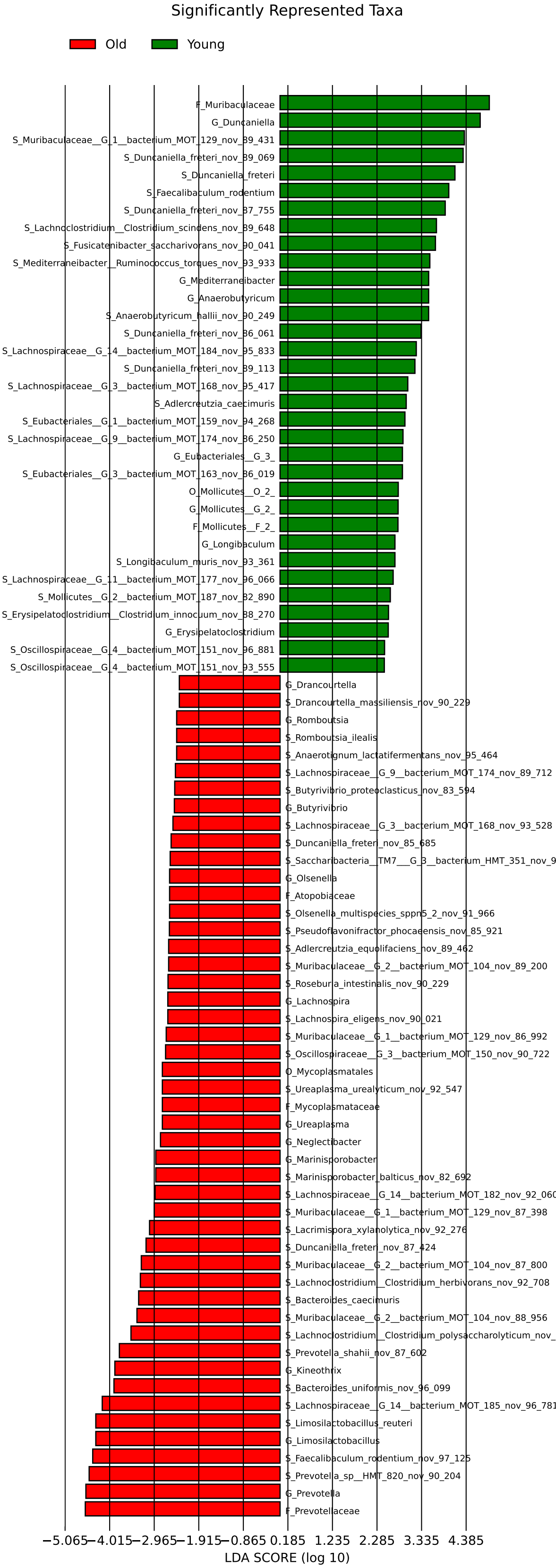

Significantly Represented Taxa
Old
Young
F_Muribaculaceae
G_Duncaniella
S_Muribaculaceae__G_1__bacterium_MOT_129_nov_89_431
S_Duncaniella_freteri_nov_89_069
S_Duncaniella_freteri
S_Faecalibaculum_rodentium
S_Duncaniella_freteri_nov_87_755
S_Lachnoclostridium__Clostridium_scindens_nov_89_648
S_Fusicatenibacter_saccharivorans_nov_90_041
S_Mediterraneibacter__Ruminococcus_torques_nov_93_933
G_Mediterraneibacter
G_Anaerobutyricum
S_Anaerobutyricum_hallii_nov_90_249
S_Duncaniella_freteri_nov_86_061
S_Lachnospiraceae__G_14__bacterium_MOT_184_nov_95_833
S_Duncaniella_freteri_nov_89_113
S_Lachnospiraceae__G_3__bacterium_MOT_168_nov_95_417
S_Adlercreutzia_caecimuris
S_Eubacteriales__G_1__bacterium_MOT_159_nov_94_268
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_86_250
G_Eubacteriales__G_3_
S_Eubacteriales__G_3__bacterium_MOT_163_nov_86_019
O_Mollicutes__O_2_
G_Mollicutes__G_2_
F_Mollicutes__F_2_
G_Longibaculum
S_Longibaculum_muris_nov_93_361
S_Lachnospiraceae__G_11__bacterium_MOT_177_nov_96_066
S_Mollicutes__G_2__bacterium_MOT_187_nov_82_890
S_Erysipelatoclostridium__Clostridium_innocuum_nov_88_270
G_Erysipelatoclostridium
S_Oscillospiraceae__G_4__bacterium_MOT_151_nov_96_881
S_Oscillospiraceae__G_4__bacterium_MOT_151_nov_93_555
G_Drancourtella
S_Drancourtella_massiliensis_nov_90_229
G_Romboutsia
S_Romboutsia_ilealis
S_Anaerotignum_lactatifermentans_nov_95_464
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_89_712
S_Butyrivibrio_proteoclasticus_nov_83_594
G_Butyrivibrio
S_Lachnospiraceae__G_3__bacterium_MOT_168_nov_93_528
S_Duncaniella_freteri_nov_85_685
S_Saccharibacteria__TM7___G_3__bacterium_HMT_351_nov_92_4
G_Olsenella
F_Atopobiaceae
S_Olsenella_multispecies_sppn5_2_nov_91_966
S_Pseudoflavonifractor_phocaeensis_nov_85_921
S_Adlercreutzia_equolifaciens_nov_89_462
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_89_200
S_Roseburia_intestinalis_nov_90_229
G_Lachnospira
S_Lachnospira_eligens_nov_90_021
S_Muribaculaceae__G_1__bacterium_MOT_129_nov_86_992
S_Oscillospiraceae__G_3__bacterium_MOT_150_nov_90_722
O_Mycoplasmatales
S_Ureaplasma_urealyticum_nov_92_547
F_Mycoplasmataceae
G_Ureaplasma
G_Neglectibacter
G_Marinisporobacter
S_Marinisporobacter_balticus_nov_82_692
S_Lachnospiraceae__G_14__bacterium_MOT_182_nov_92_060
S_Muribaculaceae__G_1__bacterium_MOT_129_nov_87_398
S_Lacrimispora_xylanolytica_nov_92_276
S_Duncaniella_freteri_nov_87_424
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_87_800
S_Lachnoclostridium__Clostridium_herbivorans_nov_92_708
S_Bacteroides_caecimuris
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_88_956
S_Lachnoclostridium__Clostridium_polysaccharolyticum_nov_86_2
S_Prevotella_shahii_nov_87_602
G_Kineothrix
S_Bacteroides_uniformis_nov_96_099
S_Lachnospiraceae__G_14__bacterium_MOT_185_nov_96_781
S_Limosilactobacillus_reuteri
G_Limosilactobacillus
S_Faecalibaculum_rodentium_nov_97_125
S_Prevotella_sp__HMT_820_nov_90_204
G_Prevotella
F_Prevotellaceae
−5.065
−4.015
−2.965
−1.915
−0.865
0.185
1.235
2.285
3.335
4.385
LDA SCORE (log 10)