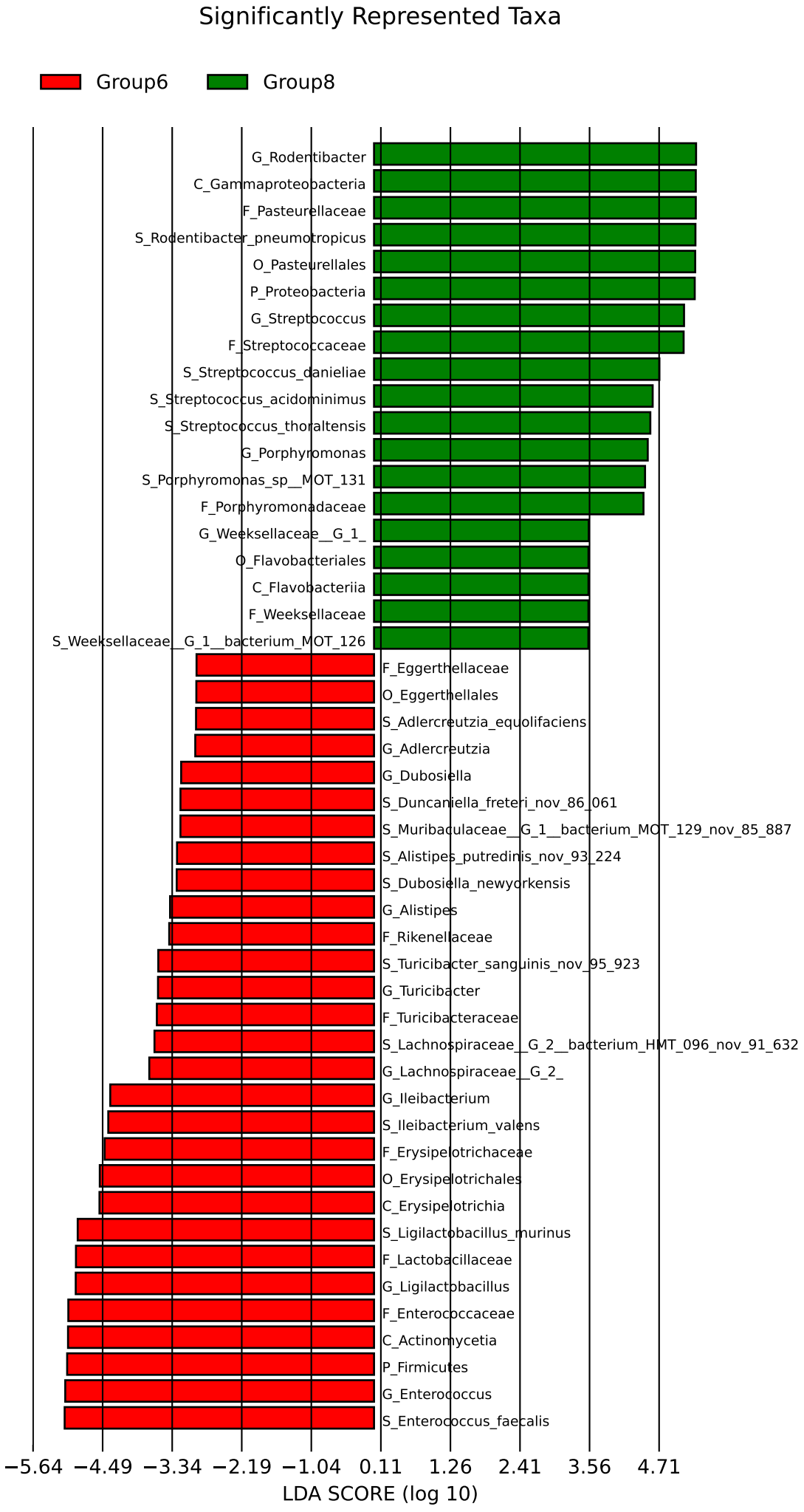
Significantly Represented Taxa
Group6
Group8
G_Rodentibacter
C_Gammaproteobacteria
F_Pasteurellaceae
S_Rodentibacter_pneumotropicus
O_Pasteurellales
P_Proteobacteria
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_danieliae
S_Streptococcus_acidominimus
S_Streptococcus_thoraltensis
G_Porphyromonas
S_Porphyromonas_sp__MOT_131
F_Porphyromonadaceae
G_Weeksellaceae__G_1_
O_Flavobacteriales
C_Flavobacteriia
F_Weeksellaceae
S_Weeksellaceae__G_1__bacterium_MOT_126
F_Eggerthellaceae
O_Eggerthellales
S_Adlercreutzia_equolifaciens
G_Adlercreutzia
G_Dubosiella
S_Duncaniella_freteri_nov_86_061
S_Muribaculaceae__G_1__bacterium_MOT_129_nov_85_887
S_Alistipes_putredinis_nov_93_224
S_Dubosiella_newyorkensis
G_Alistipes
F_Rikenellaceae
S_Turicibacter_sanguinis_nov_95_923
G_Turicibacter
F_Turicibacteraceae
S_Lachnospiraceae__G_2__bacterium_HMT_096_nov_91_632
G_Lachnospiraceae__G_2_
G_Ileibacterium
S_Ileibacterium_valens
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
S_Ligilactobacillus_murinus
F_Lactobacillaceae
G_Ligilactobacillus
F_Enterococcaceae
C_Actinomycetia
P_Firmicutes
G_Enterococcus
S_Enterococcus_faecalis
−5.64
−4.49
−3.34
−2.19
−1.04
0.11
1.26
2.41
3.56
4.71
LDA SCORE (log 10)