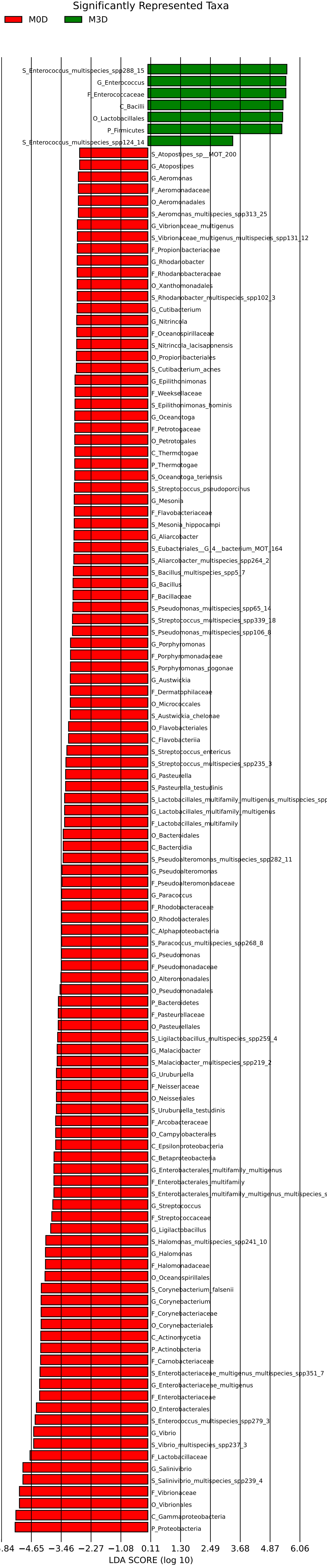
Significantly Represented Taxa
M0D
M3D
S_Enterococcus_multispecies_spp288_15
G_Enterococcus
F_Enterococcaceae
C_Bacilli
O_Lactobacillales
P_Firmicutes
S_Enterococcus_multispecies_spp124_14
S_Atopostipes_sp__MOT_200
G_Atopostipes
G_Aeromonas
F_Aeromonadaceae
O_Aeromonadales
S_Aeromonas_multispecies_spp313_25
G_Vibrionaceae_multigenus
S_Vibrionaceae_multigenus_multispecies_spp131_12
F_Propionibacteriaceae
G_Rhodanobacter
F_Rhodanobacteraceae
O_Xanthomonadales
S_Rhodanobacter_multispecies_spp102_3
G_Cutibacterium
G_Nitrincola
F_Oceanospirillaceae
S_Nitrincola_lacisaponensis
O_Propionibacteriales
S_Cutibacterium_acnes
G_Epilithonimonas
F_Weeksellaceae
S_Epilithonimonas_hominis
G_Oceanotoga
F_Petrotogaceae
O_Petrotogales
C_Thermotogae
P_Thermotogae
S_Oceanotoga_teriensis
S_Streptococcus_pseudoporcinus
G_Mesonia
F_Flavobacteriaceae
S_Mesonia_hippocampi
G_Aliarcobacter
S_Eubacteriales__G_4__bacterium_MOT_164
S_Aliarcobacter_multispecies_spp264_2
S_Bacillus_multispecies_spp5_7
G_Bacillus
F_Bacillaceae
S_Pseudomonas_multispecies_spp65_14
S_Streptococcus_multispecies_spp339_18
S_Pseudomonas_multispecies_spp106_8
G_Porphyromonas
F_Porphyromonadaceae
S_Porphyromonas_pogonae
G_Austwickia
F_Dermatophilaceae
O_Micrococcales
S_Austwickia_chelonae
O_Flavobacteriales
C_Flavobacteriia
S_Streptococcus_entericus
S_Streptococcus_multispecies_spp235_3
G_Pasteurella
S_Pasteurella_testudinis
S_Lactobacillales_multifamily_multigenus_multispecies_spp168_
G_Lactobacillales_multifamily_multigenus
F_Lactobacillales_multifamily
O_Bacteroidales
C_Bacteroidia
S_Pseudoalteromonas_multispecies_spp282_11
G_Pseudoalteromonas
F_Pseudoalteromonadaceae
G_Paracoccus
F_Rhodobacteraceae
O_Rhodobacterales
C_Alphaproteobacteria
S_Paracoccus_multispecies_spp268_8
G_Pseudomonas
F_Pseudomonadaceae
O_Alteromonadales
O_Pseudomonadales
P_Bacteroidetes
F_Pasteurellaceae
O_Pasteurellales
S_Ligilactobacillus_multispecies_spp259_4
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
G_Streptococcus
F_Streptococcaceae
G_Ligilactobacillus
S_Halomonas_multispecies_spp241_10
G_Halomonas
F_Halomonadaceae
O_Oceanospirillales
S_Corynebacterium_falsenii
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
C_Actinomycetia
P_Actinobacteria
F_Carnobacteriaceae
S_Enterobacteriaceae_multigenus_multispecies_spp351_7
G_Enterobacteriaceae_multigenus
F_Enterobacteriaceae
O_Enterobacterales
S_Enterococcus_multispecies_spp279_3
G_Vibrio
S_Vibrio_multispecies_spp237_3
F_Lactobacillaceae
G_Salinivibrio
S_Salinivibrio_multispecies_spp239_4
F_Vibrionaceae
O_Vibrionales
C_Gammaproteobacteria
P_Proteobacteria
−5.84
−4.65
−3.46
−2.27
−1.08
0.11
1.30
2.49
3.68
4.87
6.06
LDA SCORE (log 10)