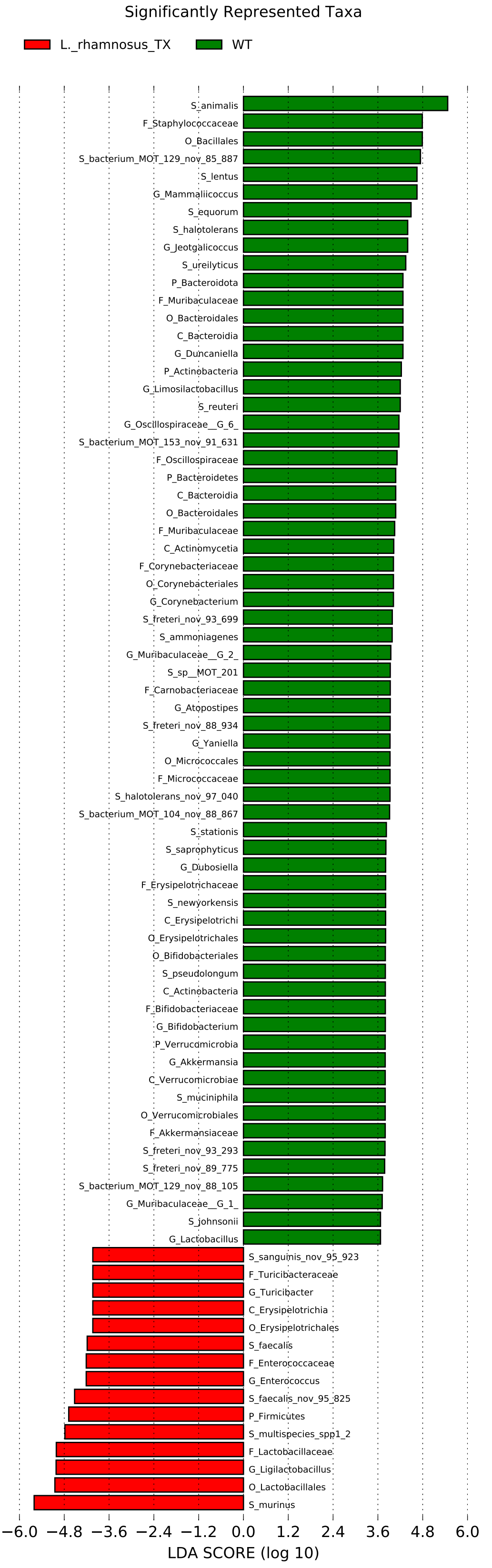
Significantly Represented Taxa
L._rhamnosus_TX
WT
S_animalis
F_Staphylococcaceae
O_Bacillales
S_bacterium_MOT_129_nov_85_887
S_lentus
G_Mammaliicoccus
S_equorum
S_halotolerans
G_Jeotgalicoccus
S_ureilyticus
P_Bacteroidota
F_Muribaculaceae
O_Bacteroidales
C_Bacteroidia
G_Duncaniella
P_Actinobacteria
G_Limosilactobacillus
S_reuteri
G_Oscillospiraceae__G_6_
S_bacterium_MOT_153_nov_91_631
F_Oscillospiraceae
P_Bacteroidetes
C_Bacteroidia
O_Bacteroidales
F_Muribaculaceae
C_Actinomycetia
F_Corynebacteriaceae
O_Corynebacteriales
G_Corynebacterium
S_freteri_nov_93_699
S_ammoniagenes
G_Muribaculaceae__G_2_
S_sp__MOT_201
F_Carnobacteriaceae
G_Atopostipes
S_freteri_nov_88_934
G_Yaniella
O_Micrococcales
F_Micrococcaceae
S_halotolerans_nov_97_040
S_bacterium_MOT_104_nov_88_867
S_stationis
S_saprophyticus
G_Dubosiella
F_Erysipelotrichaceae
S_newyorkensis
C_Erysipelotrichi
O_Erysipelotrichales
O_Bifidobacteriales
S_pseudolongum
C_Actinobacteria
F_Bifidobacteriaceae
G_Bifidobacterium
P_Verrucomicrobia
G_Akkermansia
C_Verrucomicrobiae
S_muciniphila
O_Verrucomicrobiales
F_Akkermansiaceae
S_freteri_nov_93_293
S_freteri_nov_89_775
S_bacterium_MOT_129_nov_88_105
G_Muribaculaceae__G_1_
S_johnsonii
G_Lactobacillus
S_sanguinis_nov_95_923
F_Turicibacteraceae
G_Turicibacter
C_Erysipelotrichia
O_Erysipelotrichales
S_faecalis
F_Enterococcaceae
G_Enterococcus
S_faecalis_nov_95_825
P_Firmicutes
S_multispecies_spp1_2
F_Lactobacillaceae
G_Ligilactobacillus
O_Lactobacillales
S_murinus
−6.0
−4.8
−3.6
−2.4
−1.2
0.0
1.2
2.4
3.6
4.8
6.0
LDA SCORE (log 10)