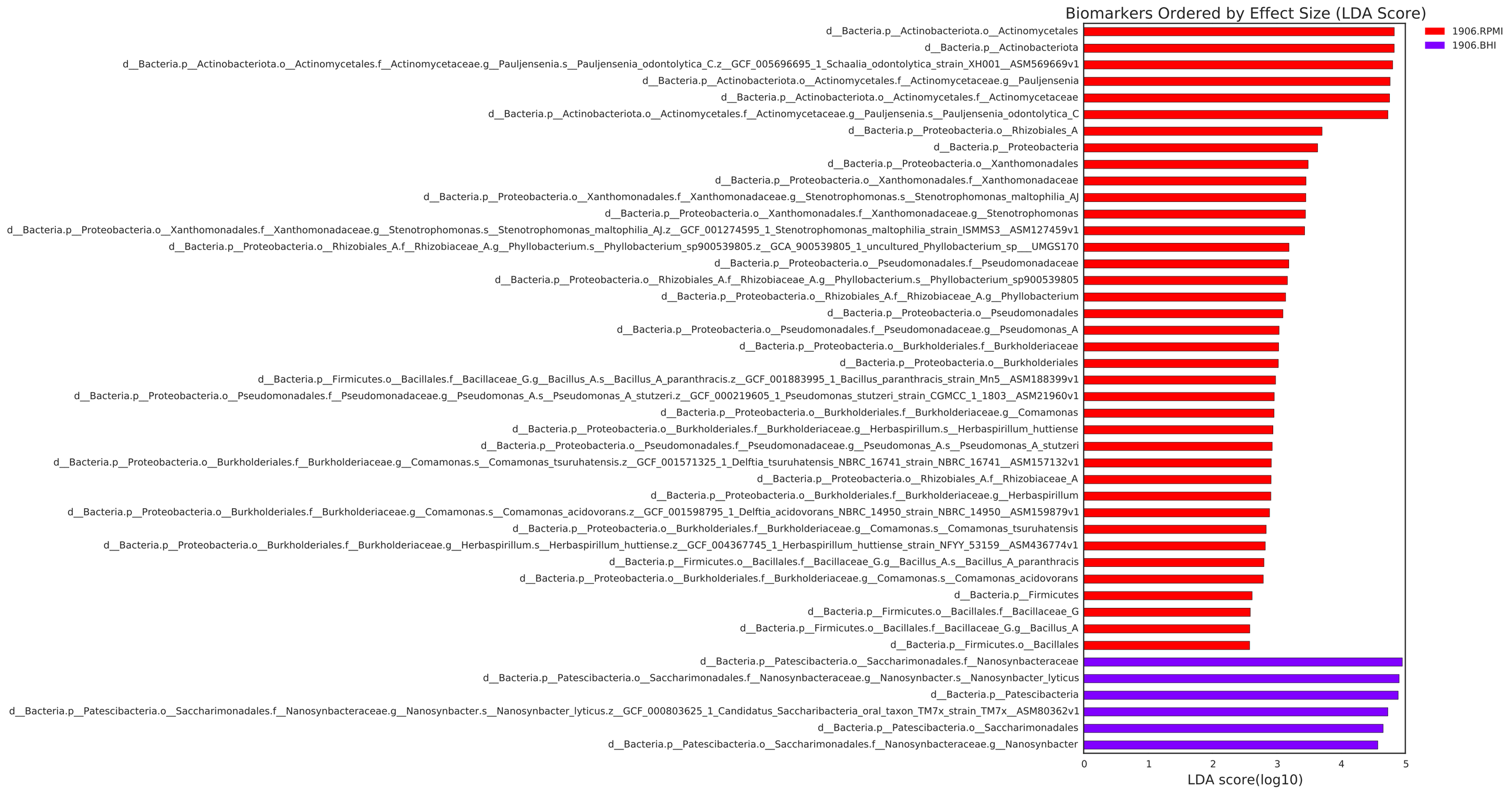

Biomarkers Ordered by Effect Size (LDA Score)
1906.RPMI
1906.BHI
d__Bacteria.p__Actinobacteriota.o__Actinomycetales
d__Bacteria.p__Actinobacteriota
d__Bacteria.p__Actinobacteriota.o__Actinomycetales.f__Actinomycetaceae.g__Pauljensenia.s__Pauljensenia_odontolytica_C.z__GCF_005696695_1_Schaalia_odontolytica_strain_XH001__ASM569669v1
d__Bacteria.p__Actinobacteriota.o__Actinomycetales.f__Actinomycetaceae.g__Pauljensenia
d__Bacteria.p__Actinobacteriota.o__Actinomycetales.f__Actinomycetaceae
d__Bacteria.p__Actinobacteriota.o__Actinomycetales.f__Actinomycetaceae.g__Pauljensenia.s__Pauljensenia_odontolytica_C
d__Bacteria.p__Proteobacteria.o__Rhizobiales_A
d__Bacteria.p__Proteobacteria
d__Bacteria.p__Proteobacteria.o__Xanthomonadales
d__Bacteria.p__Proteobacteria.o__Xanthomonadales.f__Xanthomonadaceae
d__Bacteria.p__Proteobacteria.o__Xanthomonadales.f__Xanthomonadaceae.g__Stenotrophomonas.s__Stenotrophomonas_maltophilia_AJ
d__Bacteria.p__Proteobacteria.o__Xanthomonadales.f__Xanthomonadaceae.g__Stenotrophomonas
d__Bacteria.p__Proteobacteria.o__Xanthomonadales.f__Xanthomonadaceae.g__Stenotrophomonas.s__Stenotrophomonas_maltophilia_AJ.z__GCF_001274595_1_Stenotrophomonas_maltophilia_strain_ISMMS3__ASM127459v1
d__Bacteria.p__Proteobacteria.o__Rhizobiales_A.f__Rhizobiaceae_A.g__Phyllobacterium.s__Phyllobacterium_sp900539805.z__GCA_900539805_1_uncultured_Phyllobacterium_sp___UMGS170
d__Bacteria.p__Proteobacteria.o__Pseudomonadales.f__Pseudomonadaceae
d__Bacteria.p__Proteobacteria.o__Rhizobiales_A.f__Rhizobiaceae_A.g__Phyllobacterium.s__Phyllobacterium_sp900539805
d__Bacteria.p__Proteobacteria.o__Rhizobiales_A.f__Rhizobiaceae_A.g__Phyllobacterium
d__Bacteria.p__Proteobacteria.o__Pseudomonadales
d__Bacteria.p__Proteobacteria.o__Pseudomonadales.f__Pseudomonadaceae.g__Pseudomonas_A
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae
d__Bacteria.p__Proteobacteria.o__Burkholderiales
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__Proteobacteria.o__Pseudomonadales.f__Pseudomonadaceae.g__Pseudomonas_A.s__Pseudomonas_A_stutzeri.z__GCF_000219605_1_Pseudomonas_stutzeri_strain_CGMCC_1_1803__ASM21960v1
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Comamonas
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Herbaspirillum.s__Herbaspirillum_huttiense
d__Bacteria.p__Proteobacteria.o__Pseudomonadales.f__Pseudomonadaceae.g__Pseudomonas_A.s__Pseudomonas_A_stutzeri
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Comamonas.s__Comamonas_tsuruhatensis.z__GCF_001571325_1_Delftia_tsuruhatensis_NBRC_16741_strain_NBRC_16741__ASM157132v1
d__Bacteria.p__Proteobacteria.o__Rhizobiales_A.f__Rhizobiaceae_A
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Herbaspirillum
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Comamonas.s__Comamonas_acidovorans.z__GCF_001598795_1_Delftia_acidovorans_NBRC_14950_strain_NBRC_14950__ASM159879v1
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Comamonas.s__Comamonas_tsuruhatensis
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Herbaspirillum.s__Herbaspirillum_huttiense.z__GCF_004367745_1_Herbaspirillum_huttiense_strain_NFYY_53159__ASM436774v1
d__Bacteria.p__Firmicutes.o__Bacillales.f__Bacillaceae_G.g__Bacillus_A.s__Bacillus_A_paranthracis
d__Bacteria.p__Proteobacteria.o__Burkholderiales.f__Burkholderiaceae.g__Comamonas.s__Comamonas_acidovorans
d__Bacteria.p__Firmicutes
d__Bacteria.p__Firmicutes.o__Bacillales.f__Bacillaceae_G
d__Bacteria.p__Firmicutes.o__Bacillales.f__Bacillaceae_G.g__Bacillus_A
d__Bacteria.p__Firmicutes.o__Bacillales
d__Bacteria.p__Patescibacteria.o__Saccharimonadales.f__Nanosynbacteraceae
d__Bacteria.p__Patescibacteria.o__Saccharimonadales.f__Nanosynbacteraceae.g__Nanosynbacter.s__Nanosynbacter_lyticus
d__Bacteria.p__Patescibacteria
d__Bacteria.p__Patescibacteria.o__Saccharimonadales.f__Nanosynbacteraceae.g__Nanosynbacter.s__Nanosynbacter_lyticus.z__GCF_000803625_1_Candidatus_Saccharibacteria_oral_taxon_TM7x_strain_TM7x__ASM80362v1
d__Bacteria.p__Patescibacteria.o__Saccharimonadales
d__Bacteria.p__Patescibacteria.o__Saccharimonadales.f__Nanosynbacteraceae.g__Nanosynbacter
0
1
2
3
4
5
LDA score(log10)