Figure S2 Correlation analysis of hub lncRNAs and genes. (A) The lncRNA-mRNA correlation network. Based on the correlation coefficient > 0.6, $p < 0.05$ as the cut-off values, the closer the distance is, the higher the correlation coefficient. **(B)** Venn analysis for co-expressed genes of lnc-TRIM28-14 and genes in magenta module. **(C)** Venn analysis showing that 6 candidate genes exactly included in the intersection in **B**. **(D)** PCA plot showing the correlation between GCPM-related biomarkers.