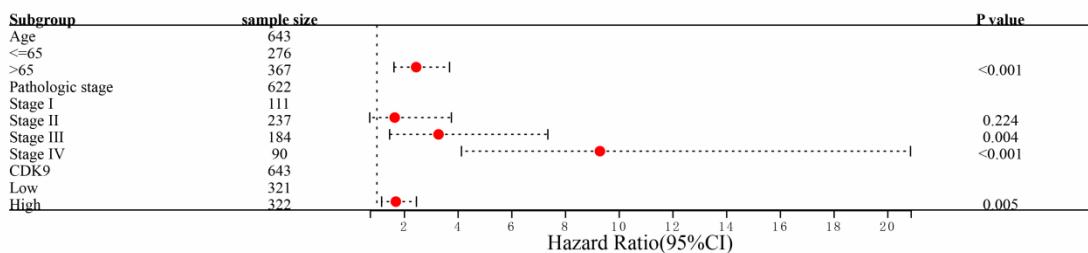


Supplementary Figure 1 Protein-protein interaction networks of cyclin-dependent kinase 9. CDK9: Cyclin-dependent kinase 9.

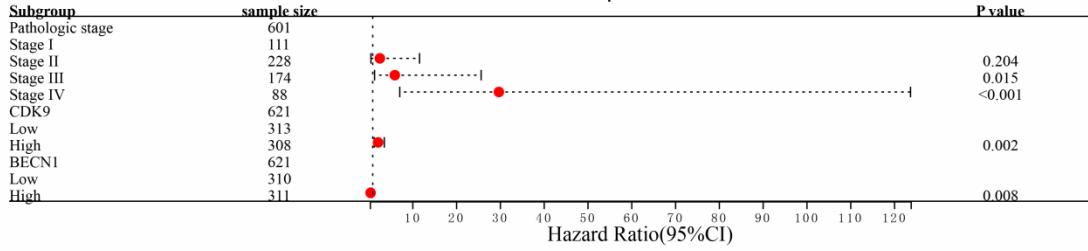
A

## OS forest plots



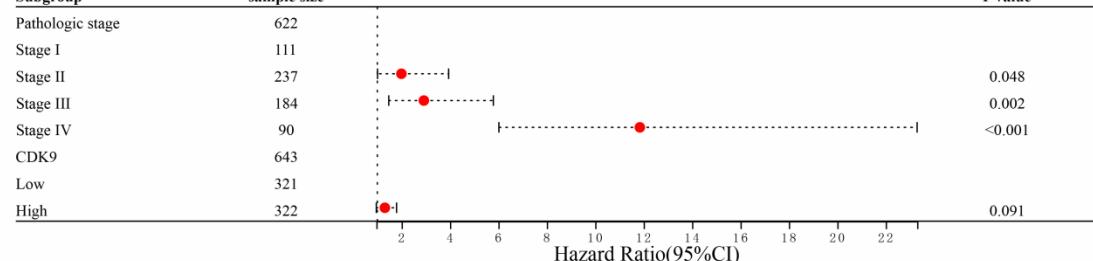
B

## DSS forest plots



C

## PFI forest plots



**Supplementary Figure 2 Multivariate analysis of overall survival, disease-specific survival, and progression-free interval.** A: Overall survival (OS); B: Forest map of disease-specific survival (DSS); C: Forest map of progression-free interval (PFI). CI: Confidence interval; CDK9: Cyclin-dependent kinase 9.

**Supplementary Table 1** Clinicopathological characteristics of the sample group from the public database

Characteristic	Category	Samples from TCGA, n = 644	Paired samples from TCGA, n = 50
n		644	50
Age in yr		68.36 (58.667,76.249)	73.367 (61.441,80.462)
T stage	T1	20 (3.12)	2 (4)
	T2	111 (17.32)	7 (14)
	T3	436 (68.02)	35 (70)
	T4	74 (11.54)	6 (12)
N stage	N0	368 (57.5)	34 (68)
	N1	153 (23.91)	8 (16)
	N2	119 (18.59)	8 (16)
M stage	M0	475 (84.22)	35 (81.395)
	M1	89 (15.78)	8 (18.605)
Sex	Female	301 (46.74)	27 (54)
	Male	343 (53.26)	23 (46)
Tumor tissue site	Colon adenocarcinoma	477 (74.07)	39 (79.592)
	Rectum adenocarcinoma	163 (25.31)	7 (14.286)
	Colon mucinous adenocarcinoma	Not Application	2 (4.082)
	Rectal mucinous adenocarcinoma	Not Application	1 (2.04)
Vital status	Dead	129 (20.03)	14 (28)
	Alive	515 (79.97)	36 (72)

Data are n (%). TGCA: The Cancer Genome Atlas.

**Supplementary Table 2** Clinicopathological characteristics of the sample group from our hospital

Characteristic	Category	Number of samples with metastasis, n = 19	Number of paired samples with liver cancer, n = 24
Age in yr		65 (53, 73)	67 (57, 75)
T stage	T1	0 (0)	2 (8.33)
	T2	0 (0)	3 (12.50)
	T3	14 (73.68)	7 (29.17)
	T4	5 (26.32)	12 (50.00)
N stage	N0	11 (57.89)	0 (0)
	N1	5 (26.32)	9 (37.50)
	N2	3 (15.79)	15 (62.50)
M stage	M0	0 (0)	0 (0)
	M1	19 (100)	24 (100)
Sex	Female	11 (57.89)	13 (54.17)
	Male	8 (42.11)	11 (45.83)
Tumor site	Right CRC	7 (36.84)	9 (37.50)
	Left CRC	12 (63.16)	15 (62.50)

Data are n (%) or median [interquartile range (IQR)]. CRC: Colorectal cancer.

**Supplementary Table 3 Univariate and multivariate analysis of overall survival, disease-specific survival, and disease-free survival for cyclin-dependent kinase 9, ABCG2, and BECN1 expression**

Char acteri stics	Overall Survival				Disease-Specific Survival				Disease Free Survival			
	Tot al, n	Univariate		Multivariate	Tot al, n	Univariate		Multivariate	Tot al, n	Univariate		Multivariate
	Hazard ratio	P valu	Hazard ratio	P valu		Hazard ratio	P valu	Hazard ratio	P valu		Hazard ratio	P valu
	643		621			643		643			643	
	(95%CI)	e	(95%CI)	e		(95%CI)	e	(95%CI)	e		(95%CI)	e
Age in yr												
≤ 65	276	Referen ce			273	Referen ce			276	Referen ce		
> 65	367	1.939 (1.320-2. 849)	< 0.00	2.462 (1.635-3. 705)	< 0.00	348	1.421 (0.894-2. 1)	0.13 7		367	1.006 (0.737-1. 371)	0.97 2
Patho logic	622				601				622			

## stage

	111	Referen				111	Referen				111	Referen			
		ce					ce					ce			
II	237	1.735	0.18	1.663	0.22	228	2.741	0.18	2.631	0.20	237	2.009	0.04	1.994	0.04
		(0.765-3.	7	(0.733-3.	4		(0.618-1	5	(0.592-1	4		(1.015-3.	5	(1.007-3.	8
		933)		774)			2.165)		1.682)			974)		945)	
III	184	3.263	0.00	3.295	0.00	174	6.688	0.01	6.042	0.01	184	3.028	0.00	2.921	0.00
		(1.462-7.	4	(1.474-7.	4		(1.571-2		(1.418-2	5		(1.529-5.	1	(1.473-5.	2
		283)		366)			8.477)		5.751)			998)		792)	
IV	90	8.065	<	9.305	<	88	27.388	<	29.800	<	90	11.749	<	11.844	<
		(3.605-1	0.00	(4.149-2	0.00		(6.598-1	0.00	(7.173-1	0.00		(5.973-2	0.00	(6.022-2	0.00
		8.043)	1	0.871)	1		13.686)	1	23.801)	1		3.108)	1	3.298)	1
Sex	643					621					643				
Fema	301	Referen				290	Referen				301	Referen			
le		ce					ce					ce			
Male	342	1.054	0.76			331	1.207	0.41			342	1.217	0.21		
		(0.744-1.	9				(0.769-1.	2				(0.892-1.	6		

		491)		895)		660)
CDK	643		621		643	
9						
Low	321	Referen		313	Referen	
		ce		ce		ce
High	322	1.525	0.01	1.705	0.00	308
		(1.073-2.	8	(1.176-2.	5	
						2.336
					<	2.213
						0.00
						322
						1.313
						0.08
						1.311
						0.09
						(0.965-1.
						3
						(0.958-1.
						1
						168)
						472)
						760)
						1
						638)
						786)
						795)
ABC	643		621		643	
G2						
Low	321	Referen		307	Referen	
		ce		ce		ce
High	322	1.068	0.70		314	1.006
		(0.756-1.	9			0.97
						(0.645-1.
						9
						(0.862-1.
						4
						510)
						570)
BEC	643		621		643	
N1						

	Low	322	Referen		310	Referen		322	Referen
			ce			ce			ce
High	321	0.783	0.16		311	0.497	0.00	0.513	0.00
		(0.554-1.	8			(0.312-0.	3	(0.313-0.	8
		108)			793)		842)		342)

CDK9: Cyclin-dependent kinase 9; CI: Confidence interval.