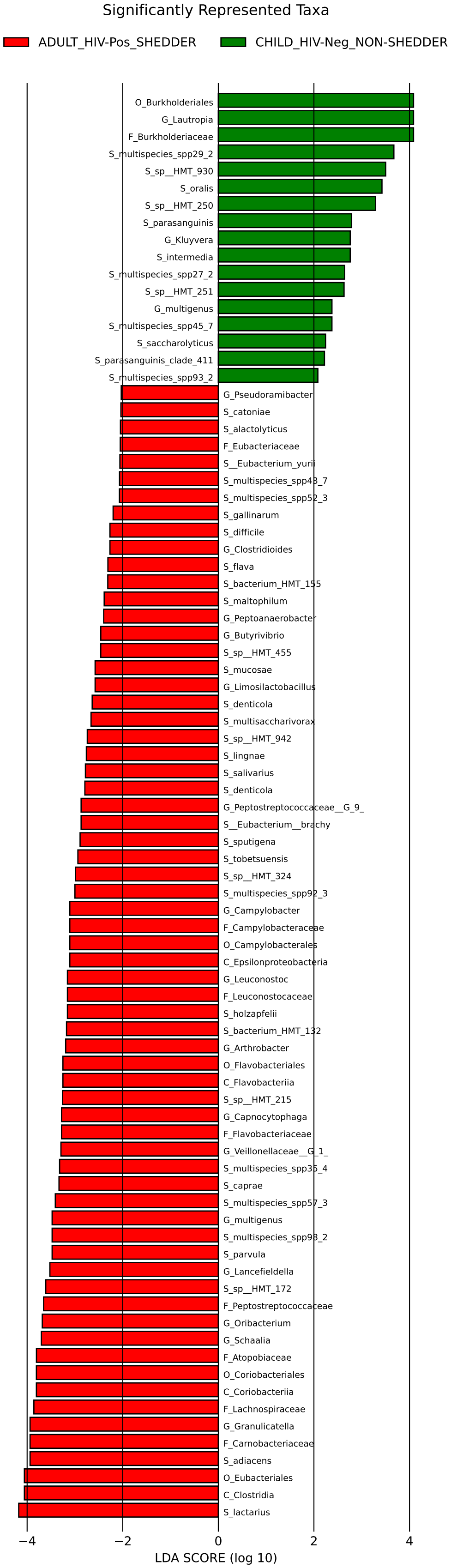
Significantly Represented Taxa
ADULT_HIV-Pos_SHEDDER
CHILD_HIV-Neg_NON-SHEDDER
O_Burkholderiales
G_Lautropia
F_Burkholderiaceae
S_multispecies_spp29_2
S_sp__HMT_930
S_oralis
S_sp__HMT_250
S_parasanguinis
G_Kluyvera
S_intermedia
S_multispecies_spp27_2
S_sp__HMT_251
G_multigenus
S_multispecies_spp45_7
S_saccharolyticus
S_parasanguinis_clade_411
S_multispecies_spp93_2
G_Pseudoramibacter
S_catoniae
S_alactolyticus
F_Eubacteriaceae
S__Eubacterium_yurii
S_multispecies_spp43_7
S_multispecies_spp52_3
S_gallinarum
S_difficile
G_Clostridioides
S_flava
S_bacterium_HMT_155
S_maltophilum
G_Peptoanaerobacter
G_Butyrivibrio
S_sp__HMT_455
S_mucosae
G_Limosilactobacillus
S_denticola
S_multisaccharivorax
S_sp__HMT_942
S_lingnae
S_salivarius
S_denticola
G_Peptostreptococcaceae__G_9_
S__Eubacterium__brachy
S_sputigena
S_tobetsuensis
S_sp__HMT_324
S_multispecies_spp92_3
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
G_Leuconostoc
F_Leuconostocaceae
S_holzapfelii
S_bacterium_HMT_132
G_Arthrobacter
O_Flavobacteriales
C_Flavobacteriia
S_sp__HMT_215
G_Capnocytophaga
F_Flavobacteriaceae
G_Veillonellaceae__G_1_
S_multispecies_spp35_4
S_caprae
S_multispecies_spp57_3
G_multigenus
S_multispecies_spp98_2
S_parvula
G_Lancefieldella
S_sp__HMT_172
F_Peptostreptococcaceae
G_Oribacterium
G_Schaalia
F_Atopobiaceae
O_Coriobacteriales
C_Coriobacteriia
F_Lachnospiraceae
G_Granulicatella
F_Carnobacteriaceae
S_adiacens
O_Eubacteriales
C_Clostridia
S_lactarius
−4
−2
0
2
4
LDA SCORE (log 10)