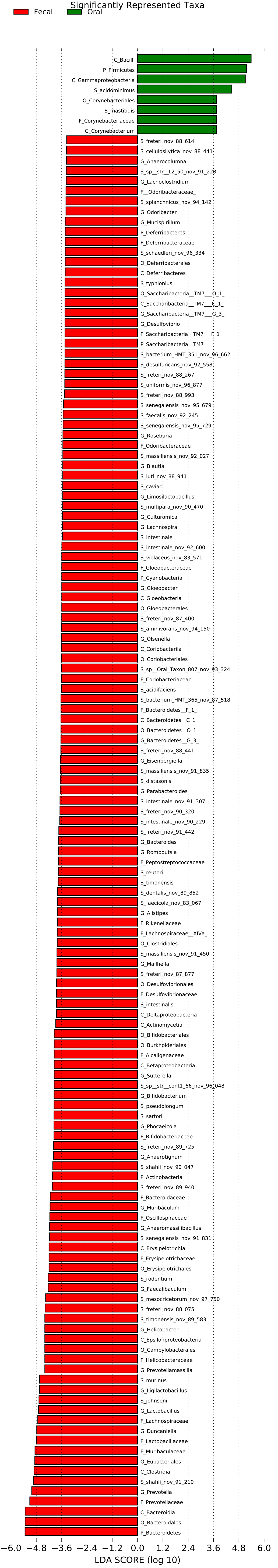
Significantly Represented Taxa
Fecal
Oral
C_Bacilli
P_Firmicutes
C_Gammaproteobacteria
S_acidominimus
O_Corynebacteriales
S_mastitidis
F_Corynebacteriaceae
G_Corynebacterium
S_freteri_nov_88_614
S_cellulosilytica_nov_88_441
G_Anaerocolumna
S_sp__str__L2_50_nov_91_228
G_Lacnoclostridium
F__Odoribacteraceae_
S_splanchnicus_nov_94_142
G_Odoribacter
G_Mucispirillum
P_Deferribacteres
F_Deferribacteraceae
S_schaedleri_nov_96_334
O_Deferribacterales
C_Deferribacteres
S_typhlonius
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
G_Saccharibacteria__TM7___G_3_
G_Desulfovibrio
F_Saccharibacteria__TM7___F_1_
P_Saccharibacteria__TM7_
S_bacterium_HMT_351_nov_96_662
S_desulfuricans_nov_92_558
S_freteri_nov_88_267
S_uniformis_nov_96_877
S_freteri_nov_88_993
S_senegalensis_nov_95_679
S_faecalis_nov_92_245
S_senegalensis_nov_95_729
G_Roseburia
F_Odoribacteraceae
S_massiliensis_nov_92_027
G_Blautia
S_luti_nov_88_941
S_caviae
G_Limosilactobacillus
S_multipara_nov_90_470
G_Culturomica
G_Lachnospira
S_intestinale
S_intestinale_nov_92_600
S_violaceus_nov_83_571
F_Gloeobacteraceae
P_Cyanobacteria
G_Gloeobacter
C_Gloeobacteria
O_Gloeobacterales
S_freteri_nov_87_400
S_aminivorans_nov_94_150
G_Olsenella
C_Coriobacteriia
O_Coriobacteriales
S_sp__Oral_Taxon_807_nov_93_324
F_Coriobacteriaceae
S_acidifaciens
S_bacterium_HMT_365_nov_87_518
F_Bacteroidetes__F_1_
C_Bacteroidetes__C_1_
O_Bacteroidetes__O_1_
G_Bacteroidetes__G_3_
S_freteri_nov_88_441
G_Eisenbergiella
S_massiliensis_nov_91_835
S_distasonis
G_Parabacteroides
S_intestinale_nov_91_307
S_freteri_nov_90_320
S_intestinale_nov_90_229
S_freteri_nov_91_442
G_Bacteroides
G_Romboutsia
F_Peptostreptococcaceae
S_reuteri
S_timonensis
S_dentalis_nov_89_852
S_faecicola_nov_83_067
G_Alistipes
F_Rikenellaceae
F_Lachnospiraceae__XIVa_
O_Clostridiales
S_massiliensis_nov_91_450
G_Mailhella
S_freteri_nov_87_877
O_Desulfovibrionales
F_Desulfovibrionaceae
S_intestinalis
C_Deltaproteobacteria
C_Actinomycetia
O_Bifidobacteriales
O_Burkholderiales
F_Alcaligenaceae
C_Betaproteobacteria
G_Sutterella
S_sp__str__cont1_66_nov_96_048
G_Bifidobacterium
S_pseudolongum
S_sartorii
G_Phocaeicola
F_Bifidobacteriaceae
S_freteri_nov_89_725
G_Anaerotignum
S_shahii_nov_90_047
P_Actinobacteria
S_freteri_nov_89_940
F_Bacteroidaceae
G_Muribaculum
F_Oscillospiraceae
G_Anaeromassilibacillus
S_senegalensis_nov_91_831
C_Erysipelotrichia
F_Erysipelotrichaceae
O_Erysipelotrichales
S_rodentium
G_Faecalibaculum
S_mesocricetorum_nov_97_750
S_freteri_nov_88_075
S_timonensis_nov_89_583
G_Helicobacter
C_Epsilonproteobacteria
O_Campylobacterales
F_Helicobacteraceae
G_Prevotellamassilia
S_murinus
G_Ligilactobacillus
S_johnsonii
G_Lactobacillus
F_Lachnospiraceae
G_Duncaniella
F_Lactobacillaceae
F_Muribaculaceae
O_Eubacteriales
C_Clostridia
S_shahii_nov_91_210
G_Prevotella
F_Prevotellaceae
C_Bacteroidia
O_Bacteroidales
P_Bacteroidetes
−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)