**Supplementary figure 2: Functional uptake of three different doses of cobomarsen in ABC-DLBCL cell lines.**

Cells were treated for 96h with 2.5μM, 5μMand 10μMcobomarsenandoligonucleotide control. The graphs show changes in the expression of miR-155 target genes, *BACH1, INPP5D, TP53INP1* and *HIVEP2* assessed by qRT-PCR and each gene was normalized to *GAPDH*. Each bar represents the mean change in expression of four miR-155 direct target genes by three different concentrations of cobomarsen relative to the oligonucleotide treated cells in **A. U2932, B. OCI-LY3 and C. RCK8.** Results are the meanof three biological replicates (±SD). Statistical analysis was performed by 2-way ANOVA, Tukey’s multiple comparisons between the mean of three biological replicates treated with cobomarsen versus the oligonucleotide control for each indicated concentration. :\*\*p<0.01, \*\*\*p<0.001 and \*\*\*\*p<0.0001.