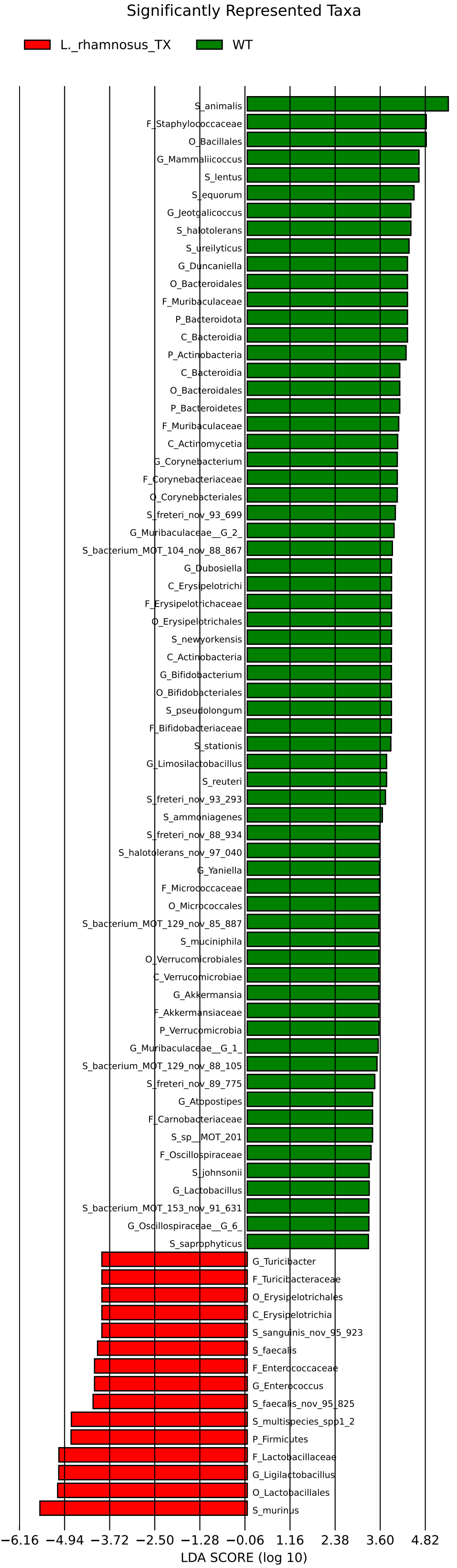
Significantly Represented Taxa
L._rhamnosus_TX
WT
S_animalis
F_Staphylococcaceae
O_Bacillales
G_Mammaliicoccus
S_lentus
S_equorum
G_Jeotgalicoccus
S_halotolerans
S_ureilyticus
G_Duncaniella
O_Bacteroidales
F_Muribaculaceae
P_Bacteroidota
C_Bacteroidia
P_Actinobacteria
C_Bacteroidia
O_Bacteroidales
P_Bacteroidetes
F_Muribaculaceae
C_Actinomycetia
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
S_freteri_nov_93_699
G_Muribaculaceae__G_2_
S_bacterium_MOT_104_nov_88_867
G_Dubosiella
C_Erysipelotrichi
F_Erysipelotrichaceae
O_Erysipelotrichales
S_newyorkensis
C_Actinobacteria
G_Bifidobacterium
O_Bifidobacteriales
S_pseudolongum
F_Bifidobacteriaceae
S_stationis
G_Limosilactobacillus
S_reuteri
S_freteri_nov_93_293
S_ammoniagenes
S_freteri_nov_88_934
S_halotolerans_nov_97_040
G_Yaniella
F_Micrococcaceae
O_Micrococcales
S_bacterium_MOT_129_nov_85_887
S_muciniphila
O_Verrucomicrobiales
C_Verrucomicrobiae
G_Akkermansia
F_Akkermansiaceae
P_Verrucomicrobia
G_Muribaculaceae__G_1_
S_bacterium_MOT_129_nov_88_105
S_freteri_nov_89_775
G_Atopostipes
F_Carnobacteriaceae
S_sp__MOT_201
F_Oscillospiraceae
S_johnsonii
G_Lactobacillus
S_bacterium_MOT_153_nov_91_631
G_Oscillospiraceae__G_6_
S_saprophyticus
G_Turicibacter
F_Turicibacteraceae
O_Erysipelotrichales
C_Erysipelotrichia
S_sanguinis_nov_95_923
S_faecalis
F_Enterococcaceae
G_Enterococcus
S_faecalis_nov_95_825
S_multispecies_spp1_2
P_Firmicutes
F_Lactobacillaceae
G_Ligilactobacillus
O_Lactobacillales
S_murinus
−6.16
−4.94
−3.72
−2.50
−1.28
−0.06
1.16
2.38
3.60
4.82
LDA SCORE (log 10)