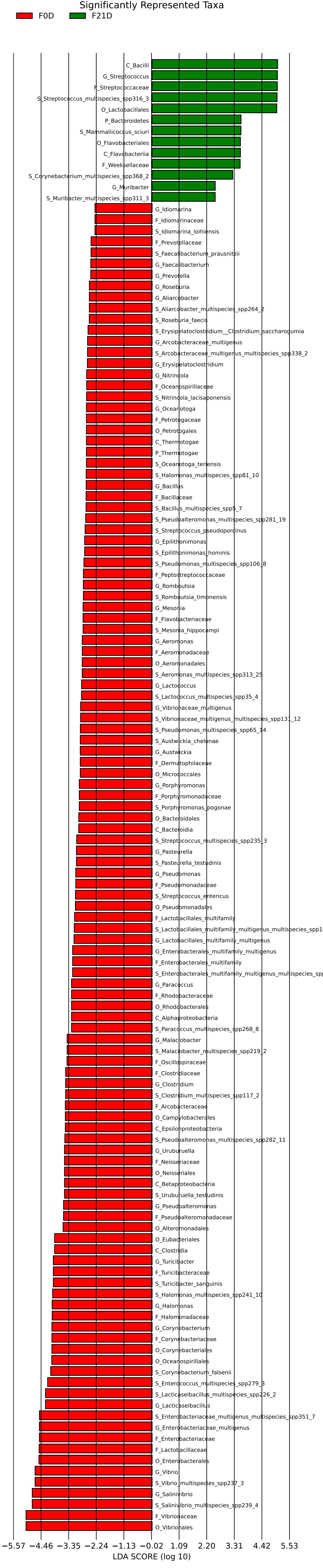
Significantly Represented Taxa
F0D
F21D
C_Bacilli
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_multispecies_spp316_3
O_Lactobacillales
P_Bacteroidetes
S_Mammaliicoccus_sciuri
O_Flavobacteriales
C_Flavobacteriia
F_Weeksellaceae
S_Corynebacterium_multispecies_spp368_2
G_Muribacter
S_Muribacter_multispecies_spp311_3
G_Idiomarina
F_Idiomarinaceae
S_Idiomarina_loihiensis
F_Prevotellaceae
S_Faecalibacterium_prausnitzii
G_Faecalibacterium
G_Prevotella
G_Roseburia
G_Aliarcobacter
S_Aliarcobacter_multispecies_spp264_2
S_Roseburia_faecis
S_Erysipelatoclostridium__Clostridium_saccharogumia
G_Arcobacteraceae_multigenus
S_Arcobacteraceae_multigenus_multispecies_spp338_2
G_Erysipelatoclostridium
G_Nitrincola
F_Oceanospirillaceae
S_Nitrincola_lacisaponensis
G_Oceanotoga
F_Petrotogaceae
O_Petrotogales
C_Thermotogae
P_Thermotogae
S_Oceanotoga_teriensis
S_Halomonas_multispecies_spp81_10
G_Bacillus
F_Bacillaceae
S_Bacillus_multispecies_spp5_7
S_Pseudoalteromonas_multispecies_spp281_19
S_Streptococcus_pseudoporcinus
G_Epilithonimonas
S_Epilithonimonas_hominis
S_Pseudomonas_multispecies_spp106_8
F_Peptostreptococcaceae
G_Romboutsia
S_Romboutsia_timonensis
G_Mesonia
F_Flavobacteriaceae
S_Mesonia_hippocampi
G_Aeromonas
F_Aeromonadaceae
O_Aeromonadales
S_Aeromonas_multispecies_spp313_25
G_Lactococcus
S_Lactococcus_multispecies_spp35_4
G_Vibrionaceae_multigenus
S_Vibrionaceae_multigenus_multispecies_spp131_12
S_Pseudomonas_multispecies_spp65_14
S_Austwickia_chelonae
G_Austwickia
F_Dermatophilaceae
O_Micrococcales
G_Porphyromonas
F_Porphyromonadaceae
S_Porphyromonas_pogonae
O_Bacteroidales
C_Bacteroidia
S_Streptococcus_multispecies_spp235_3
G_Pasteurella
S_Pasteurella_testudinis
G_Pseudomonas
F_Pseudomonadaceae
S_Streptococcus_entericus
O_Pseudomonadales
F_Lactobacillales_multifamily
S_Lactobacillales_multifamily_multigenus_multispecies_spp168
G_Lactobacillales_multifamily_multigenus
G_Enterobacterales_multifamily_multigenus
F_Enterobacterales_multifamily
S_Enterobacterales_multifamily_multigenus_multispecies_spp54
G_Paracoccus
F_Rhodobacteraceae
O_Rhodobacterales
C_Alphaproteobacteria
S_Paracoccus_multispecies_spp268_8
G_Malaciobacter
S_Malaciobacter_multispecies_spp219_2
F_Oscillospiraceae
F_Clostridiaceae
G_Clostridium
S_Clostridium_multispecies_spp117_2
F_Arcobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Pseudoalteromonas_multispecies_spp282_11
G_Uruburuella
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
S_Uruburuella_testudinis
G_Pseudoalteromonas
F_Pseudoalteromonadaceae
O_Alteromonadales
O_Eubacteriales
C_Clostridia
G_Turicibacter
F_Turicibacteraceae
S_Turicibacter_sanguinis
S_Halomonas_multispecies_spp241_10
G_Halomonas
F_Halomonadaceae
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
O_Oceanospirillales
S_Corynebacterium_falsenii
S_Enterococcus_multispecies_spp279_3
S_Lacticaseibacillus_multispecies_spp226_2
G_Lacticaseibacillus
S_Enterobacteriaceae_multigenus_multispecies_spp351_7
G_Enterobacteriaceae_multigenus
F_Enterobacteriaceae
F_Lactobacillaceae
O_Enterobacterales
G_Vibrio
S_Vibrio_multispecies_spp237_3
G_Salinivibrio
S_Salinivibrio_multispecies_spp239_4
F_Vibrionaceae
O_Vibrionales
−5.57
−4.46
−3.35
−2.24
−1.13
−0.02
1.09
2.20
3.31
4.42
5.53
LDA SCORE (log 10)