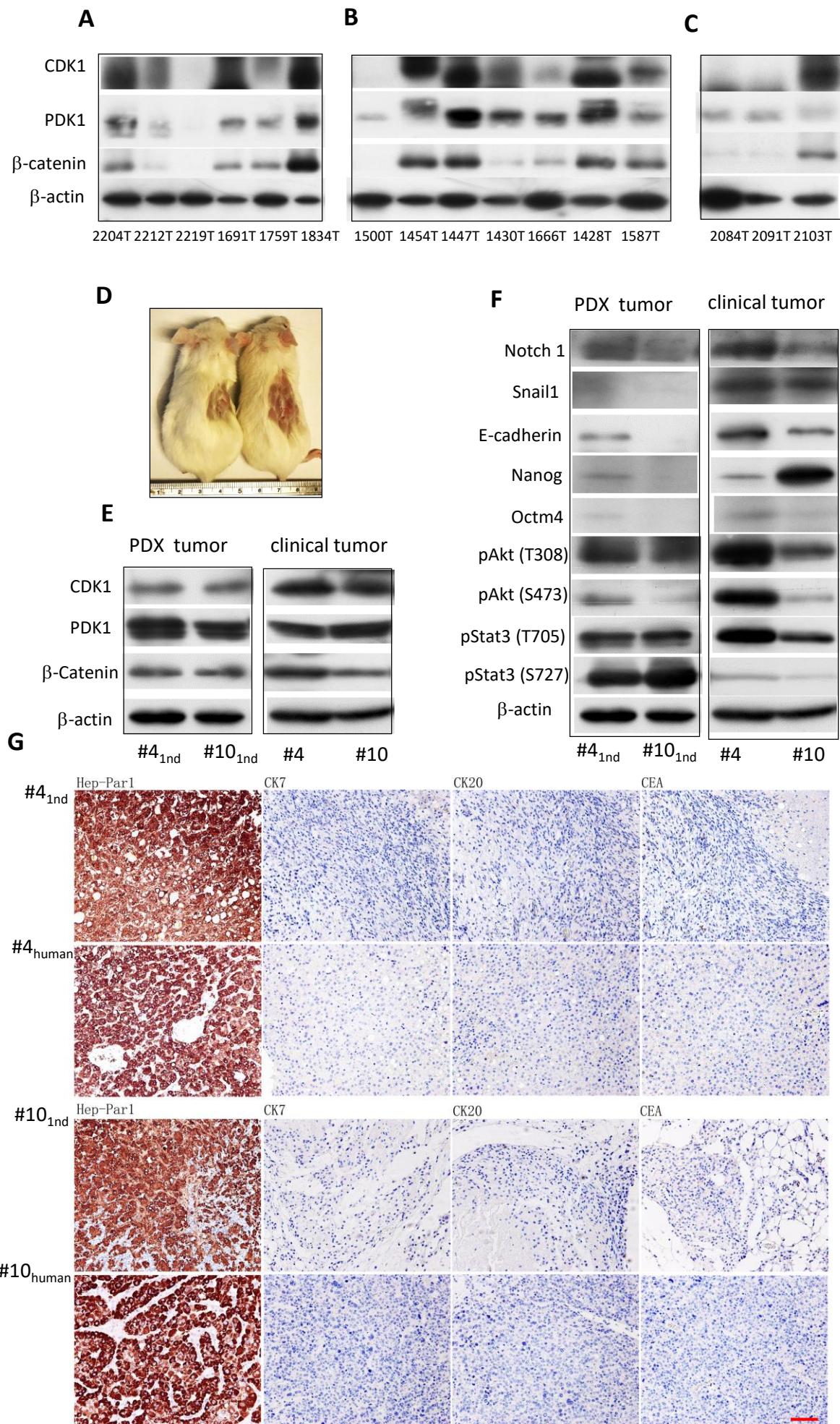
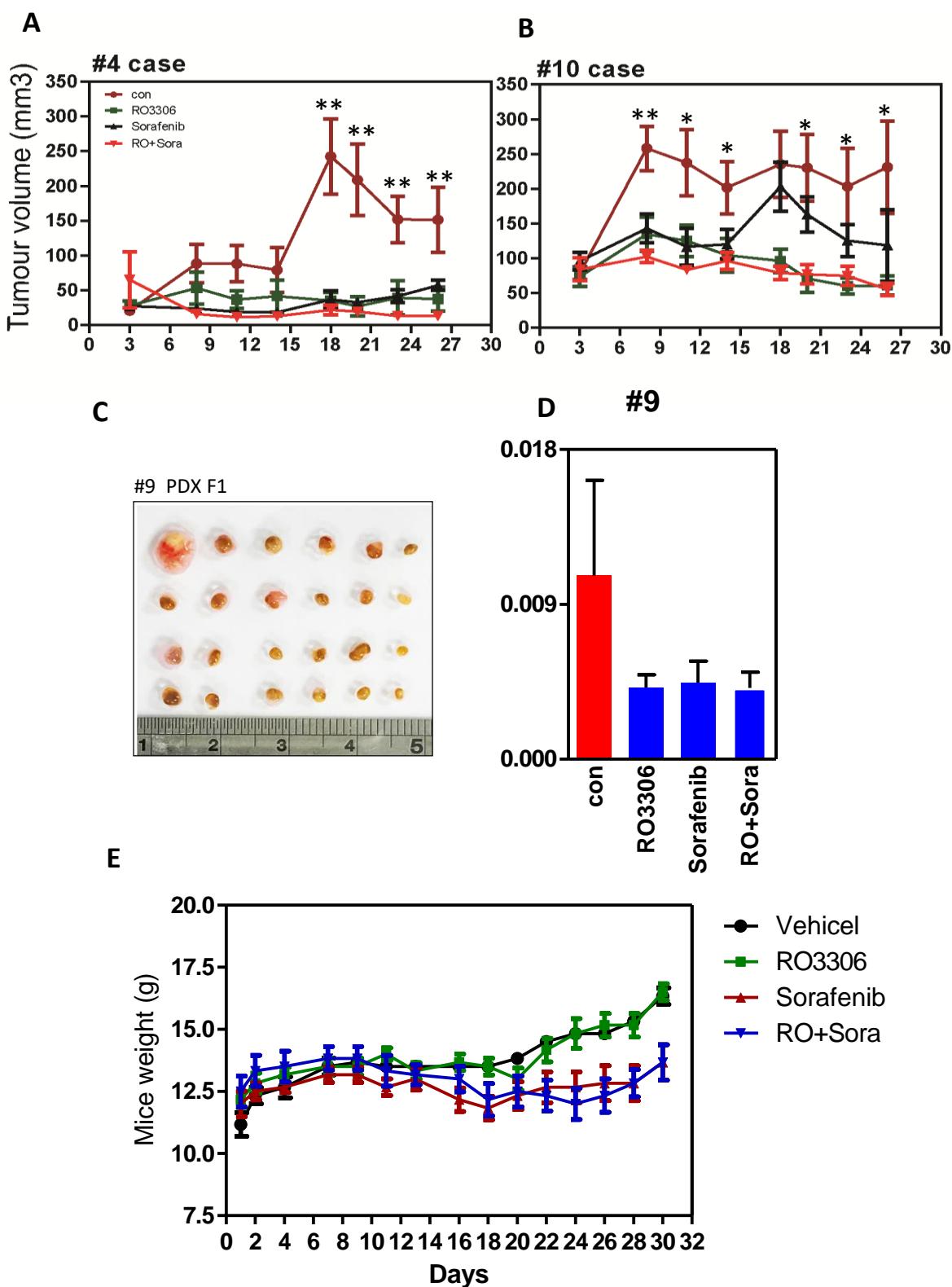


Supplementary figure 1



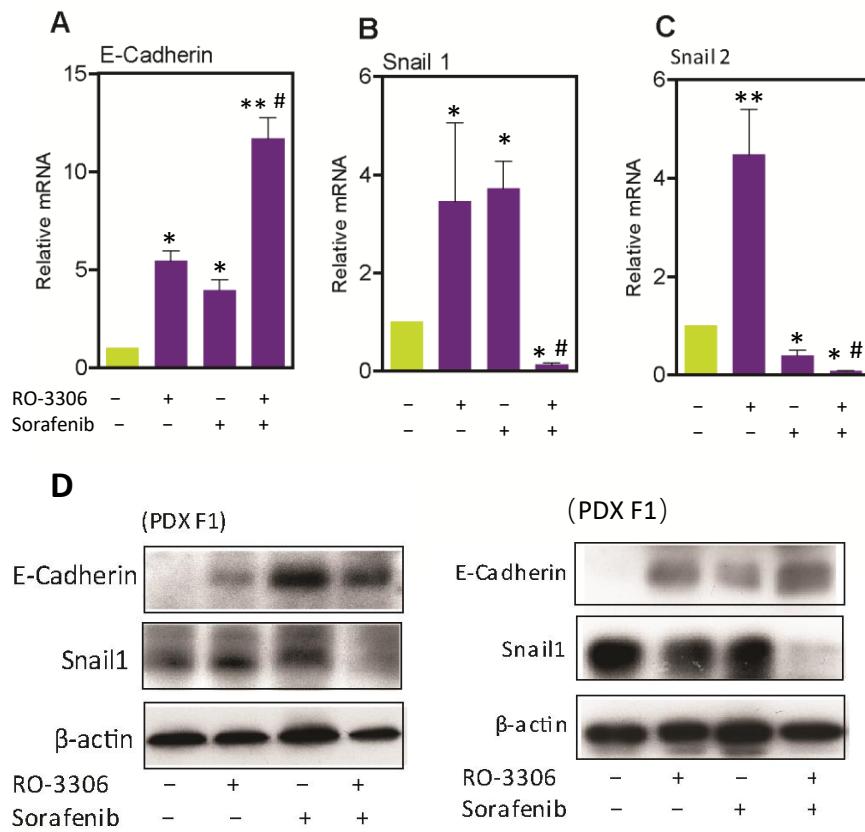
**Supplementary Figure.1 F1 PDX tumor grafts resemble the patient tumors from where they are derived.** A-C, The corresponding expression of CDK1, PDK1 and β-catenin in clinical tumor tissues. D, Image represented the mouse model of F1 PDX tumors. E-F, In comparison of the protein expression levels, PDX tumors highly mimic the paired clinical tumors analysed by western blot. G, Hep Par1, cytokeratin 7 (CK7), cytokeratin 20 (CK20) and CEA staining for F1 PDX tumors and paired clinical tissues. Scale bar represent 100 μm.

Supplementary figure 2



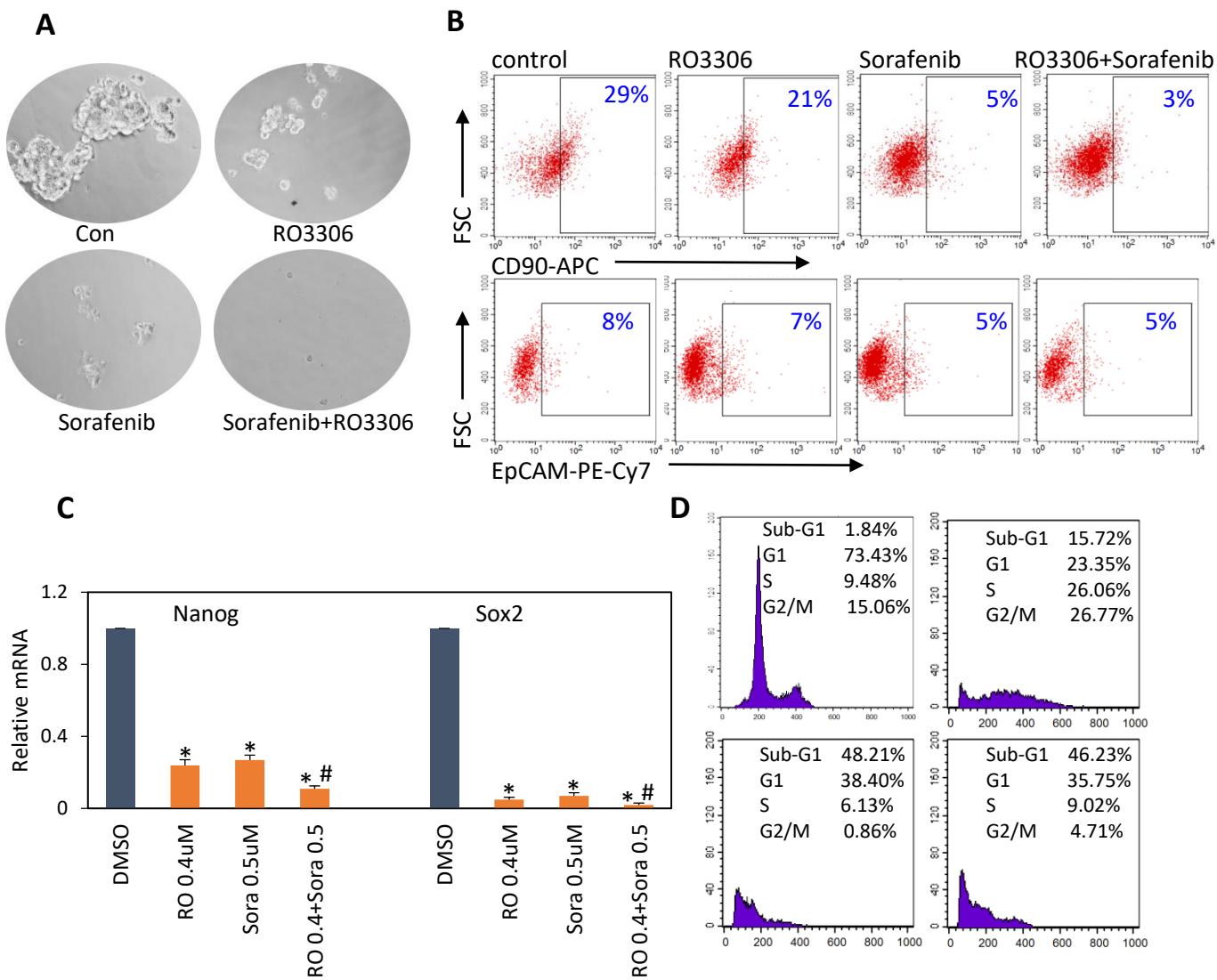
**Supplementary Figure.2 Effectiveness of RO3306 and sorafenib on PDX tumors.** A-B, The growth curves of HCC case #4 and #10 PDX models, respectively. C-D, The no inhibition effect of indicated treatments on HCC case #9 PDX models, due to the low tumor progression process. E, The tumor weight during the treatments from day 1 to day 30 with no obvious body weight change. (\*, p < 0.05; \*\*, p < 0.01 compared to control.)

Supplementary figure 3



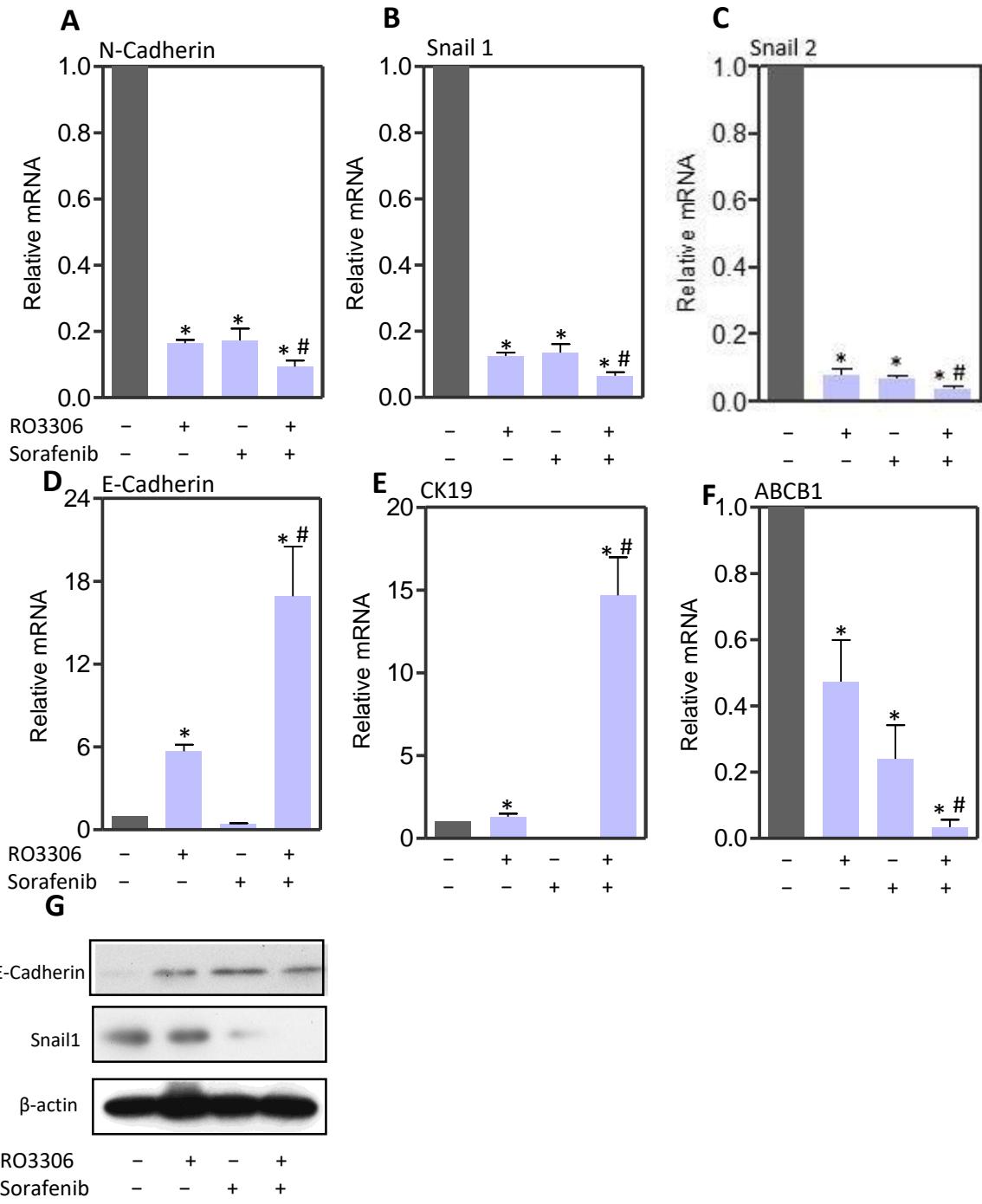
**Supplementary Figure 3 The combined effect on the EMT transition.** A-C, The synergistic effect of increased E-Cadherin and decreased Snail1 and Snail 2 on F1 PDX models. D, The combined treatment could decrease Snail1 and upregulate E-Cadherin analysed by western blot (left: case #4; right: PDX case #10) . (\*, p < 0.05; \*\*, p < 0.01 compared to control and #, p < 0.05 compared to single agent treatments.)

### Supplementary figure 4



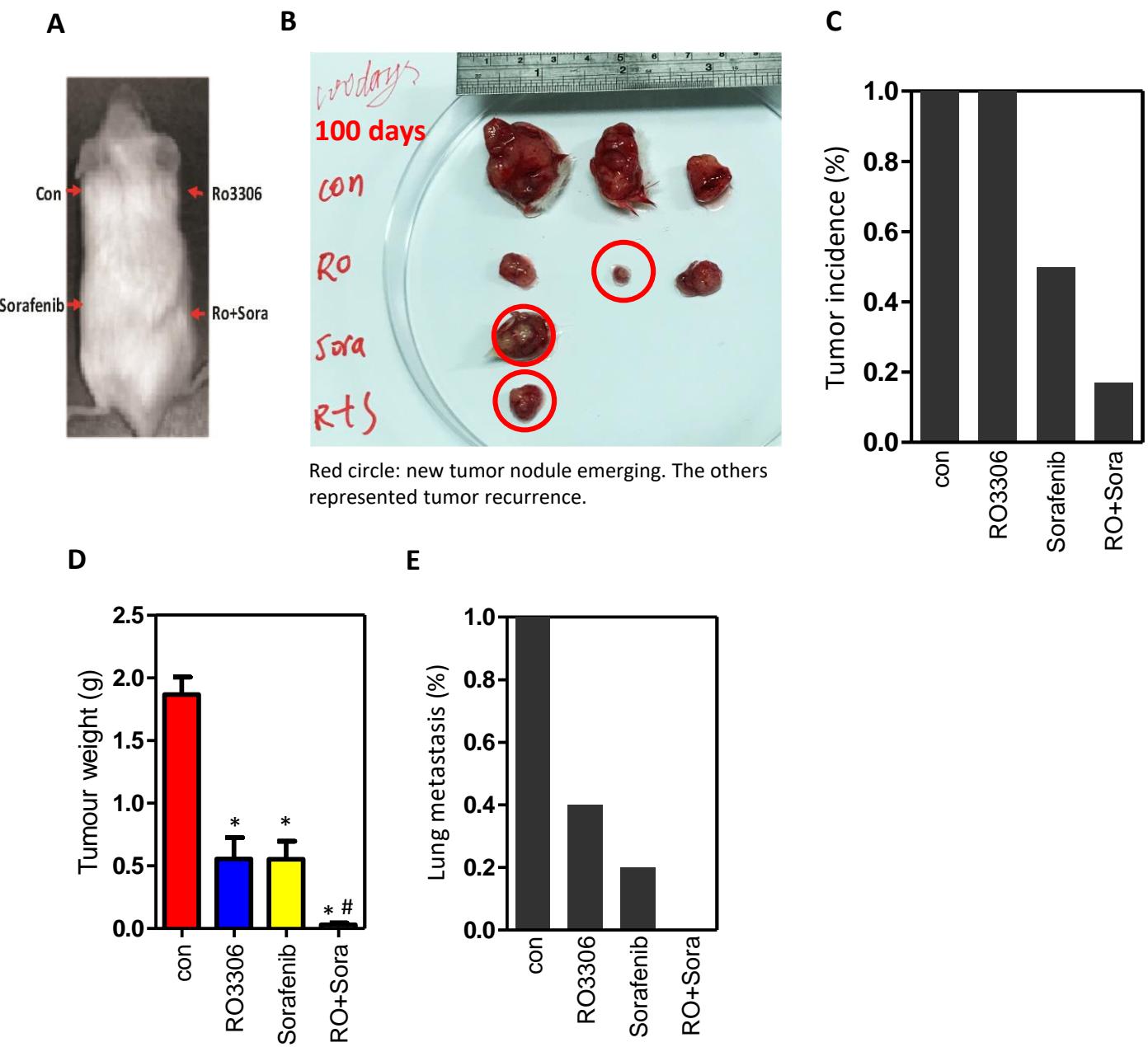
**Supplementary Figure.4 Anticancer effect of RO3306 and sorafenib combination or alone on 97H sphere cells.** A, The morphology of sphere formation in indicated treatment groups. B, The suppression effect on CD90 and EpCAM CSCs. C, Low dose of RO3306 (4  $\mu$ M) and sorafenib (2.5  $\mu$ M) decreased the stemness-related genes Nanog and Sox2. D, Low dose RO3306 and sorafenib combination for 72h promoted CSCs enter into a Sub-G1 phase. (\*, p < 0.01 compared to control. #, p < 0.05 compared to single agent treatments).

Supplementary figure 5



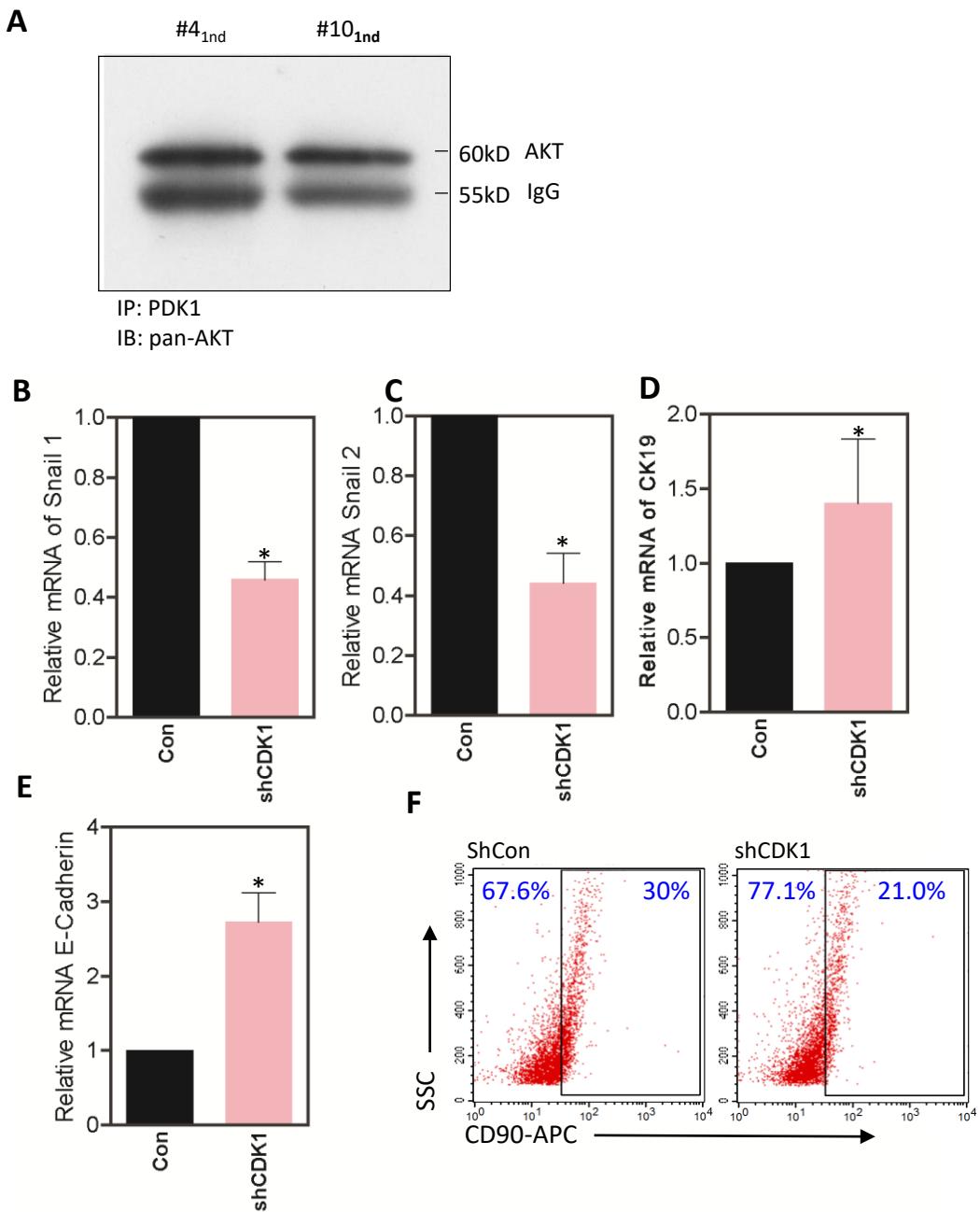
**Supplementary Figure.5 RO3306 and sorafenib alone or combination on the mRNA levels of EMT in 97H sphere cells.** A-E, The synergistic effect of downregulation N-Cadherin, Snail1 and Snail2, whereas the E-Cadherin and CK19 were upregulated. F, The synergistic effect on ABCB1 drugs resistance gene. G, Western blot analysis of E-Cadherin and Snail 1 in indicated groups. (\*, p < 0.01 compared to control. #, p < 0.05 compared to single agent treatments).

Supplementary figure 6



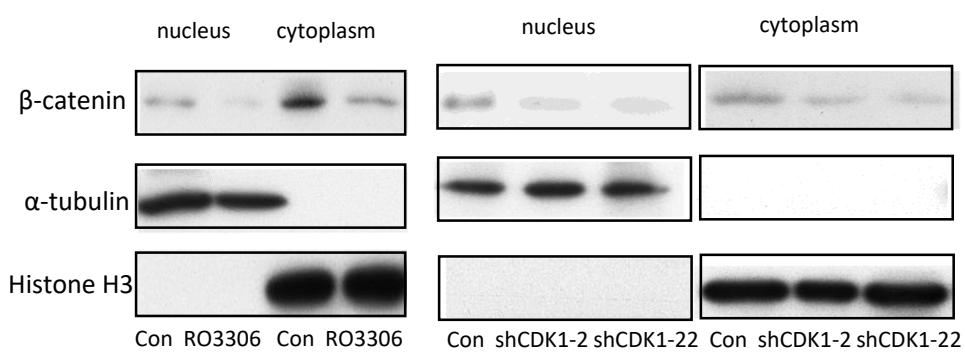
**Supplementary Figure.6 The combined effect on CSCs tumorigenicity after 100 days.** A, The indicated points of the pretreated-CSCs injection. B, Images of recurrence tumor and new emerging tumor after 100 days. C, Statistical comparison of tumor incidence (%) in various pretreat-groups after 100 days without further treatment. D, Tumor weight of 97H CSC-derived orthotopic tumor models in indicated groups. E, The percentage of lung metastasis 100%, 40%, 20% and 0% in four indicated treatments labeled above. (\*, p < 0.01 compared to control. #, p < 0.05 compared to single agent treatments).

Supplementary figure 7



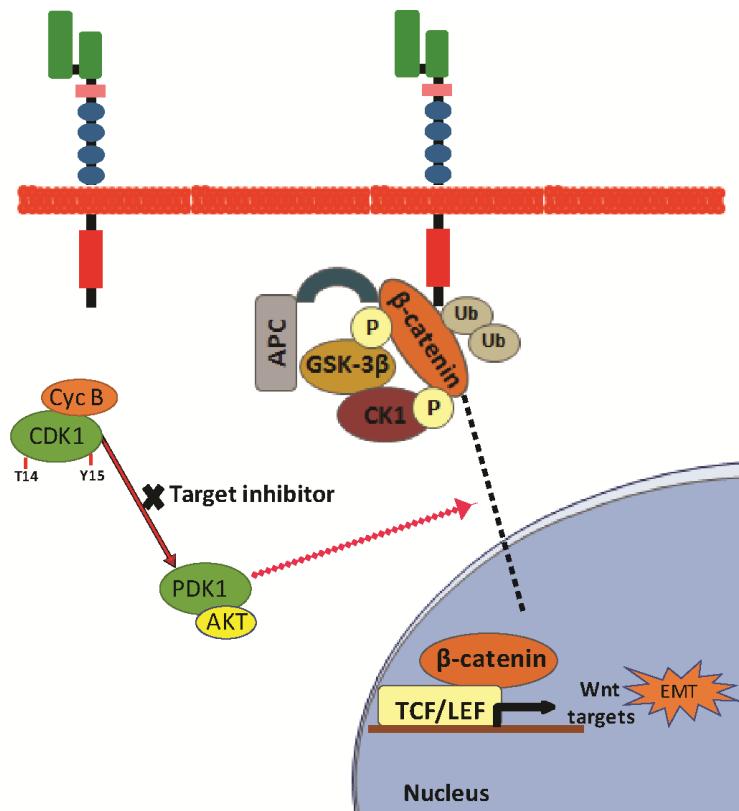
**Supplementary Figure.7 The interaction of PDK1 and AKT, and CDK1 silencing effect on 97H sphere cells.** A, The Co-IP analysis the interaction between PDK1 and AKT. B-E, Knockdown CDK1 reverse EMT process. F, The CD90+ CSCs population of shCDK1 knockdown compared with the scramble group after 24h. (\*, p < 0.01 compared to control).

Supplementary figure 8



**Supplementary Figure. 8 The expression level of β-catenin in cytoplasm and nucleus of various groups and CDK1 Knockdown.** A, Level of β-catenin in cytoplasm and nucleus were measured by western blot analysis. Decreased β-catenin was detected both in cytoplasm and nucleus. A-tubulin and Histone H3 as control loading.

Supplementary figure 9



**Supplementary Figure.9 The CDK1-PDK1-β-Catenin axis.** A schematic diagram illustrates the proposed CDK1-PDK1-β-Catenin associated pathways and its downstream targets EMT relates to HCC progression and metastasis.

**Table S1**

Table S1: Primer sequences

Primer	Sequence (5' to 3') F: forward; R: reverse
NANOG	F: CAAAGGCAAACAACCCACTT R: TCTGCTGGAGGCTGAGGTAT
OCT4	F: CTCACCCTGGGGTTCTATT R: CTCCAGGTTGCCTCTCACTC
SOX2	F: GCTGCGAACAGTCAGACAGA R: ACCTCCCCTCCAAGGTAGG
CDH1	F: CGACCCAACCCAAGAACATCA R: AGGCTGTGCCTTACAGA
CDH2	F: GACAATGCCCTCAAGTGTT R: CCATTAAGCCGAGTGATGGT
CK19	F: TTTGAGACGGAACAGGCTCT R: AATCCACCTCCACACTGACC
SNAIL1	F: CACTATGCCGCGCTCTTC R: GGTCGTAGGGCTGCTGGAA
SNAIL2	F: GAGCATTGCAGACAGGTCA R: GCTTCGGAGTGAAGAAATGC
ABCB1	F: GCCTGGCAGCTGGAAGACAAATAC R: ATGGCCAAAATCACAAGGGTTAGC
shRNA-Scramble1	AGCGGGATGTGCTTATGCAGGATTCCACGAGTGGAA TCCTGCATAAGCACATCC
shRNA-CDK1-2	AGCGGGTCAGTACATGGATTCTCACTCGAGAGTGAA AGAATCCATGTACTGACC
shRNA-Scramble11	AAAAGGATGTGCTTATGCAGGATTCCACTCGTGGAA TCCTGCATAAGCACATCC
shRNA-CDK1-22	AAAAGGTCACTACATGGATTCTCACTCTCGAGTGAA AGAATCCATGTACTGACC

Table S2. Correlation between clinicopathological parameters and CDK1 high or low expression

Parameters	Category	Cases (n=39)	CDK1 level		<i>P</i>
			low	high	
Sex	Male	32	19	13	0.139
	Female	7	2	5	
Age	≥60	22	12	10	0.921
	<60	17	9	8	
Tumor UICC7 stage	I-II	30	17	13	0.519
	III-V	9	4	5	
Tumor size	≤ 5 mm	16	9	7	0.802
	> 5 mm	23	12	11	
Tumor nodules (no.)	1-2 nodeuls	31	15	16	0.178
	≥ 3 nodeuls	8	6	2	
Venous infiltration	absent	22	13	9	0.455
	present	17	8	9	
HBsAg	positive	34	18	16	0.768
	negative	5	3	2	
AFP level	low than 20ng/ml	16	9	7	0.802
	high than 20ng/ml	23	12	11	
One year recurrence	no recurrence	29	19	10	0.013*
	recurrence	10	2	8	
Five year recurrence	no recurrence	21	15	6	0.017*
	recurrence	18	6	12	

Table S3. Correlation between clinicopathological parameters and CDK1 and PDK1 related

Parameters	Category	Cases (n=39)	CDK1 and PDK1 related		<i>P</i>
			NO	YES	
Sex	Male	28	15	13	0.790
	Female	5	3	2	
Age	≥60	19	10	9	0.797
	<60	14	8	6	
Tumor UICC7 stage	I-II	25	14	11	0.767
	III-V	8	4	4	
Tumor size	≤ 5 mm	14	9	5	0.335
	> 5 mm	19	9	10	
Tumor nodules (no.)	1-2 nodeuls	31	15	16	0.604
	≥ 3 nodeuls	8	6	2	
Venous infiltration	absent	25	13	12	0.898
	present	8	5	3	
HBsAg	positive	28	16	12	0.478
	negative	5	2	3	
AFP level	low than 20ng/ml	13	9	4	0.172
	high than 20ng/ml	20	9	11	
One year recurrence	no recurrence	24	15	9	0.134
	recurrence	9	3	6	
Five year recurrence	no recurrence	18	13	5	0.025*
	recurrence	15	5	10	

**Table S4.** The association of clinicopathological parameters, CDK1, PDK1 and CDK1 related PDK1 expression with HCC overall survival and disease-free survival using univariate and multivariate analysis

Clinical parameters	Overall survival						Disease free survival					
	Univariate analysis			multivariate analysis			Univariate analysis			multivariate analysis		
	HR	95%CI	p value	HR	95%CI	p value	HR	95%CI	p value	HR	95%CI	p value
Sex (male vs. female)	3.08 (0.92-10.36)		0.069				1.93 (0.70-5.35)		0.204			
Age (<60 vs. ≥60)	1.37 (0.44-4.24)		0.590				1.58 (0.68-3.67)		0.283			
HBsAg (pres vs. abs)	0.74 (0.16-3.38)		0.698				0.74 (0.24-2.21)		0.586			
AFP level (≤20 vs. ≥20)	1.39 (0.42-4.63)		0.589				2.07 (0.80-5.32)		0.132			
Tumor UIICC7 stage (I,II and III,IV,V)	5.84 (1.84-18.50)		<b>0.003**</b>	4.01 (1.16-13.90)		<b>0.029*</b>	4.33 (1.74-10.758)		0.002**	4.49 (1.50-13.36)		<b>0.007**</b>
Tumor size (<5 vs. ≥5)	2.44 (0.66-9.05)		0.181				1.91 (0.77-4.74)		0.160			
Tumor nodules (no.) (≤1 vs. ≥2)	2.15 (0.68-6.79)		0.191				1.54 (0.65-3.68)		0.328			
Venous infiltration (pres vs. abs)	3.15 (0.95-10.50)		0.061				3.16 (1.29-7.75)		0.012*	3.32 (1.11-9.99)		<b>0.032*</b>
CDK1 (low or high)	4.20 (1.11-15.90)		<b>0.023*</b>	1.99 (0.37-10.69)		0.424	2.08 (0.81-5.38)		0.130			
PDK1 (Low or high)	4.20 (1.11-15.90)		<b>0.035*</b>	2.61 (0.52-13.21)		0.247	2.31 (0.95-5.58)		0.064			
CDK1 related PDK1 (yes or no)	1.04 (0.28-3.86)		0.96				2.91 (1.06-7.96)		0.038**	4.62 (1.47-14.54)		<b>0.009**</b>

Univariate and multivariate analysis by cox proportional hazard regression model was used to analyze the risk factors associated with the survival. HR, hazard ratio; CI, confidence interval. \*, p < 0.05; \*\*, p < 0.01.