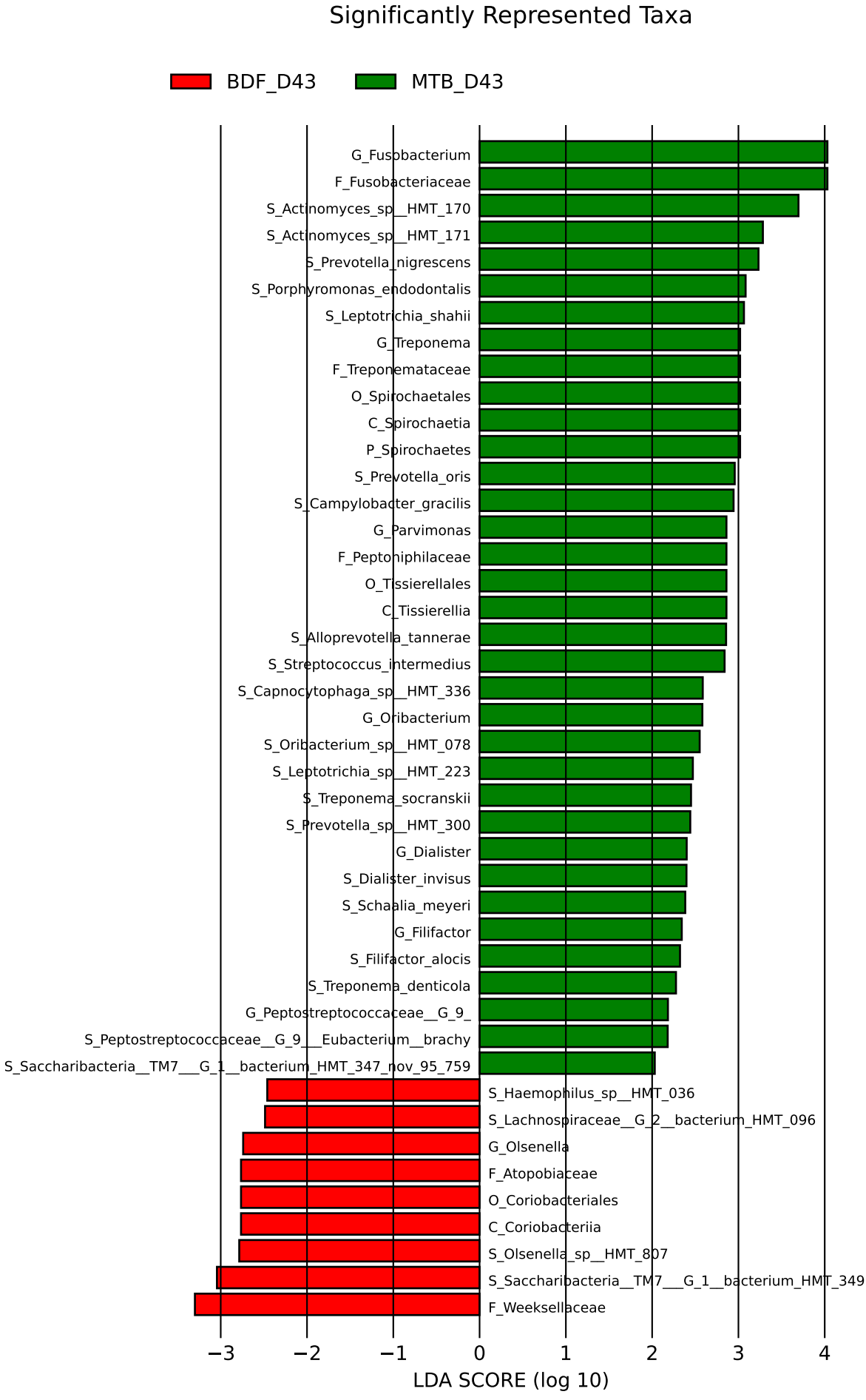
Significantly Represented Taxa
BDF_D43
MTB_D43
G_Fusobacterium
F_Fusobacteriaceae
S_Actinomyces_sp__HMT_170
S_Actinomyces_sp__HMT_171
S_Prevotella_nigrescens
S_Porphyromonas_endodontalis
S_Leptotrichia_shahii
G_Treponema
F_Treponemataceae
O_Spirochaetales
C_Spirochaetia
P_Spirochaetes
S_Prevotella_oris
S_Campylobacter_gracilis
G_Parvimonas
F_Peptoniphilaceae
O_Tissierellales
C_Tissierellia
S_Alloprevotella_tannerae
S_Streptococcus_intermedius
S_Capnocytophaga_sp__HMT_336
G_Oribacterium
S_Oribacterium_sp__HMT_078
S_Leptotrichia_sp__HMT_223
S_Treponema_socranskii
S_Prevotella_sp__HMT_300
G_Dialister
S_Dialister_invisus
S_Schaalia_meyeri
G_Filifactor
S_Filifactor_alocis
S_Treponema_denticola
G_Peptostreptococcaceae__G_9_
S_Peptostreptococcaceae__G_9___Eubacterium__brachy
S_Saccharibacteria__TM7___G_1__bacterium_HMT_347_nov_95_759
S_Haemophilus_sp__HMT_036
S_Lachnospiraceae__G_2__bacterium_HMT_096
G_Olsenella
F_Atopobiaceae
O_Coriobacteriales
C_Coriobacteriia
S_Olsenella_sp__HMT_807
S_Saccharibacteria__TM7___G_1__bacterium_HMT_349
F_Weeksellaceae
−3
−2
−1
0
1
2
3
4
LDA SCORE (log 10)