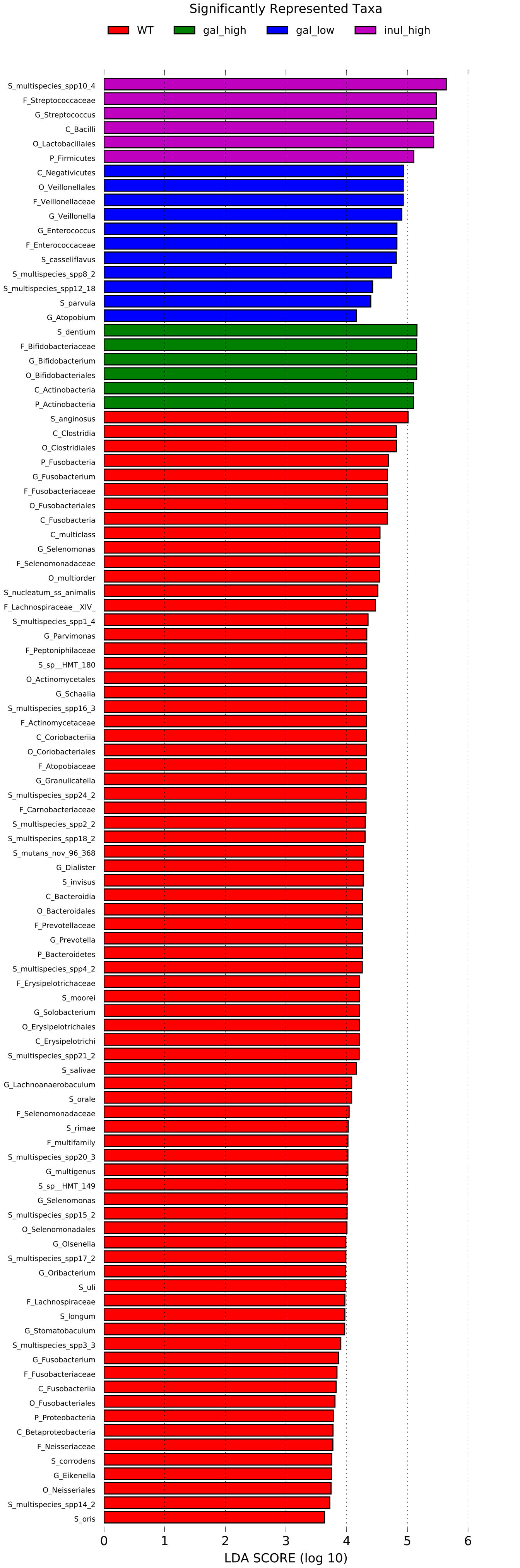
Significantly Represented Taxa
WT
gal_high
gal_low
inul_high
S_multispecies_spp10_4
F_Streptococcaceae
G_Streptococcus
C_Bacilli
O_Lactobacillales
P_Firmicutes
C_Negativicutes
O_Veillonellales
F_Veillonellaceae
G_Veillonella
G_Enterococcus
F_Enterococcaceae
S_casseliflavus
S_multispecies_spp8_2
S_multispecies_spp12_18
S_parvula
G_Atopobium
S_dentium
F_Bifidobacteriaceae
G_Bifidobacterium
O_Bifidobacteriales
C_Actinobacteria
P_Actinobacteria
S_anginosus
C_Clostridia
O_Clostridiales
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
C_multiclass
G_Selenomonas
F_Selenomonadaceae
O_multiorder
S_nucleatum_ss_animalis
F_Lachnospiraceae__XIV_
S_multispecies_spp1_4
G_Parvimonas
F_Peptoniphilaceae
S_sp__HMT_180
O_Actinomycetales
G_Schaalia
S_multispecies_spp16_3
F_Actinomycetaceae
C_Coriobacteriia
O_Coriobacteriales
F_Atopobiaceae
G_Granulicatella
S_multispecies_spp24_2
F_Carnobacteriaceae
S_multispecies_spp2_2
S_multispecies_spp18_2
S_mutans_nov_96_368
G_Dialister
S_invisus
C_Bacteroidia
O_Bacteroidales
F_Prevotellaceae
G_Prevotella
P_Bacteroidetes
S_multispecies_spp4_2
F_Erysipelotrichaceae
S_moorei
G_Solobacterium
O_Erysipelotrichales
C_Erysipelotrichi
S_multispecies_spp21_2
S_salivae
G_Lachnoanaerobaculum
S_orale
F_Selenomonadaceae
S_rimae
F_multifamily
S_multispecies_spp20_3
G_multigenus
S_sp__HMT_149
G_Selenomonas
S_multispecies_spp15_2
O_Selenomonadales
G_Olsenella
S_multispecies_spp17_2
G_Oribacterium
S_uli
F_Lachnospiraceae
S_longum
G_Stomatobaculum
S_multispecies_spp3_3
G_Fusobacterium
F_Fusobacteriaceae
C_Fusobacteriia
O_Fusobacteriales
P_Proteobacteria
C_Betaproteobacteria
F_Neisseriaceae
S_corrodens
G_Eikenella
O_Neisseriales
S_multispecies_spp14_2
S_oris
0
1
2
3
4
5
6
LDA SCORE (log 10)