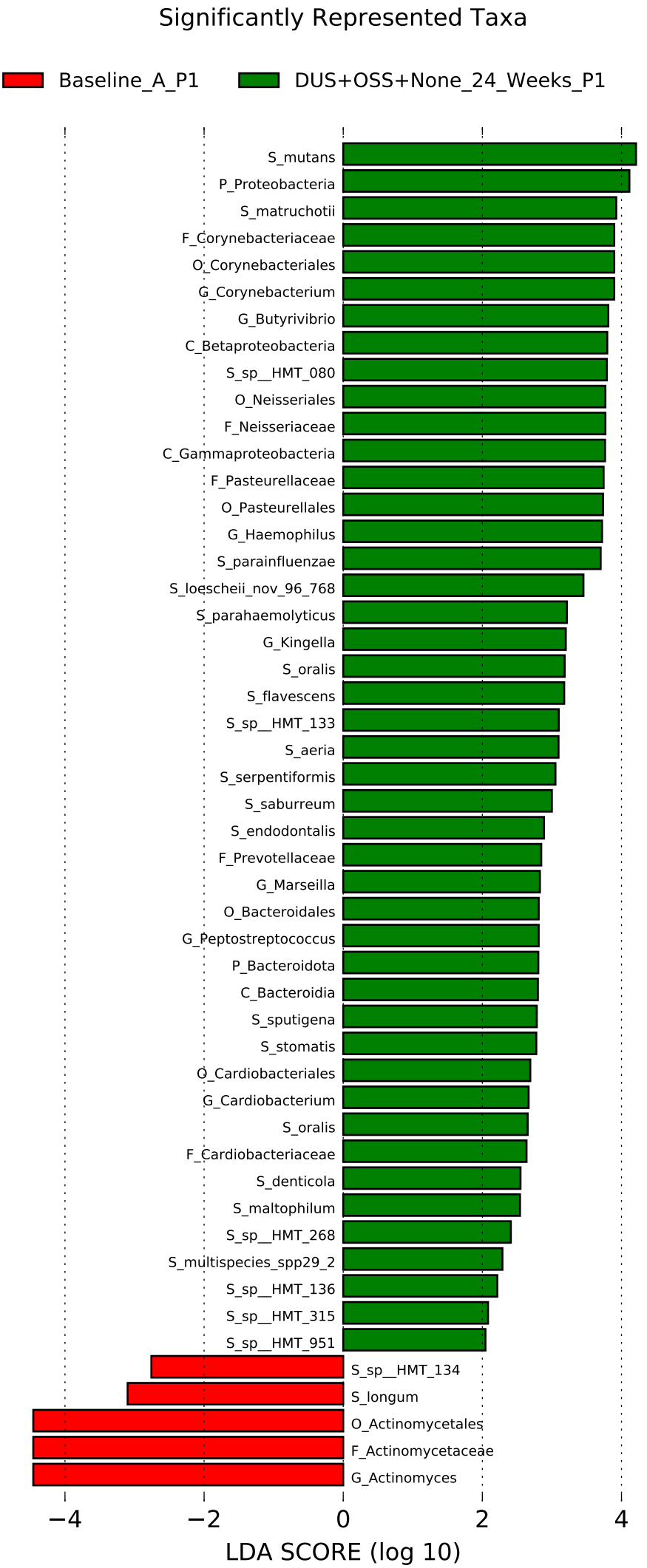
Significantly Represented Taxa
Baseline_A_P1
DUS+OSS+None_24_Weeks_P1
S_mutans
P_Proteobacteria
S_matruchotii
F_Corynebacteriaceae
O_Corynebacteriales
G_Corynebacterium
G_Butyrivibrio
C_Betaproteobacteria
S_sp__HMT_080
O_Neisseriales
F_Neisseriaceae
C_Gammaproteobacteria
F_Pasteurellaceae
O_Pasteurellales
G_Haemophilus
S_parainfluenzae
S_loescheii_nov_96_768
S_parahaemolyticus
G_Kingella
S_oralis
S_flavescens
S_sp__HMT_133
S_aeria
S_serpentiformis
S_saburreum
S_endodontalis
F_Prevotellaceae
G_Marseilla
O_Bacteroidales
G_Peptostreptococcus
P_Bacteroidota
C_Bacteroidia
S_sputigena
S_stomatis
O_Cardiobacteriales
G_Cardiobacterium
S_oralis
F_Cardiobacteriaceae
S_denticola
S_maltophilum
S_sp__HMT_268
S_multispecies_spp29_2
S_sp__HMT_136
S_sp__HMT_315
S_sp__HMT_951
S_sp__HMT_134
S_longum
O_Actinomycetales
F_Actinomycetaceae
G_Actinomyces
−4
−2
0
2
4
LDA SCORE (log 10)