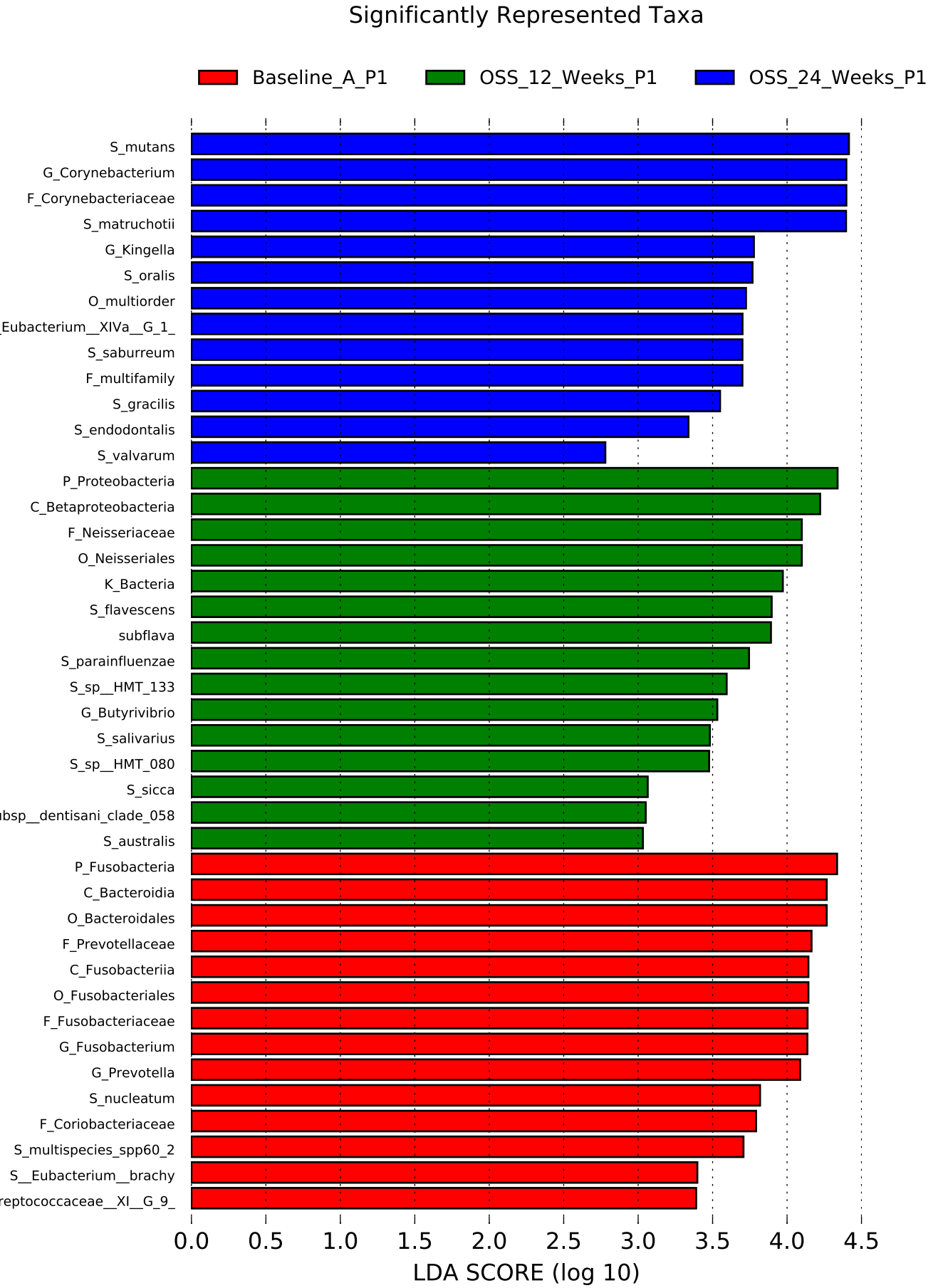
Significantly Represented Taxa
Baseline_A_P1
OSS_12_Weeks_P1
OSS_24_Weeks_P1
S_mutans
G_Corynebacterium
F_Corynebacteriaceae
S_matruchotii
G_Kingella
S_oralis
O_multiorder
_Eubacterium__XIVa__G_1_
S_saburreum
F_multifamily
S_gracilis
S_endodontalis
S_valvarum
P_Proteobacteria
C_Betaproteobacteria
F_Neisseriaceae
O_Neisseriales
K_Bacteria
S_flavescens
subflava
S_parainfluenzae
S_sp__HMT_133
G_Butyrivibrio
S_salivarius
S_sp__HMT_080
S_sicca
ubsp__dentisani_clade_058
S_australis
P_Fusobacteria
C_Bacteroidia
O_Bacteroidales
F_Prevotellaceae
C_Fusobacteriia
O_Fusobacteriales
F_Fusobacteriaceae
G_Fusobacterium
G_Prevotella
S_nucleatum
F_Coriobacteriaceae
S_multispecies_spp60_2
S__Eubacterium__brachy
reptococcaceae__XI__G_9_
0.0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
4.0
4.5
LDA SCORE (log 10)