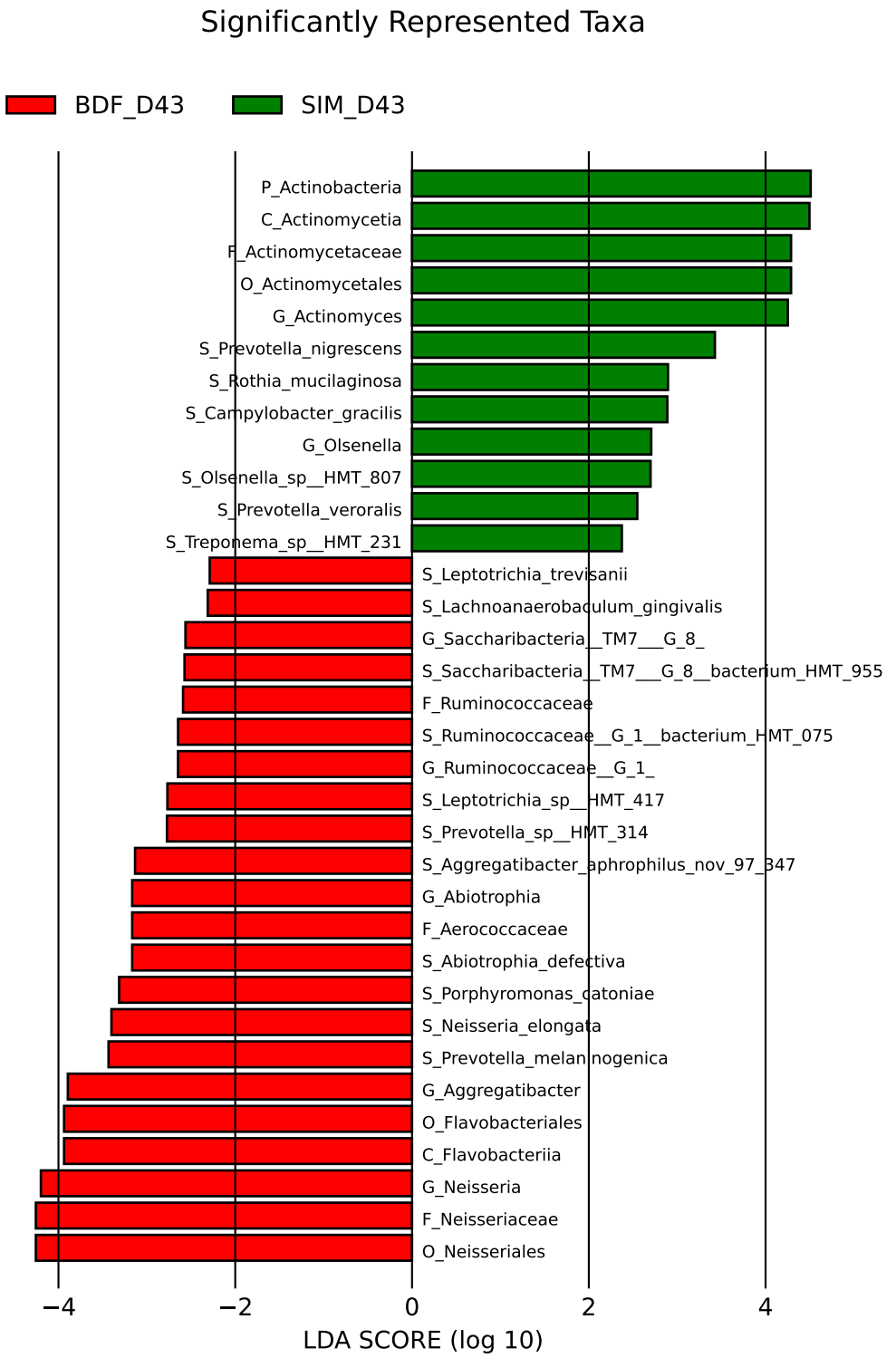
Significantly Represented Taxa
BDF_D43
SIM_D43
P_Actinobacteria
C_Actinomycetia
F_Actinomycetaceae
O_Actinomycetales
G_Actinomyces
S_Prevotella_nigrescens
S_Rothia_mucilaginosa
S_Campylobacter_gracilis
G_Olsenella
S_Olsenella_sp__HMT_807
S_Prevotella_veroralis
S_Treponema_sp__HMT_231
S_Leptotrichia_trevisanii
S_Lachnoanaerobaculum_gingivalis
G_Saccharibacteria__TM7___G_8_
S_Saccharibacteria__TM7___G_8__bacterium_HMT_955
F_Ruminococcaceae
S_Ruminococcaceae__G_1__bacterium_HMT_075
G_Ruminococcaceae__G_1_
S_Leptotrichia_sp__HMT_417
S_Prevotella_sp__HMT_314
S_Aggregatibacter_aphrophilus_nov_97_347
G_Abiotrophia
F_Aerococcaceae
S_Abiotrophia_defectiva
S_Porphyromonas_catoniae
S_Neisseria_elongata
S_Prevotella_melaninogenica
G_Aggregatibacter
O_Flavobacteriales
C_Flavobacteriia
G_Neisseria
F_Neisseriaceae
O_Neisseriales
−4
−2
0
2
4
LDA SCORE (log 10)