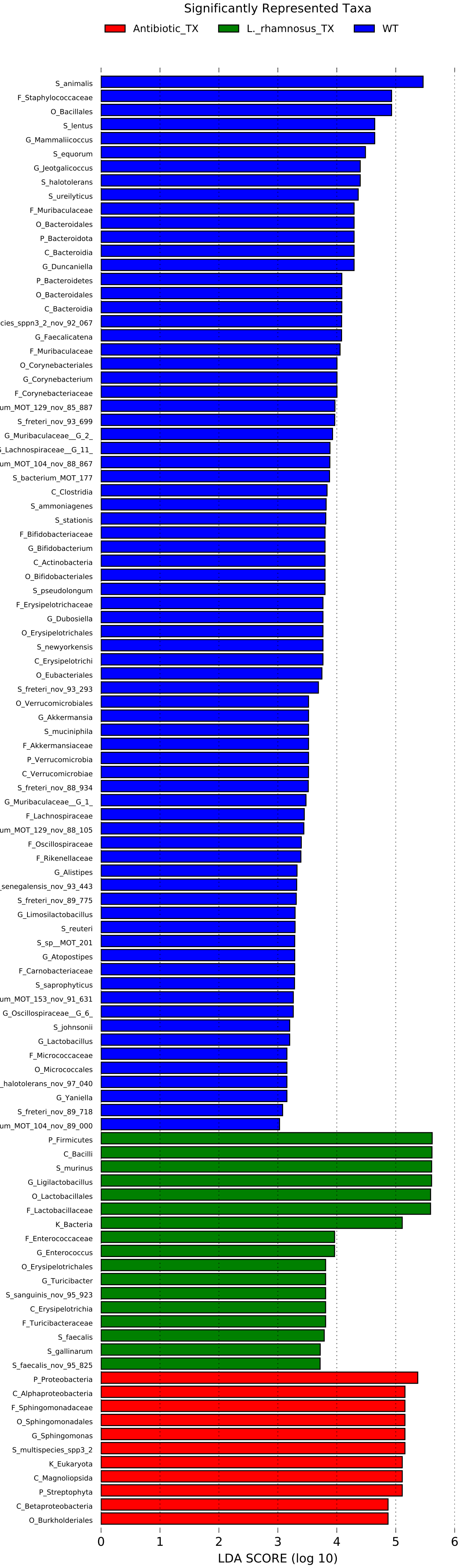
Significantly Represented Taxa
Antibiotic_TX
L._rhamnosus_TX
WT
S_animalis
F_Staphylococcaceae
O_Bacillales
S_lentus
G_Mammaliicoccus
S_equorum
G_Jeotgalicoccus
S_halotolerans
S_ureilyticus
F_Muribaculaceae
O_Bacteroidales
P_Bacteroidota
C_Bacteroidia
G_Duncaniella
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
cies_sppn3_2_nov_92_067
G_Faecalicatena
F_Muribaculaceae
O_Corynebacteriales
G_Corynebacterium
F_Corynebacteriaceae
um_MOT_129_nov_85_887
S_freteri_nov_93_699
G_Muribaculaceae__G_2_
_Lachnospiraceae__G_11_
um_MOT_104_nov_88_867
S_bacterium_MOT_177
C_Clostridia
S_ammoniagenes
S_stationis
F_Bifidobacteriaceae
G_Bifidobacterium
C_Actinobacteria
O_Bifidobacteriales
S_pseudolongum
F_Erysipelotrichaceae
G_Dubosiella
O_Erysipelotrichales
S_newyorkensis
C_Erysipelotrichi
O_Eubacteriales
S_freteri_nov_93_293
O_Verrucomicrobiales
G_Akkermansia
S_muciniphila
F_Akkermansiaceae
P_Verrucomicrobia
C_Verrucomicrobiae
S_freteri_nov_88_934
G_Muribaculaceae__G_1_
F_Lachnospiraceae
um_MOT_129_nov_88_105
F_Oscillospiraceae
F_Rikenellaceae
G_Alistipes
_senegalensis_nov_93_443
S_freteri_nov_89_775
G_Limosilactobacillus
S_reuteri
S_sp__MOT_201
G_Atopostipes
F_Carnobacteriaceae
S_saprophyticus
um_MOT_153_nov_91_631
G_Oscillospiraceae__G_6_
S_johnsonii
G_Lactobacillus
F_Micrococcaceae
O_Micrococcales
_halotolerans_nov_97_040
G_Yaniella
S_freteri_nov_89_718
um_MOT_104_nov_89_000
P_Firmicutes
C_Bacilli
S_murinus
G_Ligilactobacillus
O_Lactobacillales
F_Lactobacillaceae
K_Bacteria
F_Enterococcaceae
G_Enterococcus
O_Erysipelotrichales
G_Turicibacter
S_sanguinis_nov_95_923
C_Erysipelotrichia
F_Turicibacteraceae
S_faecalis
S_gallinarum
S_faecalis_nov_95_825
P_Proteobacteria
C_Alphaproteobacteria
F_Sphingomonadaceae
O_Sphingomonadales
G_Sphingomonas
S_multispecies_spp3_2
K_Eukaryota
C_Magnoliopsida
P_Streptophyta
C_Betaproteobacteria
O_Burkholderiales
0
1
2
3
4
5
6
LDA SCORE (log 10)