# SUPPLEMENTAL FIGURES

## Supplemental Figure 1: PCA of TCGA-UCEC gene expression levels



Using R package “DESeq2” for PCA of TCGA-UCEC gene expression levels. Normal group =35, Cancer group = 544.

## Supplemental Figure 2：The KM curves for the gene high-expression and low-expression groups in the construction of the model



## Supplemental Figure 3: COX analysis was used to verify miRNAs related to UCEC patients

