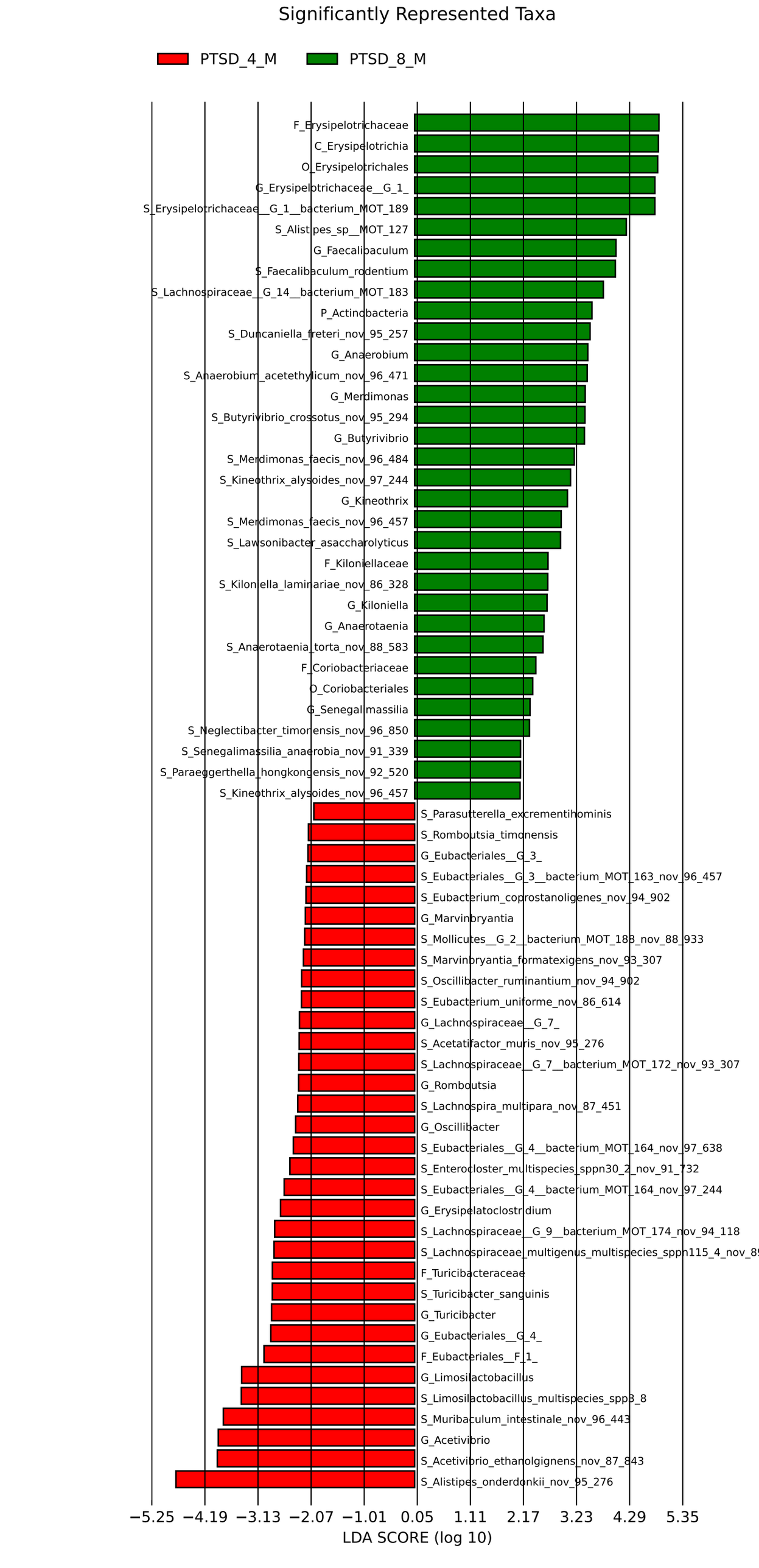
Significantly Represented Taxa
PTSD_4_M
PTSD_8_M
F_Erysipelotrichaceae
C_Erysipelotrichia
O_Erysipelotrichales
G_Erysipelotrichaceae__G_1_
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
S_Alistipes_sp__MOT_127
G_Faecalibaculum
S_Faecalibaculum_rodentium
S_Lachnospiraceae__G_14__bacterium_MOT_183
P_Actinobacteria
S_Duncaniella_freteri_nov_95_257
G_Anaerobium
S_Anaerobium_acetethylicum_nov_96_471
G_Merdimonas
S_Butyrivibrio_crossotus_nov_95_294
G_Butyrivibrio
S_Merdimonas_faecis_nov_96_484
S_Kineothrix_alysoides_nov_97_244
G_Kineothrix
S_Merdimonas_faecis_nov_96_457
S_Lawsonibacter_asaccharolyticus
F_Kiloniellaceae
S_Kiloniella_laminariae_nov_86_328
G_Kiloniella
G_Anaerotaenia
S_Anaerotaenia_torta_nov_88_583
F_Coriobacteriaceae
O_Coriobacteriales
G_Senegalimassilia
S_Neglectibacter_timonensis_nov_96_850
S_Senegalimassilia_anaerobia_nov_91_339
S_Paraeggerthella_hongkongensis_nov_92_520
S_Kineothrix_alysoides_nov_96_457
S_Parasutterella_excrementihominis
S_Romboutsia_timonensis
G_Eubacteriales__G_3_
S_Eubacteriales__G_3__bacterium_MOT_163_nov_96_457
S_Eubacterium_coprostanoligenes_nov_94_902
G_Marvinbryantia
S_Mollicutes__G_2__bacterium_MOT_188_nov_88_933
S_Marvinbryantia_formatexigens_nov_93_307
S_Oscillibacter_ruminantium_nov_94_902
S_Eubacterium_uniforme_nov_86_614
G_Lachnospiraceae__G_7_
S_Acetatifactor_muris_nov_95_276
S_Lachnospiraceae__G_7__bacterium_MOT_172_nov_93_307
G_Romboutsia
S_Lachnospira_multipara_nov_87_451
G_Oscillibacter
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_638
S_Enterocloster_multispecies_sppn30_2_nov_91_732
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_244
G_Erysipelatoclostridium
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_94_118
S_Lachnospiraceae_multigenus_multispecies_sppn115_4_nov_89
F_Turicibacteraceae
S_Turicibacter_sanguinis
G_Turicibacter
G_Eubacteriales__G_4_
F_Eubacteriales__F_1_
G_Limosilactobacillus
S_Limosilactobacillus_multispecies_spp3_8
S_Muribaculum_intestinale_nov_96_443
G_Acetivibrio
S_Acetivibrio_ethanolgignens_nov_87_843
S_Alistipes_onderdonkii_nov_95_276
−5.25
−4.19
−3.13
−2.07
−1.01
0.05
1.11
2.17
3.23
4.29
5.35
LDA SCORE (log 10)