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

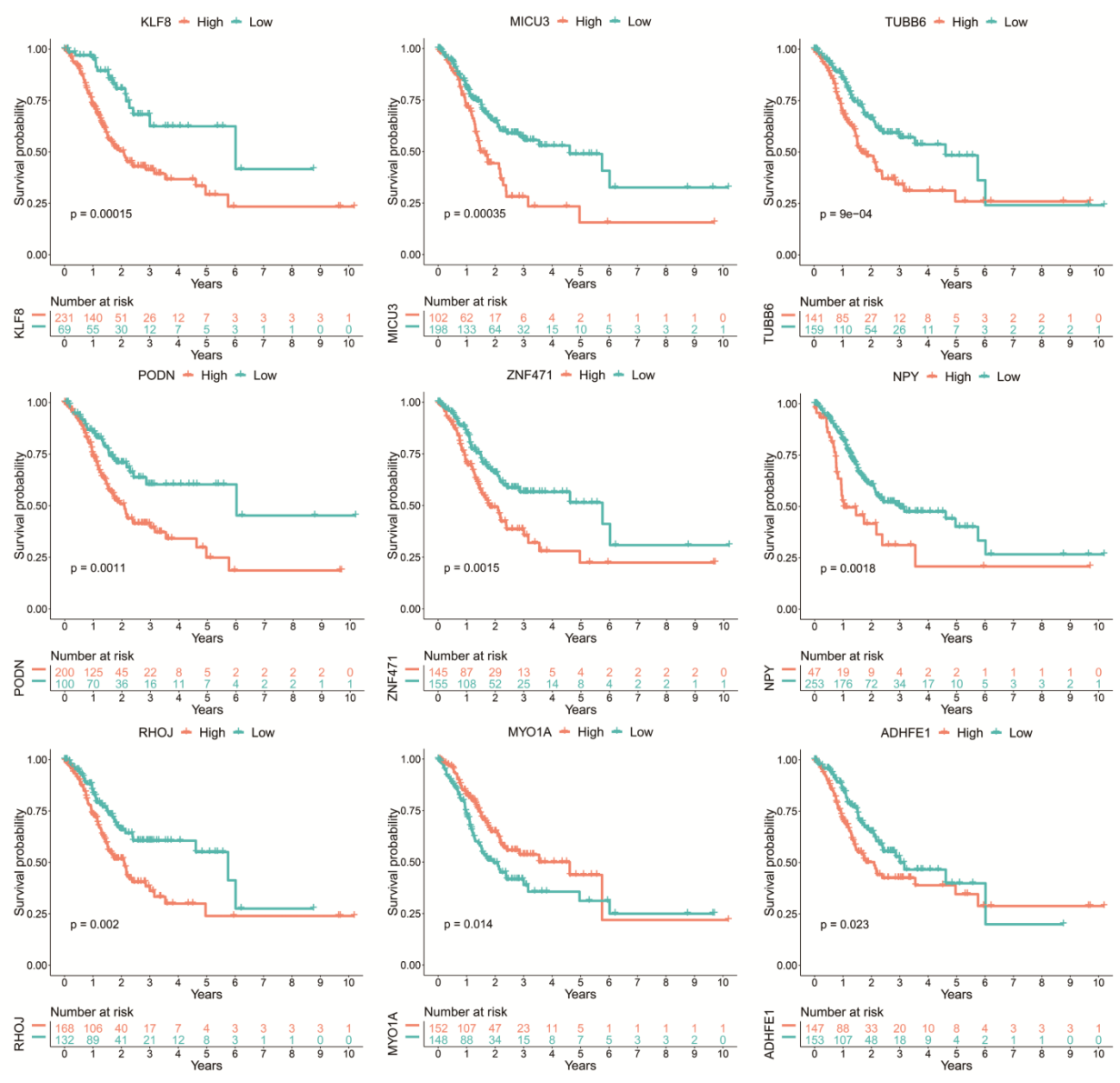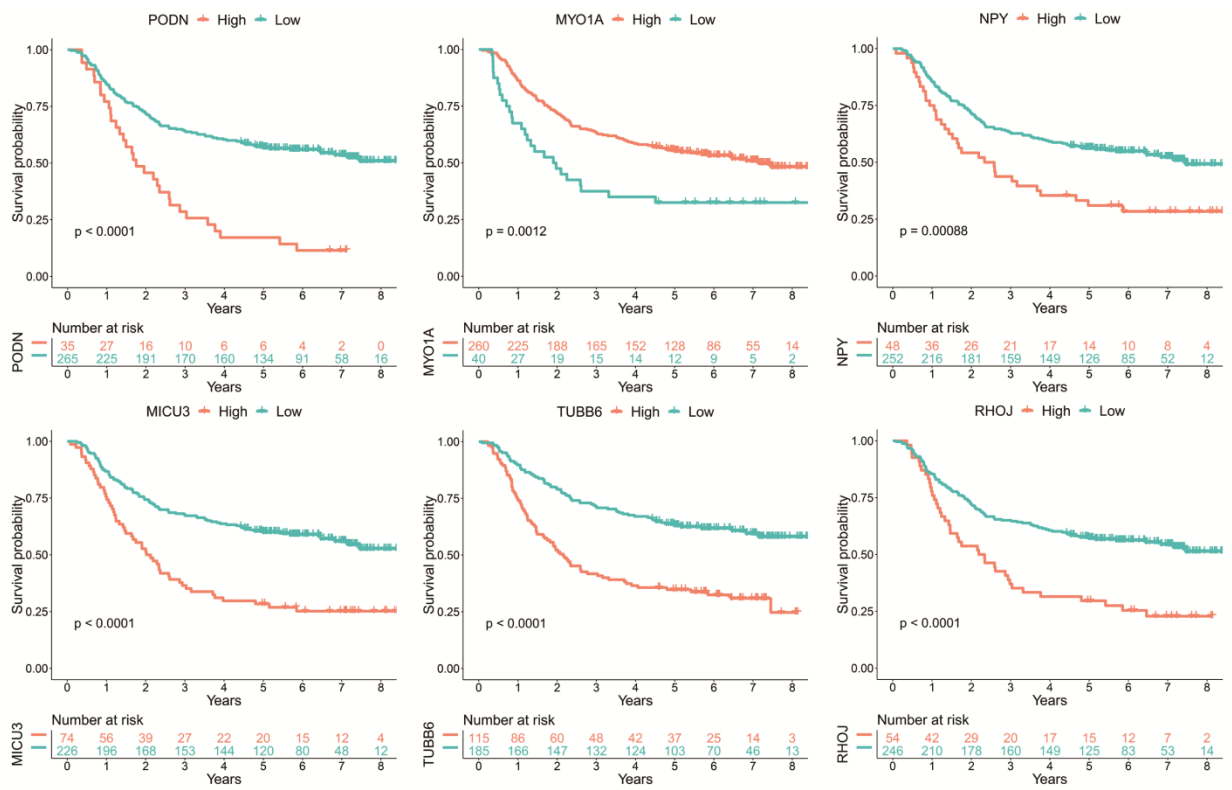K-M analysis of the thirteen candidate DNAm-driven DEGs.

Figure S1

Figure S2

Figure S3