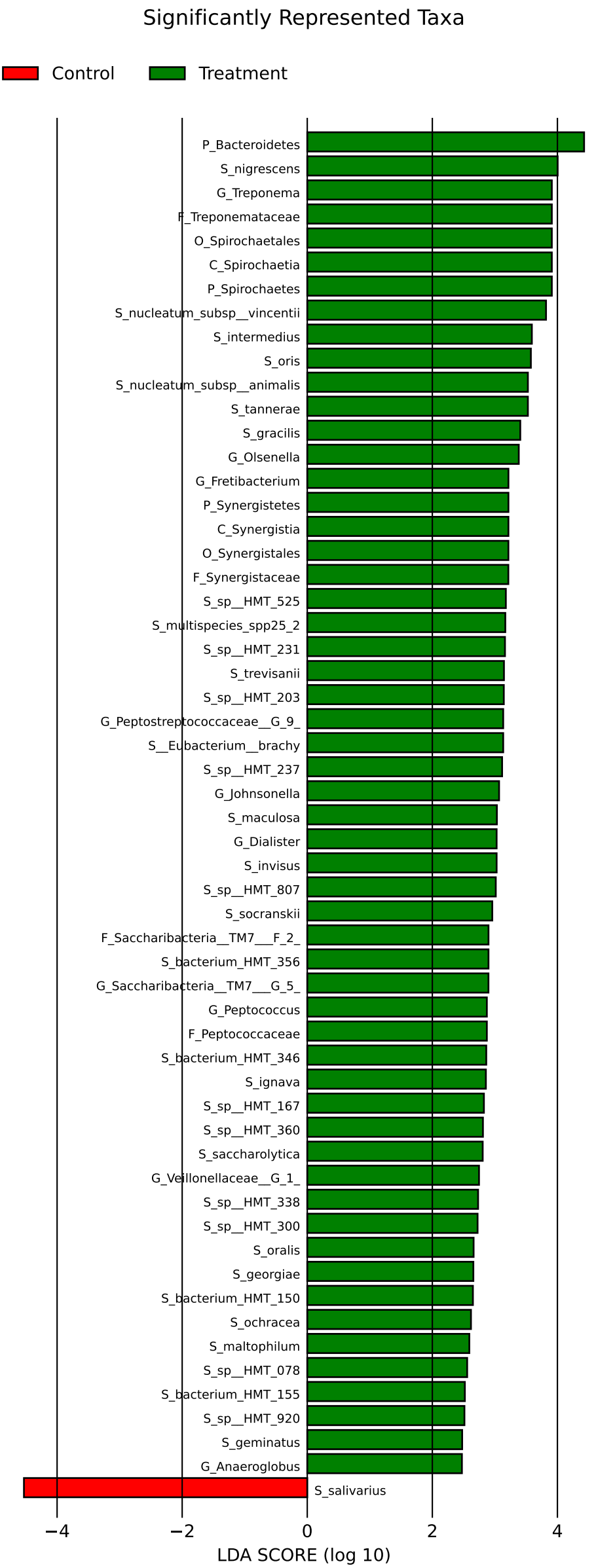
Significantly Represented Taxa
Control
Treatment
P_Bacteroidetes
S_nigrescens
G_Treponema
F_Treponemataceae
O_Spirochaetales
C_Spirochaetia
P_Spirochaetes
S_nucleatum_subsp__vincentii
S_intermedius
S_oris
S_nucleatum_subsp__animalis
S_tannerae
S_gracilis
G_Olsenella
G_Fretibacterium
P_Synergistetes
C_Synergistia
O_Synergistales
F_Synergistaceae
S_sp__HMT_525
S_multispecies_spp25_2
S_sp__HMT_231
S_trevisanii
S_sp__HMT_203
G_Peptostreptococcaceae__G_9_
S__Eubacterium__brachy
S_sp__HMT_237
G_Johnsonella
S_maculosa
G_Dialister
S_invisus
S_sp__HMT_807
S_socranskii
F_Saccharibacteria__TM7___F_2_
S_bacterium_HMT_356
G_Saccharibacteria__TM7___G_5_
G_Peptococcus
F_Peptococcaceae
S_bacterium_HMT_346
S_ignava
S_sp__HMT_167
S_sp__HMT_360
S_saccharolytica
G_Veillonellaceae__G_1_
S_sp__HMT_338
S_sp__HMT_300
S_oralis
S_georgiae
S_bacterium_HMT_150
S_ochracea
S_maltophilum
S_sp__HMT_078
S_bacterium_HMT_155
S_sp__HMT_920
S_geminatus
G_Anaeroglobus
S_salivarius
−4
−2
0
2
4
LDA SCORE (log 10)