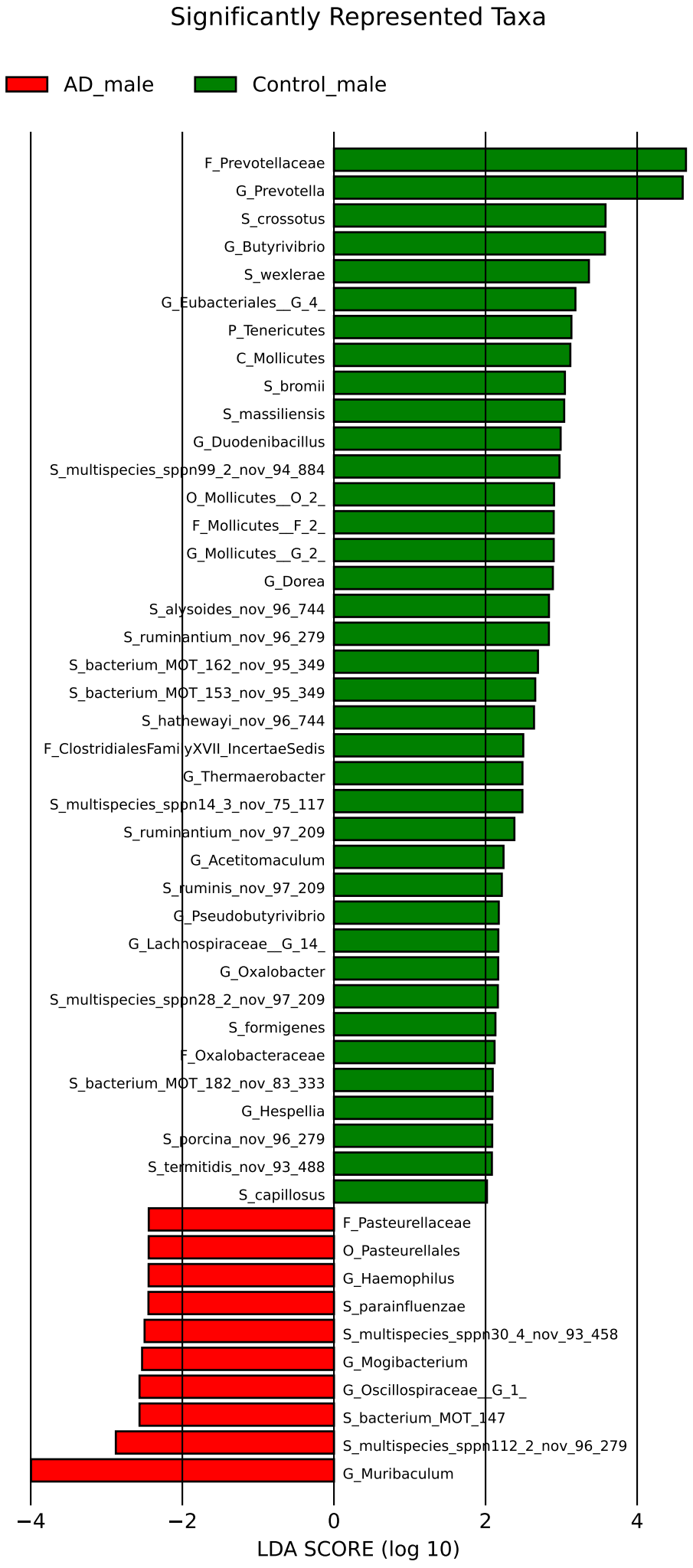
Significantly Represented Taxa
AD_male
Control_male
F_Prevotellaceae
G_Prevotella
S_crossotus
G_Butyrivibrio
S_wexlerae
G_Eubacteriales__G_4_
P_Tenericutes
C_Mollicutes
S_bromii
S_massiliensis
G_Duodenibacillus
S_multispecies_sppn99_2_nov_94_884
O_Mollicutes__O_2_
F_Mollicutes__F_2_
G_Mollicutes__G_2_
G_Dorea
S_alysoides_nov_96_744
S_ruminantium_nov_96_279
S_bacterium_MOT_162_nov_95_349
S_bacterium_MOT_153_nov_95_349
S_hathewayi_nov_96_744
F_ClostridialesFamilyXVII_IncertaeSedis
G_Thermaerobacter
S_multispecies_sppn14_3_nov_75_117
S_ruminantium_nov_97_209
G_Acetitomaculum
S_ruminis_nov_97_209
G_Pseudobutyrivibrio
G_Lachnospiraceae__G_14_
G_Oxalobacter
S_multispecies_sppn28_2_nov_97_209
S_formigenes
F_Oxalobacteraceae
S_bacterium_MOT_182_nov_83_333
G_Hespellia
S_porcina_nov_96_279
S_termitidis_nov_93_488
S_capillosus
F_Pasteurellaceae
O_Pasteurellales
G_Haemophilus
S_parainfluenzae
S_multispecies_sppn30_4_nov_93_458
G_Mogibacterium
G_Oscillospiraceae__G_1_
S_bacterium_MOT_147
S_multispecies_sppn112_2_nov_96_279
G_Muribaculum
−4
−2
0
2
4
LDA SCORE (log 10)