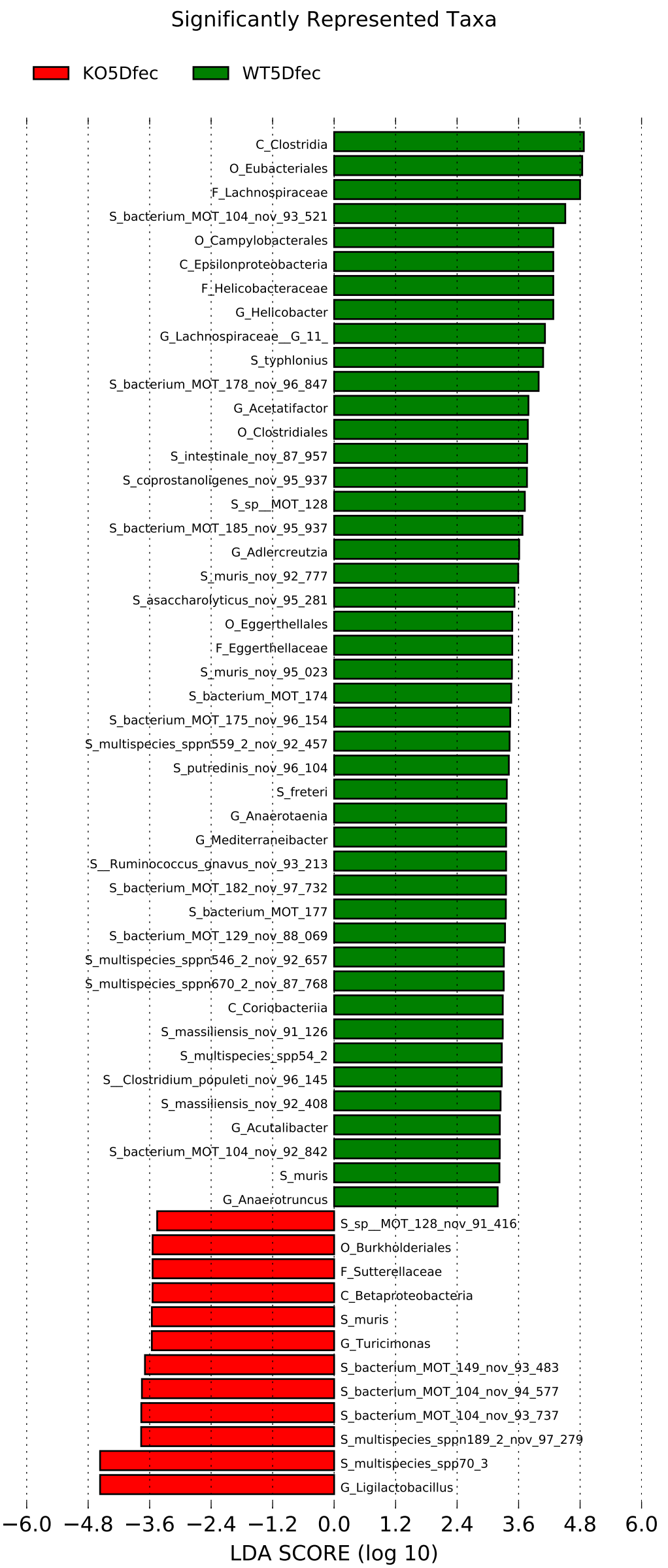
Significantly Represented Taxa
KO5Dfec
WT5Dfec
C_Clostridia
O_Eubacteriales
F_Lachnospiraceae
S_bacterium_MOT_104_nov_93_521
O_Campylobacterales
C_Epsilonproteobacteria
F_Helicobacteraceae
G_Helicobacter
G_Lachnospiraceae__G_11_
S_typhlonius
S_bacterium_MOT_178_nov_96_847
G_Acetatifactor
O_Clostridiales
S_intestinale_nov_87_957
S_coprostanoligenes_nov_95_937
S_sp__MOT_128
S_bacterium_MOT_185_nov_95_937
G_Adlercreutzia
S_muris_nov_92_777
S_asaccharolyticus_nov_95_281
O_Eggerthellales
F_Eggerthellaceae
S_muris_nov_95_023
S_bacterium_MOT_174
S_bacterium_MOT_175_nov_96_154
S_multispecies_sppn559_2_nov_92_457
S_putredinis_nov_96_104
S_freteri
G_Anaerotaenia
G_Mediterraneibacter
S__Ruminococcus_gnavus_nov_93_213
S_bacterium_MOT_182_nov_97_732
S_bacterium_MOT_177
S_bacterium_MOT_129_nov_88_069
S_multispecies_sppn546_2_nov_92_657
S_multispecies_sppn670_2_nov_87_768
C_Coriobacteriia
S_massiliensis_nov_91_126
S_multispecies_spp54_2
S__Clostridium_populeti_nov_96_145
S_massiliensis_nov_92_408
G_Acutalibacter
S_bacterium_MOT_104_nov_92_842
S_muris
G_Anaerotruncus
S_sp__MOT_128_nov_91_416
O_Burkholderiales
F_Sutterellaceae
C_Betaproteobacteria
S_muris
G_Turicimonas
S_bacterium_MOT_149_nov_93_483
S_bacterium_MOT_104_nov_94_577
S_bacterium_MOT_104_nov_93_737
S_multispecies_sppn189_2_nov_97_279
S_multispecies_spp70_3
G_Ligilactobacillus
−6.0
−4.8
−3.6
−2.4
−1.2
0.0
1.2
2.4
3.6
4.8
6.0
LDA SCORE (log 10)