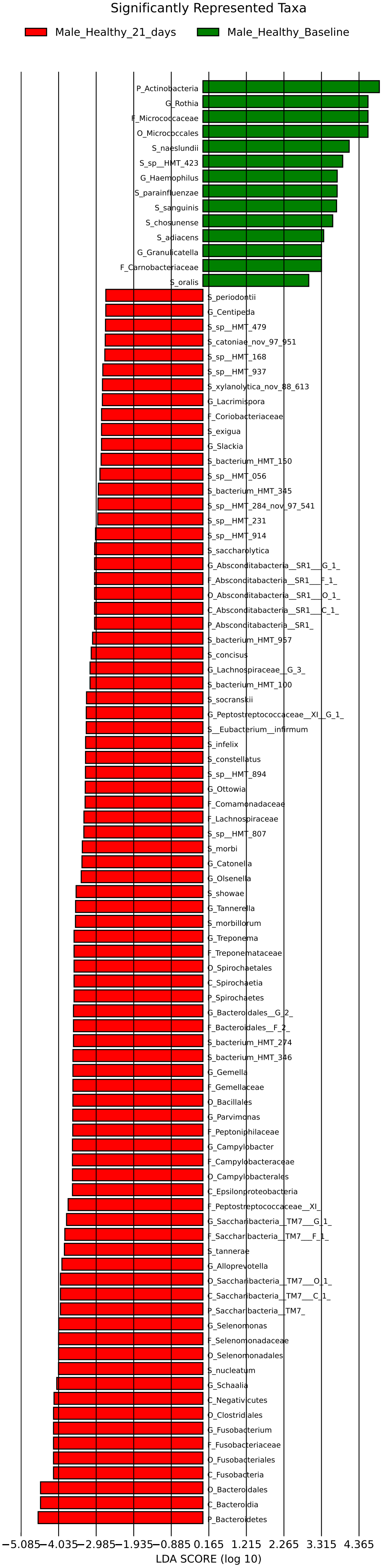
Significantly Represented Taxa
Male_Healthy_21_days
Male_Healthy_Baseline
P_Actinobacteria
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_naeslundii
S_sp__HMT_423
G_Haemophilus
S_parainfluenzae
S_sanguinis
S_chosunense
S_adiacens
G_Granulicatella
F_Carnobacteriaceae
S_oralis
S_periodontii
G_Centipeda
S_sp__HMT_479
S_catoniae_nov_97_951
S_sp__HMT_168
S_sp__HMT_937
S_xylanolytica_nov_88_613
G_Lacrimispora
F_Coriobacteriaceae
S_exigua
G_Slackia
S_bacterium_HMT_150
S_sp__HMT_056
S_bacterium_HMT_345
S_sp__HMT_284_nov_97_541
S_sp__HMT_231
S_sp__HMT_914
S_saccharolytica
G_Absconditabacteria__SR1___G_1_
F_Absconditabacteria__SR1___F_1_
O_Absconditabacteria__SR1___O_1_
C_Absconditabacteria__SR1___C_1_
P_Absconditabacteria__SR1_
S_bacterium_HMT_957
S_concisus
G_Lachnospiraceae__G_3_
S_bacterium_HMT_100
S_socranskii
G_Peptostreptococcaceae__XI__G_1_
S__Eubacterium__infirmum
S_infelix
S_constellatus
S_sp__HMT_894
G_Ottowia
F_Comamonadaceae
F_Lachnospiraceae
S_sp__HMT_807
S_morbi
G_Catonella
G_Olsenella
S_showae
G_Tannerella
S_morbillorum
G_Treponema
F_Treponemataceae
O_Spirochaetales
C_Spirochaetia
P_Spirochaetes
G_Bacteroidales__G_2_
F_Bacteroidales__F_2_
S_bacterium_HMT_274
S_bacterium_HMT_346
G_Gemella
F_Gemellaceae
O_Bacillales
G_Parvimonas
F_Peptoniphilaceae
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
F_Peptostreptococcaceae__XI_
G_Saccharibacteria__TM7___G_1_
F_Saccharibacteria__TM7___F_1_
S_tannerae
G_Alloprevotella
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
G_Selenomonas
F_Selenomonadaceae
O_Selenomonadales
S_nucleatum
G_Schaalia
C_Negativicutes
O_Clostridiales
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
−5.085
−4.035
−2.985
−1.935
−0.885
0.165
1.215
2.265
3.315
4.365
LDA SCORE (log 10)