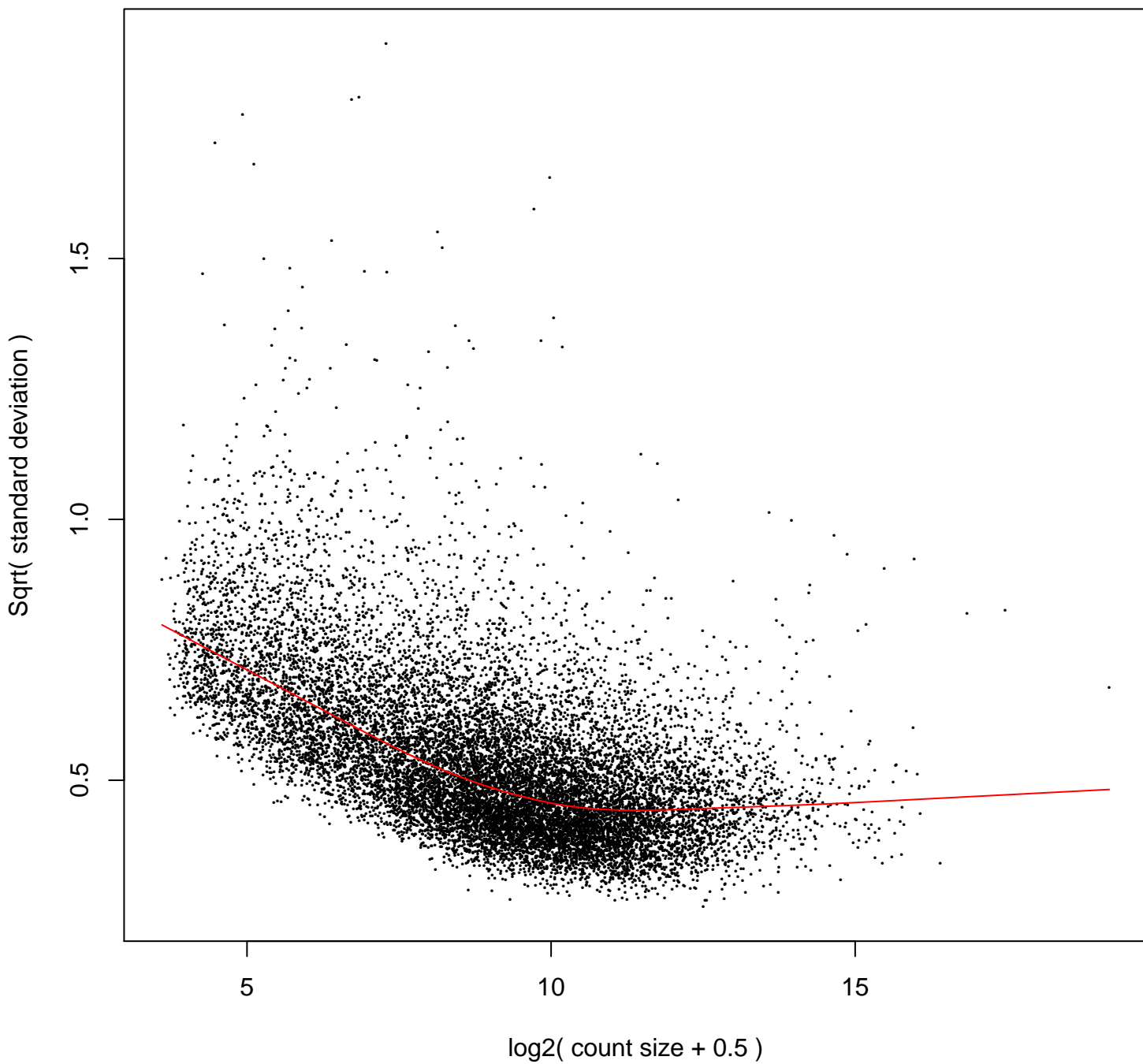
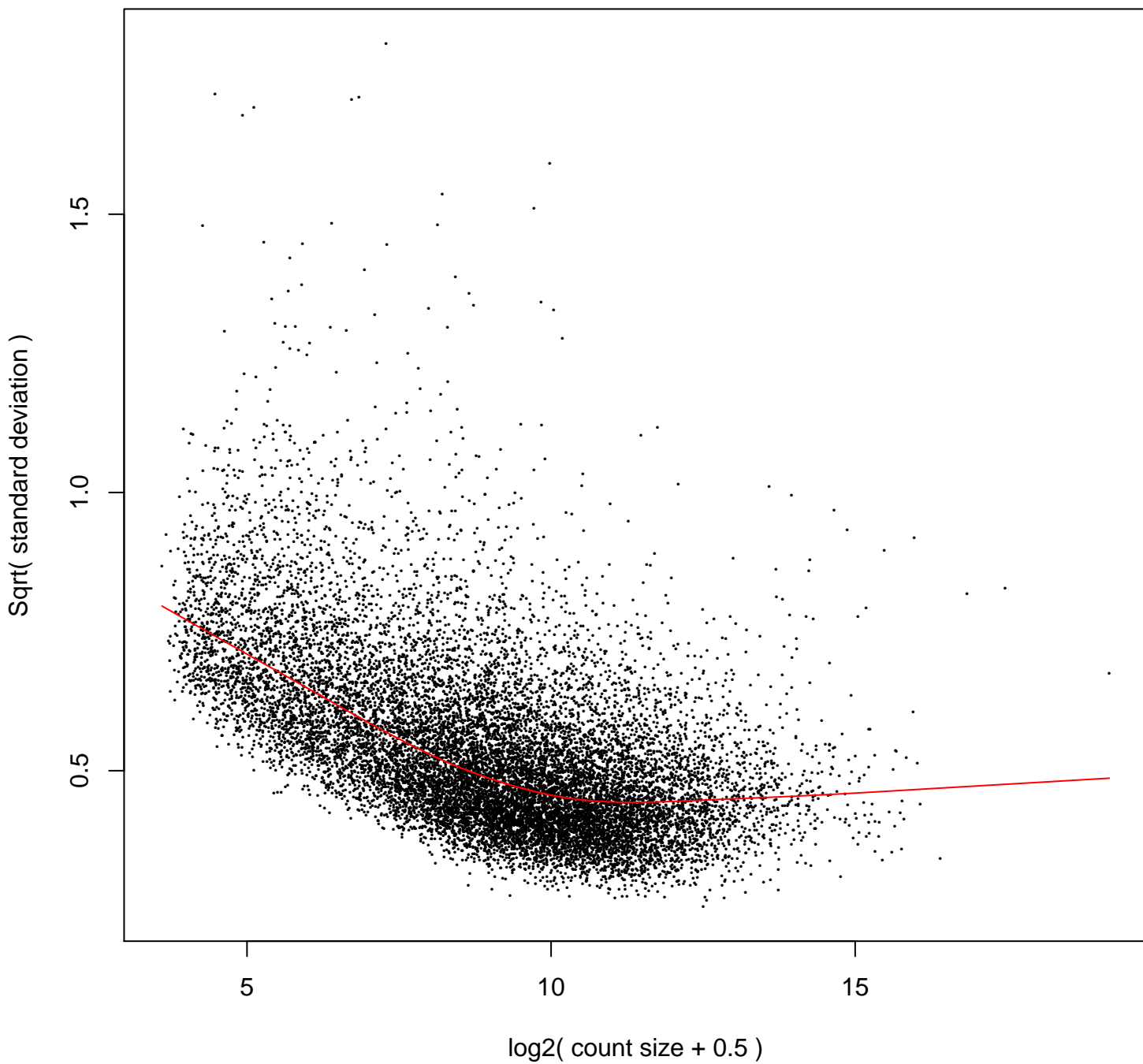


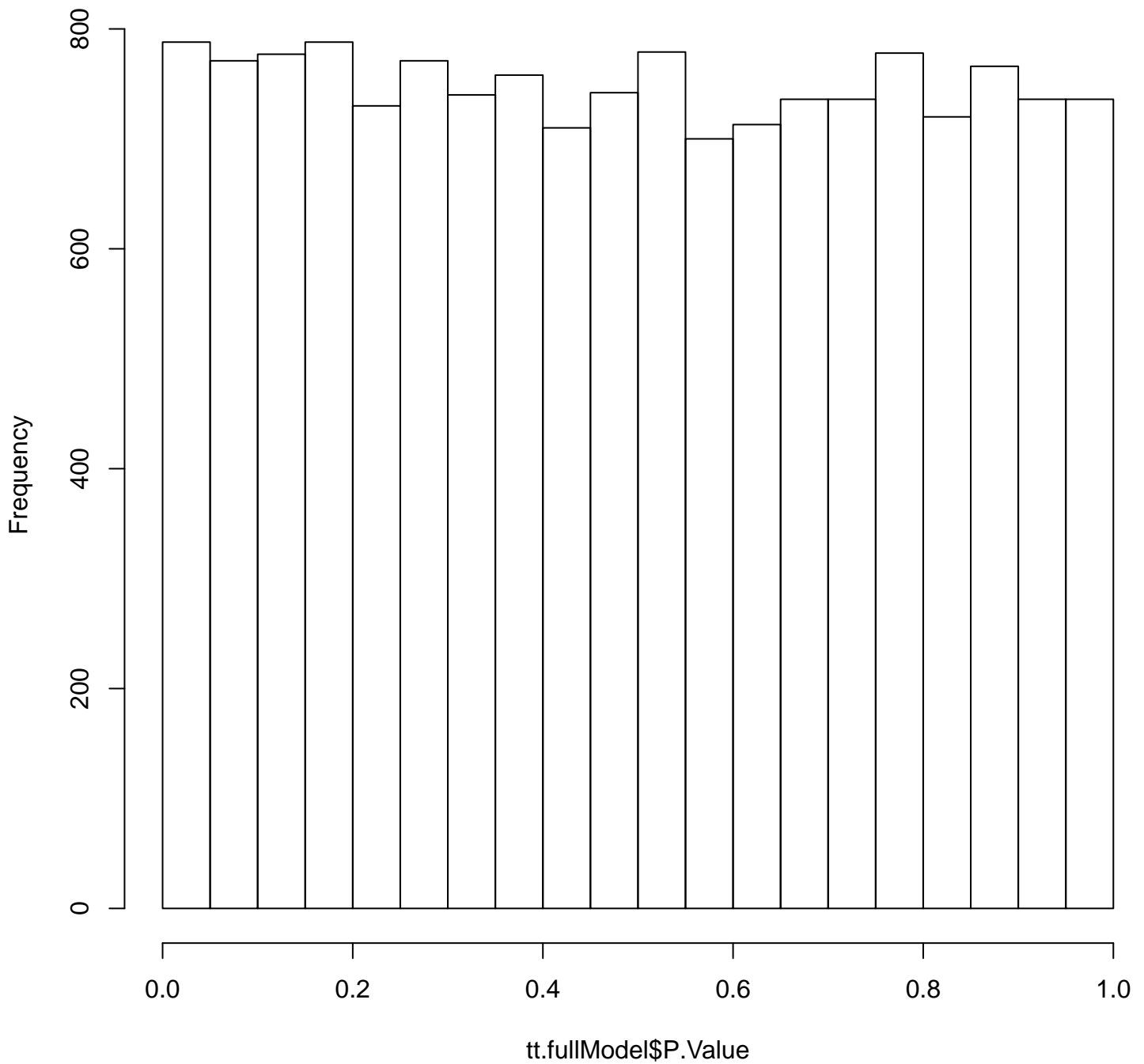
voom: Mean–variance trend



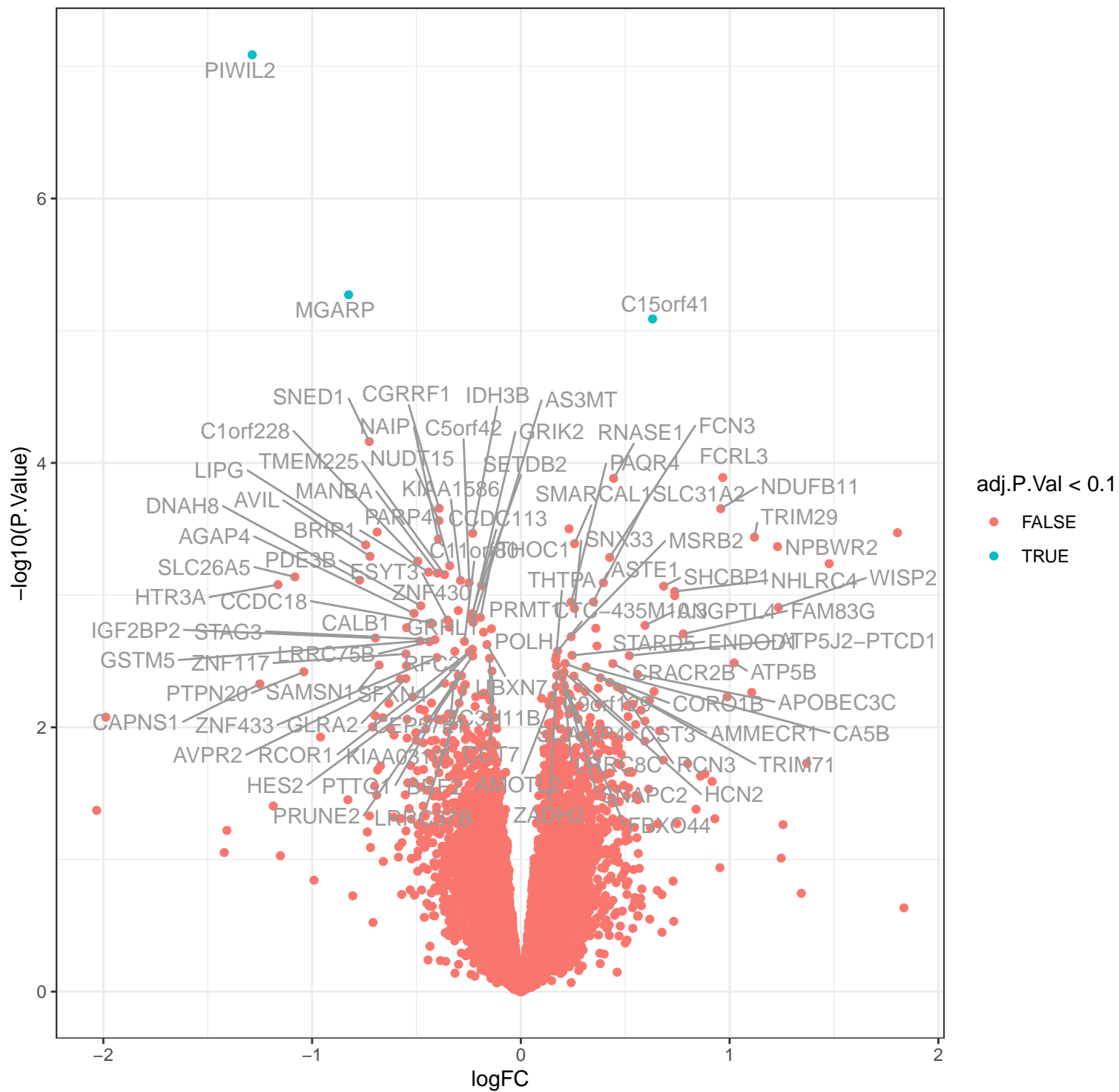
voom: Mean–variance trend



Histogram of tt.fullModel\$P.Value



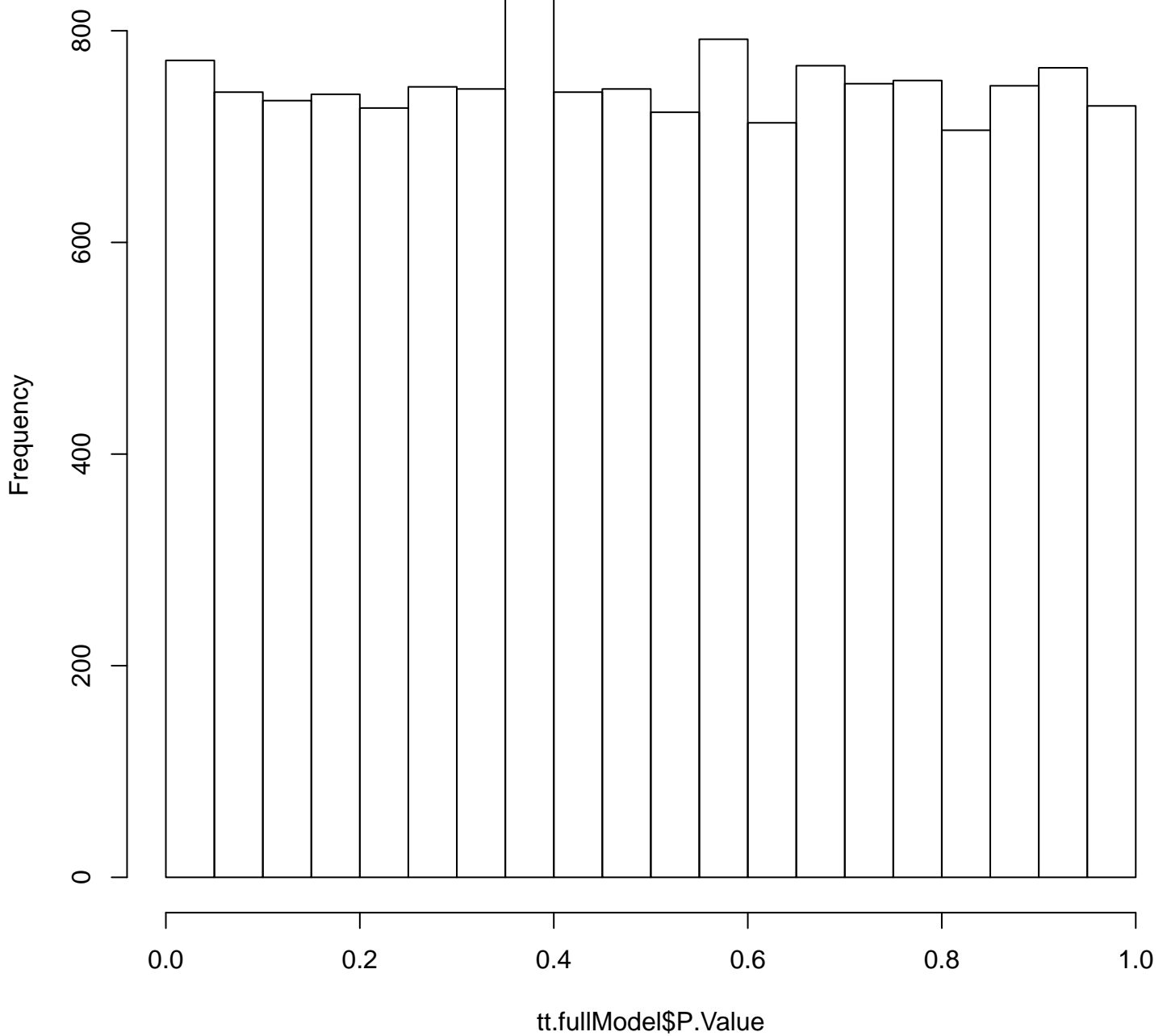
mmul CON_vs_MIA_ALL DGE Volcano Plot



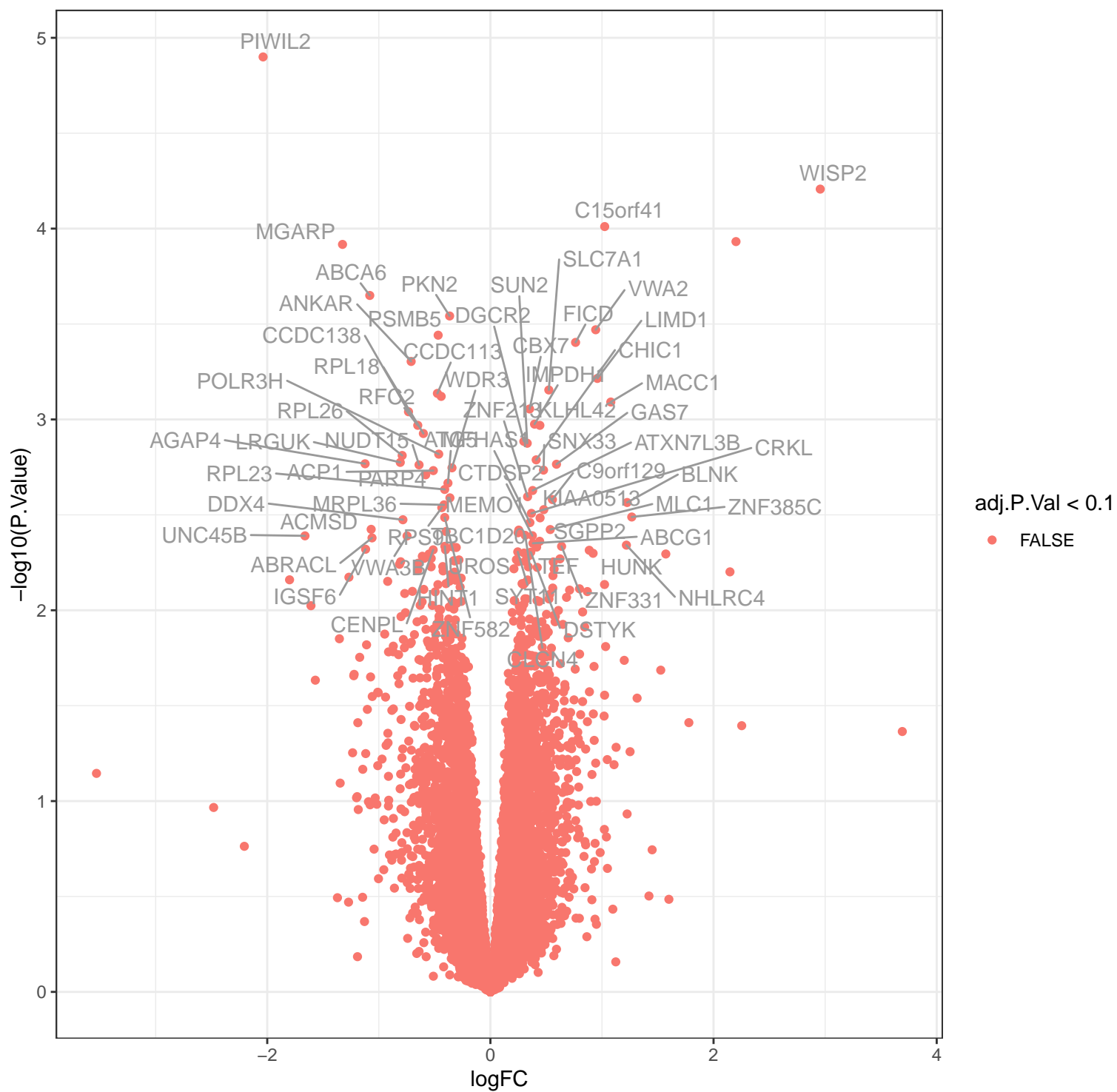
mmul CON_vs_MIA_ALL DGE Volcano Plot (after fdrtool)



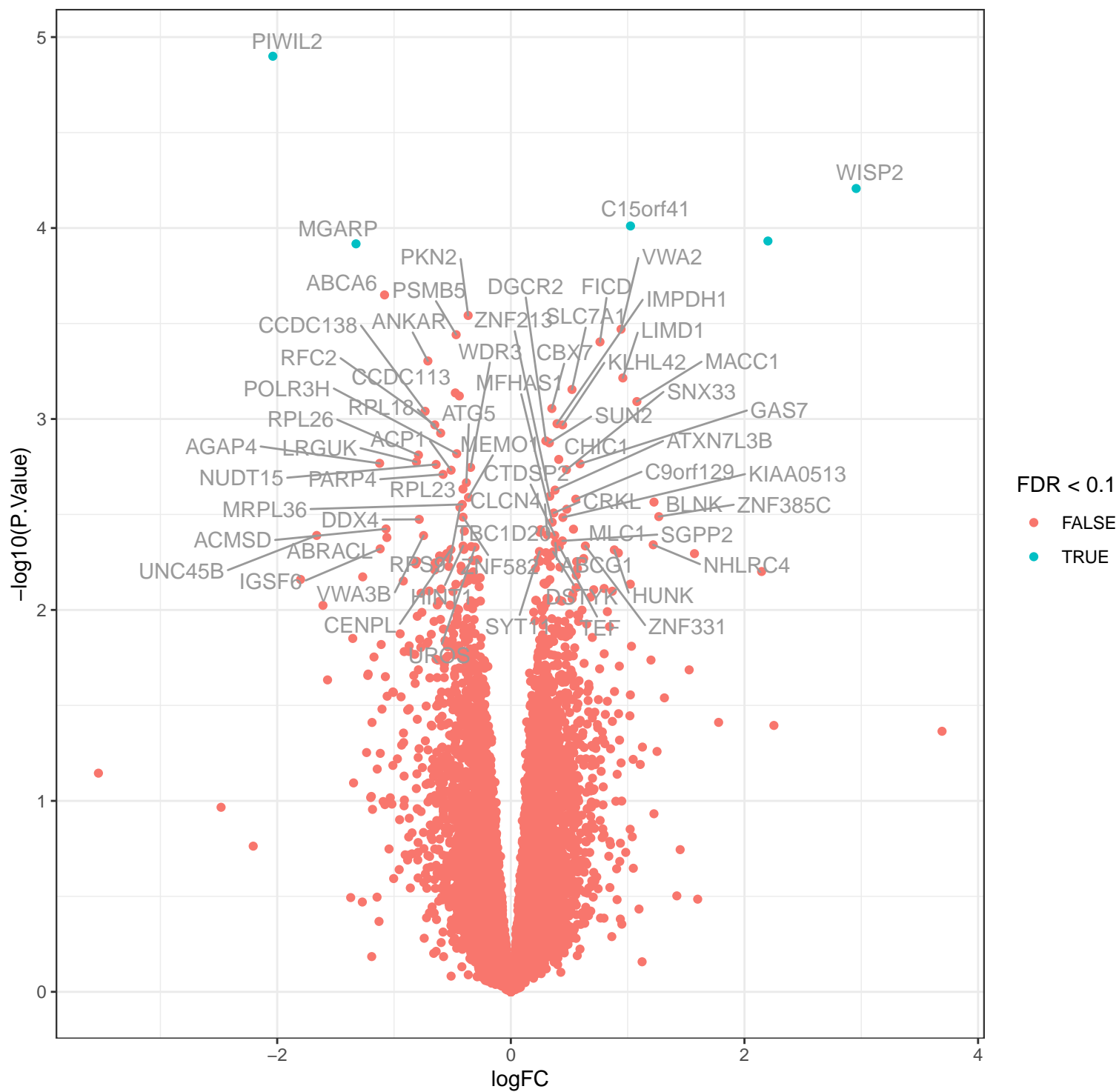
Histogram of tt.fullModel\$P.Value



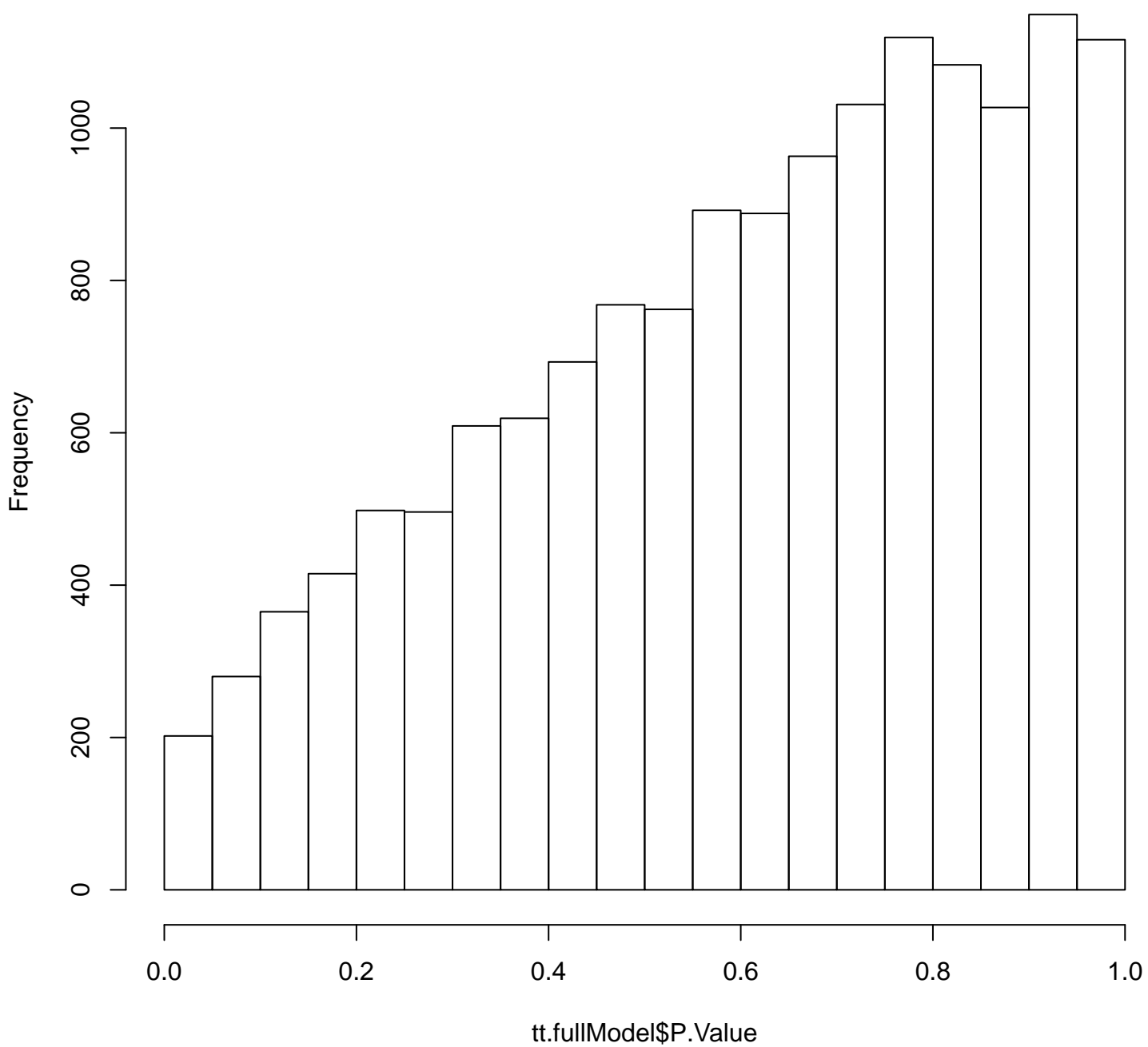
mmul CON_vs_MIA_DLPFC DGE Volcano Plot



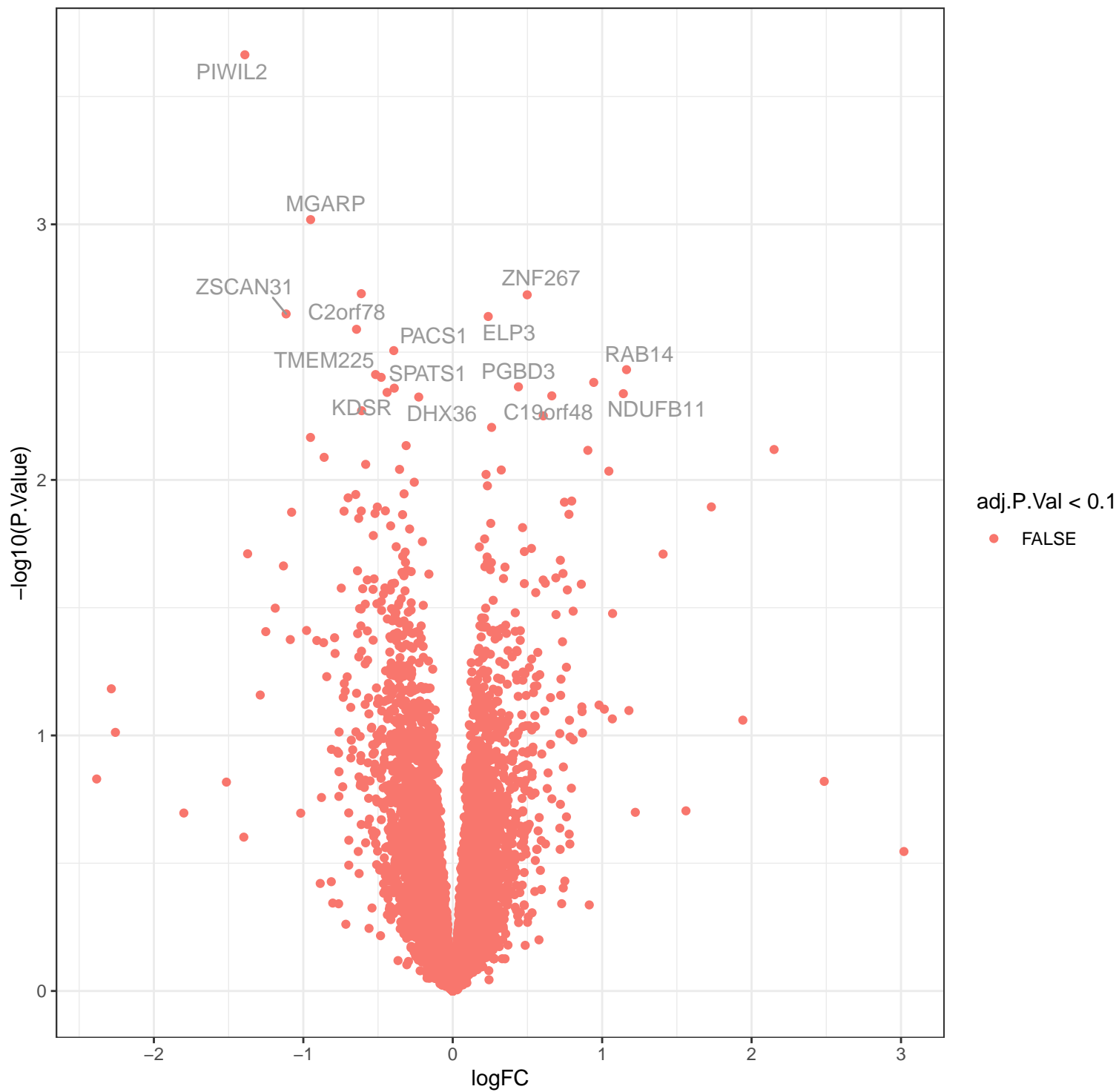
mmul CON_vs_MIA_DLPFC DGE Volcano Plot (after fdrtool)



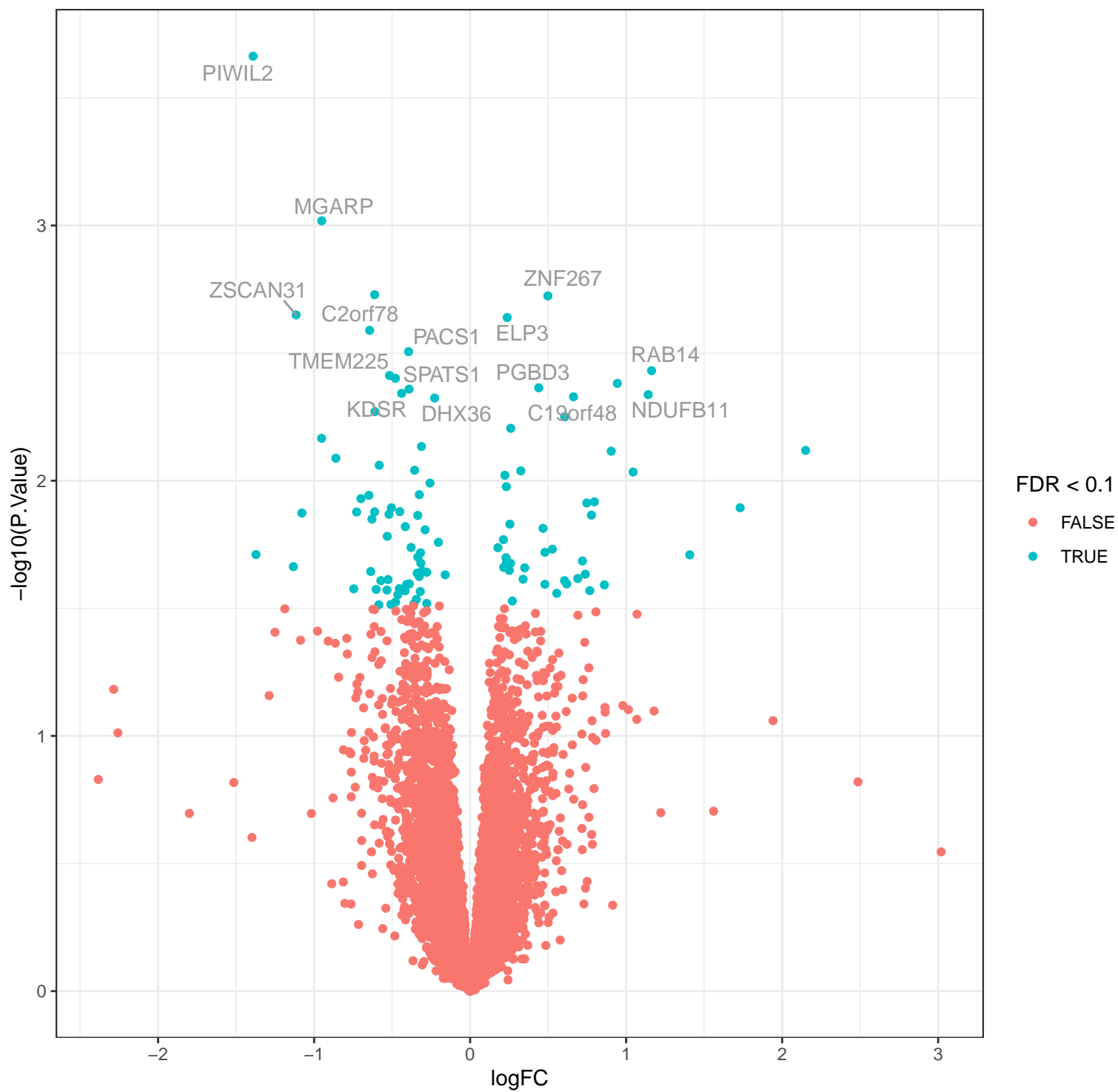
Histogram of tt.fullModel\$P.Value



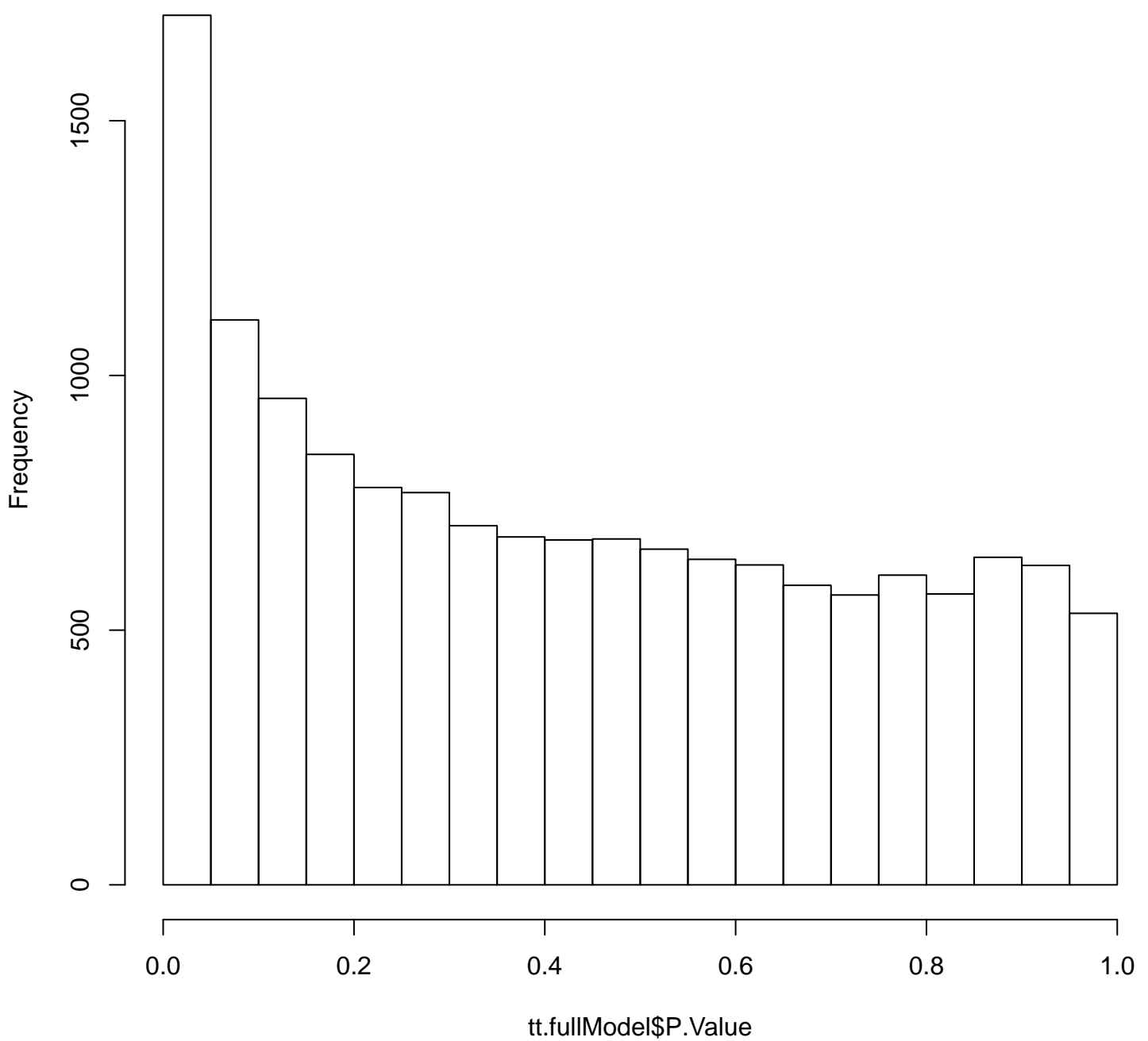
mmul CON_vs_MIA_ACC DGE Volcano Plot



mmul CON_vs_MIA_ACC DGE Volcano Plot (after fdrtool)



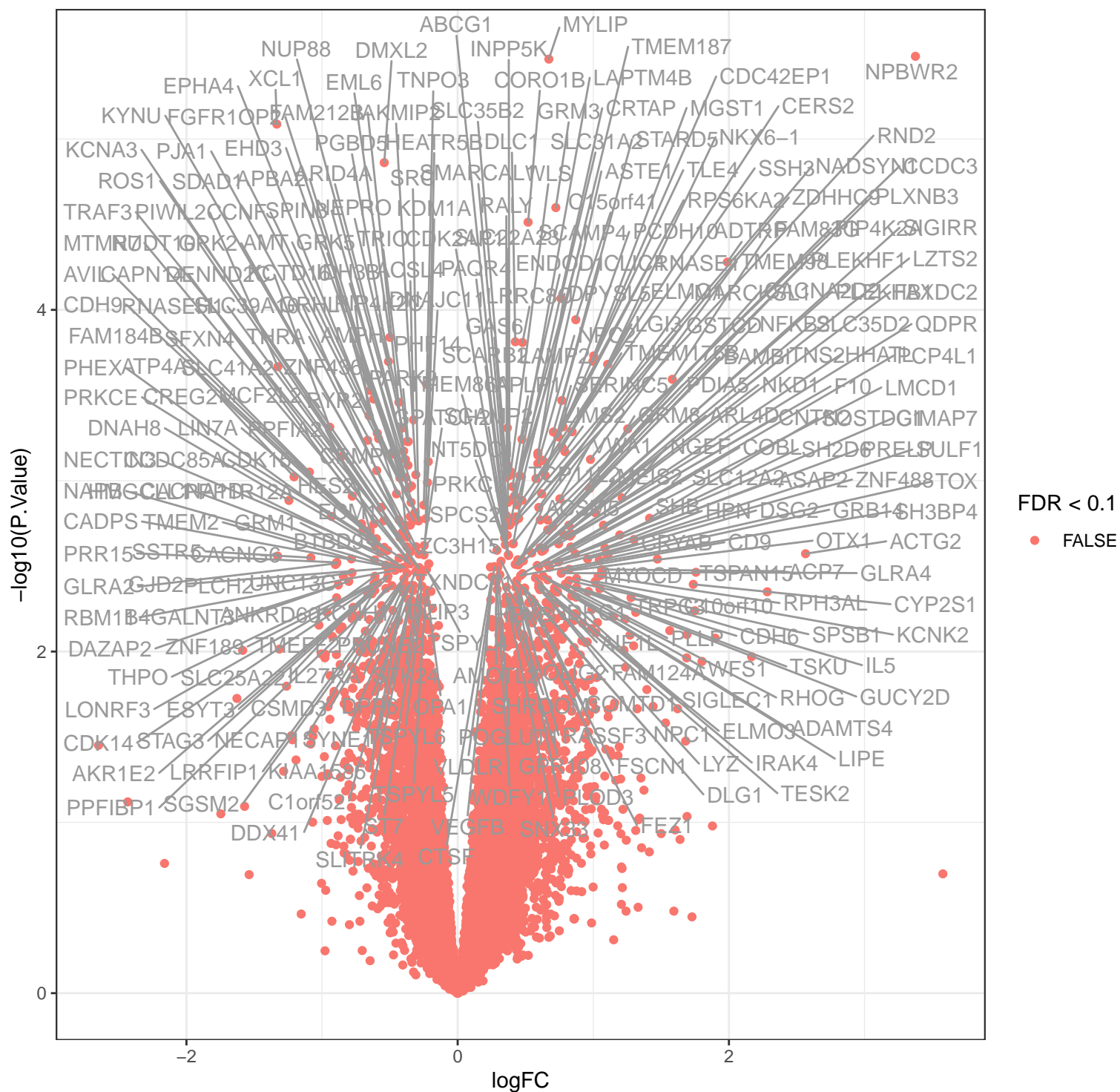
Histogram of tt.fullModel\$P.Value



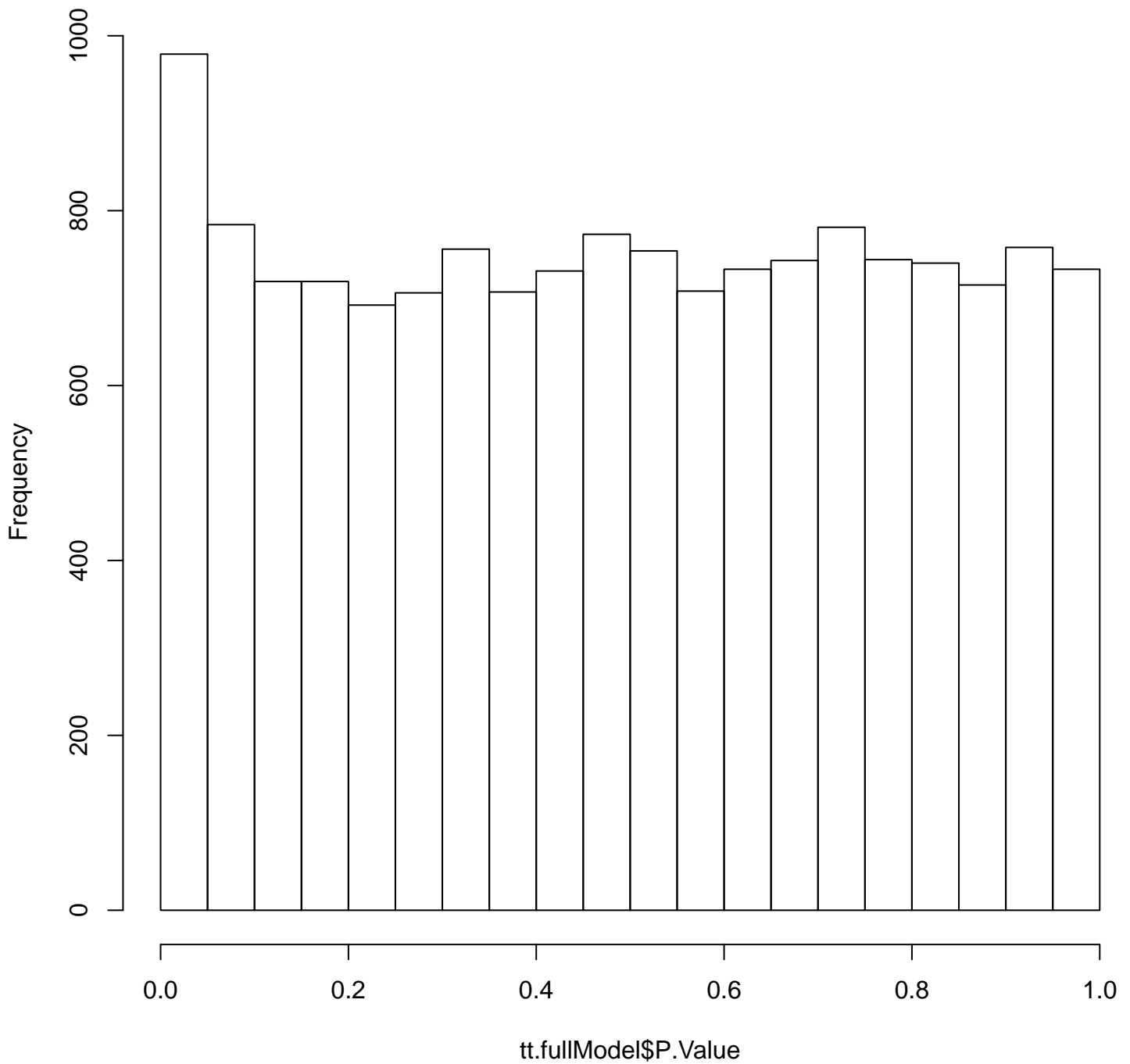
Volcano plot showing log2 fold change (logFC) on the x-axis and -log10(p-value) on the y-axis. The plot displays a large number of genes, with many labeled at the top. Red dots represent significant genes, while grey dots represent non-significant genes. The x-axis ranges from -2 to 2, and the y-axis ranges from 0 to 10. The plot shows a clear separation between significant and non-significant genes, with significant genes clustered at higher -log10(p-value) values.

- FALSE
- TRUE

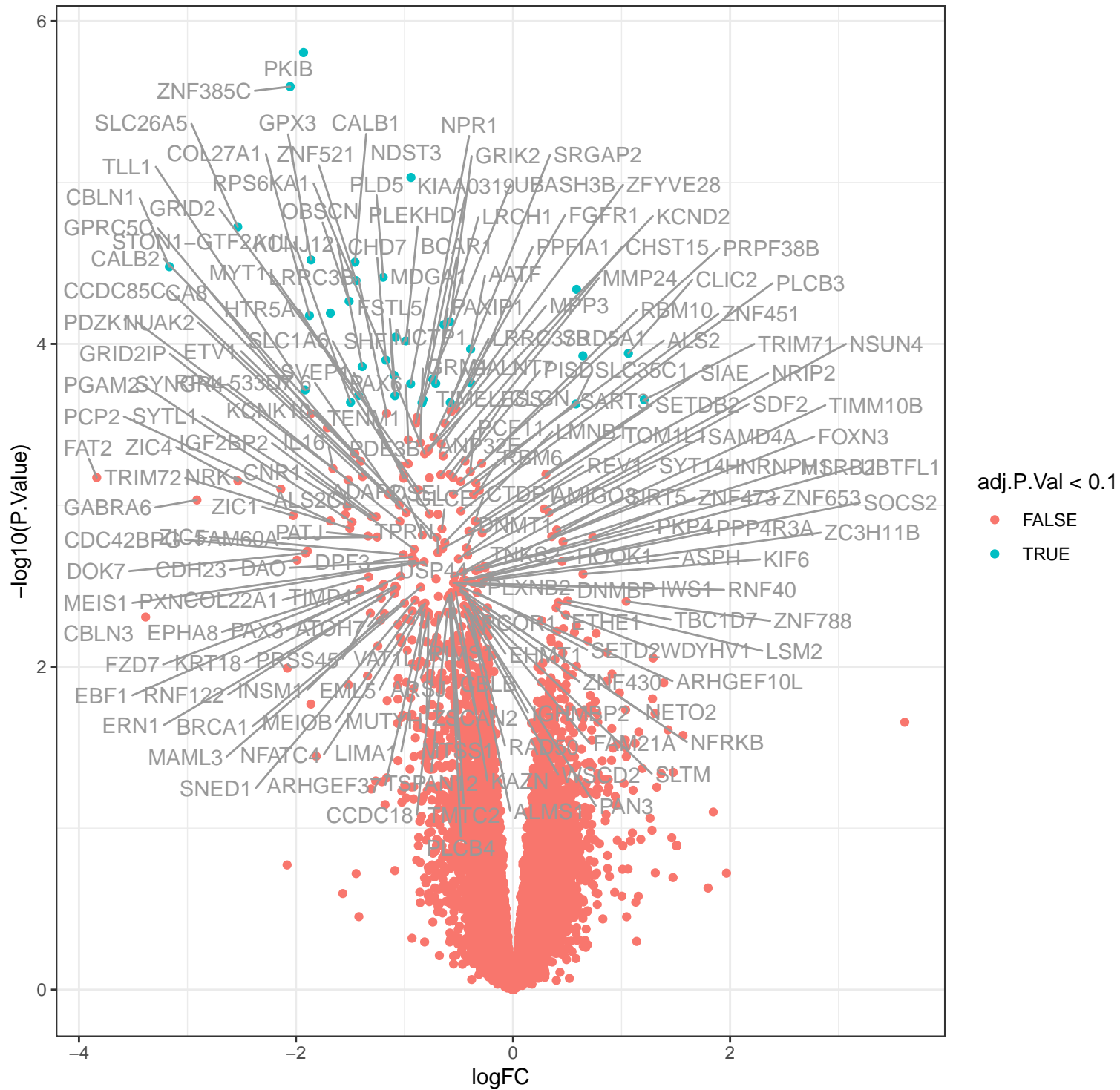
mmul CON_vs_MIA_HC DGE Volcano Plot (after fdrtool)



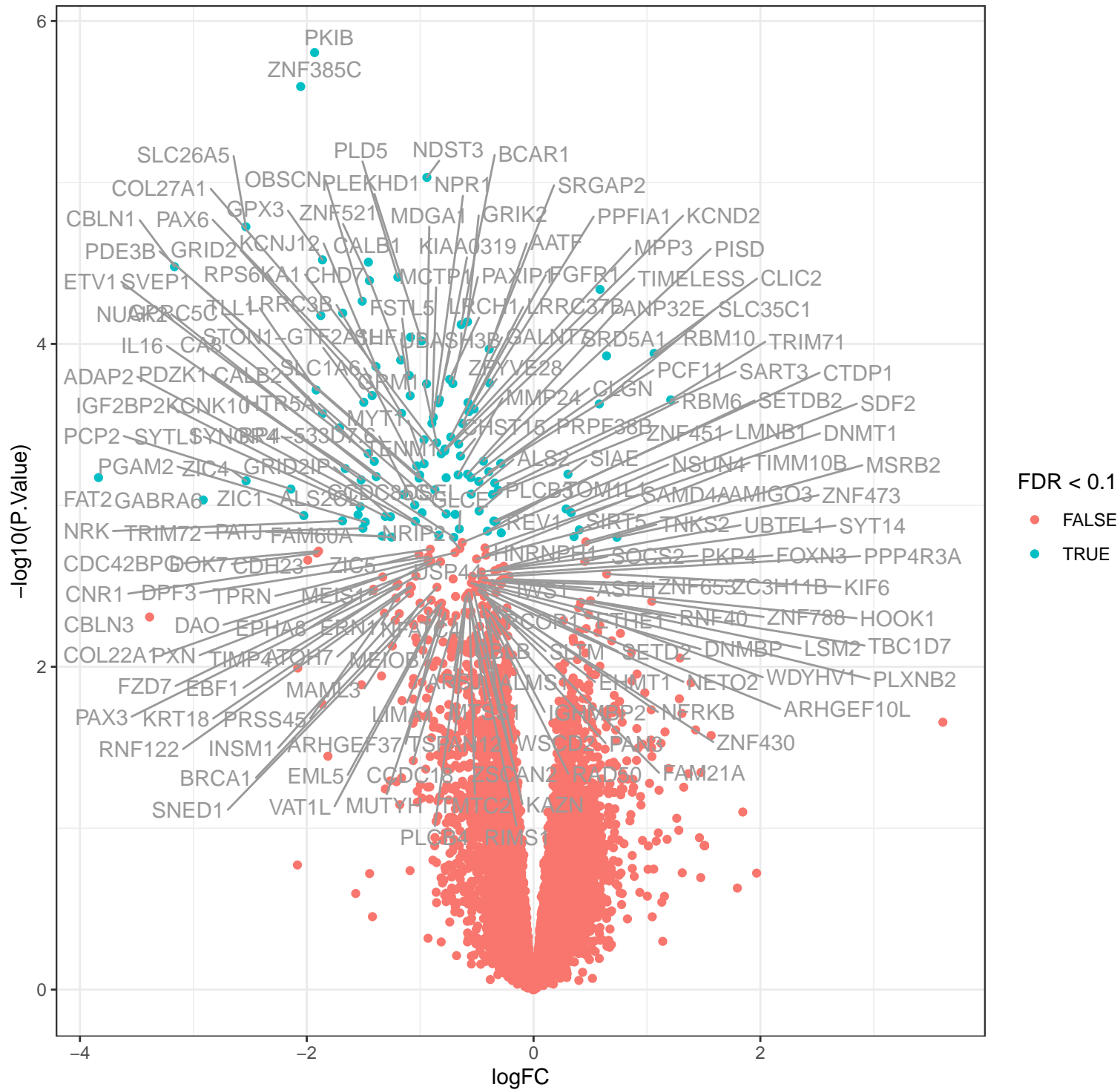
Histogram of tt.fullModel\$P.Value



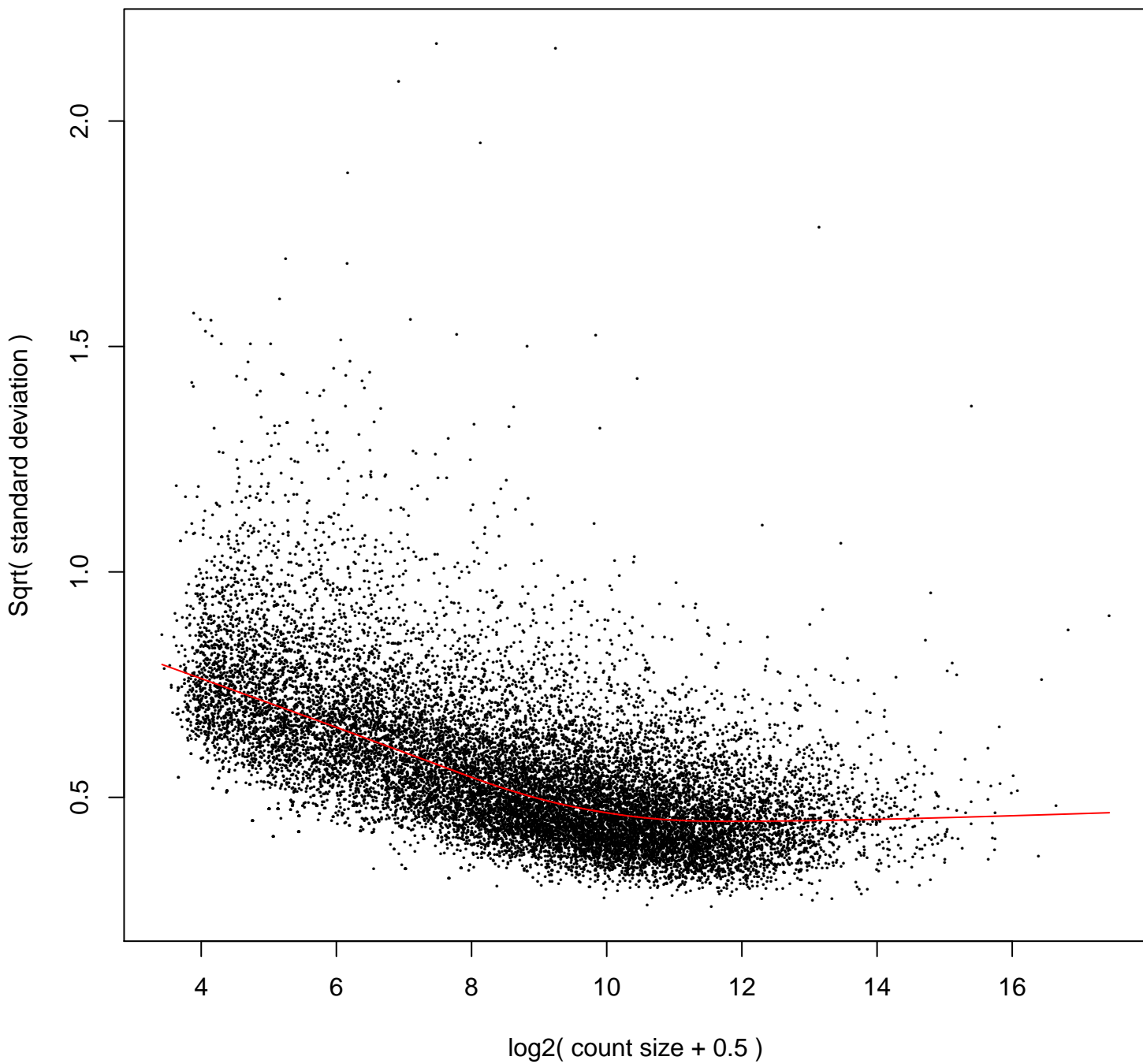
mmul CON_vs_MIA_V1 DGE Volcano Plot



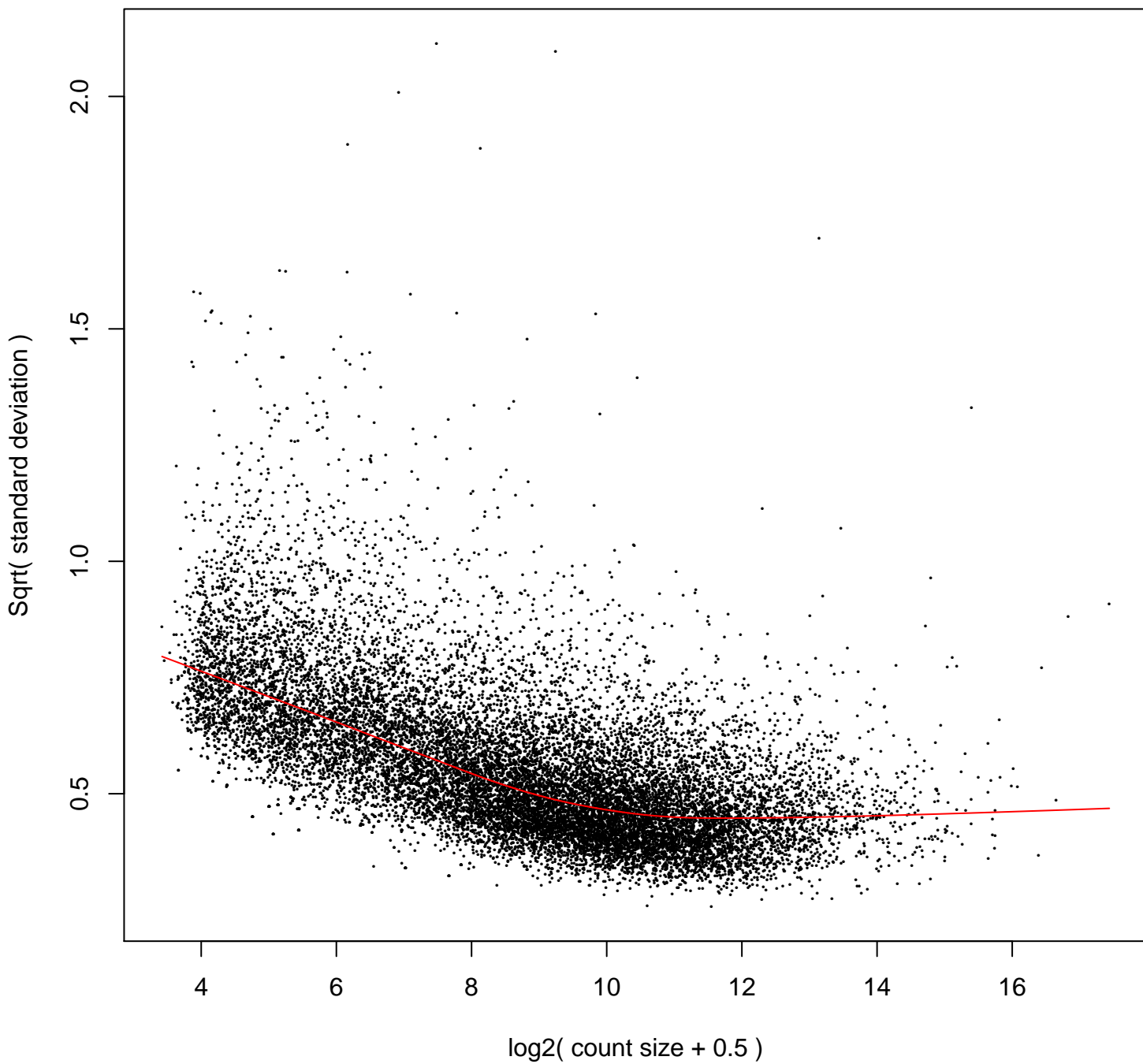
mmul CON_vs_MIA_V1 DGE Volcano Plot (after fdrtool)



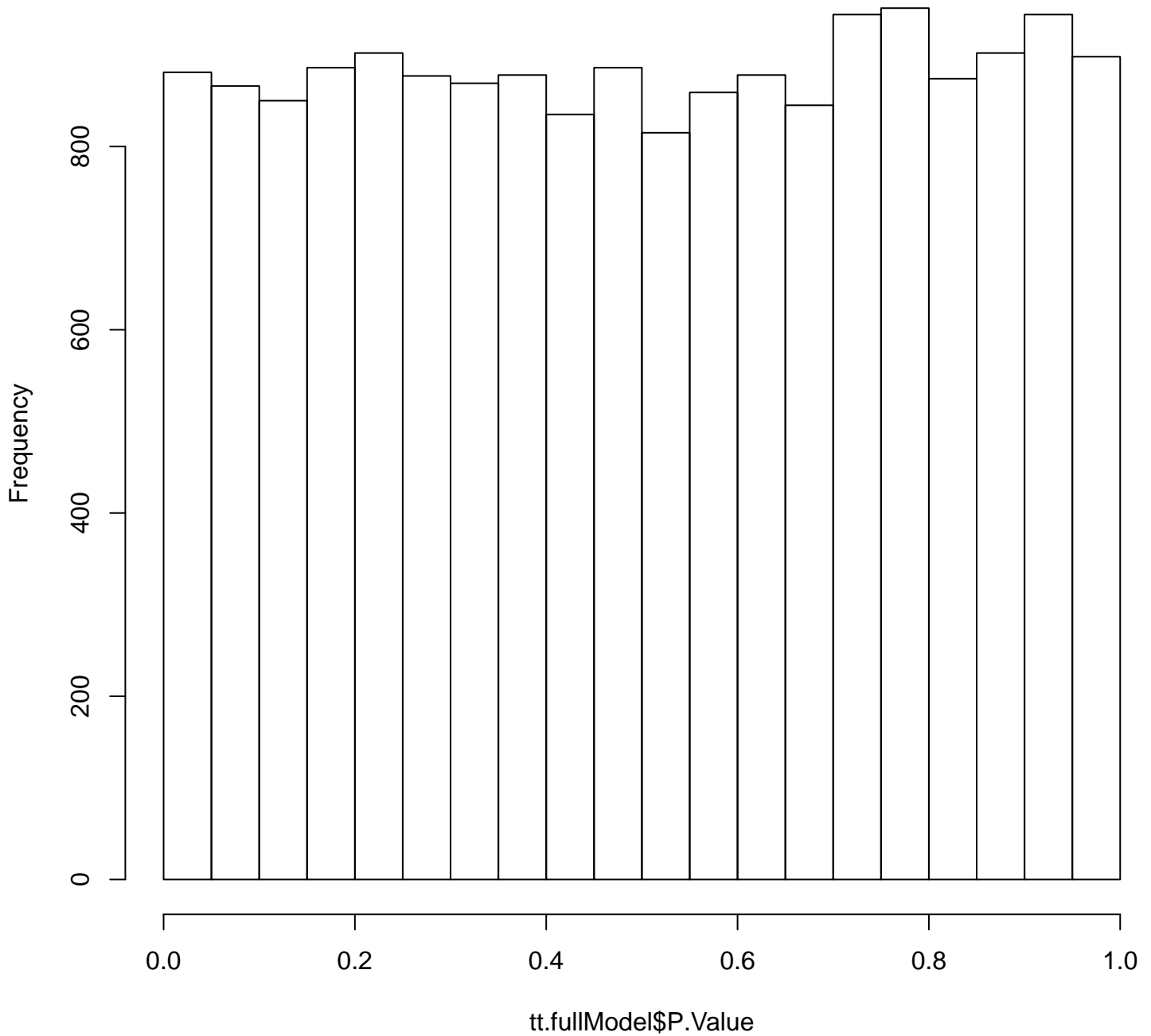
voom: Mean–variance trend



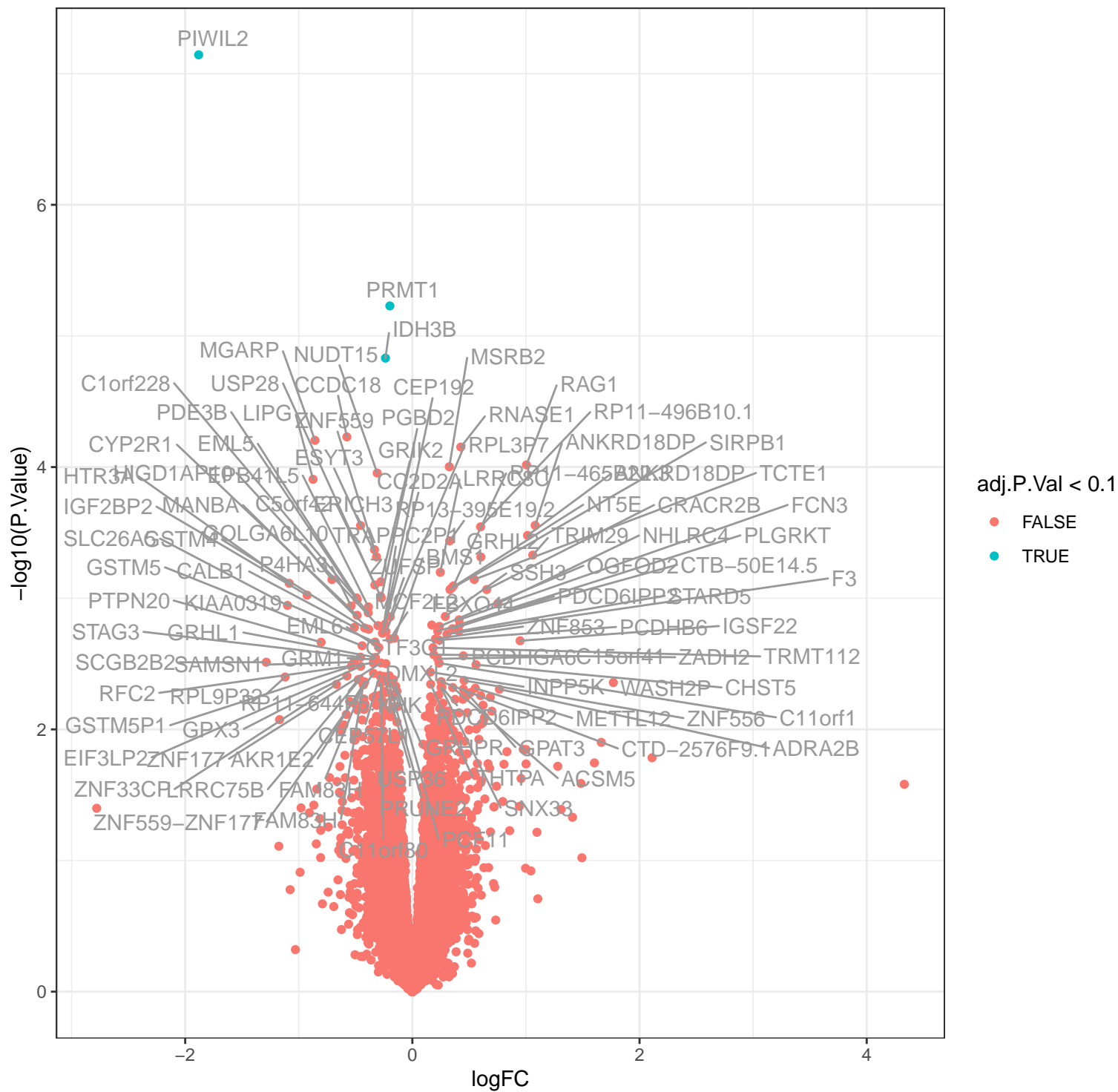
voom: Mean–variance trend



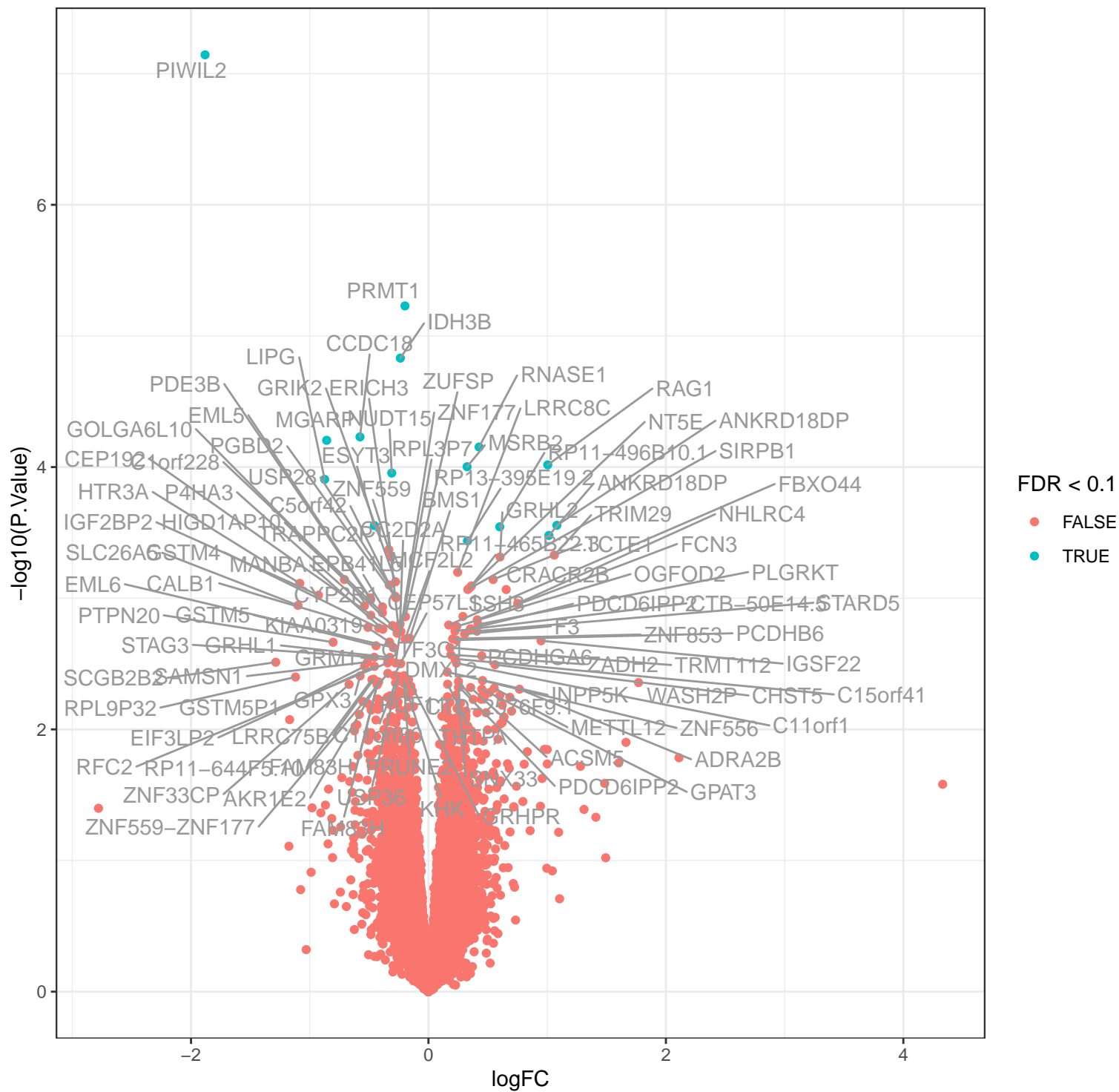
Histogram of tt.fullModel\$P.Value



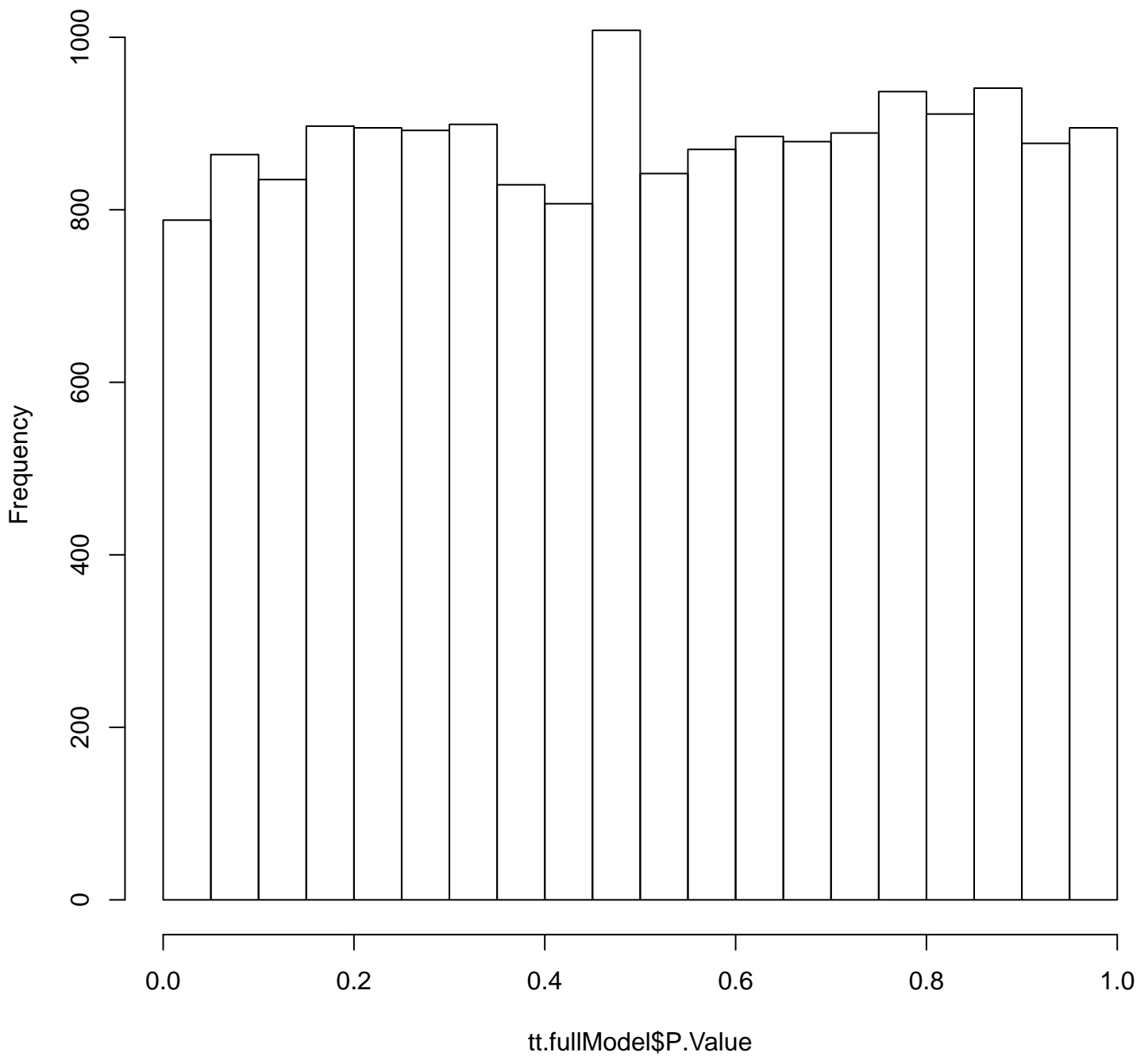
hg CON_vs_MIA_ALL DGE Volcano Plot



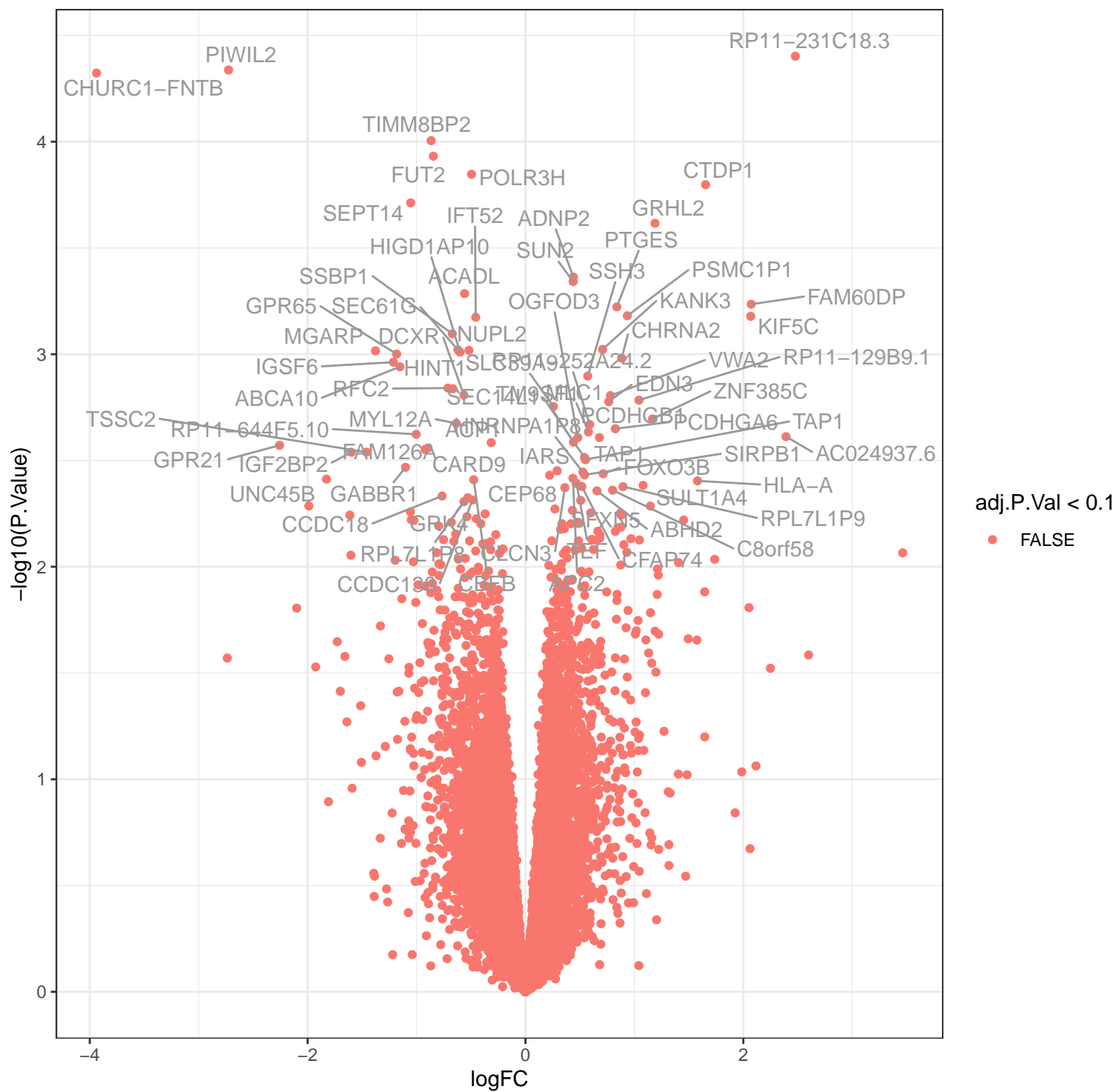
hg CON_vs_MIA_ALL DGE Volcano Plot (after fdrtool)



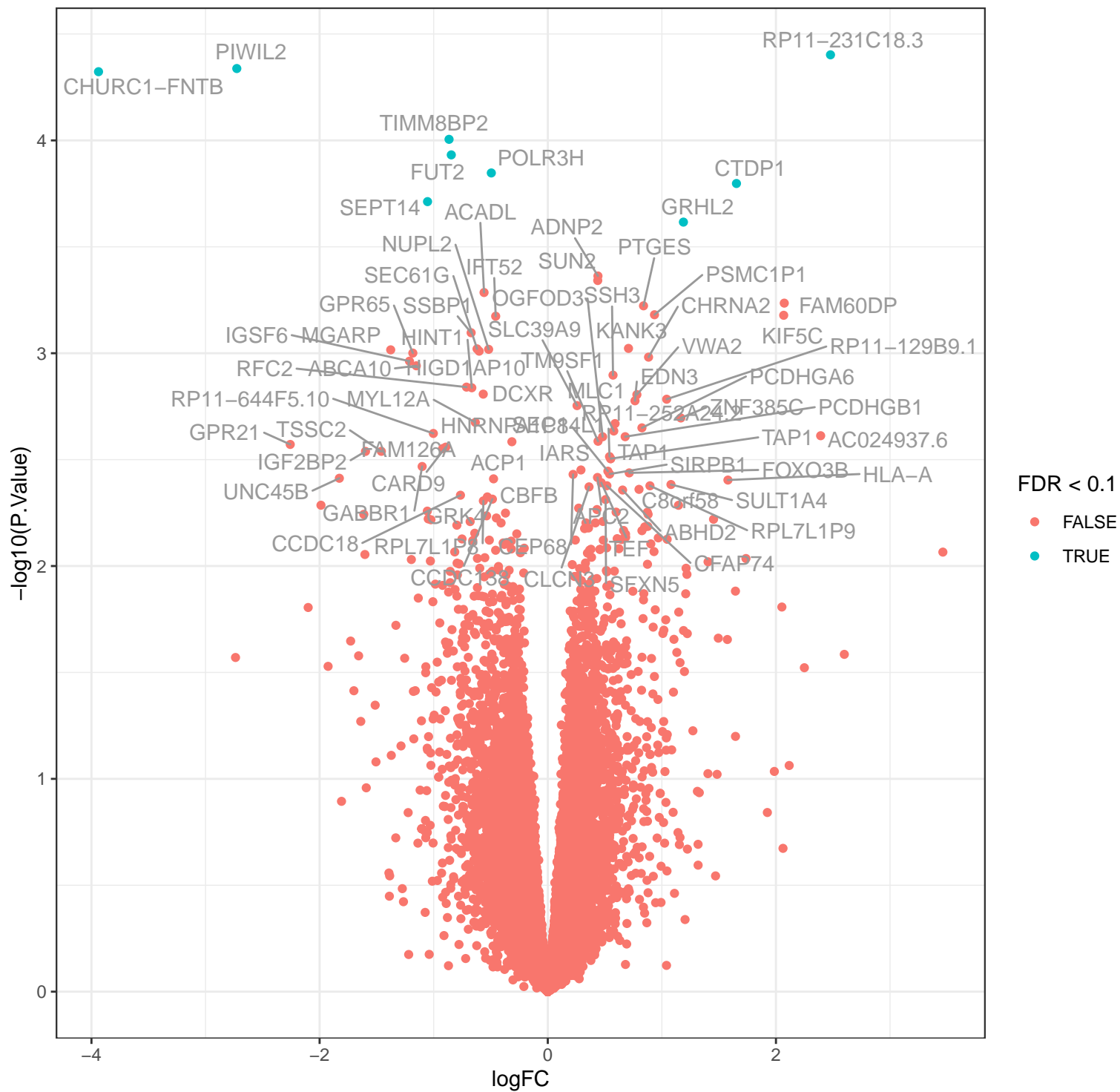
Histogram of tt.fullModel\$P.Value



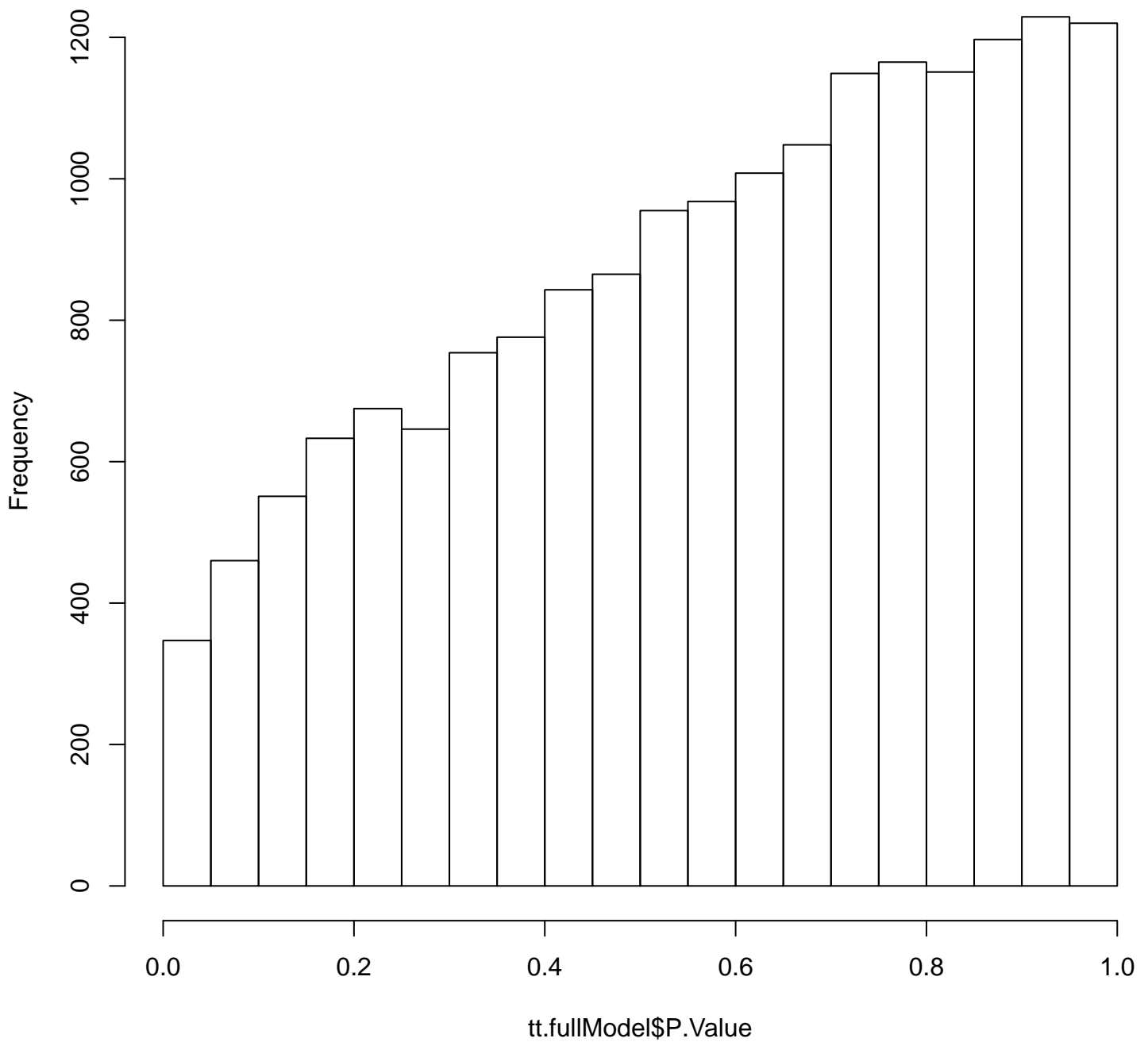
hg CON_vs_MIA_DLPFC DGE Volcano Plot



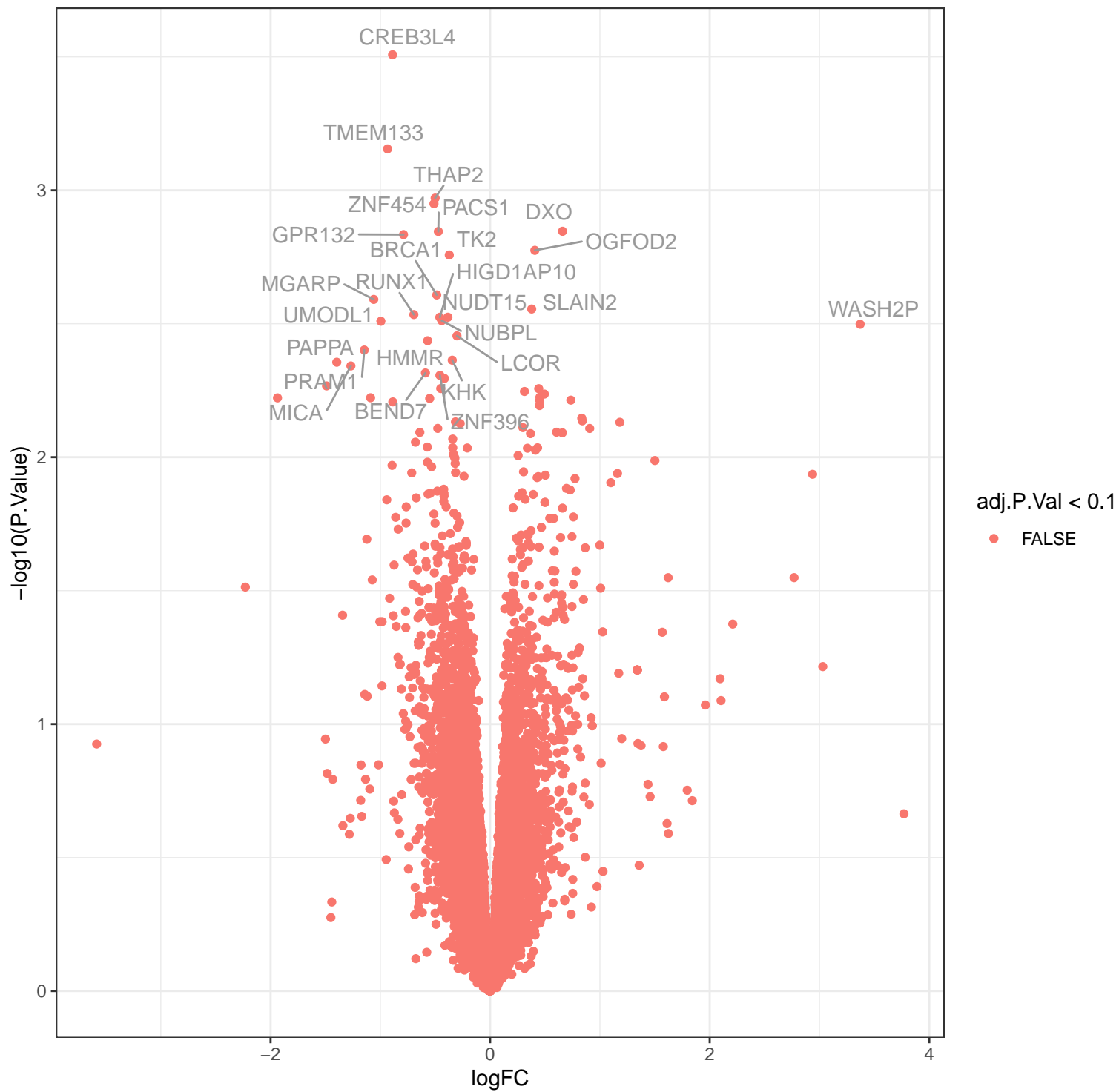
hg CON_vs_MIA_DLPFC DGE Volcano Plot (after fdrtool)



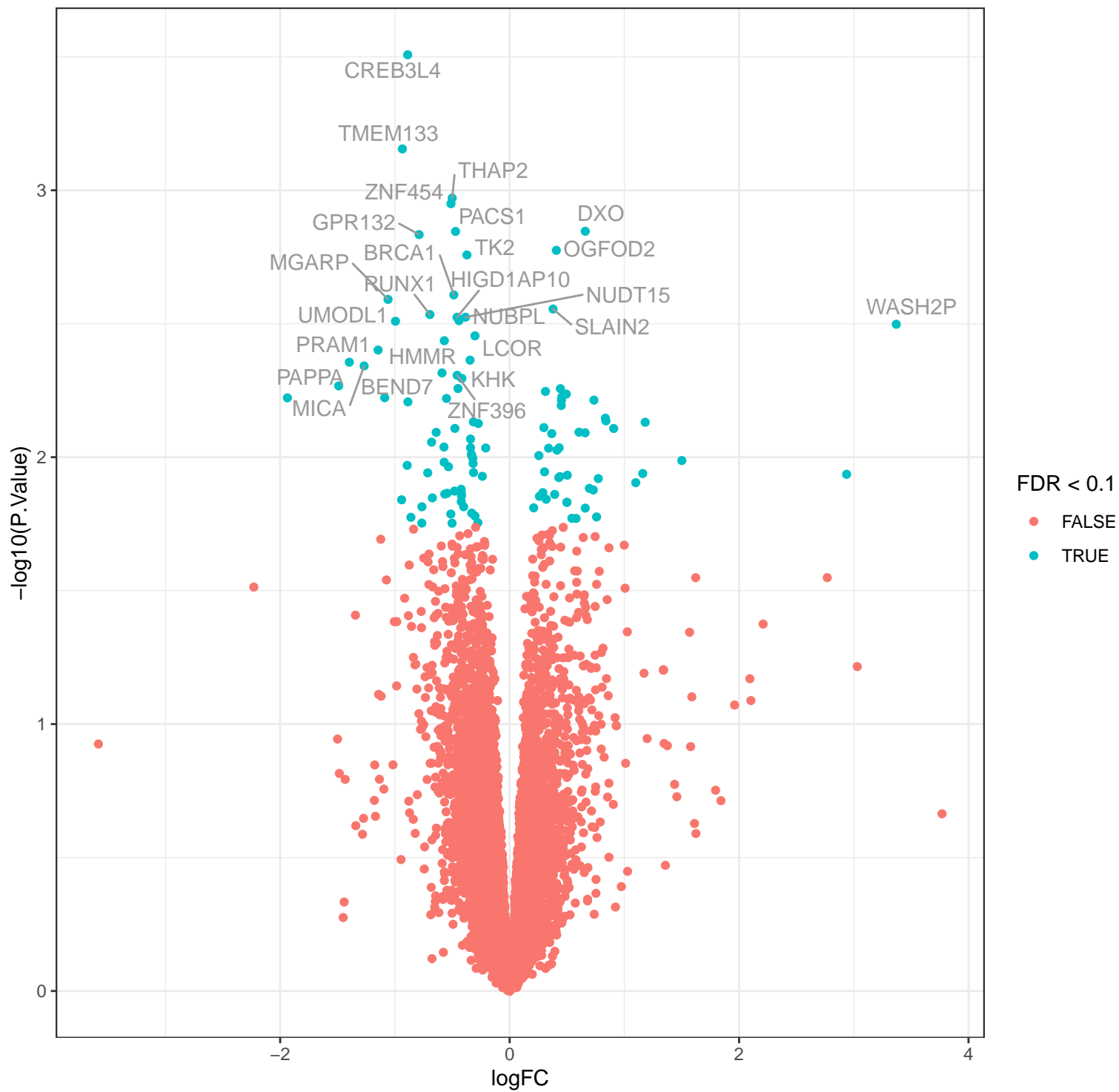
Histogram of tt.fullModel\$P.Value



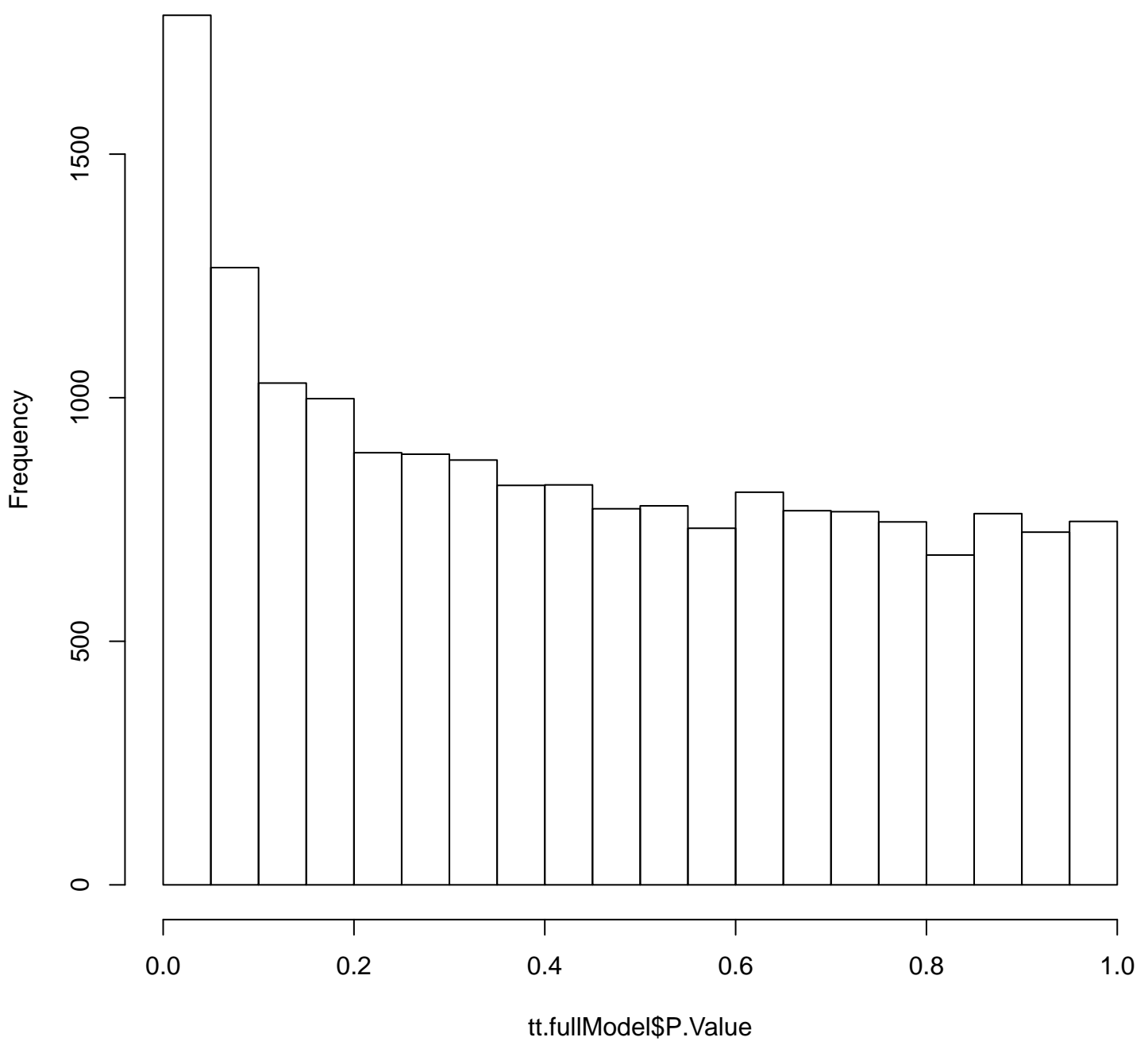
hg CON_vs_MIA_ACC DGE Volcano Plot



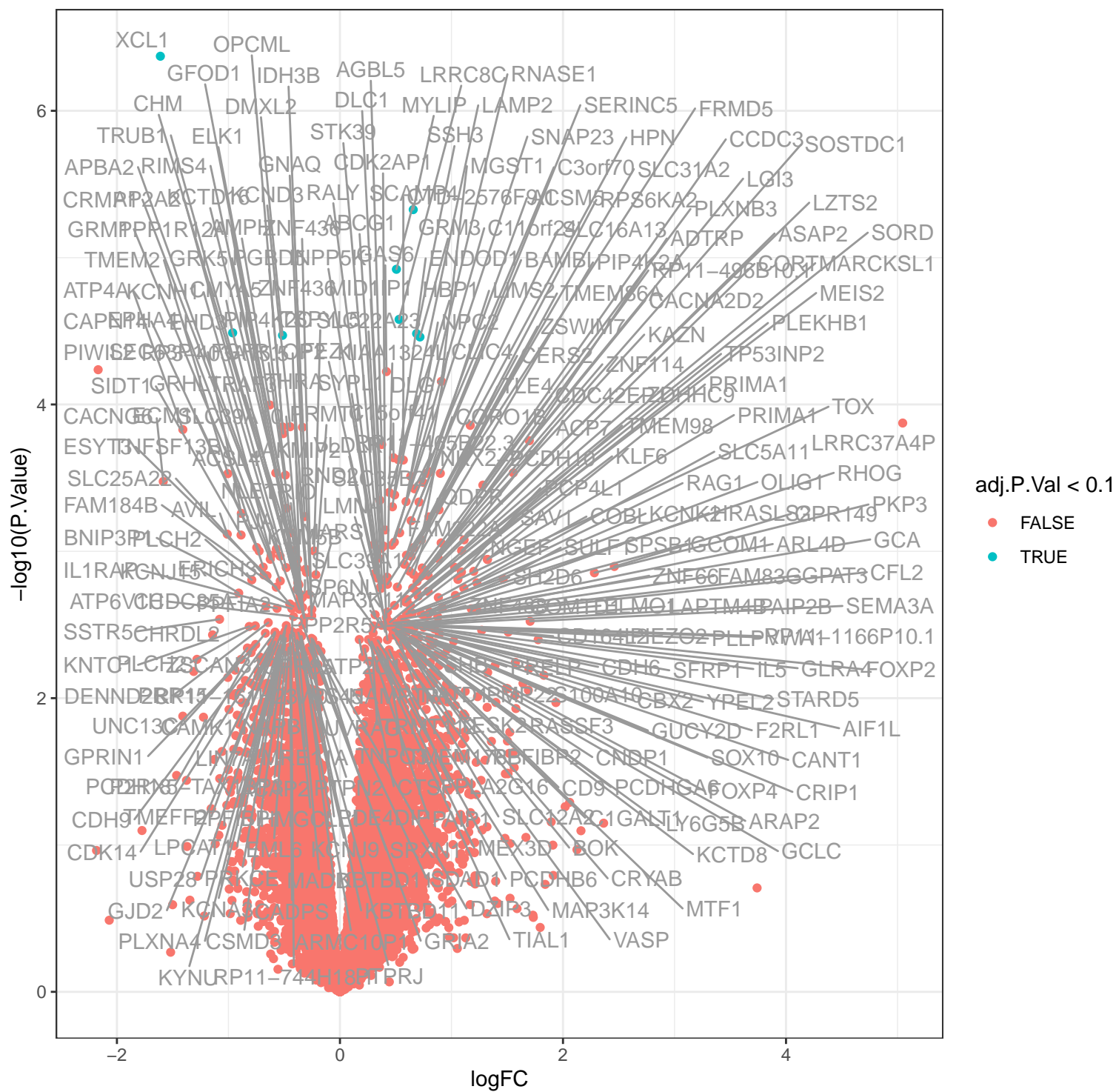
hg CON_vs_MIA_ACC DGE Volcano Plot (after fdrtool)



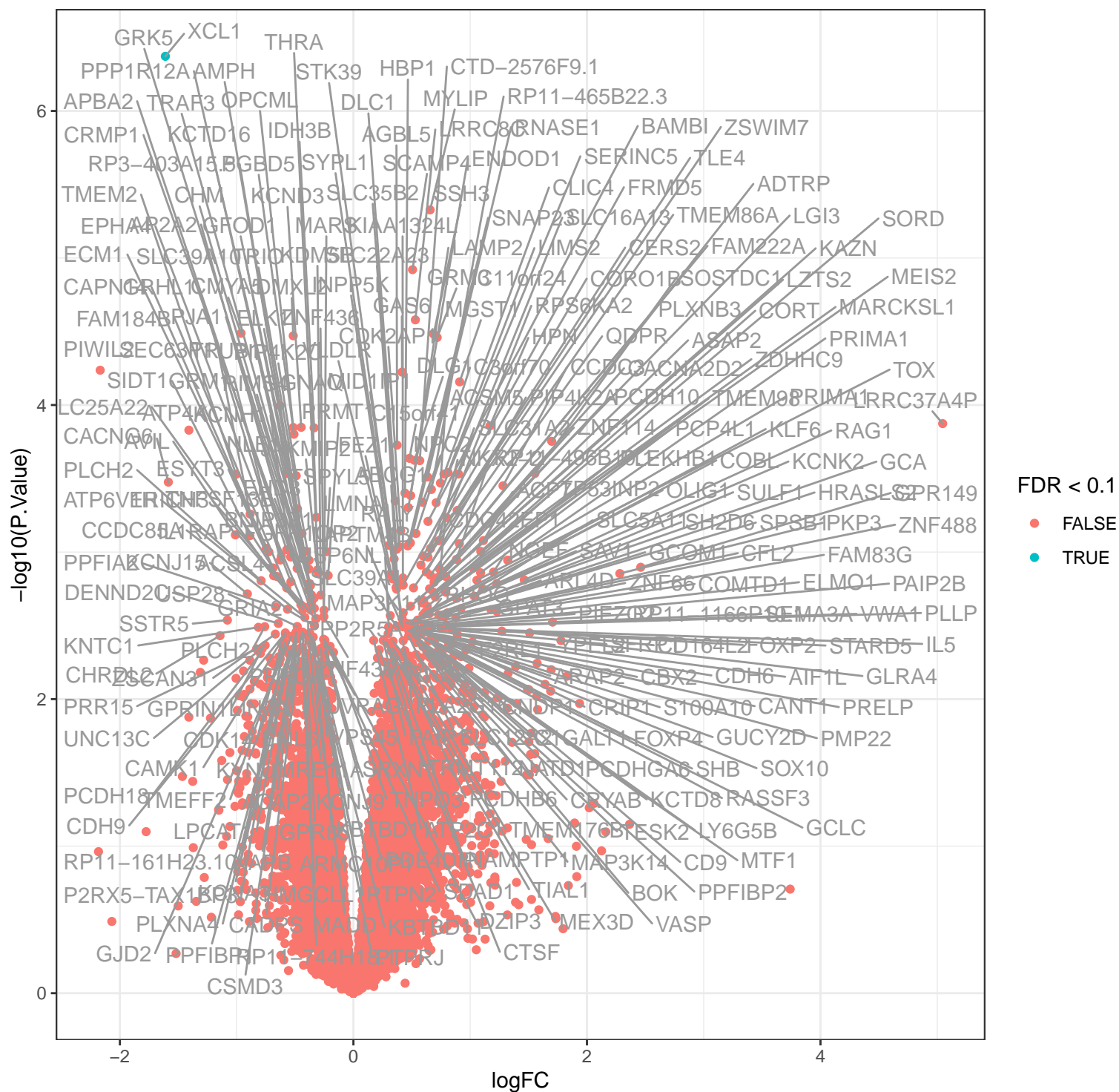
Histogram of tt.fullModel\$P.Value



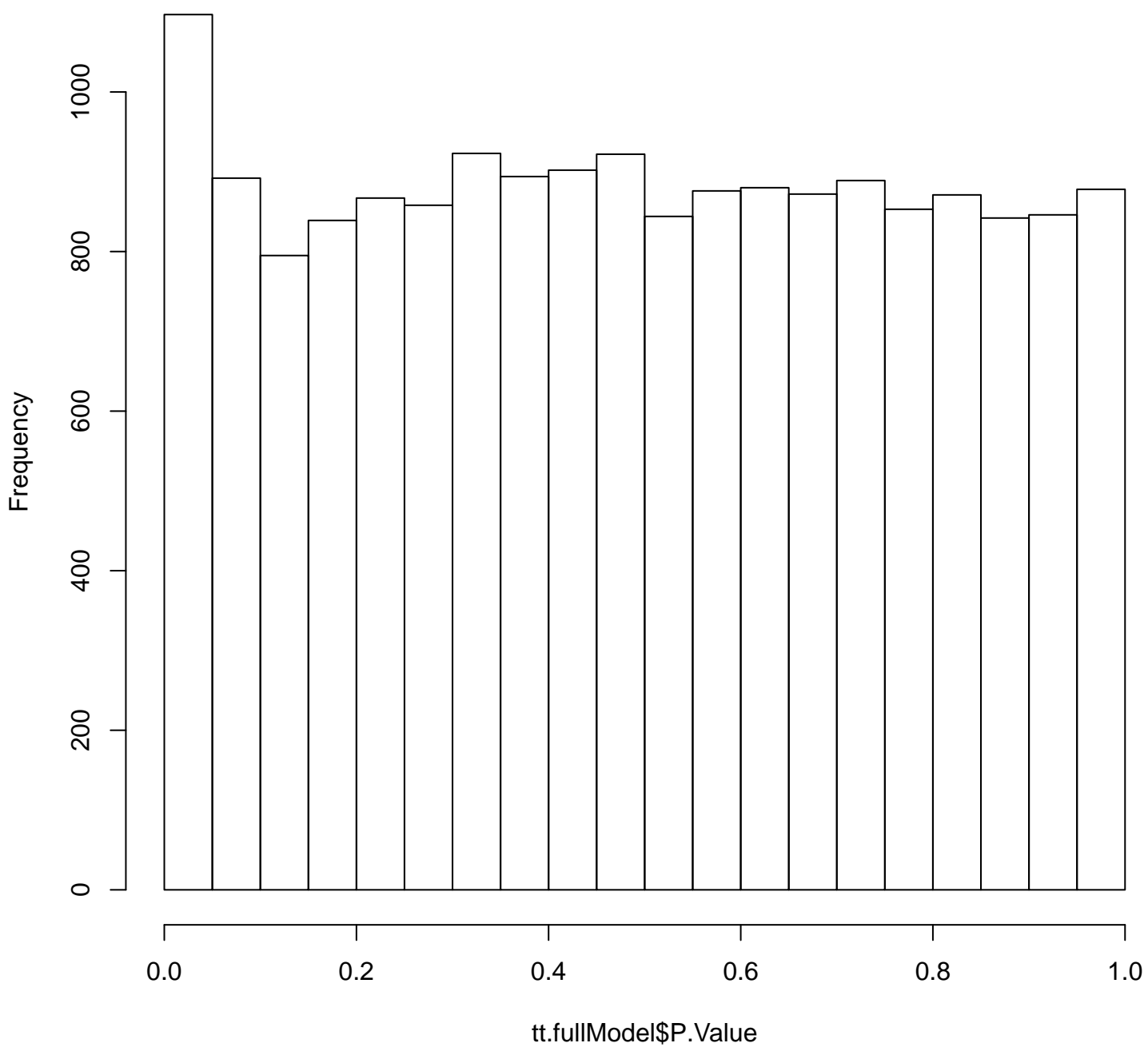
hg CON_vs_MIA_HC DGE Volcano Plot



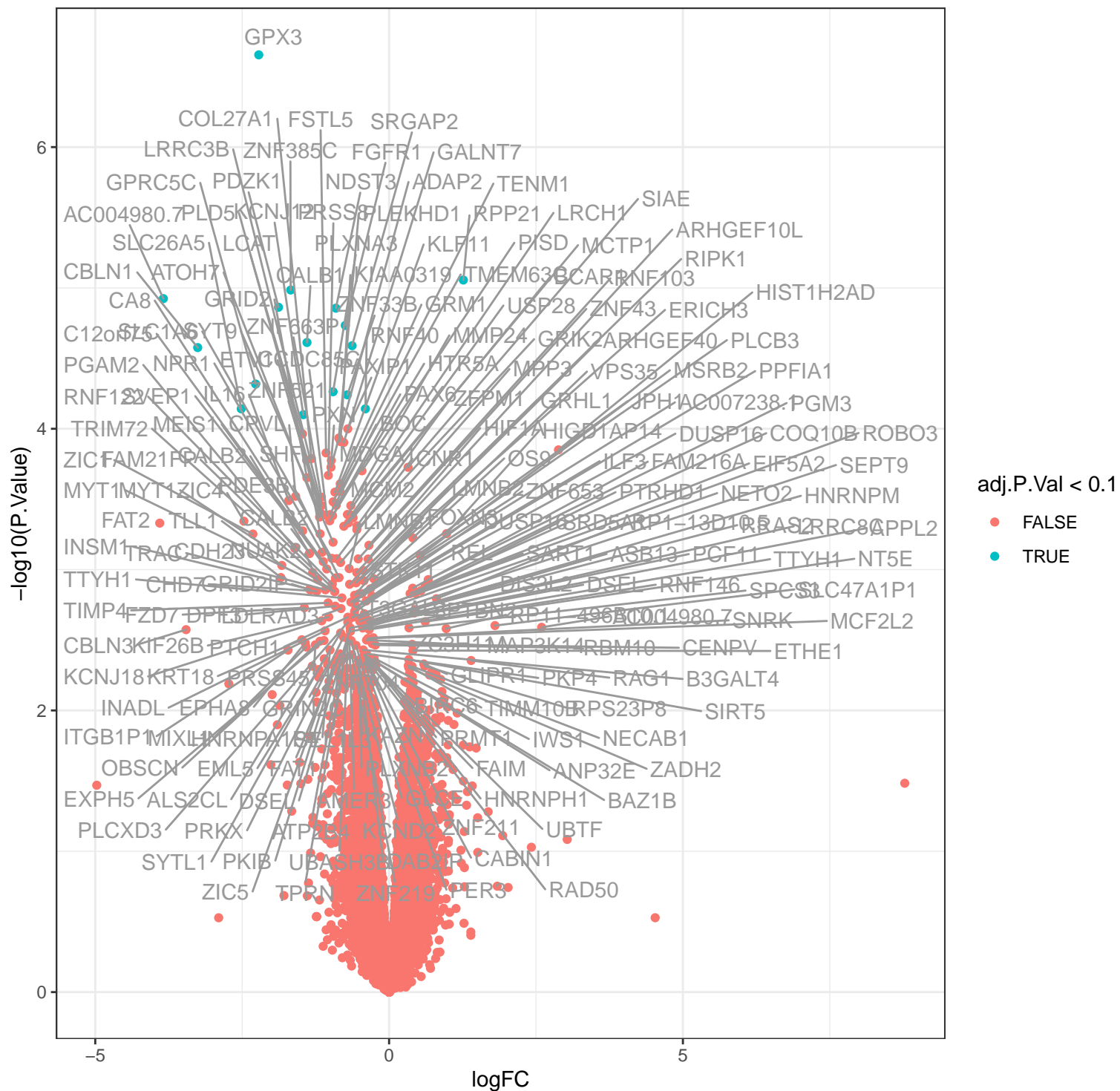
hg CON_vs_MIA_HC DGE Volcano Plot (after fdrtool)



Histogram of tt.fullModel\$P.Value



hg CON_vs_MIA_V1 DGE Volcano Plot



[illegible]

- FALSE
- TRUE