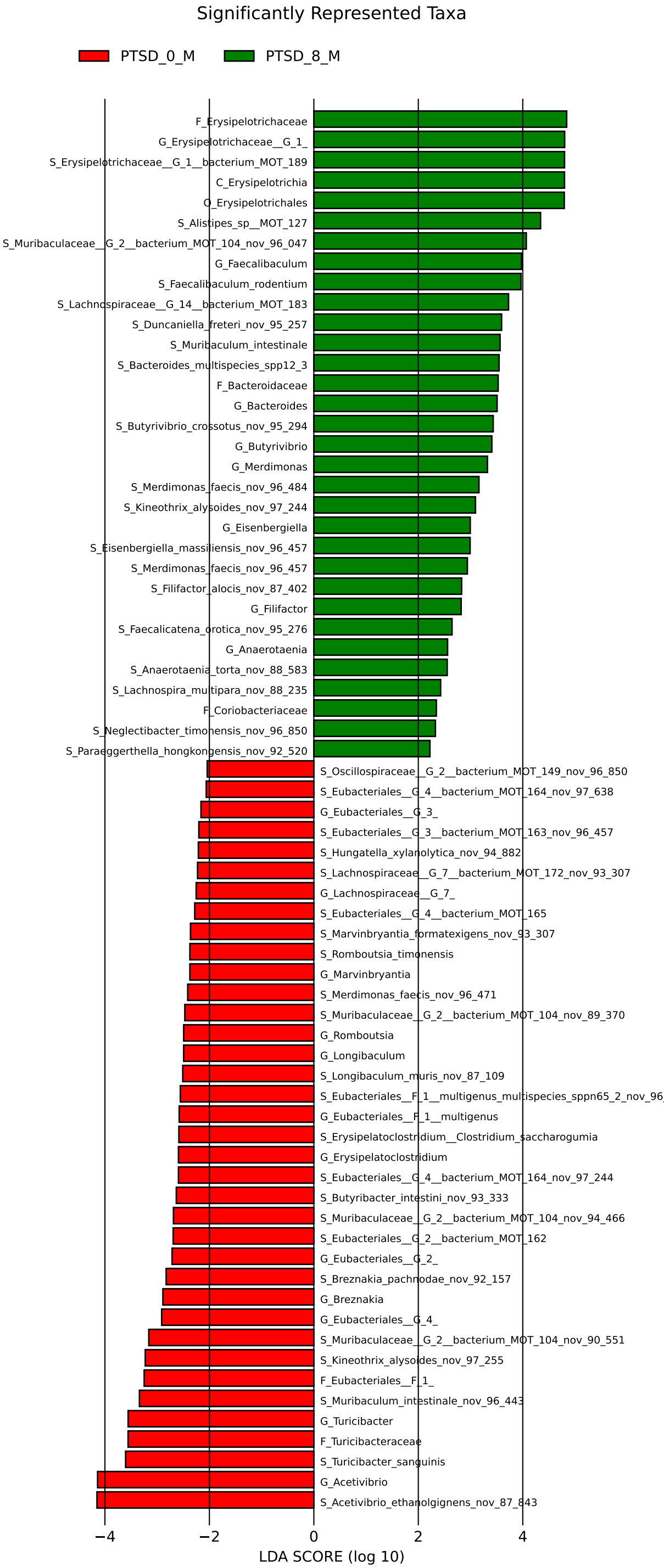
Significantly Represented Taxa
PTSD_0_M
PTSD_8_M
F_Erysipelotrichaceae
G_Erysipelotrichaceae__G_1_
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
C_Erysipelotrichia
O_Erysipelotrichales
S_Alistipes_sp__MOT_127
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_96_047
G_Faecalibaculum
S_Faecalibaculum_rodentium
S_Lachnospiraceae__G_14__bacterium_MOT_183
S_Duncaniella_freteri_nov_95_257
S_Muribaculum_intestinale
S_Bacteroides_multispecies_spp12_3
F_Bacteroidaceae
G_Bacteroides
S_Butyrivibrio_crossotus_nov_95_294
G_Butyrivibrio
G_Merdimonas
S_Merdimonas_faecis_nov_96_484
S_Kineothrix_alysoides_nov_97_244
G_Eisenbergiella
S_Eisenbergiella_massiliensis_nov_96_457
S_Merdimonas_faecis_nov_96_457
S_Filifactor_alocis_nov_87_402
G_Filifactor
S_Faecalicatena_orotica_nov_95_276
G_Anaerotaenia
S_Anaerotaenia_torta_nov_88_583
S_Lachnospira_multipara_nov_88_235
F_Coriobacteriaceae
S_Neglectibacter_timonensis_nov_96_850
S_Paraeggerthella_hongkongensis_nov_92_520
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_96_850
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_638
G_Eubacteriales__G_3_
S_Eubacteriales__G_3__bacterium_MOT_163_nov_96_457
S_Hungatella_xylanolytica_nov_94_882
S_Lachnospiraceae__G_7__bacterium_MOT_172_nov_93_307
G_Lachnospiraceae__G_7_
S_Eubacteriales__G_4__bacterium_MOT_165
S_Marvinbryantia_formatexigens_nov_93_307
S_Romboutsia_timonensis
G_Marvinbryantia
S_Merdimonas_faecis_nov_96_471
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_89_370
G_Romboutsia
G_Longibaculum
S_Longibaculum_muris_nov_87_109
S_Eubacteriales__F_1__multigenus_multispecies_sppn65_2_nov_96
G_Eubacteriales__F_1__multigenus
S_Erysipelatoclostridium__Clostridium_saccharogumia
G_Erysipelatoclostridium
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_244
S_Butyribacter_intestini_nov_93_333
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_94_466
S_Eubacteriales__G_2__bacterium_MOT_162
G_Eubacteriales__G_2_
S_Breznakia_pachnodae_nov_92_157
G_Breznakia
G_Eubacteriales__G_4_
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_90_551
S_Kineothrix_alysoides_nov_97_255
F_Eubacteriales__F_1_
S_Muribaculum_intestinale_nov_96_443
G_Turicibacter
F_Turicibacteraceae
S_Turicibacter_sanguinis
G_Acetivibrio
S_Acetivibrio_ethanolgignens_nov_87_843
−4
−2
0
2
4
LDA SCORE (log 10)