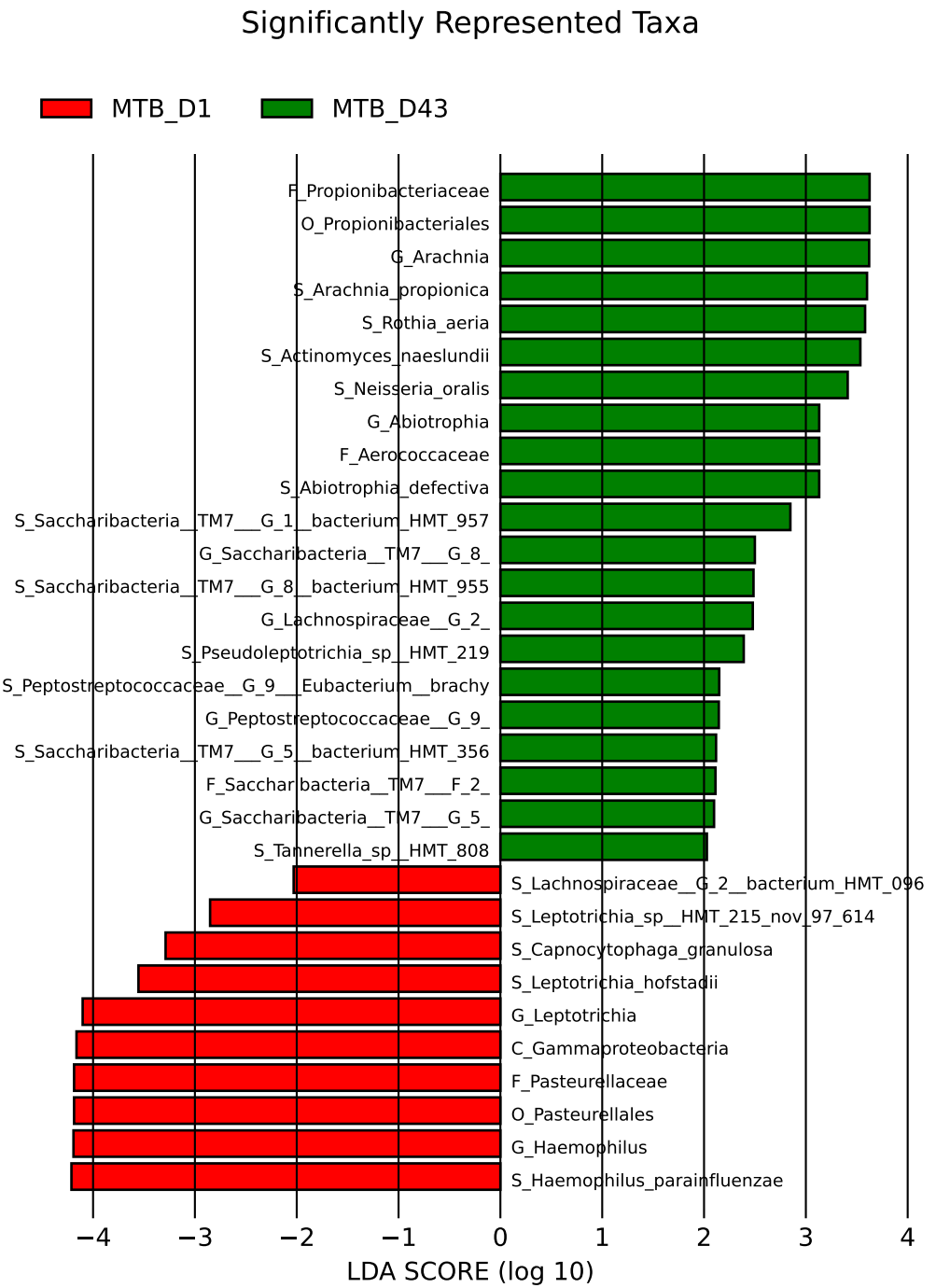
Significantly Represented Taxa
MTB_D1
MTB_D43
F_Propionibacteriaceae
O_Propionibacteriales
G_Arachnia
S_Arachnia_propionica
S_Rothia_aeria
S_Actinomyces_naeslundii
S_Neisseria_oralis
G_Abiotrophia
F_Aerococcaceae
S_Abiotrophia_defectiva
S_Saccharibacteria__TM7___G_1__bacterium_HMT_957
G_Saccharibacteria__TM7___G_8_
S_Saccharibacteria__TM7___G_8__bacterium_HMT_955
G_Lachnospiraceae__G_2_
S_Pseudoleptotrichia_sp__HMT_219
S_Peptostreptococcaceae__G_9___Eubacterium__brachy
G_Peptostreptococcaceae__G_9_
S_Saccharibacteria__TM7___G_5__bacterium_HMT_356
F_Saccharibacteria__TM7___F_2_
G_Saccharibacteria__TM7___G_5_
S_Tannerella_sp__HMT_808
S_Lachnospiraceae__G_2__bacterium_HMT_096
S_Leptotrichia_sp__HMT_215_nov_97_614
S_Capnocytophaga_granulosa
S_Leptotrichia_hofstadii
G_Leptotrichia
C_Gammaproteobacteria
F_Pasteurellaceae
O_Pasteurellales
G_Haemophilus
S_Haemophilus_parainfluenzae
−4
−3
−2
−1
0
1
2
3
4
LDA SCORE (log 10)