









Figure S9: Correlations between miRNAs and the epigenetic components in the different TCGA projects. In each box, different miRNAs and their correlations are depicted. For each miRNA the correlations within the tumors where they resulted upregulated are depicted above the midline, while those within the tumors where they resulted downregulated are depicted below the line. The pathways that resulted negatively correlated with the miRNA are listed on blue down arrows, those that resulted positively correlated are depicted on red up arrows.