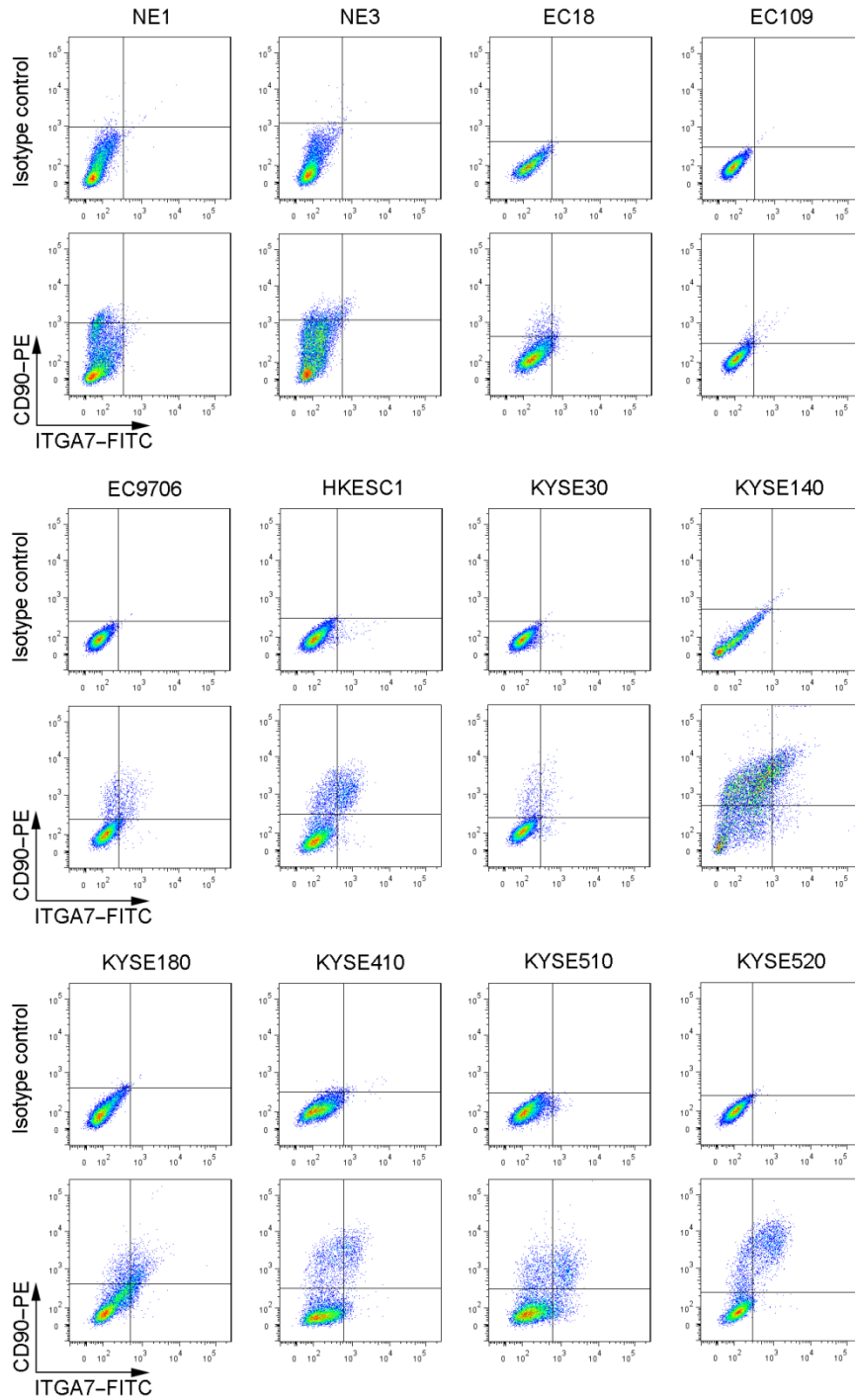
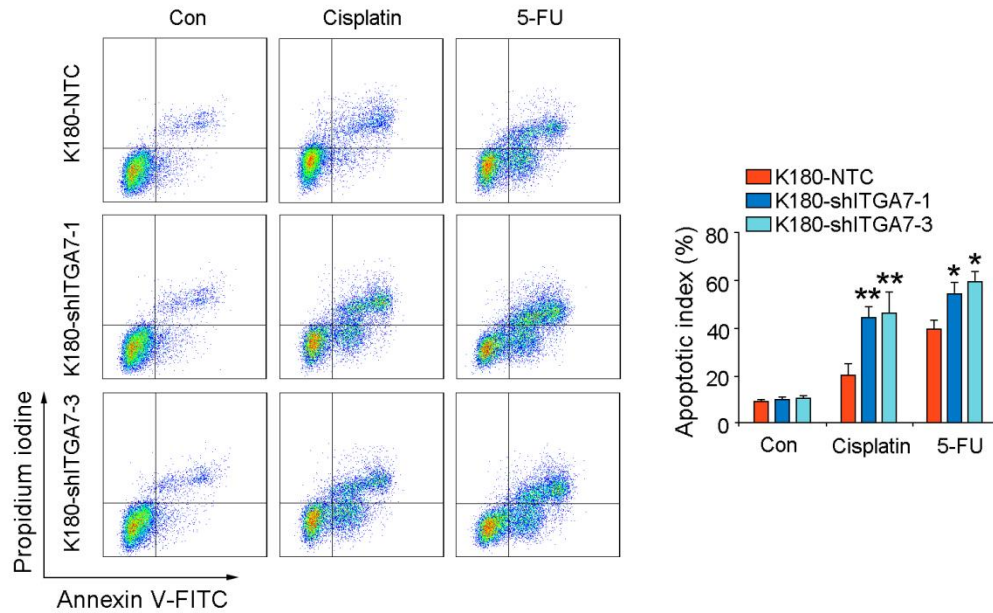


**Supplementary Figure 1. ITGA7 is sporadically expressed in ESCC.** (a) Percentage of ITGA7<sup>+</sup> cells was detected by FACS in immortalized esophageal epithelial cell line NE3 and ESCC cell lines (Red: isotype control; blue: ITGA7-PE). (b) Representative immunofluorescence images of ITGA7<sup>+</sup> cells in esophageal cell lines (green: ITGA7; blue: nuclear DAPI staining; scale bar = 50  $\mu$ m).

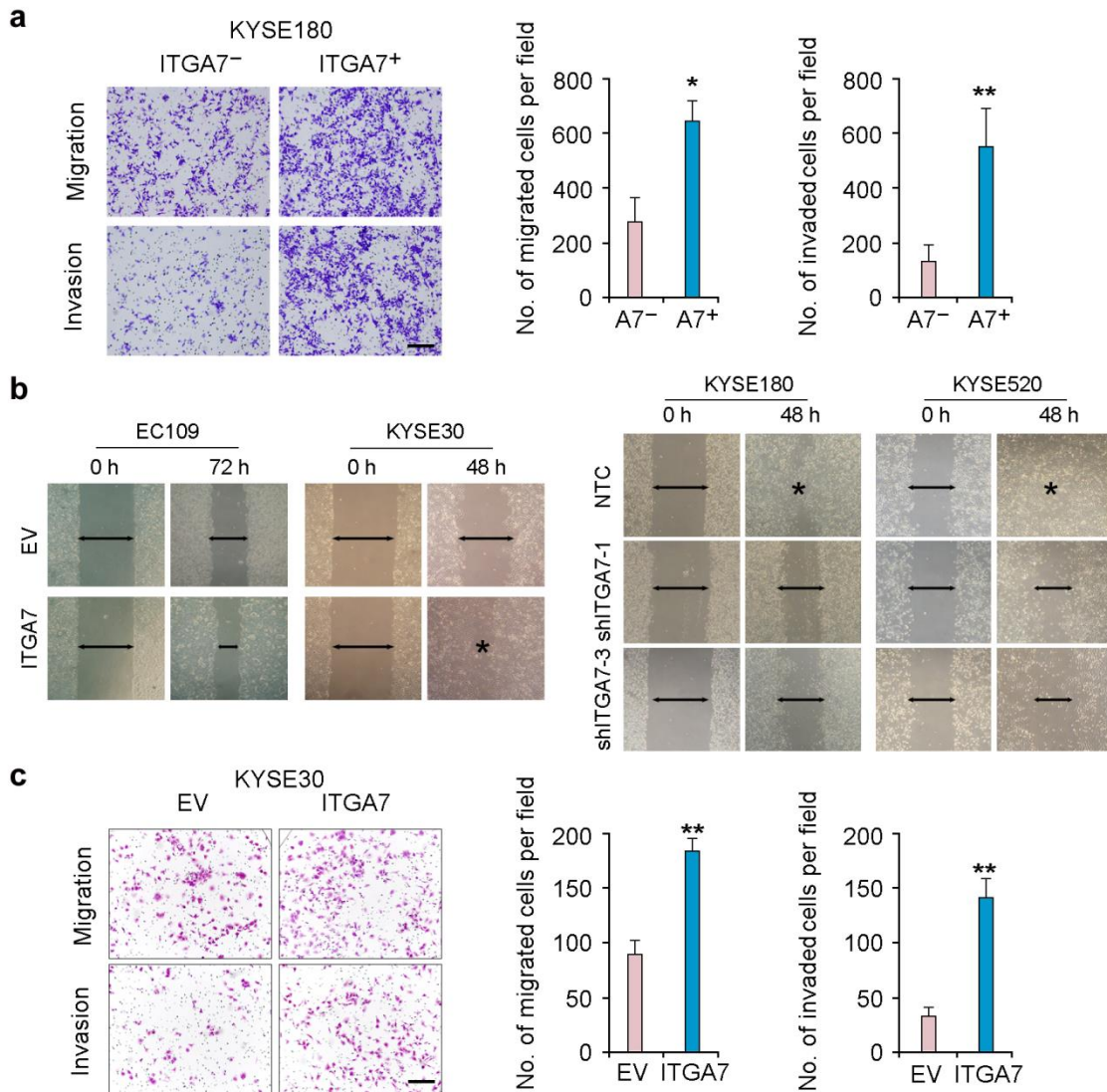


**Supplementary Figure 2.** The overlap expression of ITGA7 and CD90 in immortalized esophageal epithelial cell lines (NE1 and NE3) and ESCC cell lines.

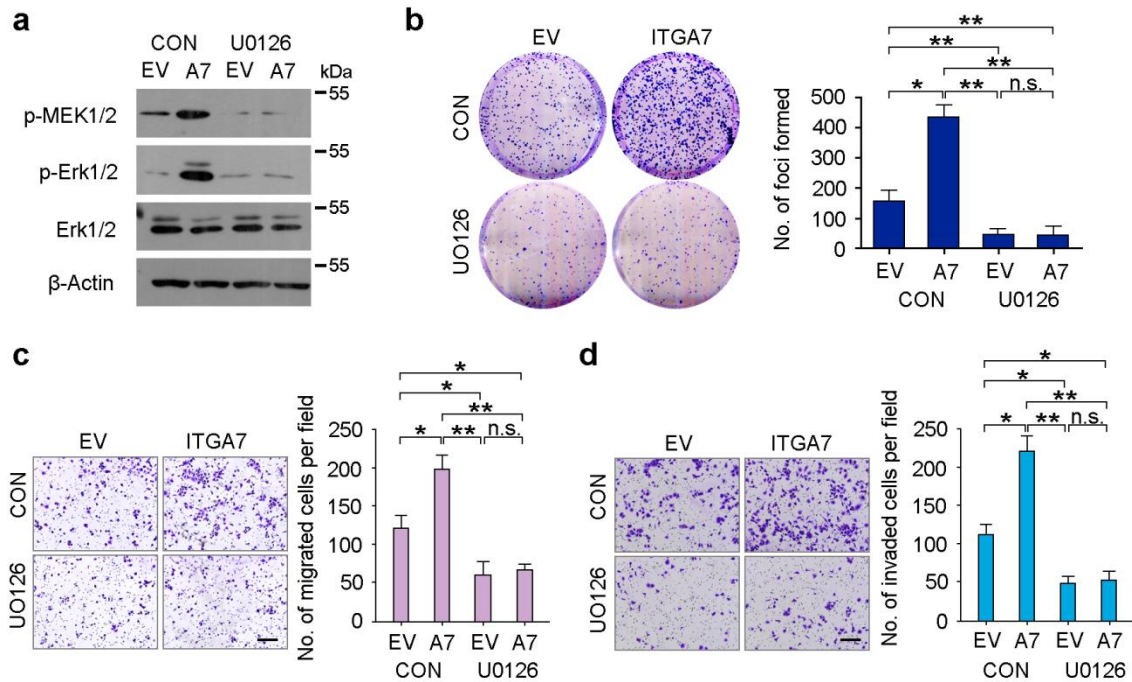


**Supplementary Figure 3. Knockdown of *ITGA7* promotes apoptosis in ESCC cells.**

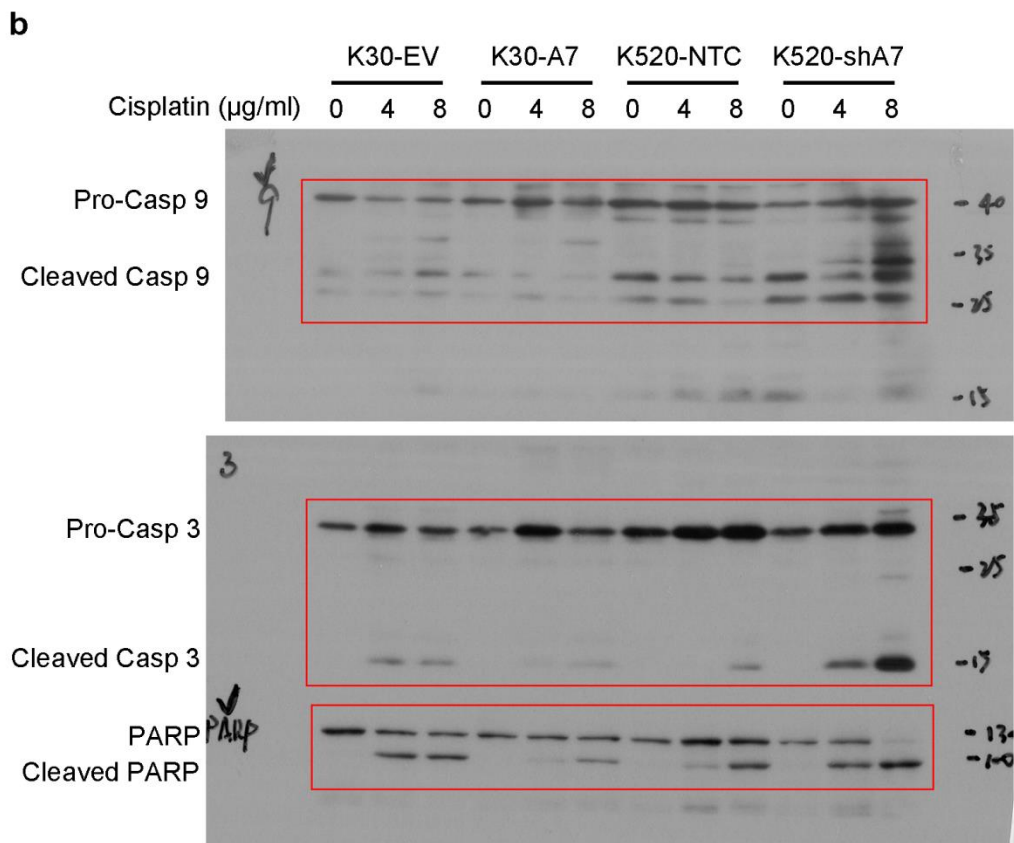
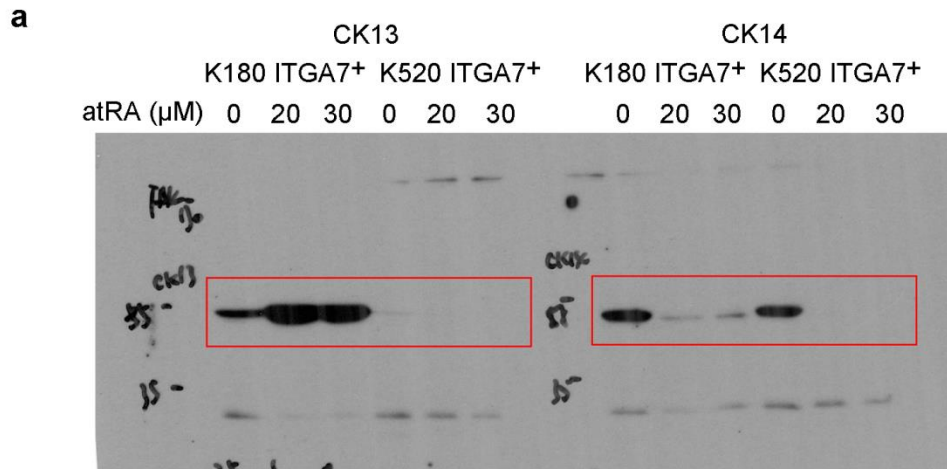
Representative of apoptosis analysis showing that sh*ITGA7*-transfected KYSE180 cells were less resistant to cisplatin and 5-FU. The apoptotic index was summarized in the bar chart and the value indicated the mean  $\pm$  s.d. of three independent experiments. Statistics: ANOVA with post hoc test. \*,  $P < 0.05$ ; \*\*,  $P < 0.001$ .



**Supplementary Figure 4. ITGA7 promotes ESCC cell migration and invasion. (a & c)** Representatives and summary of migration and invasion assays showing that ITGA7<sup>+</sup> KYSE180 cells and ITGA7-KYSE30 cells exhibited enhanced migration and invasion abilities, compare to their negative counterparts. The value expressed as the mean  $\pm$  s.d. of three independent experiments. Scale bar = 200  $\mu$ m. **(b)** ITGA7-transfected cells showed higher motility in wound-healing assay, while knockdown of ITGA7 suppressed cell motility. Statistics: (a and c) Student *t* test. \*,  $P < 0.05$ ; \*\*,  $P < 0.001$ .



**Supplementary Figure 5. MEK1/2 inhibitor suppresses ITGA7 mediated tumorigenicity and metastasis.** (a) Western blot analyses showed that the phosphorylation of MEK1/2 and downstream player Erk1/2 was significantly blocked in *ITGA7*-transfected KYSE30 cells after treatment with 20  $\mu$ M MEK1/2 inhibitor U0126 for 24 hours. Functional assays demonstrated that U0126 treatment could effectively inhibit the abilities of foci formation (b), cell migration (c; 200  $\mu$ m scale bar), and invasion (d; 200  $\mu$ m scale bar). The number of foci, migrated and invaded cells was calculated from three independent experiments and depicted as the mean  $\pm$  s.d. in the bar chart. Statistics: (b, c and d) ANOVA with post hoc test. \*,  $P < 0.05$ ; \*\*,  $P < 0.001$ ; n.s.,  $P \geq 0.05$ .



**Supplementary Figure 6. (a-b)** Full blots of Fig. 3c and Fig. 7f respectively.

**Supplementary Table 1.** Frequencies of ITGA7<sup>+</sup> cells in ESCC tumor tissues

<b>ITGA7(%)</b>	<b>No. of ESCC cases</b>
0-0.3	41
0.3-0.6	84
0.6-1	78
1-2	32
2-3	14
3-4	10
4-5	3
<b>Total</b>	<b>262</b>

**Supplementary Table 2.** Frequencies of ITGA7<sup>+</sup> and CD90<sup>+</sup> cells in esophageal cell lines

<b>Cell line</b>	<b>ITGA7<sup>+</sup>/CD90<sup>+</sup></b>	<b>ITGA7<sup>+</sup>/CD90<sup>-</sup></b>	<b>ITGA7<sup>-</sup>/CD90<sup>+</sup></b>	<b>ITGA7<sup>-</sup>/CD90<sup>-</sup></b>
NE1	0.29%	1.33%	7.04%	91.34%
NE3	1.20%	0.21%	7.30%	91.29%
EC18	2.95%	1.98%	6.47%	88.60%
EC109	0.52%	0.81%	0.97%	97.70%
EC9706	2.20%	1.53%	3.08%	93.19%
HKESC1	11.10%	1.03%	7.74%	80.13%
KYSE30	3.80%	1.02%	4.88%	90.30%
KYSE140	16.80%	1.30%	50.80%	31.10%
KYSE180	13.50%	4.26%	7.40%	74.84%
KYSE410	6.22%	2.32%	8.21%	83.25%
KYSE510	5.50%	1.40%	11.50%	81.60%
KYSE520	17.10%	1.48%	7.52%	73.90%

**Supplementary Table 3.** Overlap expression of ITGA7 and CD90

<b>Cell line</b>	<b>ITGA7<sup>+</sup><sup>a</sup></b>	<b>CD90<sup>+</sup><sup>b</sup></b>	<b>ITGA7<sup>+</sup>/CD90<sup>+</sup> in total ITGA7<sup>+</sup><sup>c</sup></b>
NE1	1.62%	7.33%	17.90%
NE3	1.41%	8.50%	85.11%
EC18	4.93%	9.42%	59.84%
EC109	1.33%	1.49%	39.10%
EC9706	3.73%	5.28%	58.98%
HKESC1	12.13%	18.84%	91.51%
KYSE30	4.82%	8.68%	78.84%
KYSE140	18.10%	67.60%	92.82%
KYSE180	17.76%	20.90%	76.01%
KYSE410	8.54%	14.43%	72.83%
KYSE510	6.90%	17.00%	79.71%
KYSE520	18.58%	24.62%	92.03%

<sup>a</sup>Calculation formula: CD90<sup>+</sup>/ITGA7<sup>+</sup>% + CD90<sup>-</sup>/ITGA7<sup>+</sup>%;

<sup>b</sup>Calculation formula: CD90<sup>+</sup>/ITGA7<sup>+</sup>% + CD90<sup>+</sup>/ITGA7<sup>-</sup>%;

<sup>c</sup>Calculation formula: CD90<sup>+</sup>/ITGA7<sup>+</sup>% / (CD90<sup>+</sup>/ITGA7<sup>+</sup>% + CD90<sup>-</sup>/ITGA7<sup>+</sup>%).



**Supplementary Table 4.** List of antibodies used in this project

<b>Antibody</b>	<b>Cat No.</b>	<b>Vendor</b>	<b>Application</b>
Mouse anti-human ITGA7	AT2566a	Abgent	WB, 1:1000; IHC FACS IF, 1:100
Rabbit anti-human ITGA7	AP18839a	Abgent	FACS IF, 1:100
PE mouse anti-human CD90	555596	BD Pharmingen	FACS, 1:20
Purified mouse anti-human CD90	555593	BD Pharmingen	IF, 1:50
Mouse anti-human CD90	ab225	Abcam	WB, 1:500
Rabbit anti-human FAK	#13009	Cell Signaling	WB, 1:1000
Rabbit anti-human FAK (Tyr397)	#8556	Cell Signaling	WB, 1:1000
Rabbit anti-human Src	#2109	Cell Signaling	WB, 1:1000
Rabbit anti-human Src (Tyr416)	#6943	Cell Signaling	WB, 1:1000
Rabbit anti-human c-Raf	#9422	Cell Signaling	WB, 1:1000
Rabbit anti-human c-Raf (Ser259)	#9421	Cell Signaling	WB, 1:1000
Rabbit anti-human MEK1/2 (Ser217/221)	#9154	Cell Signaling	WB, 1:1000
Rabbit anti-human Erk1/2	#9102	Cell Signaling	WB, 1:1000
Rabbit anti-human Erk1/2 (Thr202/204)	#9101	Cell Signaling	WB, 1:1000
Mouse anti-human $\beta$ -actin	ab6276	Abcam	WB, 1:5000
Rabbit anti-human E-Cadherin	#3195P	Cell Signaling	WB, 1:1000
Mouse anti-human $\beta$ -Catenin	#9562	Cell Signaling	WB, 1:1000
Mouse anti-human Fibronectin	ab6328	Abcam	WB, 1:500
Mouse anti-human Vimentin	#3390S	Cell Signaling	WB, 1:1000
Mouse anti-human $\alpha$ -SMA	A2547	Sigma-Aldrich	WB, 1:2000
Rabbit anti-human Akt	#9272	Cell Signaling	WB, 1:1000
Rabbit anti-human Akt (Ser473)	#9271	Cell Signaling	WB, 1:1000
Rabbit anti-human Caspase-9	#9502	Cell Signaling	WB, 1:1000
Rabbit anti-human Caspase-3	#9662	Cell Signaling	WB, 1:1000
Rabbit anti-human PARP	#9542	Cell Signaling	WB, 1:1000
Rabbit anti-human CK13	NBP2-16086	Novus biological	WB, 1:2000
Rabbit anti-human CK14	NBP1-84917	Novus biological	WB, 1:2000
Rabbit anti-human OCT3/4	#2750	Cell Signaling	WB, 1:4000
Rabbit anti-human SOX2	#3579	Cell Signaling	WB, 1:4000
Rabbit anti-human NANOG	#4903	Cell Signaling	WB, 1:4000
Goat anti-mouse IgG HRP	A4416	Sigma-Aldrich	WB, 1:10000
Goat anti-rabbit IgG HRP	A9169	Sigma-Aldrich	WB, 1:10000
Mouse IgG2a Isotype Control	M076-3	MBL	FACS, 1:200
Rabbit IgG Isotype Control	SC-3888	Santa Cruz	FACS IF, 1:100
PE mouse IgG1 Control	12-4714-82	eBioscience	FACS, 1:20
PE anti-mouse IgG2a	SC-3765	Santa Cruz	FACS, 1:100
CF <sup>TM</sup> 488 goat anti-mouse IgG	20014-1	Biotium	FACS IF, 1:5000
CF <sup>TM</sup> 488 goat anti-rabbit IgG	20012-1	Biotium	FACS IF, 1:5000

**Supplementary Table 5.** List of PCR primers for expression, cloning and BGS

<b>Gene Name</b>	<b>Primer Sequence (5'-3')</b>
<b>For Real-time PCR</b>	
<i>ITGA7</i> NM_002206.2	Forward: CTGACTCCATGTTTCGGGATCA Reverse: CACCTGTGAAGGTTTGGCG
<i>OCT3/4</i> NM_002701	Forward: CTTGCTGCAGAAGTGGGTGGAGGAA Reverse: CTGCAGTGTGGGTTTCGGGCA
<i>SOX2</i> NM_003106	Forward: AAATGGGAGGGGTGCAAAGAGGAG Reverse: CAGCTGTCATTTGCTGTGGGTGATG
<i>NANOG</i> NM_024865	Forward: AATACCTCAGCCTCCAGCAGATG Reverse: TGCGTACACCATTGCTATTCTTC
<i>NOTCH1</i> NM_017617	Forward: CCTGAGGGCTTCAAAGTGTC Reverse: CGGAACTTCTTGGTCTCCAG
<i>GAPDH</i> NM_001256799	Forward: GGAGCGAGATCCCTCCAAAAT Reverse: GGCTGTTGTCATACTTCTCATGG
<b>For full length cDNA</b>	
<i>ITGA7</i> NM_002206.2	Forward: <i>gctctaga</i> GATTTCCCTTGCATTCGCTGGGAGCTC Reverse: <i>cgggatcc</i> CTAGGCGGTGCCTGGCCCTGGAT
<b>For BGS</b>	
<i>ITGA7</i> -NCG1 +19,539 - +19,955	Forward: AGGAGGTAAGGGGAAGGGAGGG Reverse: CCCACCTCCAAACCTCACCTTC
<i>ITGA7</i> -NCG2 +13,679 - +14,162	Forward: ATTGGAGGGAGGGGGTATTGAT Reverse: TCCCCCAACTAAAACCCACAAC
<i>ITGA7</i> -NCG3 +4,167 - +4,600	Forward: GTTTTTGTTGGGTTTTTGATGA Reverse: AATATCAATTTTCTCAAACCCT

**Supplementary Table 6.** List of reagents used in this project

reagents	Cat No.	Vendor
5-aza-2'-deoxycytidine (5-aza-dC )	A3656	Sigma-Aldrich
All-trans retinoic acid (atRA)	R2625	Sigma-Aldrich
1,2,4,5-Benzenetetraamine tetrahydrochloride (Y15)	305065	Sigma-Aldrich
U0126	#9903	Cell Signaling

**Supplementary Table 7.** The shRNA sequences for lentiviral transduction

Gene	shRNA Sequence
<i>ITGA7</i> (NM_002206.2)	TRCN0000057708:CCGGCCCAGGAACCTATAATTGGAACCTCG AGTTCCAATTATAGGTCCTGGGTTTTTG TRCN0000057711:CCGGGTCCTCCATAAAGAACTTGATCTCG AGATCAAGTTCTTTATGGAGGACTTTTTG
Non-Target Control	SHC002:CCGGCAACAAGATGAAGAGCACAACCTCGAGTTGGT GCTCTTCATCTTGTTGTTTTT

## **SUPPLEMENTARY METHODS**

**Wound healing assay.** Cell mobility was studied by a scratch wound-healing assay. Briefly, cells were cultured in a 6-well plate until confluence. The culture medium was replaced by serum-free medium 24 h before wound creation. The cell layer was wounded using a sterile tip. After wounding, the medium was changed to fresh serum-free medium to remove cellular debris. Serial photographs were obtained at different time points.

**Xenograft processing and cell dissociation.** Xenografts dissected from mice were dissociated into single cells using Liberase TM (Roche) which contained a blend of collagenase I and collagenase II. Before processing, tumor tissues were rinsed with PBS once and then rinsed again with DMEM/F-12 (Gibco) supplemented with 1% penicillin/streptomycin (P/S). For every 1 cm<sup>3</sup> tissue, 5 µg per ml Liberase enzymes were added to DMEM/F-12 medium for dissociation. Tumor tissues were minced into less than 1 mm<sup>3</sup> pieces in the presence of supplemented medium in a cell culture dish under sterile condition. The cell suspension was subsequently filtered through a 100 µm cell strainer and washed with DMEM/F-12 medium. The cells were cultured in complete DMEM medium supplemented with 10% FBS and 1% P/S.