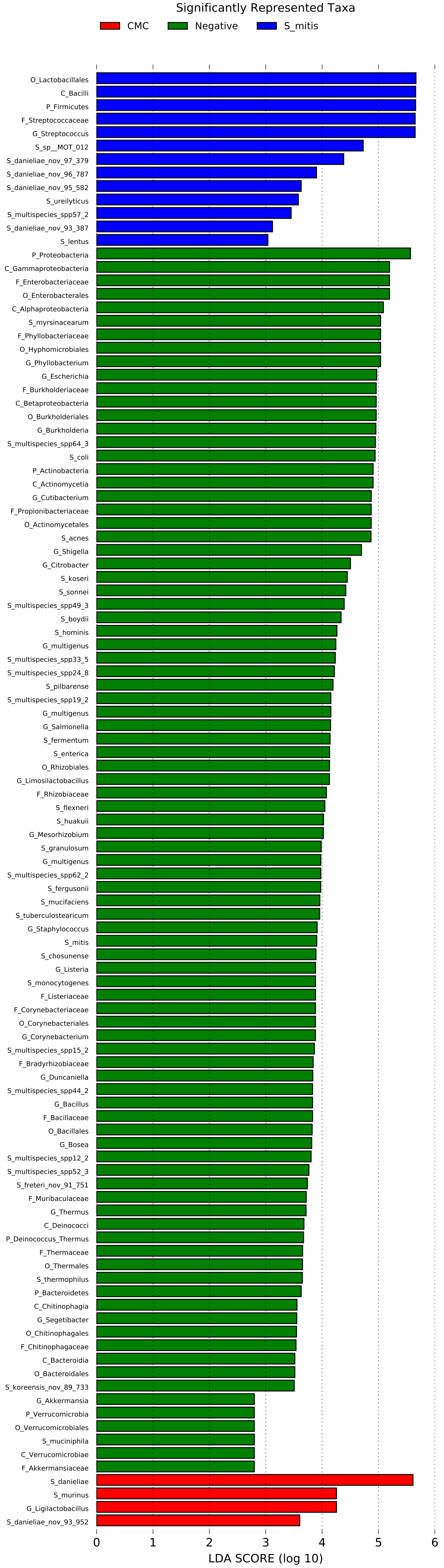
Significantly Represented Taxa
CMC
Negative
S_mitis
O_Lactobacillales
C_Bacilli
P_Firmicutes
F_Streptococcaceae
G_Streptococcus
S_sp__MOT_012
S_danieliae_nov_97_379
S_danieliae_nov_96_787
S_danieliae_nov_95_582
S_ureilyticus
S_multispecies_spp57_2
S_danieliae_nov_93_387
S_lentus
P_Proteobacteria
C_Gammaproteobacteria
F_Enterobacteriaceae
O_Enterobacterales
C_Alphaproteobacteria
S_myrsinacearum
F_Phyllobacteriaceae
O_Hyphomicrobiales
G_Phyllobacterium
G_Escherichia
F_Burkholderiaceae
C_Betaproteobacteria
O_Burkholderiales
G_Burkholderia
S_multispecies_spp64_3
S_coli
P_Actinobacteria
C_Actinomycetia
G_Cutibacterium
F_Propionibacteriaceae
O_Actinomycetales
S_acnes
G_Shigella
G_Citrobacter
S_koseri
S_sonnei
S_multispecies_spp49_3
S_boydii
S_hominis
G_multigenus
S_multispecies_spp33_5
S_multispecies_spp24_8
S_pilbarense
S_multispecies_spp19_2
G_multigenus
G_Salmonella
S_fermentum
S_enterica
O_Rhizobiales
G_Limosilactobacillus
F_Rhizobiaceae
S_flexneri
S_huakuii
G_Mesorhizobium
S_granulosum
G_multigenus
S_multispecies_spp62_2
S_fergusonii
S_mucifaciens
S_tuberculostearicum
G_Staphylococcus
S_mitis
S_chosunense
G_Listeria
S_monocytogenes
F_Listeriaceae
F_Corynebacteriaceae
O_Corynebacteriales
G_Corynebacterium
S_multispecies_spp15_2
F_Bradyrhizobiaceae
G_Duncaniella
S_multispecies_spp44_2
G_Bacillus
F_Bacillaceae
O_Bacillales
G_Bosea
S_multispecies_spp12_2
S_multispecies_spp52_3
S_freteri_nov_91_751
F_Muribaculaceae
G_Thermus
C_Deinococci
P_Deinococcus_Thermus
F_Thermaceae
O_Thermales
S_thermophilus
P_Bacteroidetes
C_Chitinophagia
G_Segetibacter
O_Chitinophagales
F_Chitinophagaceae
C_Bacteroidia
O_Bacteroidales
S_koreensis_nov_89_733
G_Akkermansia
P_Verrucomicrobia
O_Verrucomicrobiales
S_muciniphila
C_Verrucomicrobiae
F_Akkermansiaceae
S_danieliae
S_murinus
G_Ligilactobacillus
S_danieliae_nov_93_952
0
1
2
3
4
5
6
LDA SCORE (log 10)