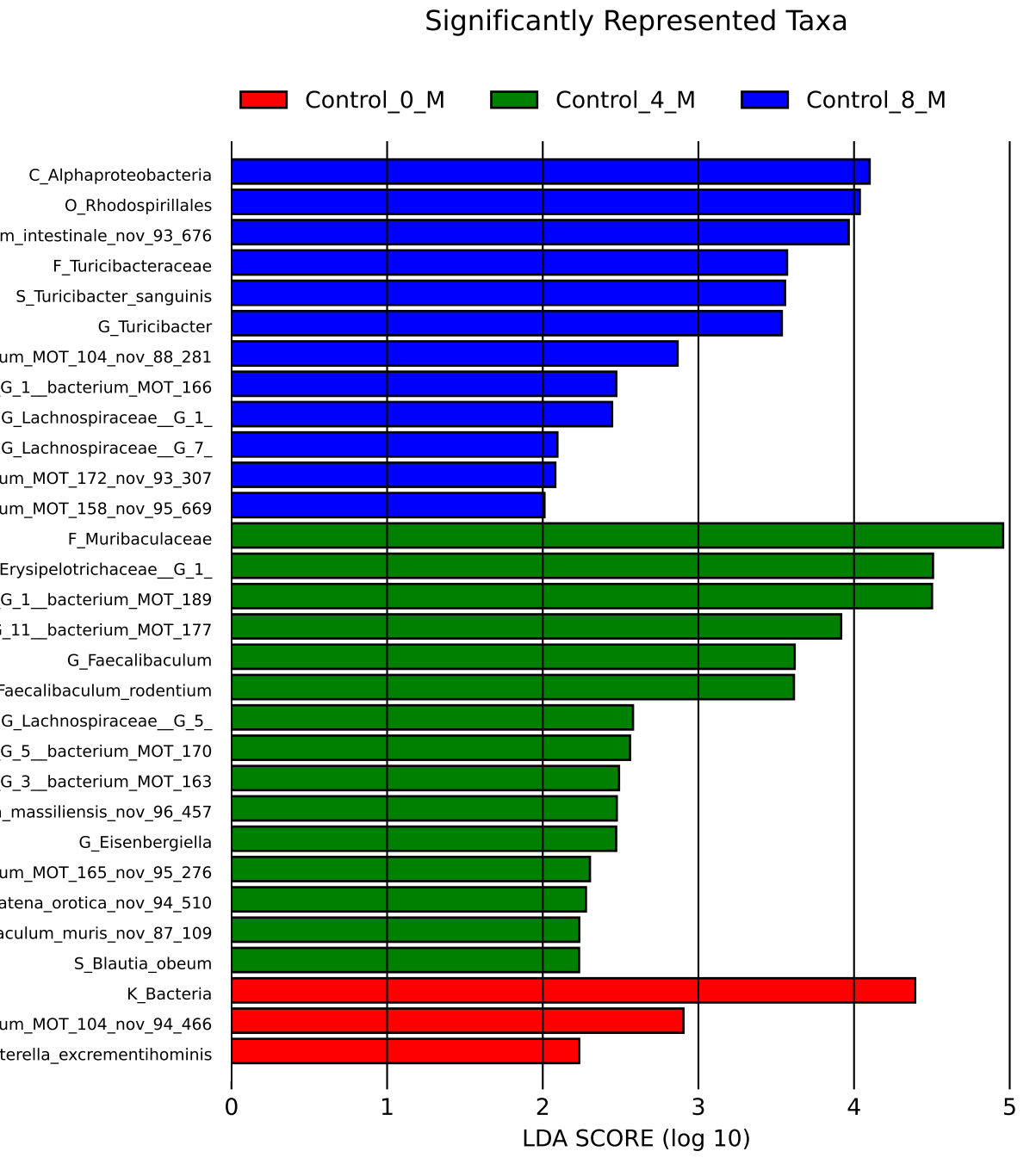
Significantly Represented Taxa
Control_0_M
Control_4_M
Control_8_M
C_Alphaproteobacteria
O_Rhodospirillales
m_intestinale_nov_93_676
F_Turicibacteraceae
S_Turicibacter_sanguinis
G_Turicibacter
um_MOT_104_nov_88_281
G_1__bacterium_MOT_166
G_Lachnospiraceae__G_1_
G_Lachnospiraceae__G_7_
um_MOT_172_nov_93_307
um_MOT_158_nov_95_669
F_Muribaculaceae
Erysipelotrichaceae__G_1_
G_1__bacterium_MOT_189
_11__bacterium_MOT_177
G_Faecalibaculum
Faecalibaculum_rodentium
G_Lachnospiraceae__G_5_
G_5__bacterium_MOT_170
G_3__bacterium_MOT_163
_massiliensis_nov_96_457
G_Eisenbergiella
um_MOT_165_nov_95_276
atena_orotica_nov_94_510
aculum_muris_nov_87_109
S_Blautia_obeum
K_Bacteria
um_MOT_104_nov_94_466
terella_excrementihominis
0
1
2
3
4
5
LDA SCORE (log 10)