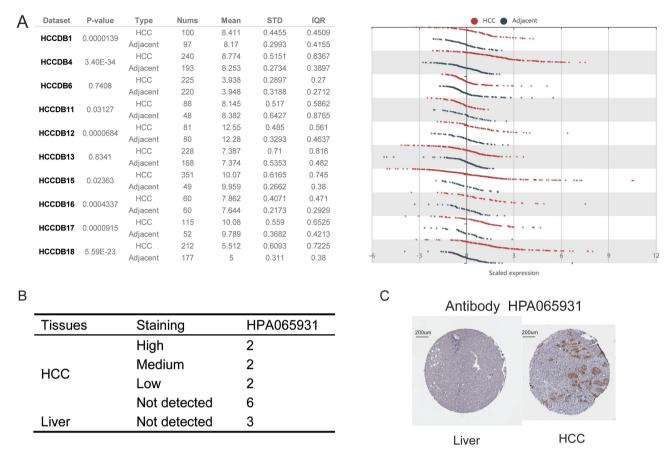
## Supplementary



**Figure S1** The expression level of RANBP1 in HCC. (A) Chart and plot showing the expression of RANBP1 in tumor tissues and the adjacent normal tissues, according to *t*-test using the Integrative Molecular Database for Hepatocellular Carcinoma (HCCDB). (B) The protein expression of RANBP1 in HCC from the HPA. (C) Distribution of RANBP1 in HCC. The staining method was immunohistochemistry staining (200 µm). Color images are available online (Open access and available on the Human Protein Atlas: https://images.proteinatlas.org/65931/162102\_A\_8\_4\_medium.jpg; https://images.proteinatlas.org/65931/162101\_B\_9\_3.jpg). RANBP1, Ranspecific binding protein 1; HCC, hepatocellular carcinoma; HPA, Human Protein Atlas.

Characteristics -	Univariate analysis		Multivariate analysis	
	HR (95% CI)	Р	HR (95% CI)	Р
T stage (T2 & T3 & T4 vs. T1)	2.384 (1.757–3.235)	<0.001	0.700 (0.092–5.335)	0.730
N stage (N1 vs. N0)	1.385 (0.342–5.611)	0.648		
M stage (M1 <i>vs.</i> M0)	3.442 (1.080–10.967)	0.037	2.272 (0.671–7.692)	0.187
Pathologic stage (stage II & stage III & stage IV <i>vs.</i> stage I)	2.310 (1.683–3.170)	<0.001	1.448 (0.191–10.968)	0.720
Histologic grade (G3&G4 vs. G1&G2)	1.154 (0.852–1.564)	0.354		
Residual tumor (R1&R2 vs. R0)	1.515 (0.841–2.730)	0.167		
Age (>60 <i>vs.</i> ≤60)	0.952 (0.710–1.275)	0.740		
Gender (male vs. female)	0.973 (0.713–1.327)	0.861		
Race (white vs. Asian & black or African American)	1.328 (0.984–1.792)	0.063	0.762 (0.492–1.179)	0.223
Child–Pugh grade (B&C <i>vs.</i> A)	1.389 (0.761–2.534)	0.285		
Fibrosis Ishak score (1/2 & 3/4 & 5/6 <i>vs.</i> 0)	1.369 (0.910–2.061)	0.132		
Adjacent hepatic tissue inflammation (mild & severe vs. none)	1.238 (0.863–1.776)	0.246		
Vascular invasion (yes <i>vs.</i> no)	1.682 (1.196–2.364)	0.003	1.631 (1.000–2.658)	0.050
RANBP1 (high <i>vs.</i> low)	1.578 (1.175–2.118)	0.002	1.805 (1.187–2.744)	0.006

Table S1 Univariate and multivariate Cox proportional hazards analysis of RANBP1 expression and progress-free interval for patients with HCC in The Cancer Genome Atlas cohort

CI, confidence interval; HR, hazard ratio; RANBP1, Ran-specific binding protein 1.

Characteristics -	Univariate analysis		Multivariate analysis	
	HR (95% CI)	Р	HR (95% CI)	Р
T stage (T2&T3&T4 vs. T1)	2.807 (1.733–4.546)	<0.001	1.857 (0.757–4.557)	0.176
N stage (N1 <i>vs.</i> N0)	3.562 (0.858–14.785)	0.080	1.010 (0.098–10.360)	0.993
M stage (M1 <i>vs.</i> M0)	5.102 (1.230–21.161)	0.025	4.563 (0.858–24.277)	0.075
Pathologic stage (stage II & stage III & stage IV <i>vs.</i> stage I)	2.887 (1.705–4.888)	<0.001	1.289 (0.142–11.687)	0.821
Histologic grade (G3 & G4 vs. G1 & G2)	1.119 (0.703–1.780)	0.636		
Residual tumor (R1 & R2 <i>vs.</i> R0)	1.640 (0.711–3.782)	0.246		
Age (>60 <i>vs.</i> ≤60)	0.880 (0.566–1.370)	0.573		
Gender (male vs. female)	0.840 (0.533–1.323)	0.452		
Race (white vs. Asian & black or African American)	1.520 (0.947–2.438)	0.083	2.266 (0.561–9.149)	0.250
Child–Pugh grade (B & C <i>vs.</i> A)	2.512 (1.103–5.724)	0.028	4.708 (1.405–15.778)	0.012
Fibrosis Ishak score (1/2 & 3/4 & 5/6 <i>vs.</i> 0)	0.921 (0.478–1.774)	0.806		
Adjacent hepatic tissue inflammation (mild & severe vs. none)	1.446 (0.791–2.645)	0.231		
Vascular invasion (yes vs. no)	1.284 (0.710–2.319)	0.408		
RANBP1 (high <i>vs.</i> low)	2.123 (1.341–3.360)	0.001	2.852 (1.178–6.903)	0.020

Table S2 Univariate and multivariate Cox proportional hazards analysis of RANBP1 expression and DSS for patients with HCC in TCGA cohort

CI, confidence interval; HR, hazard ratio.