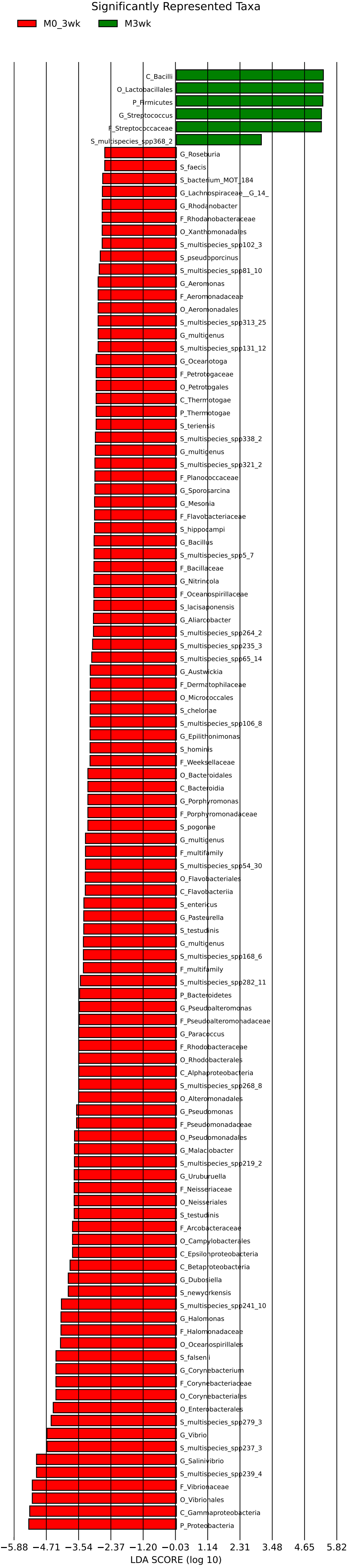
Significantly Represented Taxa
M0_3wk
M3wk
C_Bacilli
O_Lactobacillales
P_Firmicutes
G_Streptococcus
F_Streptococcaceae
S_multispecies_spp368_2
G_Roseburia
S_faecis
S_bacterium_MOT_184
G_Lachnospiraceae__G_14_
G_Rhodanobacter
F_Rhodanobacteraceae
O_Xanthomonadales
S_multispecies_spp102_3
S_pseudoporcinus
S_multispecies_spp81_10
G_Aeromonas
F_Aeromonadaceae
O_Aeromonadales
S_multispecies_spp313_25
G_multigenus
S_multispecies_spp131_12
G_Oceanotoga
F_Petrotogaceae
O_Petrotogales
C_Thermotogae
P_Thermotogae
S_teriensis
S_multispecies_spp338_2
G_multigenus
S_multispecies_spp321_2
F_Planococcaceae
G_Sporosarcina
G_Mesonia
F_Flavobacteriaceae
S_hippocampi
G_Bacillus
S_multispecies_spp5_7
F_Bacillaceae
G_Nitrincola
F_Oceanospirillaceae
S_lacisaponensis
G_Aliarcobacter
S_multispecies_spp264_2
S_multispecies_spp235_3
S_multispecies_spp65_14
G_Austwickia
F_Dermatophilaceae
O_Micrococcales
S_cheloae
S_multispecies_spp106_8
G_Epilithonimonas
S_hominis
F_Weeksellaceae
O_Bacteroidales
C_Bacteroidia
G_Porphyromonas
F_Porphyromonadaceae
S_pogonae
G_multigenus
F_multifamily
S_multispecies_spp54_30
O_Flavobacteriales
C_Flavobacteriia
S_enterica
G_Pasteurella
S_testudinis
G_multigenus
S_multispecies_spp168_6
F_multifamily
S_multispecies_spp282_11
P_Bacteroidetes
G_Pseudoalteromonas
F_Pseudoalteromonadaceae
G_Paracoccus
F_Rhodobacteraceae
O_Rhodobacterales
C_Alphaproteobacteria
S_multispecies_spp268_8
O_Alteromonadales
G_Pseudomonas
F_Pseudomonadaceae
O_Pseudomonadales
G_Malaciobacter
S_multispecies_spp219_2
G_Uruburuella
F_Neisseriaceae
O_Neisseriales
S_testudinis
F_Arcobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
C_Betaproteobacteria
G_Dubosiella
S_newyorkensis
S_multispecies_spp241_10
G_Halomonas
F_Halomonadaceae
O_Oceanospirillales
S_falsenii
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
O_Enterobacterales
S_multispecies_spp279_3
G_Vibrio
S_multispecies_spp237_3
G_Salinivibrio
S_multispecies_spp239_4
F_Vibrionaceae
O_Vibrionales
C_Gammaproteobacteria
P_Proteobacteria
−5.88
−4.71
−3.54
−2.37
−1.20
−0.03
1.14
2.31
3.48
4.65
5.82
LDA SCORE (log 10)