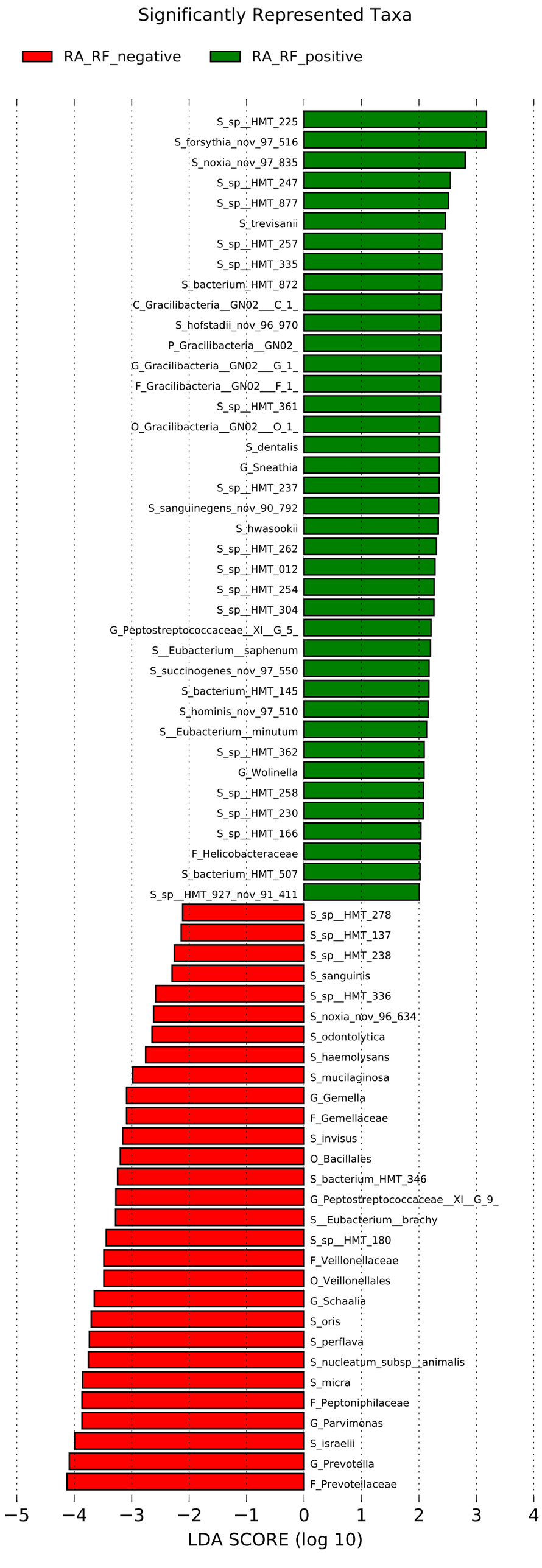
Significantly Represented Taxa
RA_RF_negative
RA_RF_positive
S_sp__HMT_225
S_forsythia_nov_97_516
S_noxia_nov_97_835
S_sp__HMT_247
S_sp__HMT_877
S_trevisanii
S_sp__HMT_257
S_sp__HMT_335
S_bacterium_HMT_872
C_Gracilibacteria__GN02___C_1_
S_hofstadii_nov_96_970
P_Gracilibacteria__GN02_
G_Gracilibacteria__GN02___G_1_
F_Gracilibacteria__GN02___F_1_
S_sp__HMT_361
O_Gracilibacteria__GN02___O_1_
S_dentalis
G_Sneathia
S_sp__HMT_237
S_sanguinegens_nov_90_792
S_hwasookii
S_sp__HMT_262
S_sp__HMT_012
S_sp__HMT_254
S_sp__HMT_304
G_Peptostreptococcaceae__XI__G_5_
S__Eubacterium__saphenum
S_succinogenes_nov_97_550
S_bacterium_HMT_145
S_hominis_nov_97_510
S__Eubacterium__minutum
S_sp__HMT_362
G_Wolinella
S_sp__HMT_258
S_sp__HMT_230
S_sp__HMT_166
F_Helicobacteraceae
S_bacterium_HMT_507
S_sp__HMT_927_nov_91_411
S_sp__HMT_278
S_sp__HMT_137
S_sp__HMT_238
S_sanguinis
S_sp__HMT_336
S_noxia_nov_96_634
S_odontolytica
S_haemolysans
S_mucilaginosa
G_Gemella
F_Gemellaceae
S_invisus
O_Bacillales
S_bacterium_HMT_346
G_Peptostreptococcaceae__XI__G_9_
S__Eubacterium__brachy
S_sp__HMT_180
F_Veillonellaceae
O_Veillonellales
G_Schaalia
S_oris
S_perflava
S_nucleatum_subsp__animalis
S_micra
F_Peptoniphilaceae
G_Parvimonas
S_israelii
G_Prevotella
F_Prevotellaceae
−5
−4
−3
−2
−1
0
1
2
3
4
LDA SCORE (log 10)