Supplementary materials

Figure S1
K-M analysis of the nine candidate DNAm-driven DEGs in the TCGA database.

Figure S2
K-M analysis of the six DNAm-driven DEGs in the GEO validation database.

Figure S3
Regression analysis between CNV and gene expression of the six DNAm-driven DEGs.

Table S1
DEGs identified by DESeq.

Table S2
Methyl Mix models for the DNAm-driven genes.

Table S3
Results of the GO analysis.

Table S4
Pathway analyses based on multiple databases.

Table S5
K-M analysis of the thirteen candidate DNAm-driven DEGs.
Figure S2