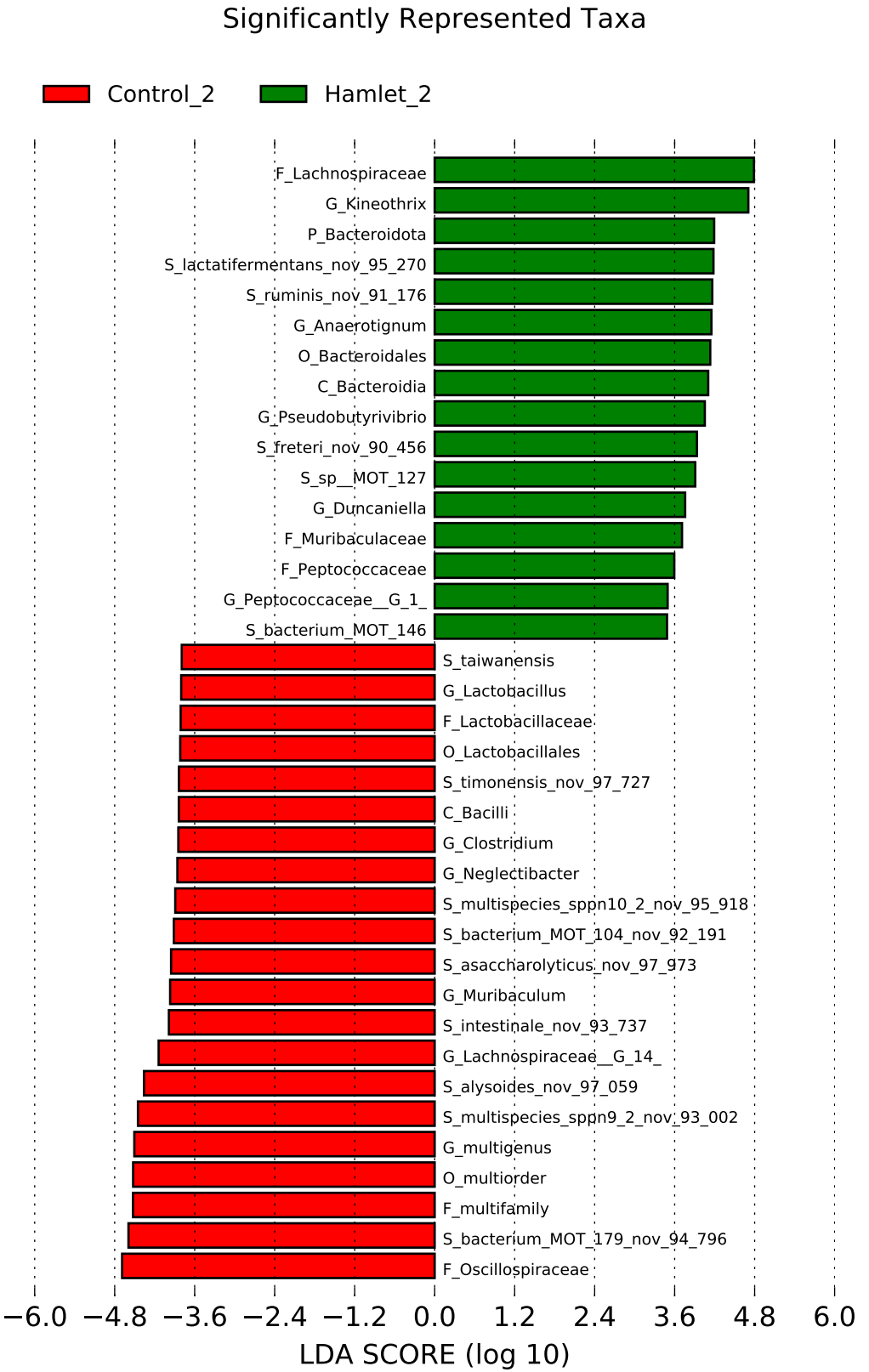
Significantly Represented Taxa
Control_2
Hamlet_2
F_Lachnospiraceae
G_Kineothrix
P_Bacteroidota
S_lactatifermentans_nov_95_270
S_ruminis_nov_91_176
G_Anaerotignum
O_Bacteroidales
C_Bacteroidia
G_Pseudobutyrivibrio
S_freteri_nov_90_456
S_sp__MOT_127
G_Duncaniella
F_Muribaculaceae
F_Peptococcaceae
G_Peptococcaceae__G_1_
S_bacterium_MOT_146
S_taiwanensis
G_Lactobacillus
F_Lactobacillaceae
O_Lactobacillales
S_timonensis_nov_97_727
C_Bacilli
G_Clostridium
G_Neglectibacter
S_multispecies_sppn10_2_nov_95_918
S_bacterium_MOT_104_nov_92_191
S_asaccharolyticus_nov_97_973
G_Muribaculum
S_intestinale_nov_93_737
G_Lachnospiraceae__G_14_
S_alysoides_nov_97_059
S_multispecies_sppn9_2_nov_93_002
G_multigenus
O_multiorder
F_multifamily
S_bacterium_MOT_179_nov_94_796
F_Oscillospiraceae
−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)