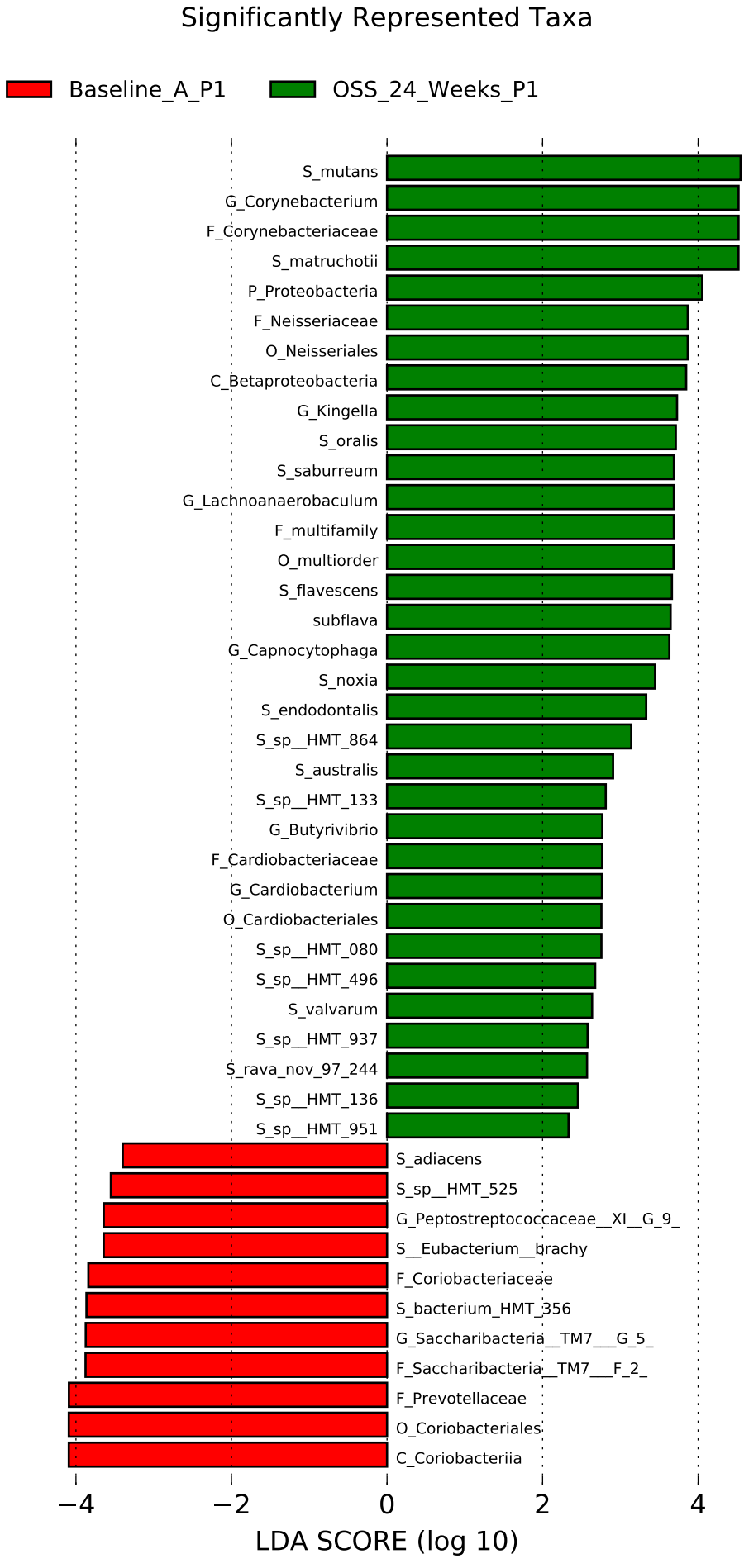
Significantly Represented Taxa
Baseline_A_P1
OSS_24_Weeks_P1
S_mutans
G_Corynebacterium
F_Corynebacteriaceae
S_matruchotii
P_Proteobacteria
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
G_Kingella
S_oralis
S_saburreum
G_Lachnoanaerobaculum
F_multifamily
O_multiorder
S_flavescens
subflava
G_Capnocytophaga
S_noxia
S_endodontalis
S_sp__HMT_864
S_australis
S_sp__HMT_133
G_Butyrivibrio
F_Cardiobacteriaceae
G_Cardiobacterium
O_Cardiobacteriales
S_sp__HMT_080
S_sp__HMT_496
S_valvarum
S_sp__HMT_937
S_rava_nov_97_244
S_sp__HMT_136
S_sp__HMT_951
S_adiacens
S_sp__HMT_525
G_Peptostreptococcaceae__XI__G_9_
S__Eubacterium__brachy
F_Coriobacteriaceae
S_bacterium_HMT_356
G_Saccharibacteria__TM7___G_5_
F_Saccharibacteria__TM7___F_2_
F_Prevotellaceae
O_Coriobacteriales
C_Coriobacteriia
−4
−2
0
2
4
LDA SCORE (log 10)