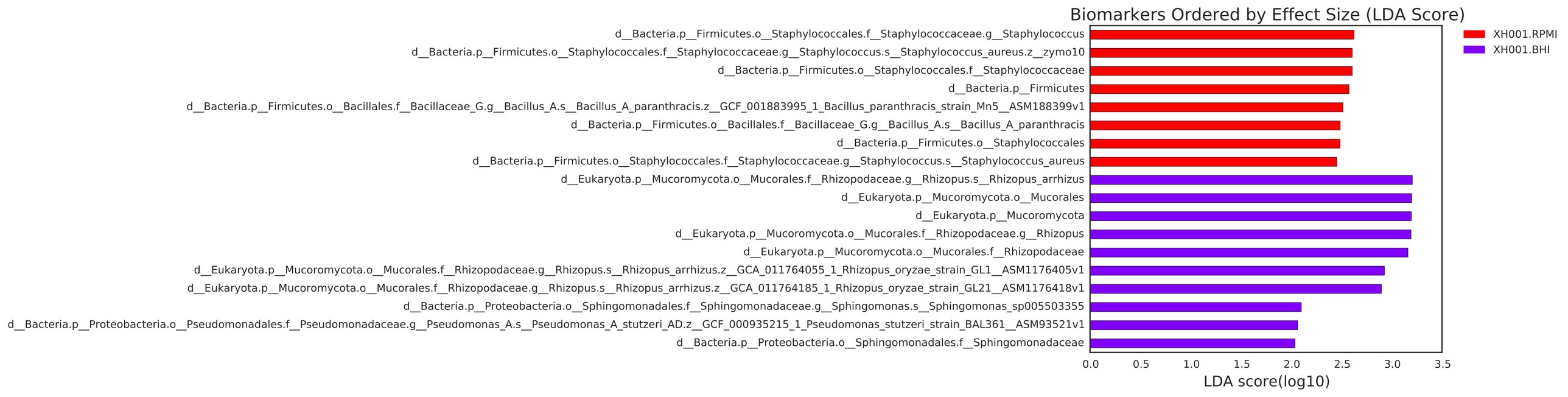

Biomarkers Ordered by Effect Size (LDA Score)
XH001.RPMI
XH001.BHI
d__Bacteria.p__Firmicutes.o__Staphylococcales.f__Staphylococcaceae.g__Staphylococcus
d__Bacteria.p__Firmicutes.o__Staphylococcales.f__Staphylococcaceae.g__Staphylococcus.s__Staphylococcus_aureus.z__zymo10
d__Bacteria.p__Firmicutes.o__Staphylococcales.f__Staphylococcaceae
d__Bacteria.p__Firmicutes
d__Bacteria.p__Firmicutes.o__Bacillales.f__Bacillaceae_G.g__Bacillus_A.s__Bacillus_A_paranthracis.z__GCF_001883995_1_Bacillus_paranthracis_strain_Mn5__ASM188399v1
d__Bacteria.p__Firmicutes.o__Bacillales.f__Bacillaceae_G.g__Bacillus_A.s__Bacillus_A_paranthracis
d__Bacteria.p__Firmicutes.o__Staphylococcales
d__Bacteria.p__Firmicutes.o__Staphylococcales.f__Staphylococcaceae.g__Staphylococcus.s__Staphylococcus_aureus
d__Eukaryota.p__Mucoromycota.o__Mucorales.f__Rhizopodaceae.g__Rhizopus.s__Rhizopus_arrhizus
d__Eukaryota.p__Mucoromycota.o__Mucorales
d__Eukaryota.p__Mucoromycota
d__Eukaryota.p__Mucoromycota.o__Mucorales.f__Rhizopodaceae.g__Rhizopus
d__Eukaryota.p__Mucoromycota.o__Mucorales.f__Rhizopodaceae
d__Eukaryota.p__Mucoromycota.o__Mucorales.f__Rhizopodaceae.g__Rhizopus.s__Rhizopus_arrhizus.z__GCA_011764055_1_Rhizopus_oryzae_strain_GL1__ASM1176405v1
d__Eukaryota.p__Mucoromycota.o__Mucorales.f__Rhizopodaceae.g__Rhizopus.s__Rhizopus_arrhizus.z__GCA_011764185_1_Rhizopus_oryzae_strain_GL21__ASM1176418v1
d__Bacteria.p__Proteobacteria.o__Sphingomonadales.f__Sphingomonadaceae.g__Sphingomonas.s__Sphingomonas_sp005503355
d__Bacteria.p__Proteobacteria.o__Pseudomonadales.f__Pseudomonadaceae.g__Pseudomonas_A.s__Pseudomonas_A_stutzeri_AD.z__GCF_000935215_1_Pseudomonas_stutzeri_strain_BAL361__ASM93521v1
d__Bacteria.p__Proteobacteria.o__Sphingomonadales.f__Sphingomonadaceae
0.0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
LDA score(log10)