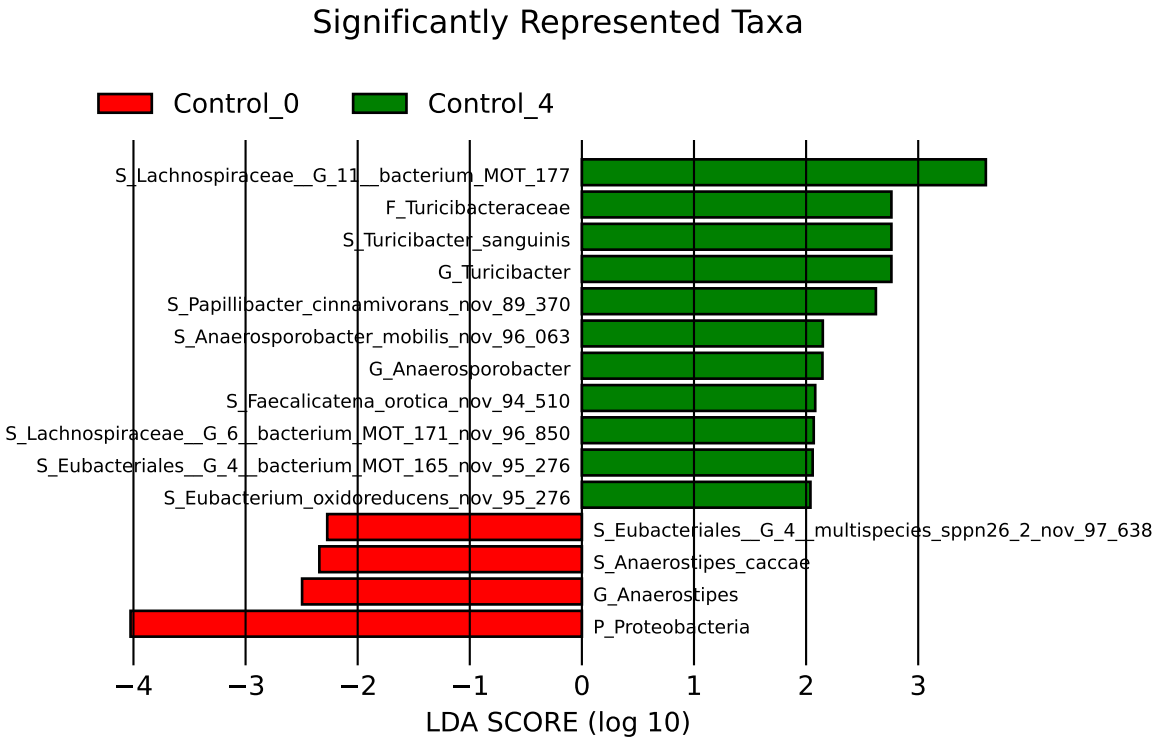
Significantly Represented Taxa
Control_0
Control_4
S_Lachnospiraceae__G_11__bacterium_MOT_177
F_Turicibacteraceae
S_Turicibacter_sanguinis
G_Turicibacter
S_Papillibacter_cinnamivorans_nov_89_370
S_Anaerosporobacter_mobilis_nov_96_063
G_Anaerosporobacter
S_Faecalicatena_orotica_nov_94_510
S_Lachnospiraceae__G_6__bacterium_MOT_171_nov_96_850
S_Eubacteriales__G_4__bacterium_MOT_165_nov_95_276
S_Eubacterium_oxidoreducens_nov_95_276
S_Eubacteriales__G_4__multispecies_sppn26_2_nov_97_638
S_Anaerostipes_caccae
G_Anaerostipes
P_Proteobacteria
−4
−3
−2
−1
0
1
2
3
LDA SCORE (log 10)