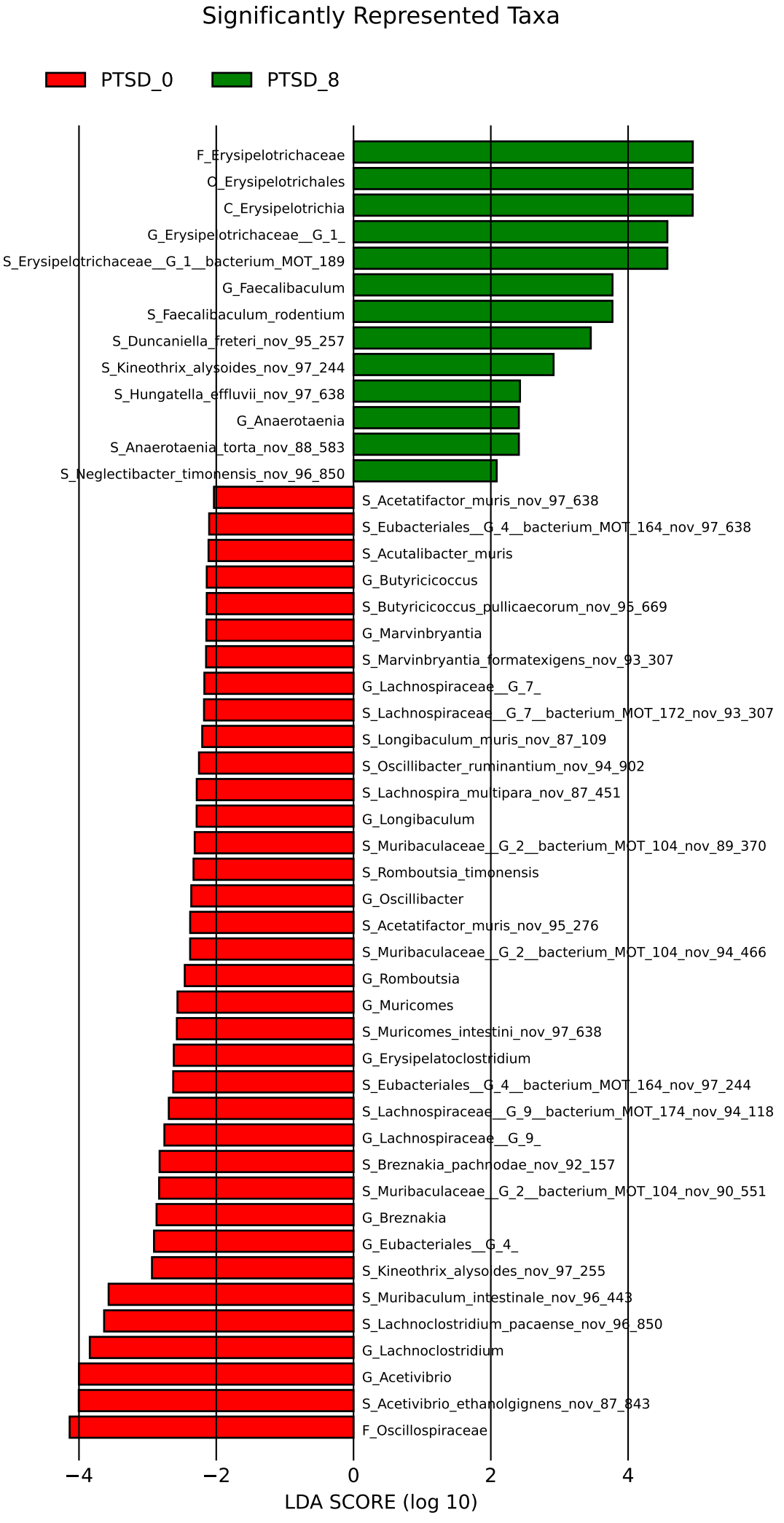
Significantly Represented Taxa
PTSD_0
PTSD_8
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
G_Erysipelotrichaceae__G_1_
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
G_Faecalibaculum
S_Faecalibaculum_rodentium
S_Duncaniella_freteri_nov_95_257
S_Kineothrix_alysoides_nov_97_244
S_Hungatella_effluvii_nov_97_638
G_Anaerotaenia
S_Anaerotaenia_torta_nov_88_583
S_Neglectibacter_timonensis_nov_96_850
S_Acetatifactor_muris_nov_97_638
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_638
S_Acutalibacter_muris
G_Butyricicoccus
S_Butyricicoccus_pullicaecorum_nov_95_669
G_Marvinbryantia
S_Marvinbryantia_formatexigens_nov_93_307
G_Lachnospiraceae__G_7_
S_Lachnospiraceae__G_7__bacterium_MOT_172_nov_93_307
S_Longibaculum_muris_nov_87_109
S_Oscillibacter_ruminantium_nov_94_902
S_Lachnospira_multipara_nov_87_451
G_Longibaculum
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_89_370
S_Romboutsia_timonensis
G_Oscillibacter
S_Acetatifactor_muris_nov_95_276
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_94_466
G_Romboutsia
G_Muricomes
S_Muricomes_intestini_nov_97_638
G_Erysipelatoclostridium
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_244
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_94_118
G_Lachnospiraceae__G_9_
S_Breznakia_pachnodae_nov_92_157
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_90_551
G_Breznakia
G_Eubacteriales__G_4_
S_Kineothrix_alysoides_nov_97_255
S_Muribaculum_intestinale_nov_96_443
S_Lachnoclostridium_pacaense_nov_96_850
G_Lachnoclostridium
G_Acetivibrio
S_Acetivibrio_ethanolgignens_nov_87_843
F_Oscillospiraceae
−4
−2
0
2
4
LDA SCORE (log 10)