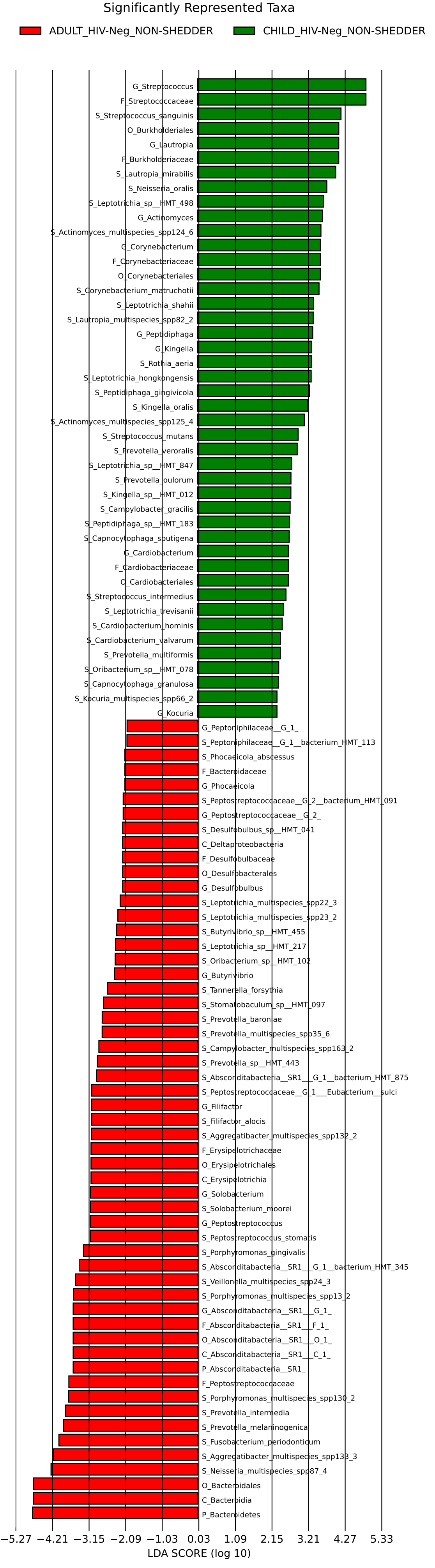
Significantly Represented Taxa
ADULT_HIV-Neg_NON-SHEDDER
CHILD_HIV-Neg_NON-SHEDDER
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_sanguinis
O_Burkholderiales
G_Lautropia
F_Burkholderiaceae
S_Lautropia_mirabilis
S_Neisseria_oralis
S_Leptotrichia_sp__HMT_498
G_Actinomyces
S_Actinomyces_multispecies_spp124_6
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
S_Corynebacterium_matruchotii
S_Leptotrichia_shahii
S_Lautropia_multispecies_spp82_2
G_Peptidiphaga
G_Kingella
S_Rothia_aeria
S_Leptotrichia_hongkongensis
S_Peptidiphaga_gingivicola
S_Kingella_oralis
S_Actinomyces_multispecies_spp125_4
S_Streptococcus_mutans
S_Prevotella_veroralis
S_Leptotrichia_sp__HMT_847
S_Prevotella_oulorum
S_Kingella_sp__HMT_012
S_Campylobacter_gracilis
S_Peptidiphaga_sp__HMT_183
S_Capnocytophaga_sputigena
G_Cardiobacterium
F_Cardiobacteriaceae
O_Cardiobacteriales
S_Streptococcus_intermedius
S_Leptotrichia_trevisanii
S_Cardiobacterium_hominis
S_Cardiobacterium_valvarum
S_Prevotella_multiformis
S_Oribacterium_sp__HMT_078
S_Capnocytophaga_granulosa
S_Kocuria_multispecies_spp66_2
G_Kocuria
G_Peptoniphilaceae__G_1_
S_Peptoniphilaceae__G_1__bacterium_HMT_113
S_Phocaeicola_abscessus
F_Bacteroidaceae
G_Phocaeicola
S_Peptostreptococcaceae__G_2__bacterium_HMT_091
G_Peptostreptococcaceae__G_2_
S_Desulfobulbus_sp__HMT_041
C_Deltaproteobacteria
F_Desulfobulbaceae
O_Desulfobacterales
G_Desulfobulbus
S_Leptotrichia_multispecies_spp22_3
S_Leptotrichia_multispecies_spp23_2
S_Butyrivibrio_sp__HMT_455
S_Leptotrichia_sp__HMT_217
S_Oribacterium_sp__HMT_102
G_Butyrivibrio
S_Tannerella_forsythia
S_Stomatobaculum_sp__HMT_097
S_Prevotella_baroniae
S_Prevotella_multispecies_spp35_6
S_Campylobacter_multispecies_spp163_2
S_Prevotella_sp__HMT_443
S_Absconditabacteria__SR1___G_1__bacterium_HMT_875
S_Peptostreptococcaceae__G_1___Eubacterium__sulci
G_Filifactor
S_Filifactor_alocis
S_Aggregatibacter_multispecies_spp132_2
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
G_Solobacterium
S_Solobacterium_moorei
G_Peptostreptococcus
S_Peptostreptococcus_stomatis
S_Porphyromonas_gingivalis
S_Absconditabacteria__SR1___G_1__bacterium_HMT_345
S_Veillonella_multispecies_spp24_3
S_Porphyromonas_multispecies_spp13_2
G_Absconditabacteria__SR1___G_1_
F_Absconditabacteria__SR1___F_1_
O_Absconditabacteria__SR1___O_1_
C_Absconditabacteria__SR1___C_1_
P_Absconditabacteria__SR1_
F_Peptostreptococcaceae
S_Porphyromonas_multispecies_spp130_2
S_Prevotella_intermedia
S_Prevotella_melaninogenica
S_Fusobacterium_periodonticum
S_Aggregatibacter_multispecies_spp133_3
S_Neisseria_multispecies_spp87_4
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
−5.27
−4.21
−3.15
−2.09
−1.03
0.03
1.09
2.15
3.21
4.27
5.33
LDA SCORE (log 10)