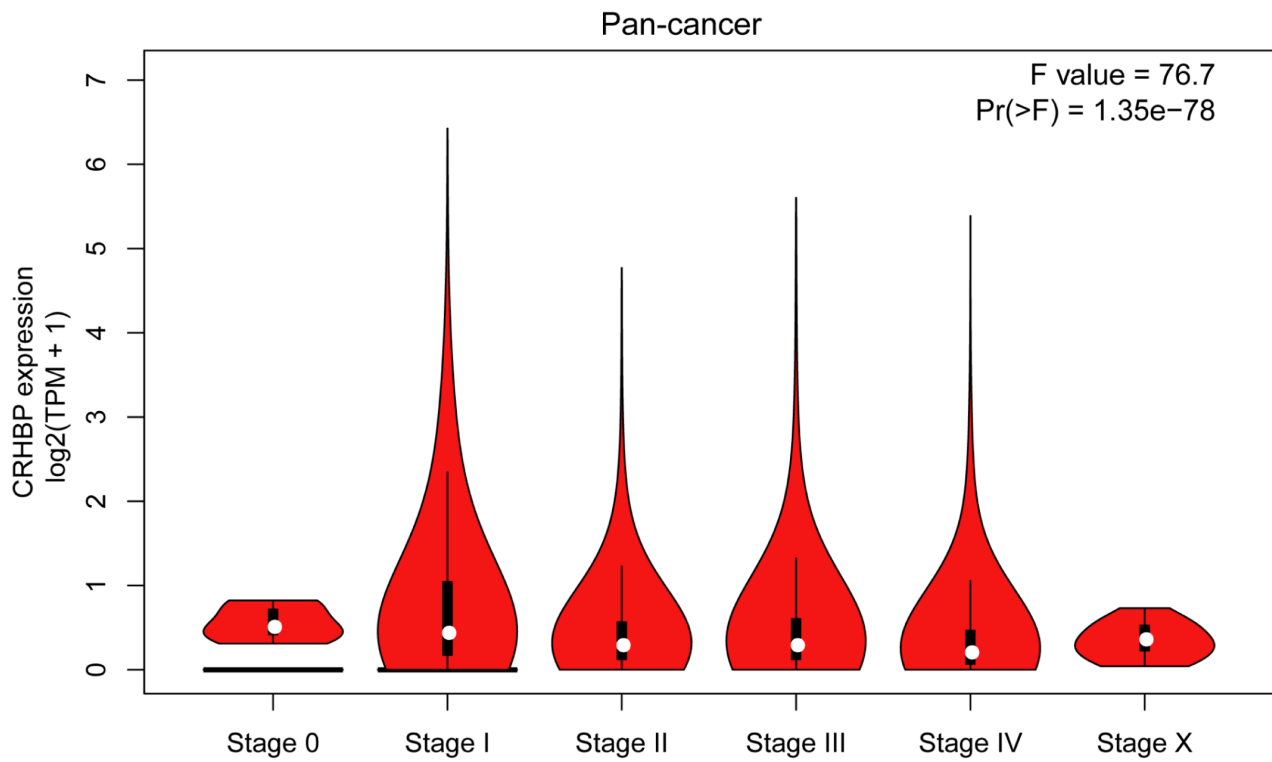
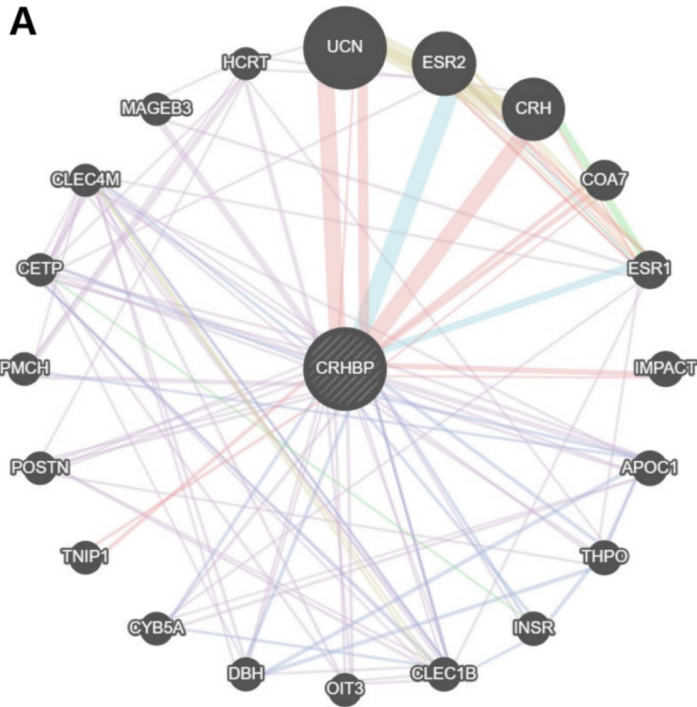


Supplementary Figure 1



Supplementary Figure 1 CRHBP expression level shown significant statistical difference between different pathological stages in the whole pan-cancer.

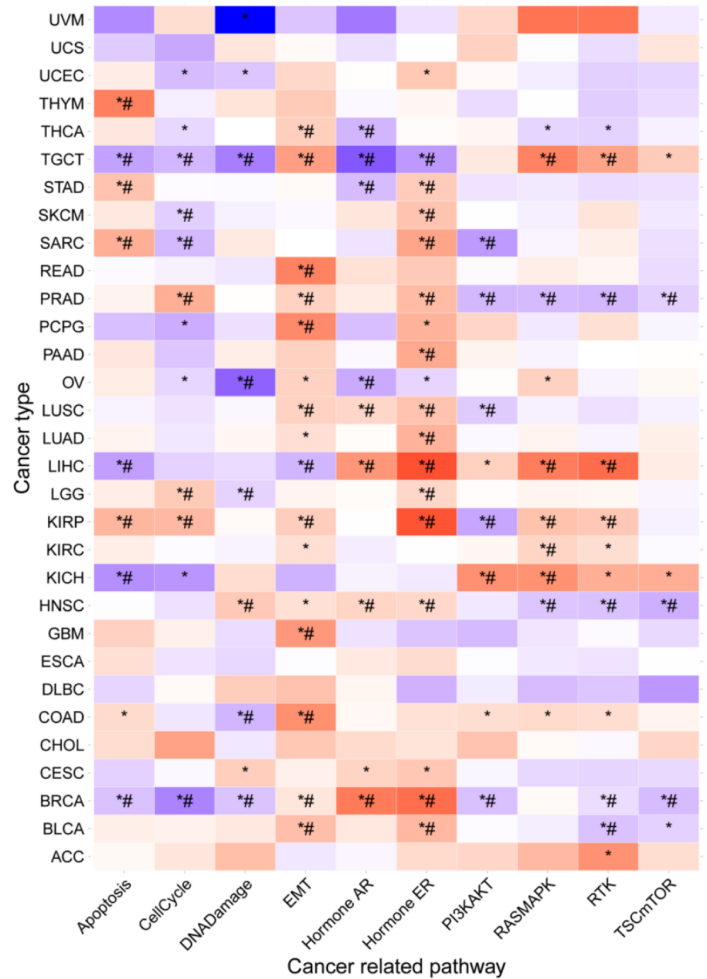
Supplementary Figure 2



Networks

- Physical Interactions
- Co-expression
- Predicted
- Co-localization
- Genetic Interactions
- Pathway
- Shared protein domains

B



*: P value ≤ 0.05; #: FDR ≤ 0.05 Spearman cor. ■ -0.6 -0.3 0.0 0.3 0.6 ■

Supplementary Figure 2 (A) Protein interaction network for CRHBP (B) Heat map reflecting the relationship between GSVA score and activity of different tumor-related pathways.

**Attachment: One-to-one correspondence
between blotting band and whole membrane**

Figure 11 A

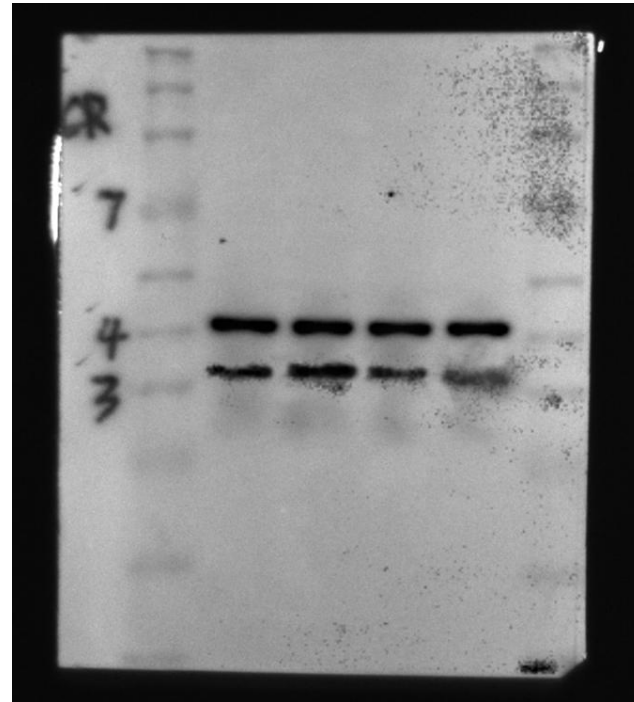
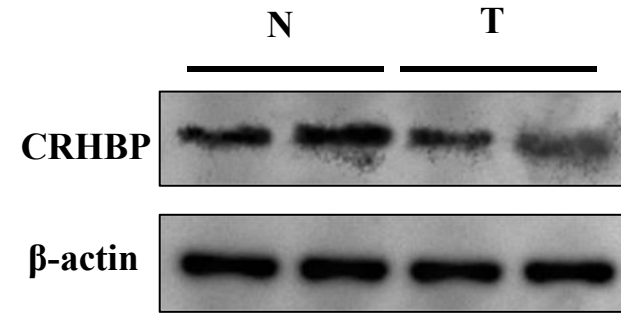
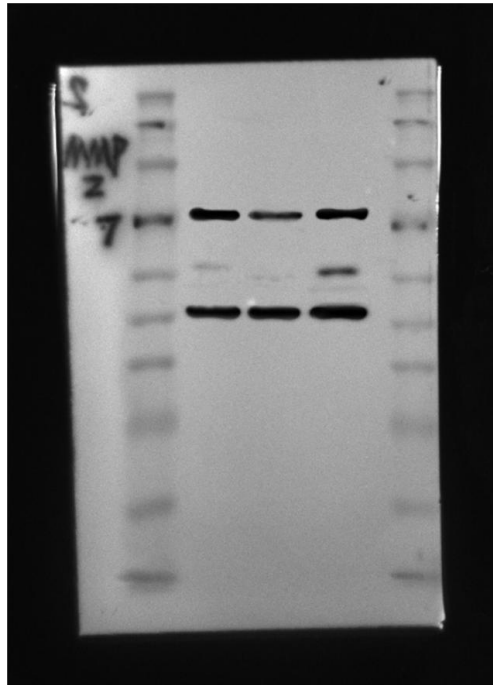
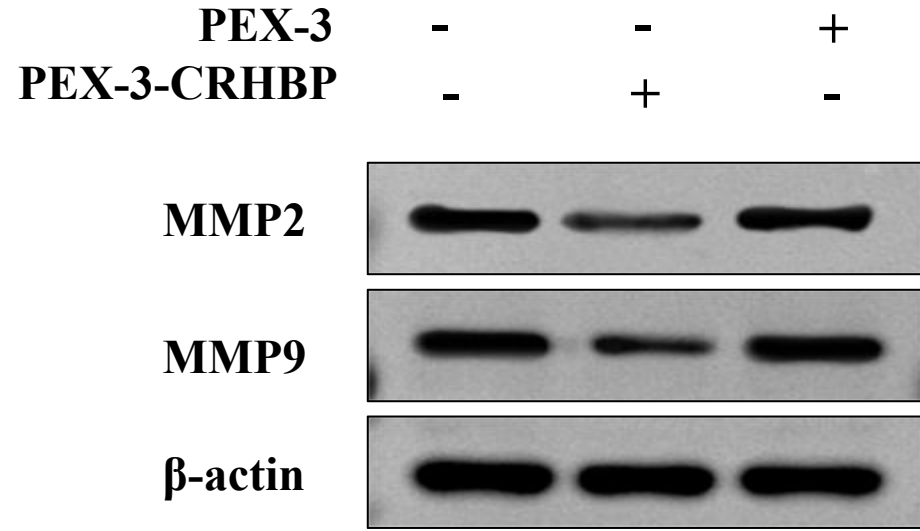
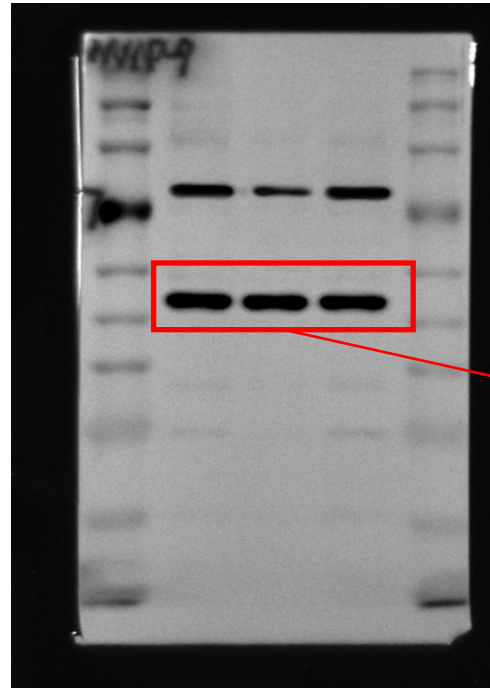


Figure 11 D



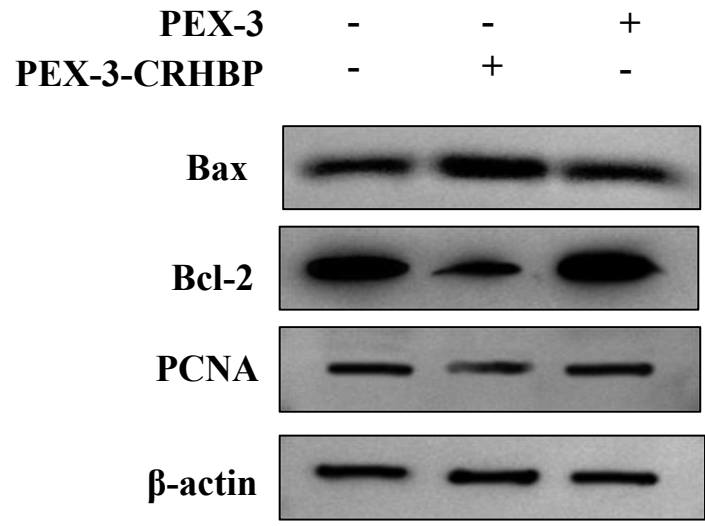
MMP2



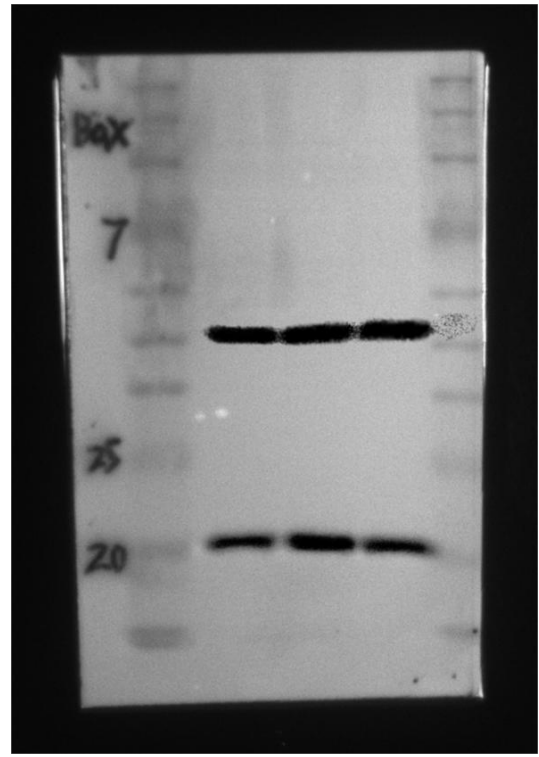
MMP9

When determining the relative expression, each group used its own internal reference protein. The internal reference proteins placed in the figure were randomly selected from the two groups.

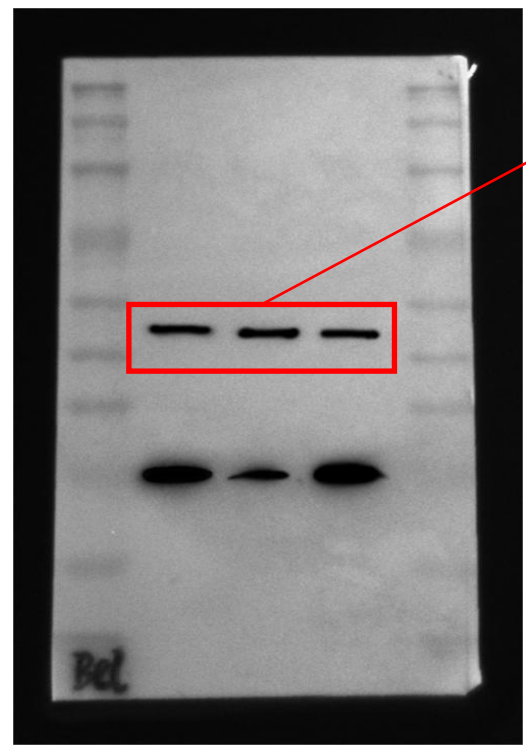
Figure 12 C



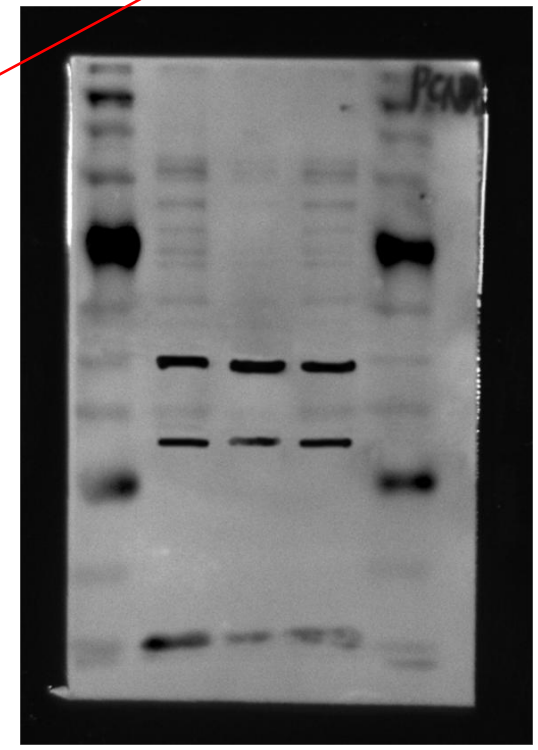
When determining the relative expression, each group used its own internal reference protein. The internal reference proteins placed in the figure were randomly selected from the three groups.



Bax



Bcl-2



PCNA