E:\重医附三院\2020在研课题\m6A regulator\submission\Revision\Figures\Supplementary Figure S1.tif**Figure S1.** Overview of this work

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**Figure S2.** **(A)** The most enriched 10 GO terms using the top 500 expressed genes in each m6A modification patterns. **(B)** GSVA analysis of each m6A clusters. **(C-F)** Overall survival curves of representative genes in each m6A modification patterns.

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**Figure S3. (A)** The survival analysis of UCSC ovarian cancer dataset which was divided by the cutoff value according risk score calculated by LASSO multivariate. **(B)** The ROC curve for evaluating the prediction efficiency of the prognostic signature.