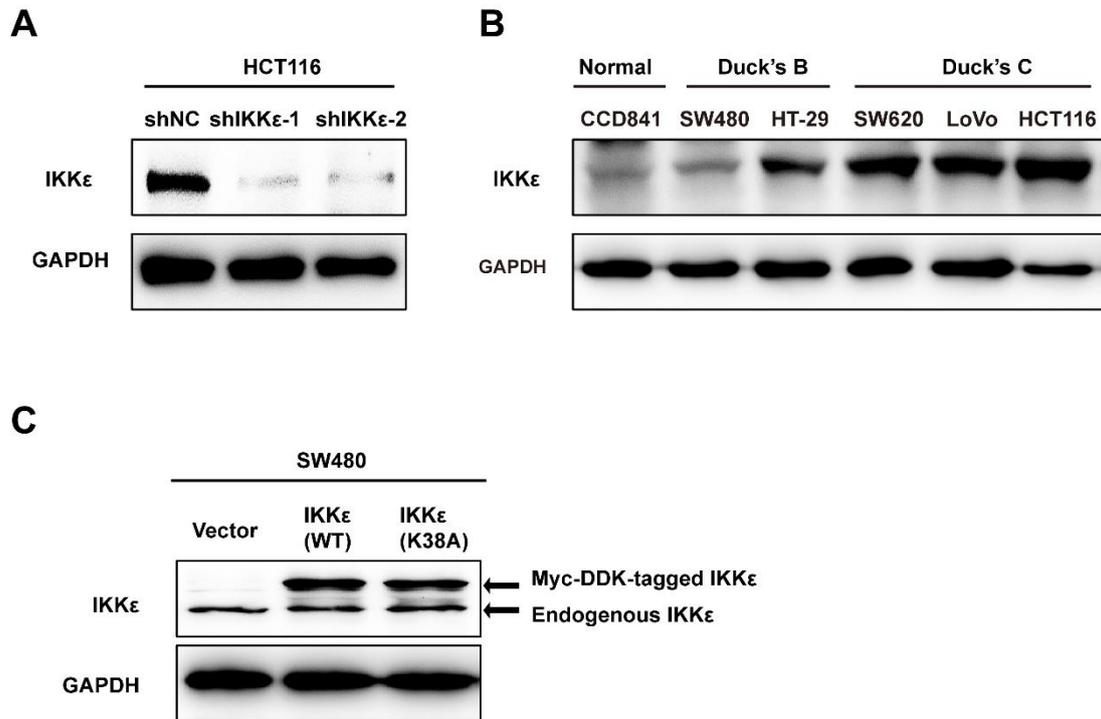
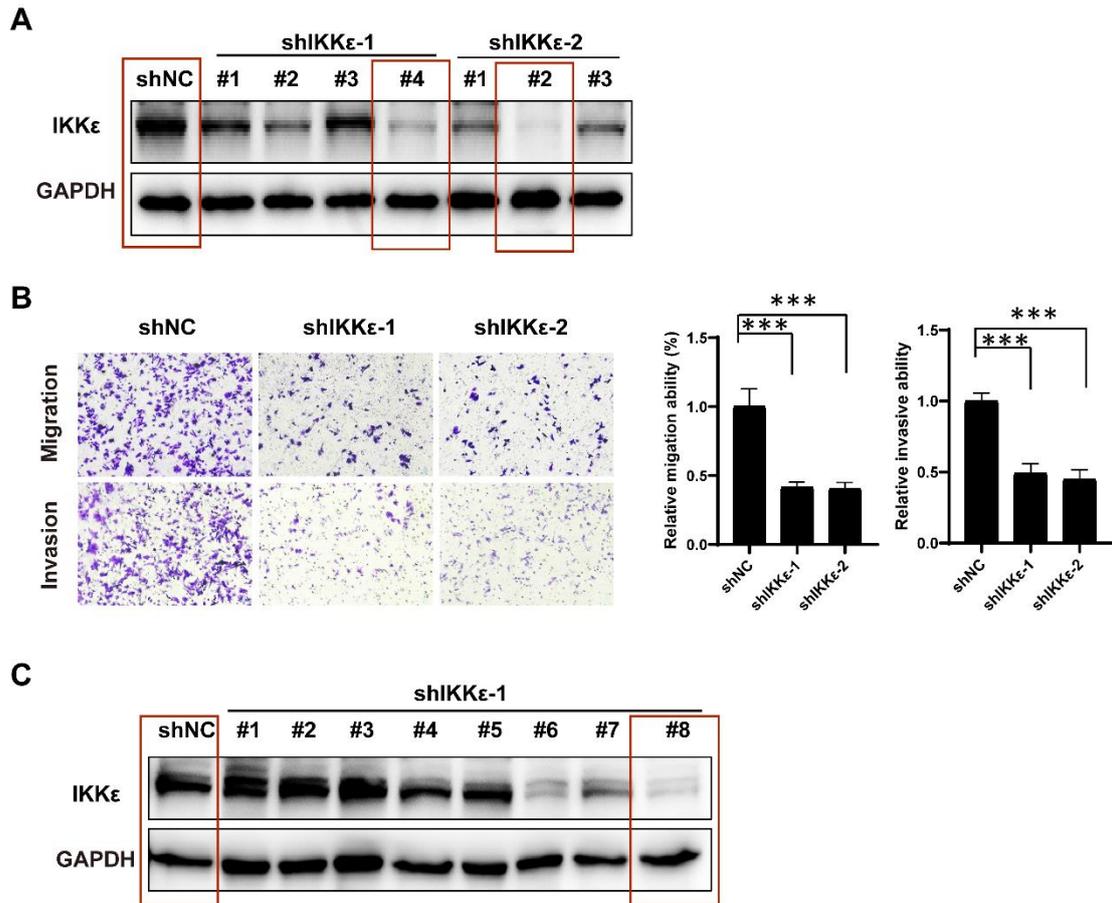


## Supplementary figures



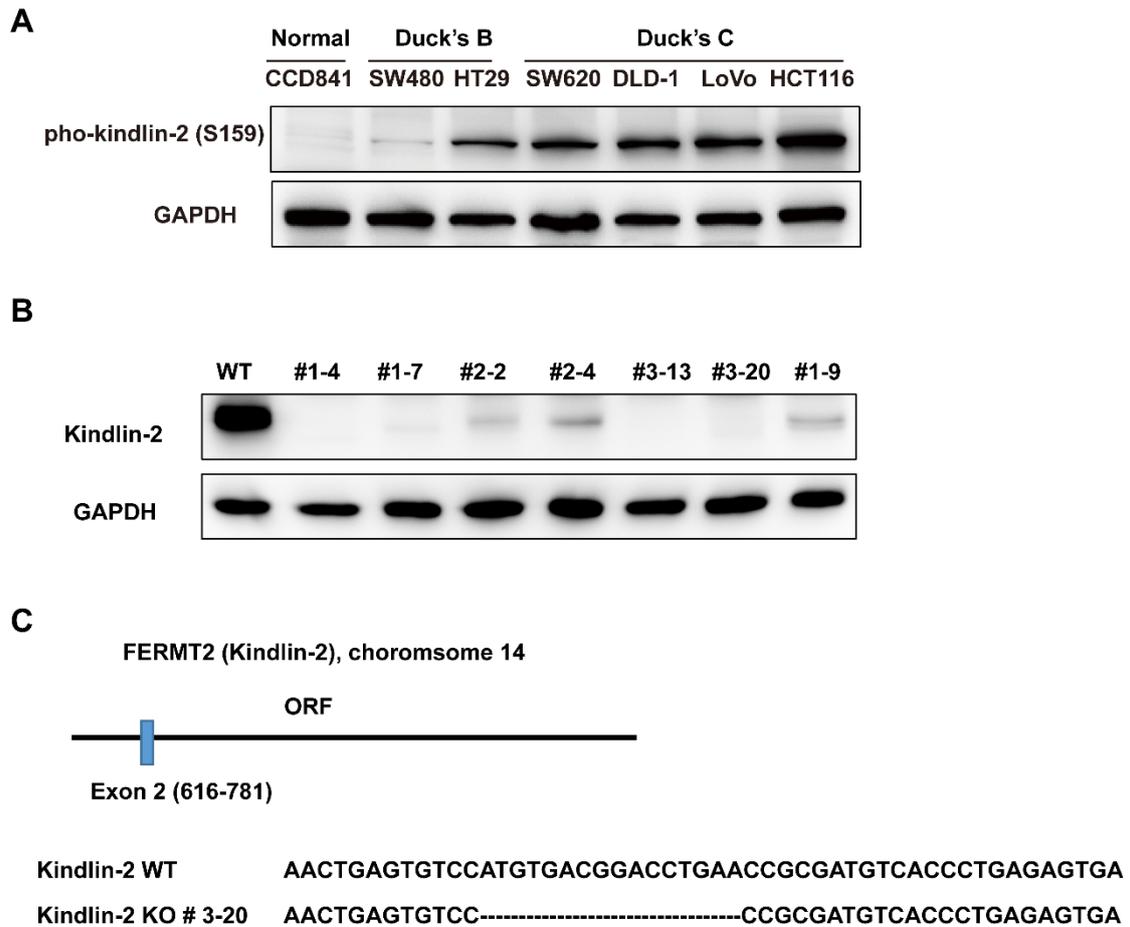
**Figure S1. Expression levels of IKKε in variety of colon cancer cell lines.**

(A) Western blot analysis of lysates from HCT116 cells stably transfected with non-targeting shRNA or shRNA targeting IKKε. (B) IKKε protein levels in the indicated cells were assessed by western blotting. (C) Western blot analysis of lysates from SW480 cells stably transfected with Myc-DDK-tagged wild-type IKKε (WT) or mutant IKKε (K38A).



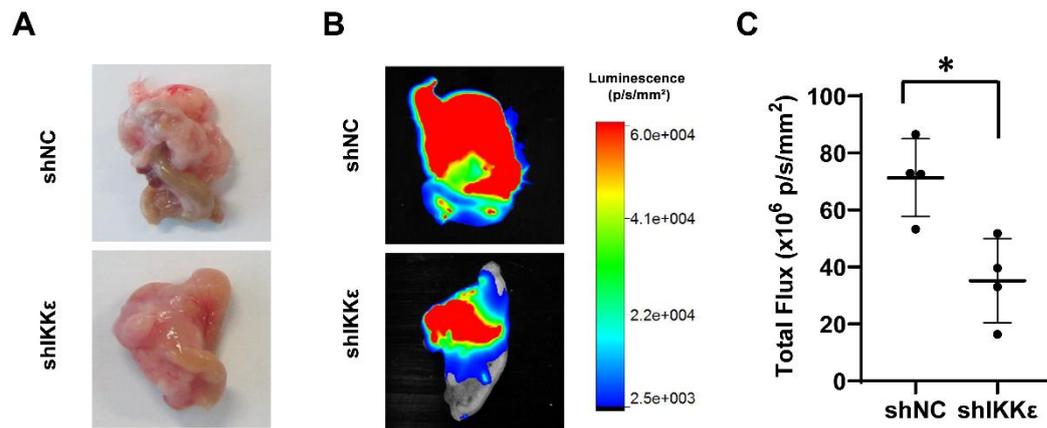
**Figure S2. Confirmation of expression levels of IKKε and the migrative/invasive ability in IKKε-knockdown in CRC cells.**

(A) Lovo cells stably expressed luciferase and IKKε-specific shRNA or control shRNA. Total protein was isolated from the indicated cells and examined for IKKε protein levels by western blotting. The red boxes highlight the cells used in the experiments. (B) Transwell invasion migration assay (top) and invasion assay (bottom) for Lovo cells stably transfected with non-targeting shRNA or shRNA targeting IKKε (left). The relative migration and invasive ability was normalized to shNC (right). All data represent the means  $\pm$  S.D. of three independent experiments ( $***P < 0.001$ ). (C) HCT116 cells stably expressed luciferase and IKKε-specific shRNA or control shRNA. Total protein was isolated from the indicated cells and examined for IKKε protein levels by western blotting. The red boxes highlight the cells used in the experiments.



**Figure S3. Confirmation of kindlin-2-knockout in HCT116 cells.**

(A) Phosphorylated-kindlin-2(S159) protein levels in the indicated cells were assessed by western blotting. (B) Kindlin-2 expression levels in HCT116 clones generated from cells that were stably transfected with the sgRNA-Cas9 knockout vector. (C) Sequence of the region of kindlin-2 that misses bases in clone #3-20 of the stable CRISPR/Cas9 knockout HCT116 cells.



**Figure S4. The growth of tumors in cecum of mice orthotopically microinjected with IKK $\epsilon$ -knockdown CRC cells was inhibited.**

(A) The gross view of tumor nodules established in cecum and (B) the bioluminescence images of cecum from mice orthotopically microinjected with HCT116 cells stably expressing shNC or shIKK $\epsilon$ . (C) Quantitative analysis of photon flux in shNC (n=4) and shIKK $\epsilon$  (n=4) mice (\* $P$ <0.05).

Supplementary table

Table. S1 Correlation between IKKε expression and clinicopathological characteristics

	variables	IKKε expression		total	χ <sup>2</sup>	p value
		low	High			
Age (year)	≤68	69	28	97	0.02	0.889
	>68	66	28	94		
	null					
T stage	T1/T2	7	2	9	0.232	0.63
	T3/T4	123	52	175		
	null					
TNM stage	I/II	78	17	95	12.8	0.000
	III/IV	51	37	88		
	null					
N stage	N0	82	20	102	10.3	0.001
	N1/N2	50	35	85		
	null					
M stage	M0	131	52	183	1.067	0.302
	M1	5	4	9		
	null					
Sex	Female	59	23	82	0.112	0.738
	Male	76	33	109		
	Null					
grade	I/II	115	41	156	3.35	0.067
	III	21	15	36		

Pearson's chi-square tests were used to analyze the correlation between IKKε and clinical features. Results were considered statistically significant at P<0.05.