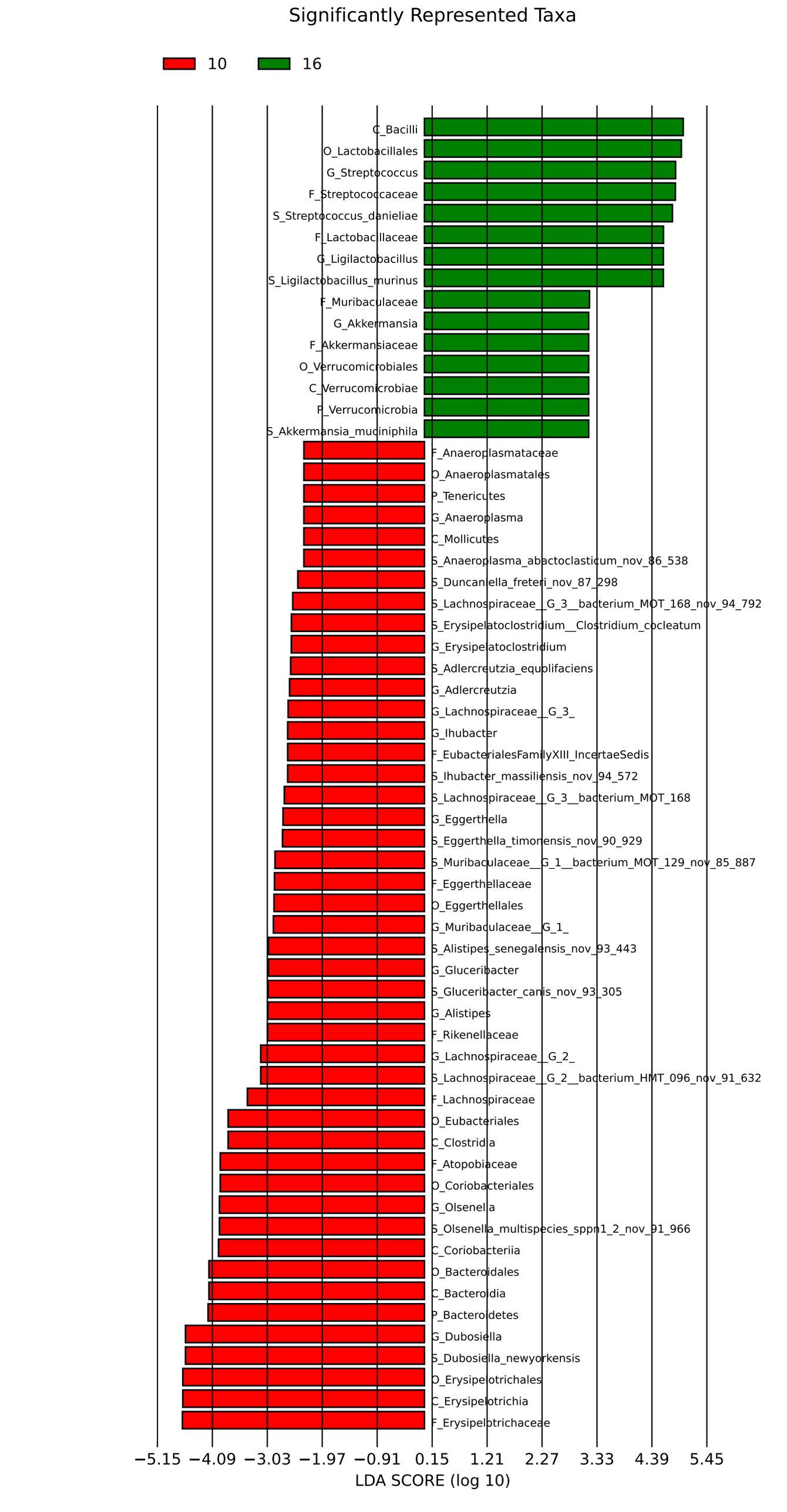

Significantly Represented Taxa
10
16
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_danieliae
F_Lactobacillaceae
G_Ligilactobacillus
S_Ligilactobacillus_murinus
F_Muribaculaceae
G_Akkermansia
F_Akkermansiaceae
O_Verrucomicrobiales
C_Verrucomicrobiae
P_Verrucomicrobia
S_Akkermansia_muciniphila
F_Anaeroplasmataceae
O_Anaeroplasmatales
P_Tenericutes
G_Anaeroplasma
C_Mollicutes
S_Anaeroplasma_abactoclasticum_nov_86_538
S_Duncaniella_freteri_nov_87_298
S_Lachnospiraceae__G_3__bacterium_MOT_168_nov_94_792
S_Erysipelatoclostridium__Clostridium_cocleatum
G_Erysipelatoclostridium
S_Adlercreutzia_equolifaciens
G_Adlercreutzia
G_Lachnospiraceae__G_3_
G_Ihubacter
F_EubacterialesFamilyXIII_IncertaeSedis
S_Ihubacter_massiliensis_nov_94_572
S_Lachnospiraceae__G_3__bacterium_MOT_168
G_Eggerthella
S_Eggerthella_timonensis_nov_90_929
S_Muribaculaceae__G_1__bacterium_MOT_129_nov_85_887
F_Eggerthellaceae
O_Eggerthellales
G_Muribaculaceae__G_1_
S_Alistipes_senegalensis_nov_93_443
G_Gluceribacter
S_Gluceribacter_canis_nov_93_305
G_Alistipes
F_Rikenellaceae
G_Lachnospiraceae__G_2_
S_Lachnospiraceae__G_2__bacterium_HMT_096_nov_91_632
F_Lachnospiraceae
O_Eubacteriales
C_Clostridia
F_Atopobiaceae
O_Coriobacteriales
G_Olsenella
S_Olsenella_multispecies_sppn1_2_nov_91_966
C_Coriobacteriia
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
G_Dubosiella
S_Dubosiella_newyorkensis
O_Erysipelotrichales
C_Erysipelotrichia
F_Erysipelotrichaceae
−5.15 −4.09 −3.03 −1.97 −0.91 0.15 1.21 2.27 3.33 4.39 5.45
LDA SCORE (log 10)