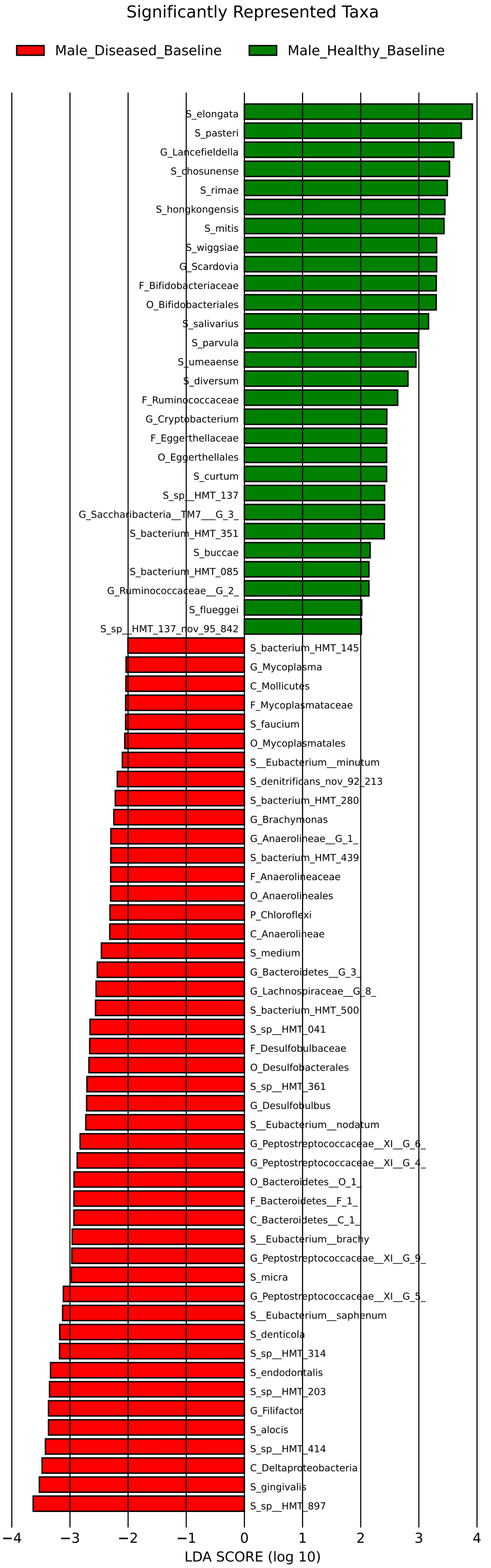
Significantly Represented Taxa
Male_Diseased_Baseline
Male_Healthy_Baseline
S_elongata
S_pasteri
G_Lancefieldella
S_chosunense
S_rimae
S_hongkongensis
S_mitis
S_wiggsiae
G_Scardovia
F_Bifidobacteriaceae
O_Bifidobacteriales
S_salivarius
S_parvula
S_umeaense
S_diversum
F_Ruminococcaceae
G_Cryptobacterium
F_Eggerthellaceae
O_Eggerthellales
S_curtum
S_sp__HMT_137
G_Saccharibacteria__TM7___G_3_
S_bacterium_HMT_351
S_buccae
S_bacterium_HMT_085
G_Ruminococcaceae__G_2_
S_flueggei
S_sp__HMT_137_nov_95_842
S_bacterium_HMT_145
G_Mycoplasma
C_Mollicutes
F_Mycoplasmataceae
S_faucium
O_Mycoplasmatales
S__Eubacterium__minutum
S_denitrificans_nov_92_213
S_bacterium_HMT_280
G_Brachymonas
G_Anaerolineae__G_1_
S_bacterium_HMT_439
F_Anaerolineaceae
O_Anaerolineales
P_Chloroflexi
C_Anaerolineae
S_medium
G_Bacteroidetes__G_3_
G_Lachnospiraceae__G_8_
S_bacterium_HMT_500
S_sp__HMT_041
F_Desulfobulbaceae
O_Desulfobacterales
S_sp__HMT_361
G_Desulfobulbus
S__Eubacterium__nodatum
G_Peptostreptococcaceae__XI__G_6_
G_Peptostreptococcaceae__XI__G_4_
O_Bacteroidetes__O_1_
F_Bacteroidetes__F_1_
C_Bacteroidetes__C_1_
S__Eubacterium__brachy
G_Peptostreptococcaceae__XI__G_9_
S_micra
G_Peptostreptococcaceae__XI__G_5_
S__Eubacterium__saphenum
S_denticola
S_sp__HMT_314
S_endodontalis
S_sp__HMT_203
G_Filifactor
S_alocis
S_sp__HMT_414
C_Deltaproteobacteria
S_gingivalis
S_sp__HMT_897
−4
−3
−2
−1
0
1
2
3
4
LDA SCORE (log 10)