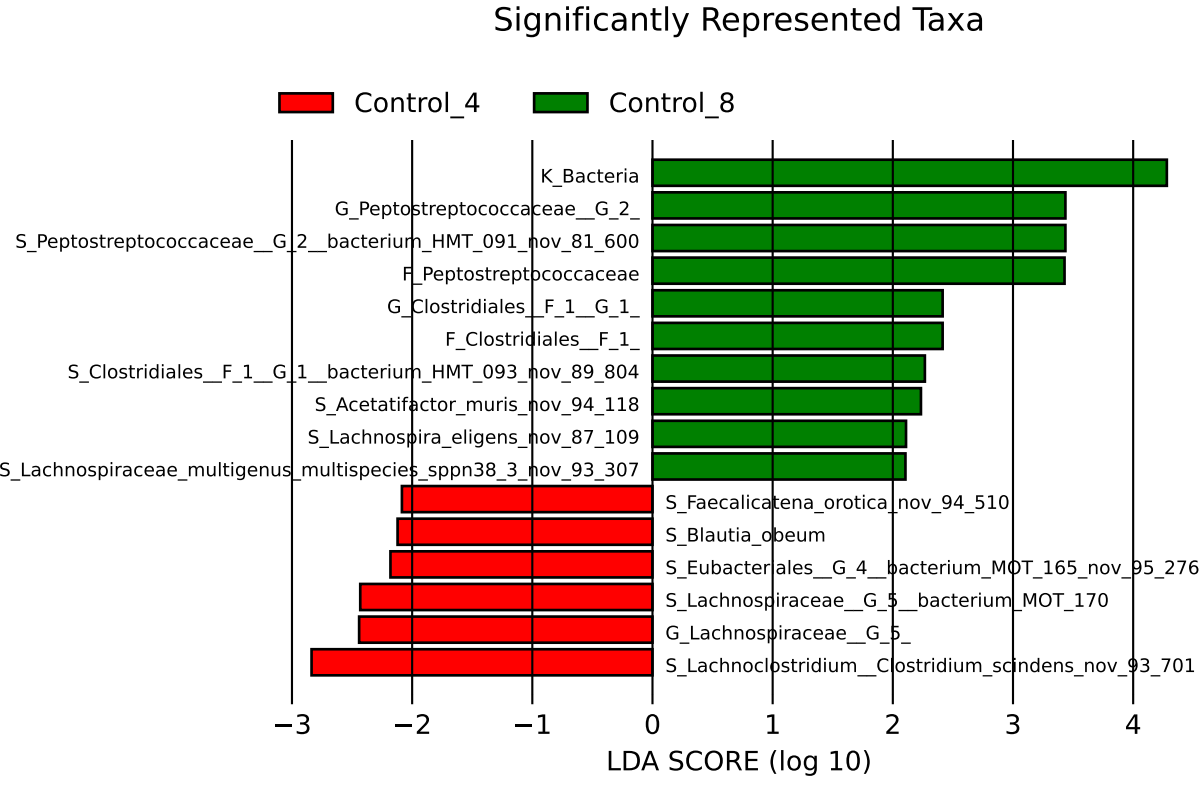
Significantly Represented Taxa
Control_4
Control_8
K_Bacteria
G_Peptostreptococcaceae__G_2_
S_Peptostreptococcaceae__G_2__bacterium_HMT_091_nov_81_600
F_Peptostreptococcaceae
G_Clostridiales__F_1__G_1_
F_Clostridiales__F_1_
S_Clostridiales__F_1__G_1__bacterium_HMT_093_nov_89_804
S_Acetatifactor_muris_nov_94_118
S_Lachnospira_eligens_nov_87_109
S_Lachnospiraceae_multigenus_multispecies_sppn38_3_nov_93_307
S_Faecalicatena_orotica_nov_94_510
S_Blautia_obeum
S_Eubacteriales__G_4__bacterium_MOT_165_nov_95_276
S_Lachnospiraceae__G_5__bacterium_MOT_170
G_Lachnospiraceae__G_5_
S_Lachnoclostridium__Clostridium_scindens_nov_93_701
−3
−2
−1
0
1
2
3
4
LDA SCORE (log 10)