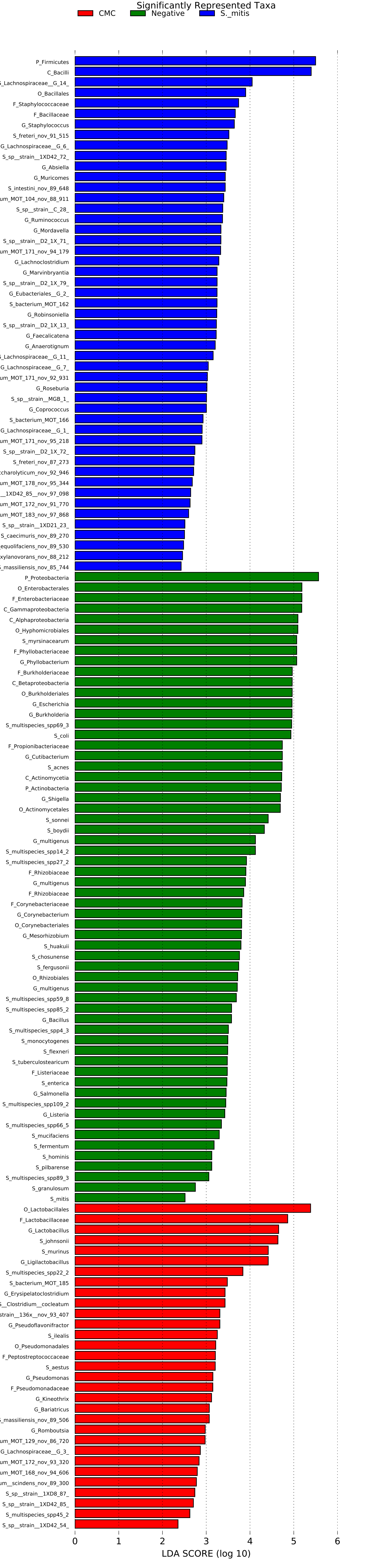
Significantly Represented Taxa
CMC
Negative
S._mitis
P_Firmicutes
C_Bacilli
G_Lachnospiraceae__G_14_
O_Bacillales
F_Staphylococcaceae
F_Bacillaceae
G_Staphylococcus
S_freteri_nov_91_515
G_Lachnospiraceae__G_6_
S_sp__strain__1XD42_72_
G_Absiella
G_Muricomes
S_intestini_nov_89_648
um_MOT_104_nov_88_911
S_sp__strain__C_28_
G_Ruminococcus
G_Mordavella
S_sp__strain__D2_1X_71_
um_MOT_171_nov_94_179
G_Lachnoclostridium
G_Marvinbryantia
S_sp__strain__D2_1X_79_
G_Eubacteriales__G_2_
S_bacterium_MOT_162
G_Robinsoniella
S_sp__strain__D2_1X_13_
G_Faecalicatena
G_Anaerotignum
G_Lachnospiraceae__G_11_
G_Lachnospiraceae__G_7_
um_MOT_171_nov_92_931
G_Roseburia
S_sp__strain__MGB_1_
G_Coprococcus
S_bacterium_MOT_166
G_Lachnospiraceae__G_1_
um_MOT_171_nov_95_218
S_sp__strain__D2_1X_72_
S_freteri_nov_87_273
ccharolyticum_nov_92_946
um_MOT_178_nov_95_344
__1XD42_85__nov_97_098
um_MOT_172_nov_91_770
um_MOT_183_nov_97_868
S_sp__strain__1XD21_23_
S_caecimuris_nov_89_270
equolifaciens_nov_89_530
xylanovorans_nov_88_212
_massiliensis_nov_85_744
P_Proteobacteria
O_Enterobacterales
F_Enterobacteriaceae
C_Gammaproteobacteria
C_Alphaproteobacteria
O_Hyphomicrobiales
S_myrsinacearum
F_Phyllobacteriaceae
G_Phyllobacterium
F_Burkholderiaceae
C_Betaproteobacteria
O_Burkholderiales
G_Escherichia
G_Burkholderia
S_multispecies_spp69_3
S_coli
F_Propionibacteriaceae
G_Cutibacterium
S_acnes
C_Actinomycetia
P_Actinobacteria
G_Shigella
O_Actinomycetales
S_sonnei
S_boydii
G_multigenus
S_multispecies_spp14_2
S_multispecies_spp27_2
F_Rhizobiaceae
G_multigenus
F_Rhizobiaceae
F_Corynebacteriaceae
G_Corynebacterium
O_Corynebacteriales
G_Mesorhizobium
S_huakuii
S_chosunense
S_fergusonii
O_Rhizobiales
G_multigenus
S_multispecies_spp59_8
S_multispecies_spp85_2
G_Bacillus
S_multispecies_spp4_3
S_monocytogenes
S_flexneri
S_tuberculostearicum
F_Listeriaceae
S_enterica
G_Salmonella
S_multispecies_spp109_2
G_Listeria
S_multispecies_spp66_5
S_mucifaciens
S_fermentum
S_hominis
S_pilbarense
S_multispecies_spp89_3
S_granulosum
S_mitis
O_Lactobacillales
F_Lactobacillaceae
G_Lactobacillus
S_johnsonii
S_murinus
G_Ligilactobacillus
S_multispecies_spp22_2
S_bacterium_MOT_185
G_Erysipelatoclostridium
S__Clostridium__cocleatum
strain__136x__nov_93_407
G_Pseudoflavonifractor
S_ilealis
O_Pseudomonadales
F_Peptostreptococcaceae
S_aestus
G_Pseudomonas
F_Pseudomonadaceae
G_Kineothrix
G_Bariatricus
_massiliensis_nov_89_506
G_Romboutsia
um_MOT_129_nov_86_720
G_Lachnospiraceae__G_3_
um_MOT_172_nov_93_320
um_MOT_168_nov_94_606
um__scindens_nov_89_300
S_sp__strain__1XD8_87_
S_sp__strain__1XD42_85_
S_multispecies_spp45_2
S_sp__strain__1XD42_54_
0
1
2
3
4
5
6
LDA SCORE (log 10)