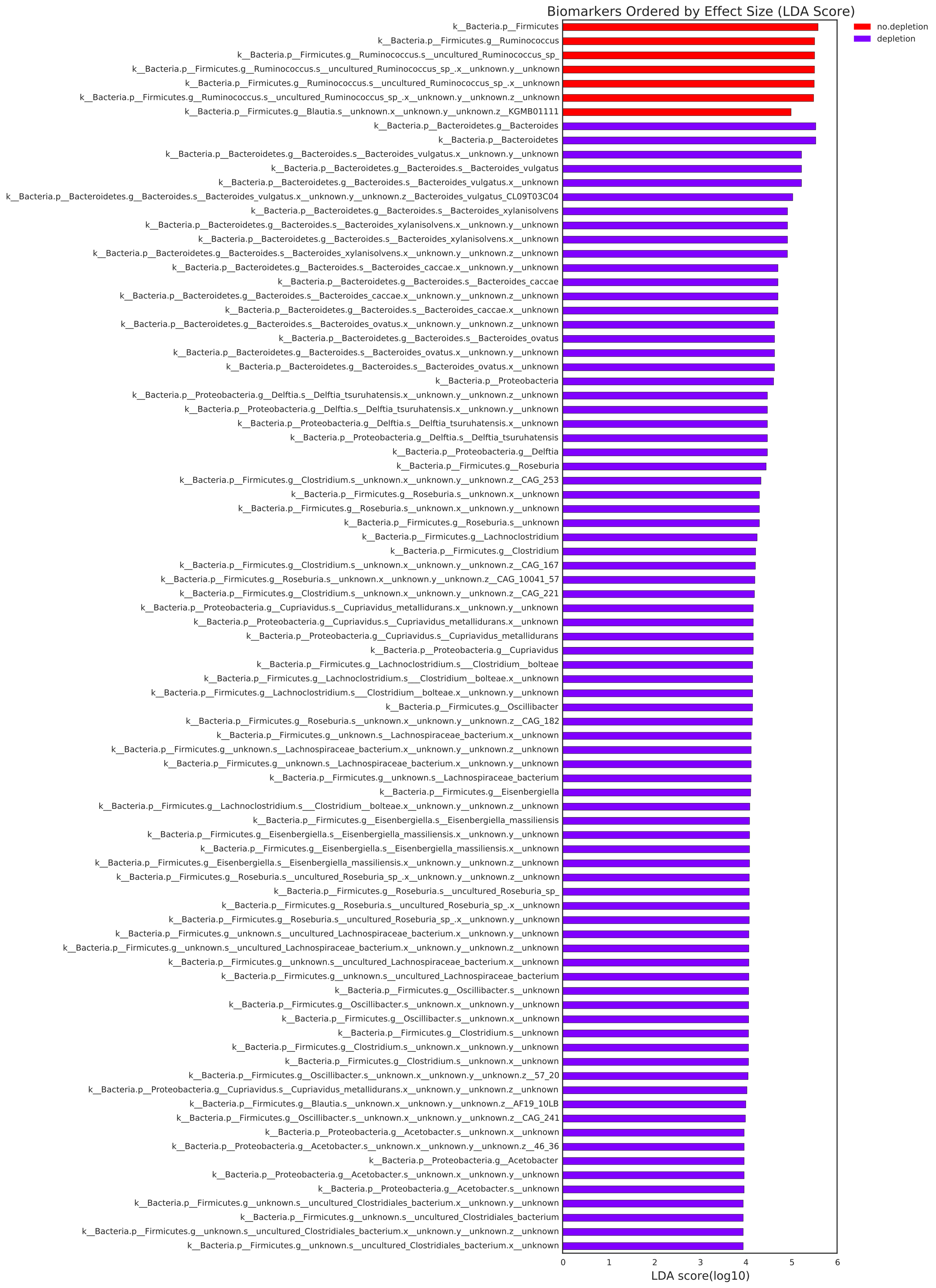
Biomarkers Ordered by Effect Size (LDA Score)
no.depletion
depletion
k__Bacteria.p__Firmicutes
k__Bacteria.p__Firmicutes.g__Ruminococcus
k__Bacteria.p__Firmicutes.g__Ruminococcus.s__uncultured_Ruminococcus_sp_
k__Bacteria.p__Firmicutes.g__Ruminococcus.s__uncultured_Ruminococcus_sp_.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__Ruminococcus.s__uncultured_Ruminococcus_sp_.x__unknown
k__Bacteria.p__Firmicutes.g__Ruminococcus.s__uncultured_Ruminococcus_sp_.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__Blautia.s__unknown.x__unknown.y__unknown.z__KGMB01111
k__Bacteria.p__Bacteroidetes.g__Bacteroides
k__Bacteria.p__Bacteroidetes
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_vulgatus.x__unknown.y__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_vulgatus
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_vulgatus.x__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_vulgatus.x__unknown.y__unknown.z__Bacteroides_vulgatus_CL09T03C04
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_xylanisolvens
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_xylanisolvens.x__unknown.y__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_xylanisolvens.x__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_xylanisolvens.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_caccae.x__unknown.y__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_caccae
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_caccae.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_caccae.x__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_ovatus.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_ovatus
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_ovatus.x__unknown.y__unknown
k__Bacteria.p__Bacteroidetes.g__Bacteroides.s__Bacteroides_ovatus.x__unknown
k__Bacteria.p__Proteobacteria
k__Bacteria.p__Proteobacteria.g__Delftia.s__Delftia_tsuruhatensis.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Proteobacteria.g__Delftia.s__Delftia_tsuruhatensis.x__unknown.y__unknown
k__Bacteria.p__Proteobacteria.g__Delftia.s__Delftia_tsuruhatensis.x__unknown
k__Bacteria.p__Proteobacteria.g__Delftia.s__Delftia_tsuruhatensis
k__Bacteria.p__Proteobacteria.g__Delftia
k__Bacteria.p__Firmicutes.g__Roseburia
k__Bacteria.p__Firmicutes.g__Clostridium.s__unknown.x__unknown.y__unknown.z__CAG_253
k__Bacteria.p__Firmicutes.g__Roseburia.s__unknown.x__unknown
k__Bacteria.p__Firmicutes.g__Roseburia.s__unknown.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__Roseburia.s__unknown
k__Bacteria.p__Firmicutes.g__Lachnoclostridium
k__Bacteria.p__Firmicutes.g__Clostridium
k__Bacteria.p__Firmicutes.g__Clostridium.s__unknown.x__unknown.y__unknown.z__CAG_167
k__Bacteria.p__Firmicutes.g__Roseburia.s__unknown.x__unknown.y__unknown.z__CAG_10041_57
k__Bacteria.p__Firmicutes.g__Clostridium.s__unknown.x__unknown.y__unknown.z__CAG_221
k__Bacteria.p__Proteobacteria.g__Cupriavidus.s__Cupriavidus_metallidurans.x__unknown.y__unknown
k__Bacteria.p__Proteobacteria.g__Cupriavidus.s__Cupriavidus_metallidurans.x__unknown
k__Bacteria.p__Proteobacteria.g__Cupriavidus.s__Cupriavidus_metallidurans
k__Bacteria.p__Proteobacteria.g__Cupriavidus
k__Bacteria.p__Firmicutes.g__Lachnoclostridium.s___Clostridium__bolteae
k__Bacteria.p__Firmicutes.g__Lachnoclostridium.s___Clostridium__bolteae.x__unknown
k__Bacteria.p__Firmicutes.g__Lachnoclostridium.s___Clostridium__bolteae.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__Oscillibacter
k__Bacteria.p__Firmicutes.g__Roseburia.s__unknown.x__unknown.y__unknown.z__CAG_182
k__Bacteria.p__Firmicutes.g__unknown.s__Lachnospiraceae_bacterium.x__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__Lachnospiraceae_bacterium.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__Lachnospiraceae_bacterium.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__Lachnospiraceae_bacterium
k__Bacteria.p__Firmicutes.g__Eisenbergiella
k__Bacteria.p__Firmicutes.g__Lachnoclostridium.s___Clostridium__bolteae.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__Eisenbergiella.s__Eisenbergiella_massiliensis
k__Bacteria.p__Firmicutes.g__Eisenbergiella.s__Eisenbergiella_massiliensis.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__Eisenbergiella.s__Eisenbergiella_massiliensis.x__unknown
k__Bacteria.p__Firmicutes.g__Eisenbergiella.s__Eisenbergiella_massiliensis.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__Roseburia.s__uncultured_Roseburia_sp_.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__Roseburia.s__uncultured_Roseburia_sp_
k__Bacteria.p__Firmicutes.g__Roseburia.s__uncultured_Roseburia_sp_.x__unknown
k__Bacteria.p__Firmicutes.g__Roseburia.s__uncultured_Roseburia_sp_.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Lachnospiraceae_bacterium.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Lachnospiraceae_bacterium.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Lachnospiraceae_bacterium.x__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Lachnospiraceae_bacterium
k__Bacteria.p__Firmicutes.g__Oscillibacter.s__unknown
k__Bacteria.p__Firmicutes.g__Oscillibacter.s__unknown.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__Oscillibacter.s__unknown.x__unknown
k__Bacteria.p__Firmicutes.g__Clostridium.s__unknown
k__Bacteria.p__Firmicutes.g__Clostridium.s__unknown.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__Clostridium.s__unknown.x__unknown
k__Bacteria.p__Firmicutes.g__Oscillibacter.s__unknown.x__unknown.y__unknown.z__57_20
k__Bacteria.p__Proteobacteria.g__Cupriavidus.s__Cupriavidus_metallidurans.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__Blautia.s__unknown.x__unknown.y__unknown.z__AF19_10LB
k__Bacteria.p__Firmicutes.g__Oscillibacter.s__unknown.x__unknown.y__unknown.z__CAG_241
k__Bacteria.p__Proteobacteria.g__Acetobacter.s__unknown.x__unknown
k__Bacteria.p__Proteobacteria.g__Acetobacter.s__unknown.x__unknown.y__unknown.z__46_36
k__Bacteria.p__Proteobacteria.g__Acetobacter
k__Bacteria.p__Proteobacteria.g__Acetobacter.s__unknown.x__unknown.y__unknown
k__Bacteria.p__Proteobacteria.g__Acetobacter.s__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Clostridiales_bacterium.x__unknown.y__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Clostridiales_bacterium
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Clostridiales_bacterium.x__unknown.y__unknown.z__unknown
k__Bacteria.p__Firmicutes.g__unknown.s__uncultured_Clostridiales_bacterium.x__unknown
0
1
2
3
4
5
6
LDA score(log10)