

FIG. S10.Promoter methylation related miRNA expression changes. Cells were treated with the DNA methyltransferase inhibitor 5'-AZA, followed by global gene expression analysis using miRNA microarrays **(A)** Venn diagram of miRNA predicted to be hypomethylated in SN12C and spontaneously hypermethylated in LM2 cells. **(B)** Venn analysis of genes predicted to be hypermethylated in SN12C and spontaneously demethylated in LM2 cells.