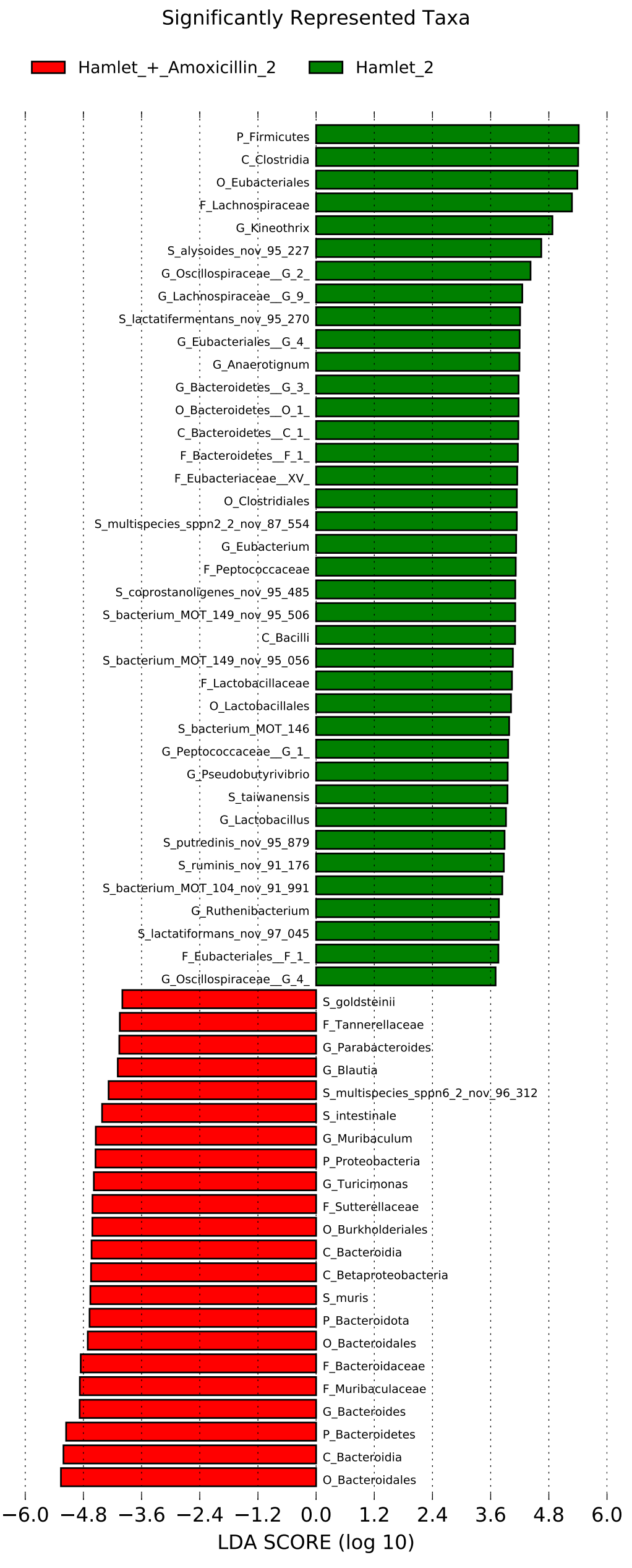
Significantly Represented Taxa
Hamlet_+_Amoxicillin_2
Hamlet_2
P_Firmicutes
C_Clostridia
O_Eubacteriales
F_Lachnospiraceae
G_Kineothrix
S_alysoides_nov_95_227
G_Oscillospiraceae__G_2_
G_Lachnospiraceae__G_9_
S_lactatifermentans_nov_95_270
G_Eubacteriales__G_4_
G_Anaerotignum
G_Bacteroidetes__G_3_
O_Bacteroidetes__O_1_
C_Bacteroidetes__C_1_
F_Bacteroidetes__F_1_
F_Eubacteriaceae__XV_
O_Clostridiales
S_multispecies_sppn2_2_nov_87_554
G_Eubacterium
F_Peptococcaceae
S_coprostanoligenes_nov_95_485
S_bacterium_MOT_149_nov_95_506
C_Bacilli
S_bacterium_MOT_149_nov_95_056
F_Lactobacillaceae
O_Lactobacillales
S_bacterium_MOT_146
G_Peptococcaceae__G_1_
G_Pseudobutyrivibrio
S_taiwanensis
G_Lactobacillus
S_putredinis_nov_95_879
S_ruminis_nov_91_176
S_bacterium_MOT_104_nov_91_991
G_Ruthenibacterium
S_lactatiformans_nov_97_045
F_Eubacteriales__F_1_
G_Oscillospiraceae__G_4_
S_goldsteinii
F_Tannerellaceae
G_Parabacteroides
G_Blautia
S_multispecies_sppn6_2_nov_96_312
S_intestinale
G_Muribaculum
P_Proteobacteria
G_Turicimonas
F_Sutterellaceae
O_Burkholderiales
C_Bacteroidia
C_Betaproteobacteria
S_muris
P_Bacteroidota
O_Bacteroidales
F_Bacteroidaceae
F_Muribaculaceae
G_Bacteroides
P_Bacteroidetes
C_Bacteroidia
O_Bacteroidales
−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)