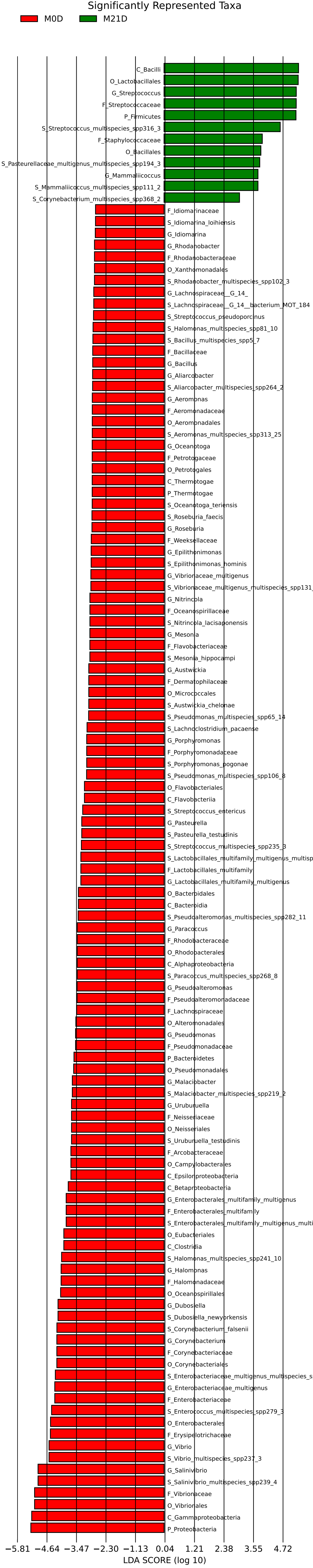
Significantly Represented Taxa
M0D
M21D
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
P_Firmicutes
S_Streptococcus_multispecies_spp316_3
F_Staphylococcaceae
O_Bacillales
S_Pasteurellaceae_multigenus_multispecies_spp194_3
G_Mammaliicoccus
S_Mammaliicoccus_multispecies_spp111_2
S_Corynebacterium_multispecies_spp368_2
F_Idiomarinaceae
S_Idiomarina_loihiensis
G_Idiomarina
G_Rhodanobacter
F_Rhodanobacteraceae
O_Xanthomonadales
S_Rhodanobacter_multispecies_spp102_3
G_Lachnospiraceae__G_14_
S_Lachnospiraceae__G_14__bacterium_MOT_184
S_Streptococcus_pseudoporcinus
S_Halomonas_multispecies_spp81_10
S_Bacillus_multispecies_spp5_7
F_Bacillaceae
G_Bacillus
G_Aliarcobacter
S_Aliarcobacter_multispecies_spp264_2
G_Aeromonas
F_Aeromonadaceae
O_Aeromonadales
S_Aeromonas_multispecies_spp313_25
G_Oceanotoga
F_Petrotogaceae
O_Petrotogales
C_Thermotogae
P_Thermotogae
S_Oceanotoga_teriensis
S_Roseburia_faecis
G_Roseburia
F_Weeksellaceae
G_Epilithonimonas
S_Epilithonimonas_hominis
G_Vibrionaceae_multigenus
S_Vibrionaceae_multigenus_multispecies_spp131_12
G_Nitrincola
F_Oceanospirillaceae
S_Nitrincola_lacisaponensis
G_Mesonia
F_Flavobacteriaceae
S_Mesonia_hippocampi
G_Austwickia
F_Dermatophilaceae
O_Micrococcales
S_Austwickia_chelonae
S_Pseudomonas_multispecies_spp65_14
S_Lachnoclostridium_pacaense
G_Porphyromonas
F_Porphyromonadaceae
S_Porphyromonas_pogonae
S_Pseudomonas_multispecies_spp106_8
O_Flavobacteriales
C_Flavobacteriia
S_Streptococcus_entericus
G_Pasteurella
S_Pasteurella_testudinis
S_Streptococcus_multispecies_spp235_3
S_Lactobacillales_multifamily_multigenus_multispecies_spp168
F_Lactobacillales_multifamily
G_Lactobacillales_multifamily_multigenus
O_Bacteroidales
C_Bacteroidia
S_Pseudoalteromonas_multispecies_spp282_11
G_Paracoccus
F_Rhodobacteraceae
O_Rhodobacterales
C_Alphaproteobacteria
S_Paracoccus_multispecies_spp268_8
G_Pseudoalteromonas
F_Pseudoalteromonadaceae
F_Lachnospiraceae
O_Alteromonadales
G_Pseudomonas
F_Pseudomonadaceae
P_Bacteroidetes
O_Pseudomonadales
G_Malaciobacter
S_Malaciobacter_multispecies_spp219_2
G_Uruburuella
F_Neisseriaceae
O_Neisseriales
S_Uruburuella_testudinis
F_Arcobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
C_Betaproteobacteria
G_Enterobacterales_multifamily_multigenus
F_Enterobacterales_multifamily
S_Enterobacterales_multifamily_multigenus_multispecies_spp54
O_Eubacteriales
C_Clostridia
S_Halomonas_multispecies_spp241_10
G_Halomonas
F_Halomonadaceae
O_Oceanospirillales
G_Dubosiella
S_Dubosiella_newyorkensis
S_Corynebacterium_falsenii
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
S_Enterobacteriaceae_multigenus_multispecies_spp351_7
G_Enterobacteriaceae_multigenus
F_Enterobacteriaceae
S_Enterococcus_multispecies_spp279_3
O_Enterobacterales
F_Erysipelotrichaceae
G_Vibrio
S_Vibrio_multispecies_spp237_3
G_Salinivibrio
S_Salinivibrio_multispecies_spp239_4
F_Vibrionaceae
O_Vibrionales
C_Gammaproteobacteria
P_Proteobacteria
−5.81
−4.64
−3.47
−2.30
−1.13
0.04
1.21
2.38
3.55
4.72
LDA SCORE (log 10)