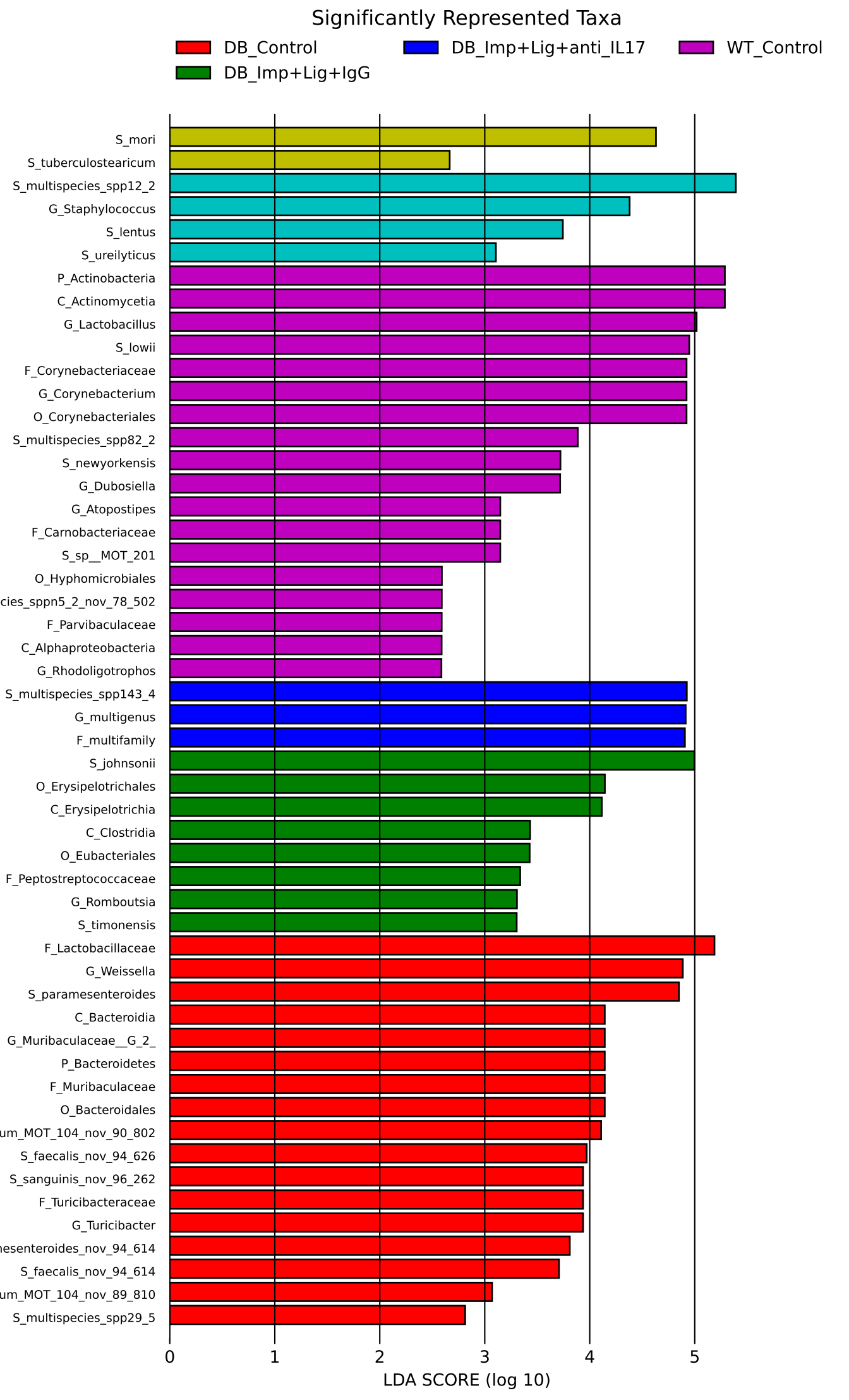
Significantly Represented Taxa
DB_Control
DB_Imp+Lig+anti_IL17
WT_Control
DB_Imp+Lig+IgG
S_mori
S_tuberculostearicum
S_multispecies_spp12_2
G_Staphylococcus
S_lentus
S_ureilyticus
P_Actinobacteria
C_Actinomycetia
G_Lactobacillus
S_lowii
F_Corynebacteriaceae
G_Corynebacterium
O_Corynebacteriales
S_multispecies_spp82_2
S_newyorkensis
G_Dubosiella
G_Atopostipes
F_Carnobacteriaceae
S_sp__MOT_201
O_Hyphomicrobiales
cies_sppn5_2_nov_78_502
F_Parvibaculaceae
C_Alphaproteobacteria
G_Rhodoligotrophos
S_multispecies_spp143_4
G_multigenus
F_multifamily
S_johnsonii
O_Erysipelotrichales
C_Erysipelotrichia
C_Clostridia
O_Eubacteriales
F_Peptostreptococcaceae
G_Romboutsia
S_timonensis
F_Lactobacillaceae
G_Weissella
S_paramesenteroides
C_Bacteroidia
G_Muribaculaceae__G_2_
P_Bacteroidetes
F_Muribaculaceae
O_Bacteroidales
um_MOT_104_nov_90_802
S_faecalis_nov_94_626
S_sanguinis_nov_96_262
F_Turicibacteraceae
G_Turicibacter
esenteroides_nov_94_614
S_faecalis_nov_94_614
um_MOT_104_nov_89_810
S_multispecies_spp29_5
0
1
2
3
4
5
LDA SCORE (log 10)