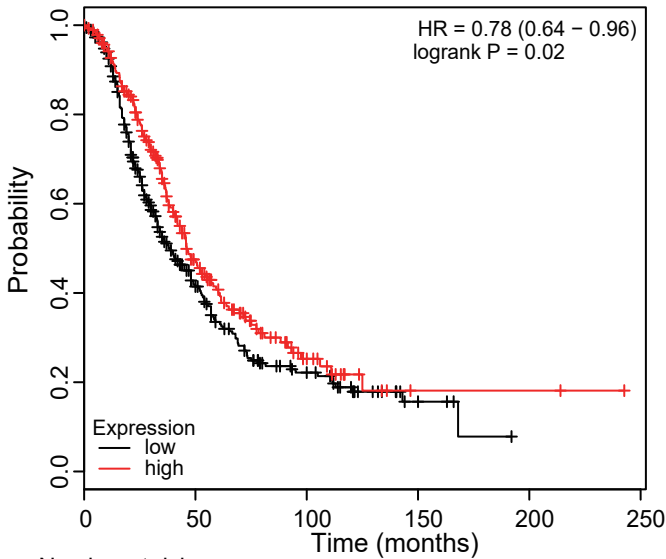
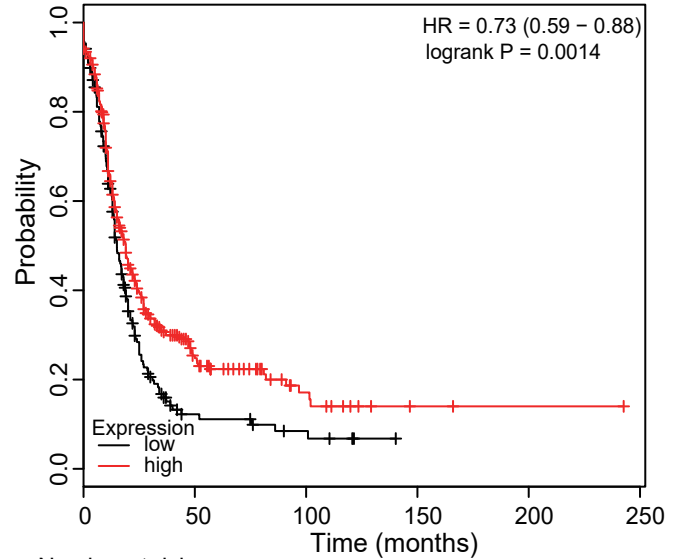
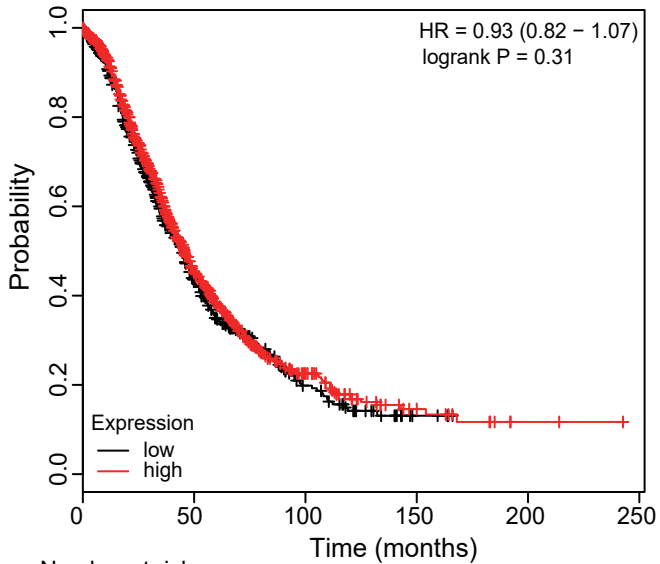


HOSP1

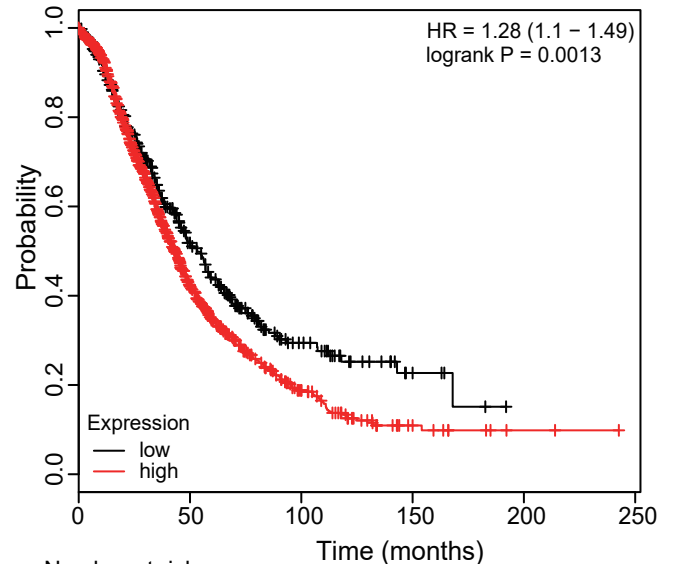
Number at risk		0	50	100	150	200	250
low	369	92	30	6	0	0	0
high	286	74	18	2	2	0	0

RHOA

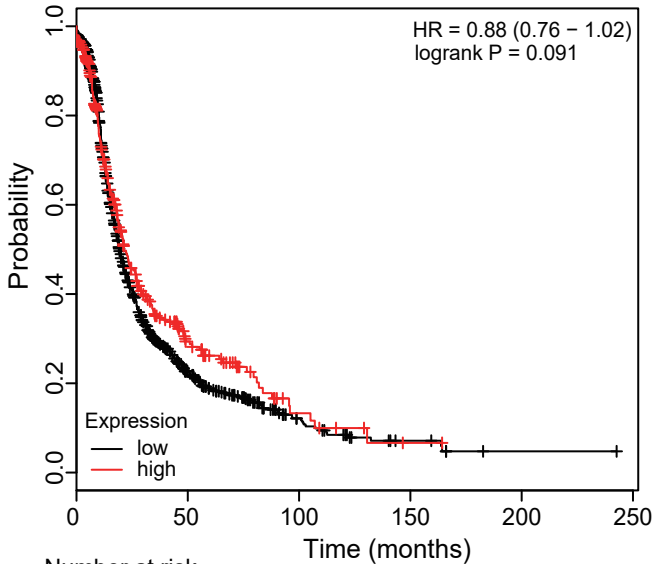
Number at risk		0	50	100	150	200	250
low	187	11	5	0	0	0	0
high	427	43	11	2	1	0	0

PRKAR1A

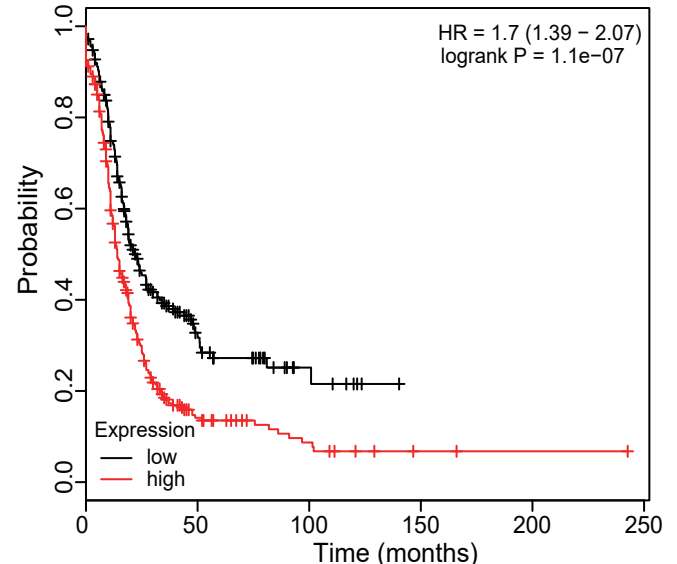
Number at risk		0	50	100	150	200	250
low	527	154	34	4	0	0	0
high	129	274	63	14	2	0	0

ANXN5

Number at risk		0	50	100	150	200	250
low	413	133	34	7	0	0	0
high	1243	295	63	11	2	0	0

CDKN3

Number at risk		0	50	100	150	200	250
low	1069	124	27	7	1	0	0
high	366	48	8	1	0	0	0

LRRN4

Number at risk		0	50	100	150	200	250
low	249	30	7	0	0	0	0
high	365	24	9	2	1	0	0

Supplementary

Figure S1. (A) Heatmap of the 115 DEGs according to the selection criteria: $p < 0.05$ and $|\logFC| > 2$.

(B) The Visualization of GO analysis of the DEGs.

Figure S2. Survival analyses of 6 candidate genes based on the Kaplan Meier Plotter database.