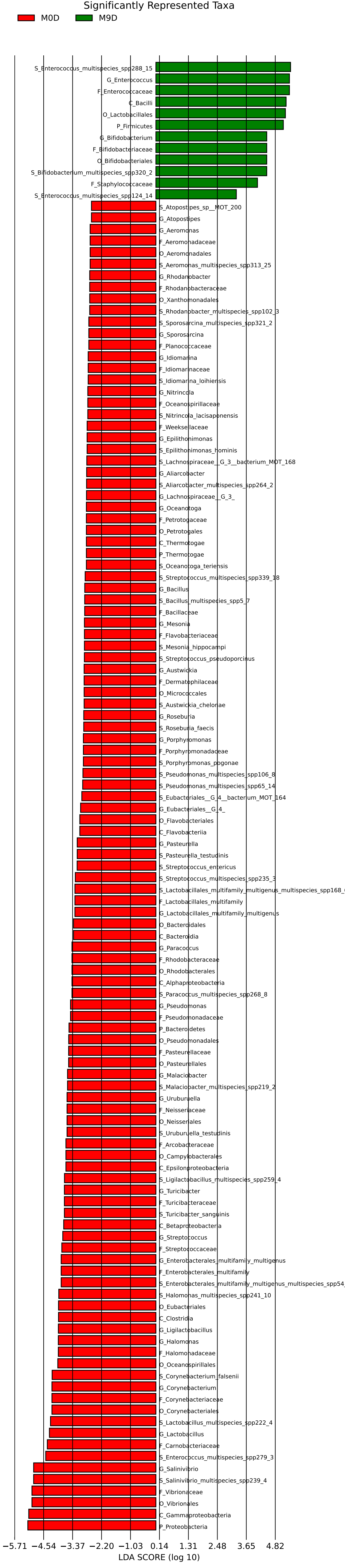
Significantly Represented Taxa
M0D
M9D
S_Enterococcus_multispecies_spp288_15
G_Enterococcus
F_Enterococcaceae
C_Bacilli
O_Lactobacillales
P_Firmicutes
G_Bifidobacterium
F_Bifidobacteriaceae
O_Bifidobacteriales
S_Bifidobacterium_multispecies_spp320_2
F_Staphylococcaceae
S_Enterococcus_multispecies_spp124_14
S_Atopostipes_sp__MOT_200
G_Atopostipes
G_Aeromonas
F_Aeromonadaceae
O_Aeromonadales
S_Aeromonas_multispecies_spp313_25
G_Rhodanobacter
F_Rhodanobacteraceae
O_Xanthomonadales
S_Rhodanobacter_multispecies_spp102_3
S_Sporosarcina_multispecies_spp321_2
G_Sporosarcina
F_Planococcaceae
G_Idiomarina
F_Idiomarinaceae
S_Idiomarina_loihiensis
G_Nitrincola
F_Oceanospirillaceae
S_Nitrincola_lacisaponensis
F_Weeksellaceae
G_Epilithonimonas
S_Epilithonimonas_hominis
S_Lachnospiraceae__G_3__bacterium_MOT_168
G_Aliarcobacter
S_Aliarcobacter_multispecies_spp264_2
G_Lachnospiraceae__G_3_
G_Oceanotoga
F_Petrotogaceae
O_Petrotogales
C_Thermotogae
P_Thermotogae
S_Oceanotoga_teriensis
S_Streptococcus_multispecies_spp339_18
G_Bacillus
S_Bacillus_multispecies_spp5_7
F_Bacillaceae
G_Mesonia
F_Flavobacteriaceae
S_Mesonia_hippocampi
S_Streptococcus_pseudoporcinus
G_Austwickia
F_Dermatophilaceae
O_Micrococcales
S_Austwickia_chelonae
G_Roseburia
S_Roseburia_faecis
G_Porphyromonas
F_Porphyromonadaceae
S_Porphyromonas_pogonae
S_Pseudomonas_multispecies_spp106_8
S_Pseudomonas_multispecies_spp65_14
S_Eubacteriales__G_4__bacterium_MOT_164
G_Eubacteriales__G_4_
O_Flavobacteriales
C_Flavobacteriia
G_Pasteurella
S_Pasteurella_testudinis
S_Streptococcus_entericus
S_Streptococcus_multispecies_spp235_3
S_Lactobacillales_multifamily_multigenus_multispecies_spp168_
F_Lactobacillales_multifamily
G_Lactobacillales_multifamily_multigenus
O_Bacteroidales
C_Bacteroidia
G_Paracoccus
F_Rhodobacteraceae
O_Rhodobacterales
C_Alphaproteobacteria
S_Paracoccus_multispecies_spp268_8
G_Pseudomonas
F_Pseudomonadaceae
P_Bacteroidetes
O_Pseudomonadales
F_Pasteurellaceae
O_Pasteurellales
G_Malaciobacter
S_Malaciobacter_multispecies_spp219_2
G_Uruburuella
F_Neisseriaceae
O_Neisseriales
S_Uruburuella_testudinis
F_Arcobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Ligilactobacillus_multispecies_spp259_4
G_Turicibacter
F_Turicibacteraceae
S_Turicibacter_sanguinis
C_Betaproteobacteria
G_Streptococcus
F_Streptococcaceae
G_Enterobacterales_multifamily_multigenus
F_Enterobacterales_multifamily
S_Enterobacterales_multifamily_multigenus_multispecies_spp54
S_Halomonas_multispecies_spp241_10
O_Eubacteriales
C_Clostridia
G_Ligilactobacillus
G_Halomonas
F_Halomonadaceae
O_Oceanospirillales
S_Corynebacterium_falsenii
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
S_Lactobacillus_multispecies_spp222_4
G_Lactobacillus
F_Carnobacteriaceae
S_Enterococcus_multispecies_spp279_3
G_Salinivibrio
S_Salinivibrio_multispecies_spp239_4
F_Vibrionaceae
O_Vibrionales
C_Gammaproteobacteria
P_Proteobacteria
−5.71
−4.54
−3.37
−2.20
−1.03
0.14
1.31
2.48
3.65
4.82
LDA SCORE (log 10)