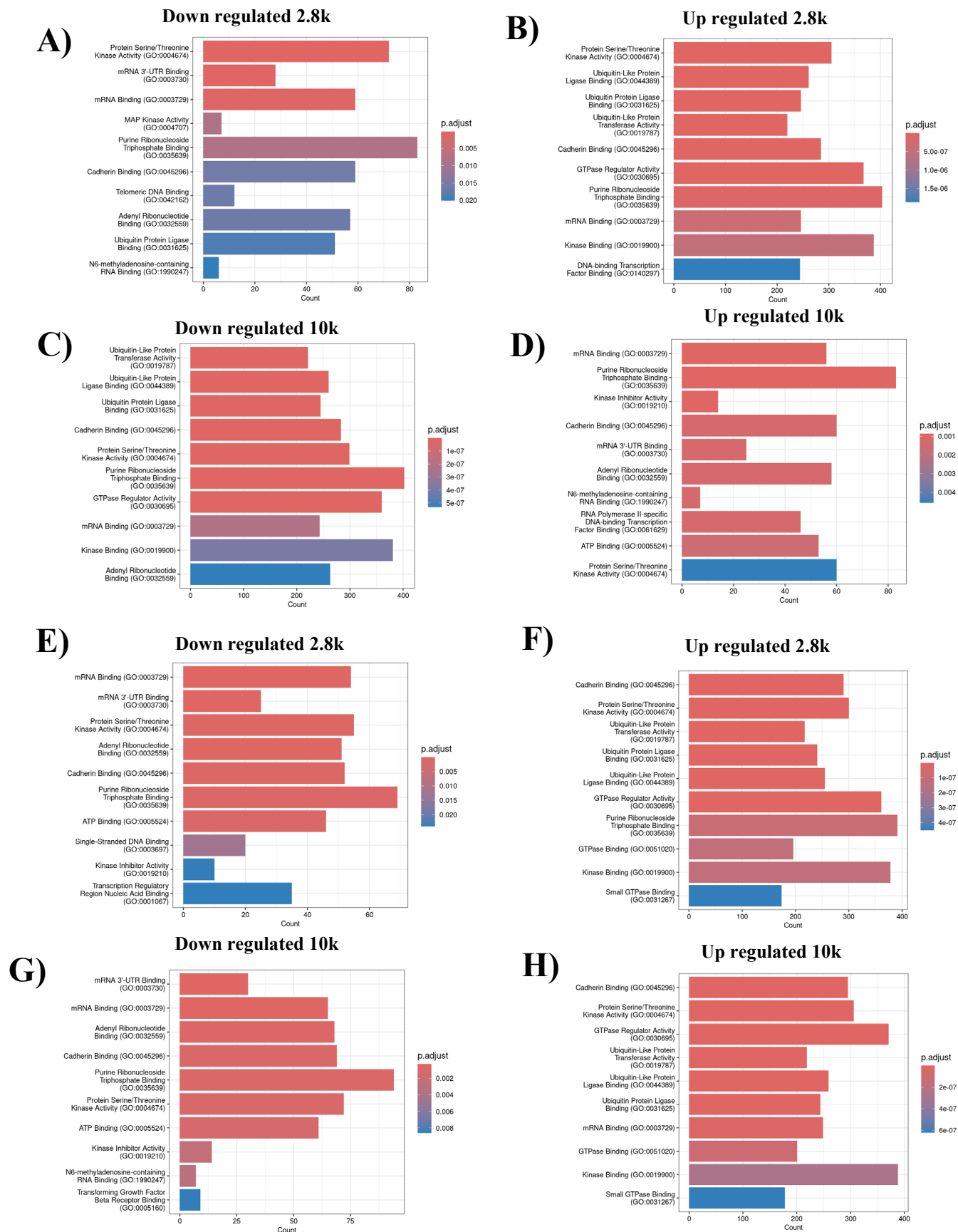


# Figure S1



**Fig S1: (A-D)** Gene set enrichment analysis (GSEA) analysis for the gene targets for the top 10 up-/downregulated miRNAs identified through differential expression analysis between Met5A EV subpopulations and VMC23 EV subpopulations.

**(E-H)** Gene set enrichment analysis (GSEA) analysis for the gene targets for the top 10 up-/downregulated miRNAs identified through differential expression analysis between Met5A EV subpopulations and MSTO EV subpopulations.