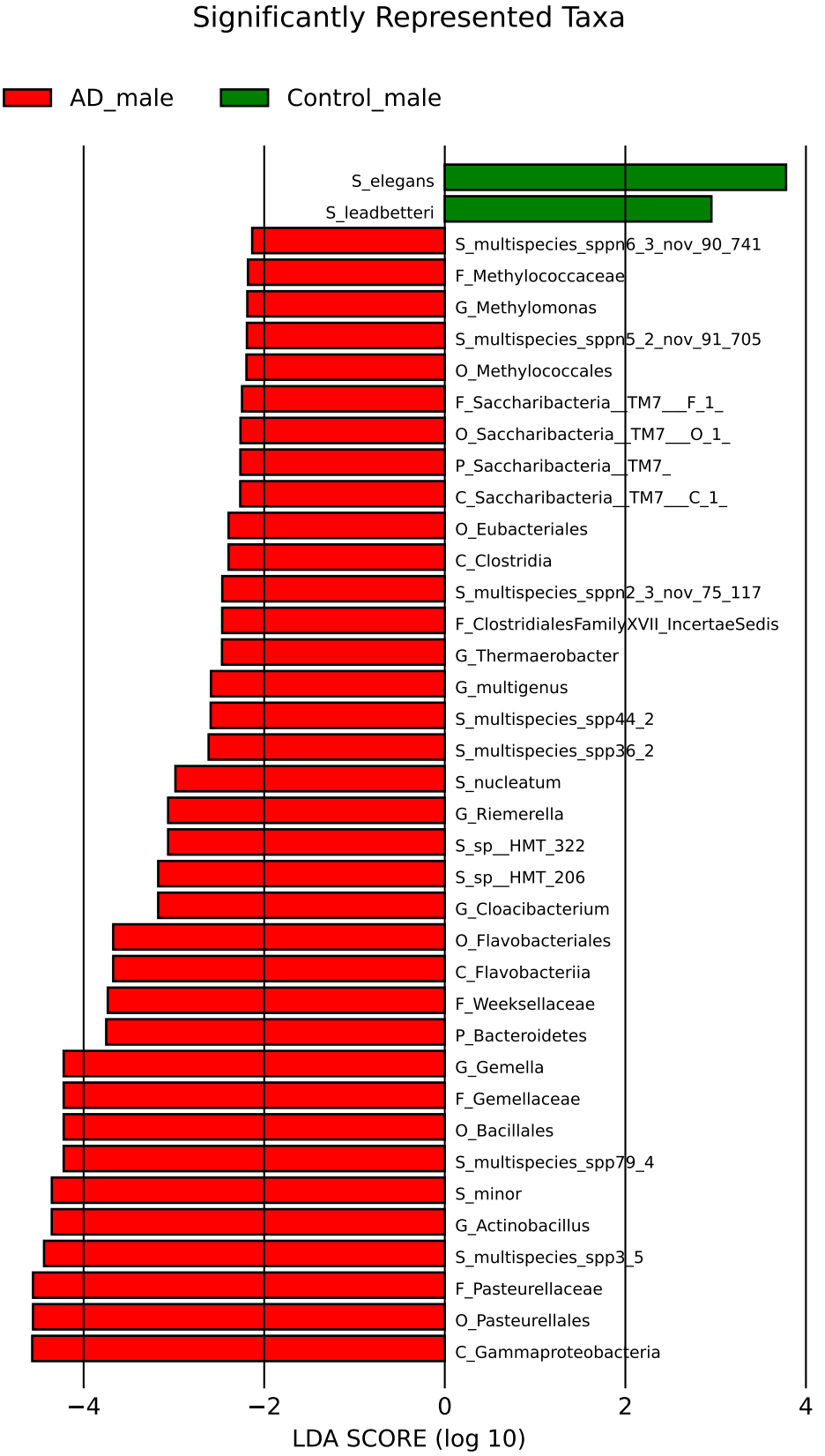
Significantly Represented Taxa
AD_male
Control_male
S_elegans
S_leadbetteri
S_multispecies_sppn6_3_nov_90_741
F_Methylococcaceae
G_Methylomonas
S_multispecies_sppn5_2_nov_91_705
O_Methylococcales
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
P_Saccharibacteria__TM7_
C_Saccharibacteria__TM7___C_1_
O_Eubacteriales
C_Clostridia
S_multispecies_sppn2_3_nov_75_117
F_ClostridialesFamilyXVII_IncertaeSedis
G_Thermaerobacter
G_multigenus
S_multispecies_spp44_2
S_multispecies_spp36_2
S_nucleatum
G_Riemerella
S_sp__HMT_322
S_sp__HMT_206
G_Cloacibacterium
O_Flavobacteriales
C_Flavobacteriia
F_Weeksellaceae
P_Bacteroidetes
G_Gemella
F_Gemellaceae
O_Bacillales
S_multispecies_spp79_4
S_minor
G_Actinobacillus
S_multispecies_spp3_5
F_Pasteurellaceae
O_Pasteurellales
C_Gammaproteobacteria
−4
−2
0
2
4
LDA SCORE (log 10)