

YAP Triggers Bladder Cancer Proliferation by Affecting the MAPK Pathway

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Background: The transcriptional regulator YAP is frequently overexpressed in human cancers, such as breast and pancreatic cancers, plays an important role in tumorigenesis and can regulate many factors affecting cancer progression. These observations encouraged us to investigate the effect of YAP expression on bladder cancer.

Methods: The changes in multiple cellular functions associated with tumor progression including cell proliferation, cell migration, cell cycle, and cell apoptosis were assessed after YAP knockdown/overexpression in bladder cancer cell lines. Additionally, Western blot was developed to verify the change of proteins caused by YAP knockdown/overexpression.

Results: YAP had relatively higher expression in bladder cancer tissues than in normal tissues. The proliferation and migration of bladder cancer cells were inhibited by YAP knockdown but were promoted by its overexpression. This promoting effect was accompanied by the increased activity of MAPK/ERK pathway.

Conclusion: Our data established that YAP is an oncogene involved in bladder cancer and thus can be a potential target for treatment.

Keywords: bladder cancer, proliferation, migration, MAPK/ERK pathway

Introduction

Bladder cancer is one of the most frequently diagnosed and lethal cancers and one of the top 10 most common cancers in men with annually increasing global incidence.^{1,2} The current standard treatment for bladder cancer is surgery combined with platinum-based chemotherapy. However, bladder cancers managed by surgery still recur and progress to locally invasive or metastatic stages. Although bladder cancer has been studied at the molecular level, its underlying mechanism is still unclear, and no effective therapeutic target is currently available.^{3,4} Therefore, investigating the mechanism of bladder cancer and screening an effective target drug for its treatment are necessary.

YAP is the key transcriptional co-activator of the Hippo pathway and controls cell proliferation, apoptosis, differentiation, and drug resistance in various cancers.^{5–7} Hippo signaling pathway is a regulator of organ size and consists of a core kinase cascade, in which MST1/2 phosphorylates and activates the Lats1/2.^{8,9} YAP also plays important roles in bladder cancer tumorigenesis, progression, and drug resistance.^{10–12} However, the underlying mechanisms on how YAP regulates bladder cancer remain unclear.

This study showed that YAP is upregulated in bladder cancer cell lines, and YAP silencing impairs the proliferating ability and migration of these cells.

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Furthermore, YAP regulates cell proliferation by influencing the MAPK/ERK pathway. In summary, YAP is required for the tumorigenesis of bladder cancer cells by regulating the MAPK/ERK pathway. Our findings suggest that targeting YAP may be a viable strategy for bladder cancer treatment.

Materials and Methods

Cell Culture

SV-HUC-1, 5637, J82, TCC-SUP, and UMUC-3 bladder cell lines were purchased from the American Type Culture Collection. SV-HUC-1 cell line was maintained in Ham's F12K medium (Thermo Scientific) supplemented with 10% fetal bovine serum, 5637 was maintained in RPMI 1640 medium (Thermo Scientific) supplemented with 10% fetal bovine serum and 5% glutamine, J82 was maintained in MEM medium (Thermo Scientific) supplemented with 10% fetal bovine serum and 5% glutamine, and TCC-SUP and UMUC-3 were maintained in MEM medium (Thermo Scientific) supplemented with 10% fetal bovine serum and 5% glutamine. These cell lines were grown at 37°C containing 5% CO₂ in a humidified chamber.

Cell Transfection

Full YAP cDNA and negative control cDNA were cloned into pc-DNA3.1. Three short-interfering RNAs targeting YAP (Si-YAP#1, Si-YAP #2, and Si-YAP#3) and a relative scrambled siRNA (Si-NC) were designed and purchased from RiboBio (Guangzhou, China). The cells were transfected with plasmids or siRNAs by using Lipofectamine 3000 (Invitrogen, USA) in accordance with the manufacturer's guidelines.

Western Blot Analysis

For Western blot analysis, the cells were harvested and lysed using RIPA lysis buffer (supplemented with 1× PMSF). Equal protein amounts (20–30 µg) were separated by SDS-PAGE and then blotted onto PVDF membranes. The blots were blocked with 5% non-fat milk for 1 h, probed with primary antibody at 4°C overnight, washed with TBS-Tween-20 three times, and finally incubated with secondary antibody for 1 h at room temperature. Immunoreactive blots were visualized with ECL Plus reagents. The following primary antibodies were purchased from Cell Signaling Technology: YAP (#14074, 1:1000), β-catenin (#8480, 1:1000), E-cadherin (#14472, 1:1000), N-cadherin (#13116, 1:1000), vimentin (#5741, 1:1000), Slug (#9585, 1:1000), P38 (#8690, 1:1000),

phosphorylated p38 (pP38) (#4511, 1:1000), ERK (#4695, 1:1000), phosphorylated ERK (pERK) (#9010, 1:1000), c-Jun NH2-terminal kinase (JNK) (#9252, 1:1000), and phosphorylated JNK (pJNK) (#9255, 1:1000).

Cell Viability Assay

The cells were seeded ($0.5-1 \times 10^4$) onto a 96-well culture plate for 12, 24, or 72 h with or without the drug to determine the number of living cells. Cell viability was evaluated using CCK8 assay. SCH772984 (MedChem Express) was reconstituted following the manufacturer's recommendations and used at the indicated doses. Absorbance was measured at the absorbance of color (450 nm) by using a microplate absorbance reader.

Apoptosis and Cell-Cycle Flow Cytometry

The percentage of apoptotic cells and the cell-cycle distribution of cells was determined using the BD Annexin V-FITC/PI Assay Kit and a Cycle Staining Kit, respectively. Both assays were according to the manufacturer's instructions. Cell apoptosis and cell-cycle analyses were performed by flow cytometry (Accuri model C6).

RNA Isolation and Quantitative Real-Time PCR

Total RNA was isolated from cells and tissues using TRIzol reagent (Invitrogen, Carlsbad, CA) in accordance with the manufacturer's instructions. Total cDNA was synthesized using a HiFiScript cDNA synthesis kit (Cwbio, Jiangsu, China). Quantitative real-time PCR was performed using the UltraSYBR mixture (CW0957, Cwbio, Jiangsu, China) in ABI 7300. mRNA expression was assessed by evaluating the threshold cycle (CT) values. Relative expression was calculated using the $\Delta\Delta CT$ method, and GAPDH mRNA was employed as an endogenous control for mRNAs. The primers were designed and synthesized in Multi Sciences (Hangzhou, China) using the following primer sequences: GAPDH forward 5'-TGTGGGCATCAATGGATTTGG -3' and GAPDH reverse 5'-ACACCATGTATTCCGGGTCAAT -3'; YAP forward 5'-TAGCCCTGCGTAGCCAGTTA-3' and YAP reverse 5'-TCATGCTTAGTCCACTGTCTGT-3'.

Statistical Analysis

Comparisons between groups were performed with multiple comparisons by one-way ANOVA. Tukey's after

one-way ANOVA was used as the post-hoc test. Student's *t* test was used for pairwise comparisons. All data were obtained from at least three independent experiments and presented as mean \pm S.D. or mean \pm SEM. In all statistical tests, * represents values of $P < 0.05$, ** represents $P < 0.01$, and NS represents not significant. Statistical analyses were performed using GraphPad Prism 7.0 (San Diego, CA, USA).

Results

YAP Functions in Human Bladder Cancer Cells

Endogenous YAP expression between bladder cancer and normal tissues by using the oncomine database (<https://www.oncomine.org>) based on existing cancer microarray data sets. As shown in Figure 1A, YAP expression in cancer tissues was relatively higher than that in normal

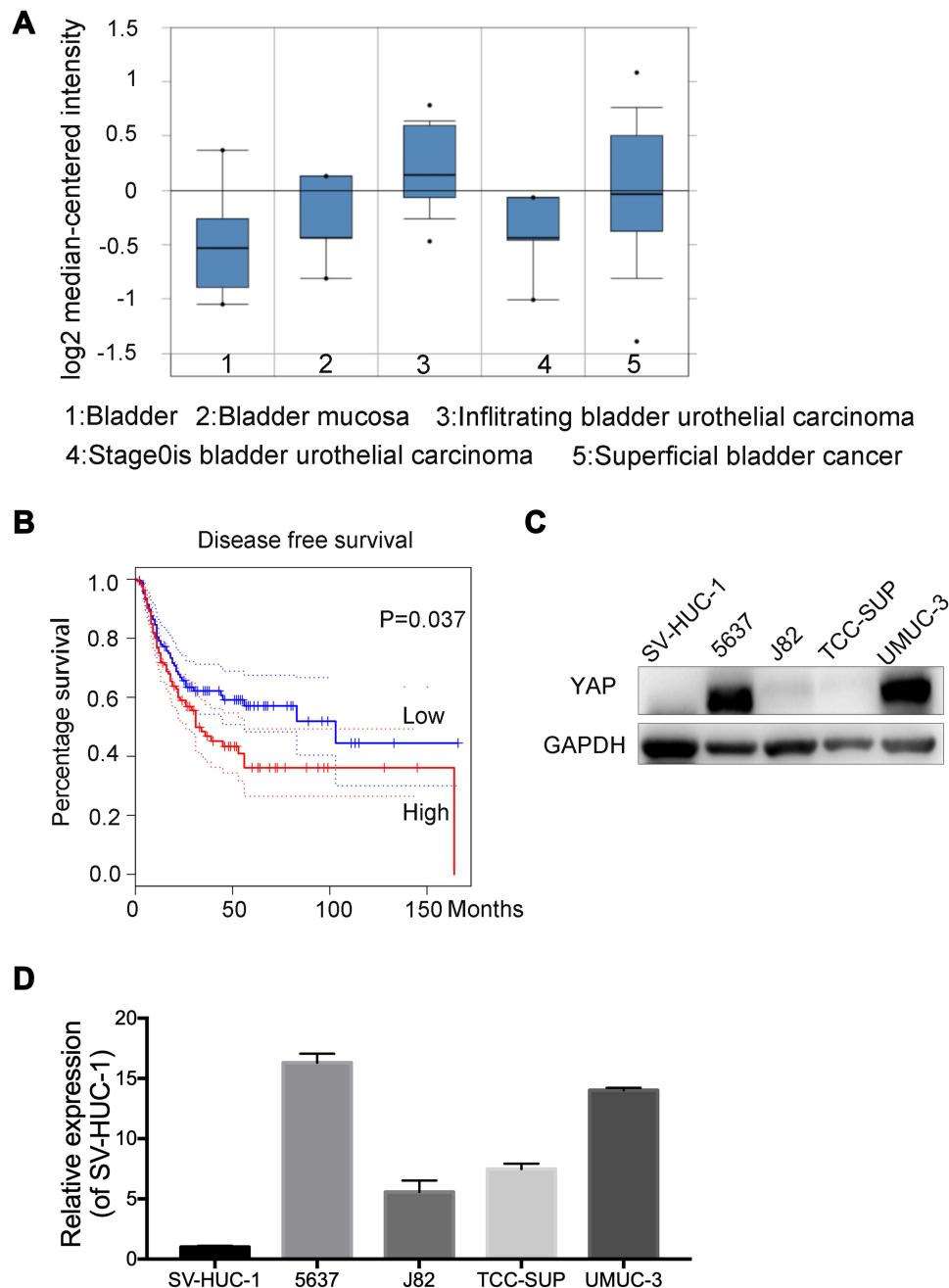


Figure 1 YAP is involved in the bladder cancer. **(A)** Oncomine databases showed that YAP was relatively highly expressed in bladder cancer. **(B)** Kaplan–Meier survival plot of bladder cancer using mRNAs of YAP from an online software KM plot. **(C and D)** Protein and mRNA expression of YAP in normal bladder cell and different bladder cancer cells.

tissues. The association of survival rate and YAP expression was further examined by using the online survival analysis software GEPIA (<http://gepia.cancer-pku.cn/>). A high YAP level was significantly associated to a low survival rate (Figure 1B). The protein and mRNA expression of YAP in normal bladder cell line and bladder cancer cell lines were then examined (Figure 1C and D). The level of YAP was higher in bladder cancer cells than in normal bladder cell line. These data strongly suggested that YAP is involved in bladder cancer progression.

YAP Promotes Cell Proliferation and Induces Cell Death in Bladder Cancer

YAP expression was relatively high in 5637 and UMUC-3 cells but low in J82 cells (Figure 1C and D). Therefore, these cell lines were selected for further experiments. Western blot and qRT-PCR analyses revealed that YAP level was altered in 5637 and UMUC-3 cells transfected with either YAP siRNA (Si-YAP #1, Si-YAP #2, and Si-YAP #3) or non-targeting control (Si-NC) (Figure 2A–C). J82 cells transfected with either YAP or vector plasmids were analyzed (Figure 2A and D). Therefore, Si-YAP#2 was employed in latter experiments. Cell proliferation assays revealed that YAP silencing attenuated the viability of 5637 cells (Figure 2E). Consistent with that in 5637 cells, YAP inhibition could reduce UMUC-3 cell proliferation (Figure 2F). However, YAP overexpression promoted the cell proliferation of J82 cells (Figure 2G). Furthermore, we determined the cell apoptosis and cell cycle whether were related with the change of YAP. The results revealed that YAP knockdown significantly induced apoptosis in 5637 and UMUC-3 cells, whereas YAP-overexpressed J82 cells inhibited apoptosis (Figure 3A). However, additional or depleted expression of YAP did not significantly influence cell cycle progression (Figure 3B). In summary, YAP promotes bladder cancer cell proliferation though inhibiting cell apoptosis.

YAP Knockdown Suppresses Cell Migration Ability

The effects of YAP on the migration ability of 5637, UMUC-3, and J82 cells were studied via Transwell assay to further investigate its biological potential in bladder cancer. The number of migrating cells increased in the YAP overexpression group but decreased in the YAP knockdown group (Figure 4A–C). The expression of EMT-related markers was also analyzed through Western

blot. E-cadherin could not be detected in 5637 and UMUC-3 cells, and vimentin was not determined in J82 cells (data not shown). These results might be related to cell migration ability. As shown in Figure 4D, YAP knockdown inhibited Slug and vimentin expression in 5637 and UMUC-3 cells. YAP overexpression for 48 h significantly inhibited E-cadherin expression but increased β -catenin, N-cadherin, and Slug expression in J82 cells (Figure 4E and F). These findings indicated that a decreased YAP expression inhibits cell migration ability.

YAP Regulates Bladder Cancer Progression by Influencing the MAPK Pathway

How YAP regulates bladder cancer cell progression was studied by controlling the expression of different target genes. The putative downstream targets of YAP were examined to further confirm the mechanisms underlying its effect on the proliferation and migration of bladder cancer cells. The activated MAPK signaling pathway is oncogenic-related and is responsible for different cellular processes, such as proliferation, differentiation, and development, in various cancers including bladder cancer.^{3,15,16} The levels of MAPK pathway-associated molecules and their phosphorylated forms were examined to determine whether YAP regulates bladder cancer proliferation via this pathway. Western blot analyses revealed that pP38, pERK, and pJNK levels were significantly high in the YAP-related high group (Figure 5A and C). In addition, YAP overexpression could induce the expression of pP38, pERK, and pJNK (Figure 5B and C). Cell proliferation in response to YAP overexpression significantly attenuated by ERK inhibitor (Figure 5D). These results indicated that YAP activates MAPK/ERK pathway and thus promotes bladder cancer proliferation.

Discussion

Experimental results revealed that YAP modulates bladder cancer. The Hippo pathway regulates cell fate and stem cell differentiation during normal organogenesis and tumorigenesis and is also involved in drug resistance.^{17–19} YAP is the major protein in the Hippo pathway, and its overexpression is involved in many diseases, especially cancer.^{10,19–22} The reciprocal antagonism between Hippo-YAP/TAZ and NF- κ B signaling is important in osteoarthritis pathogenesis and regulates matrix-degrading enzymes and cartilage degradation.²³ In breast cancer, YAP activity is maintained in a certain level by TRPS1 to regulate tumor-infiltrating

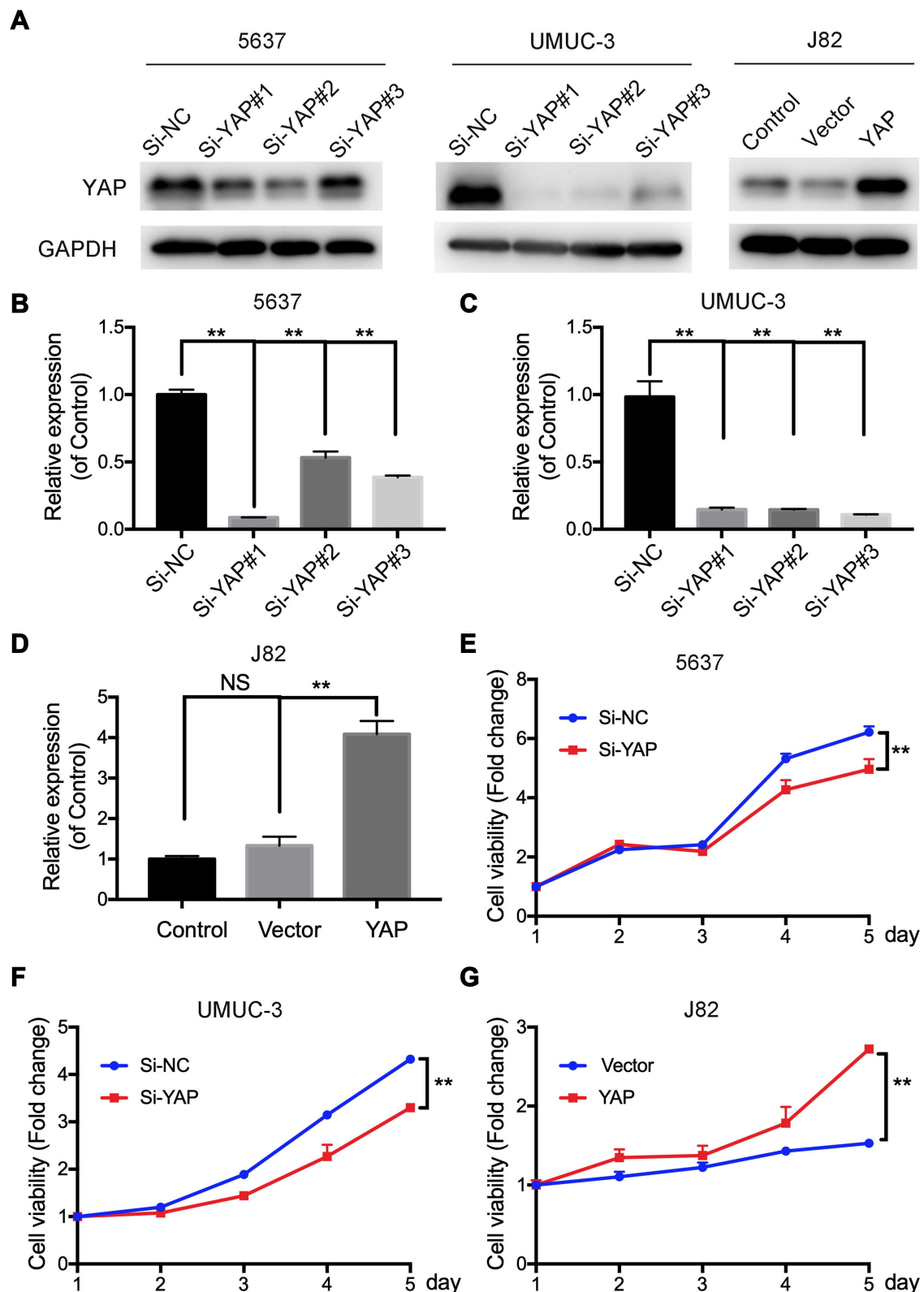
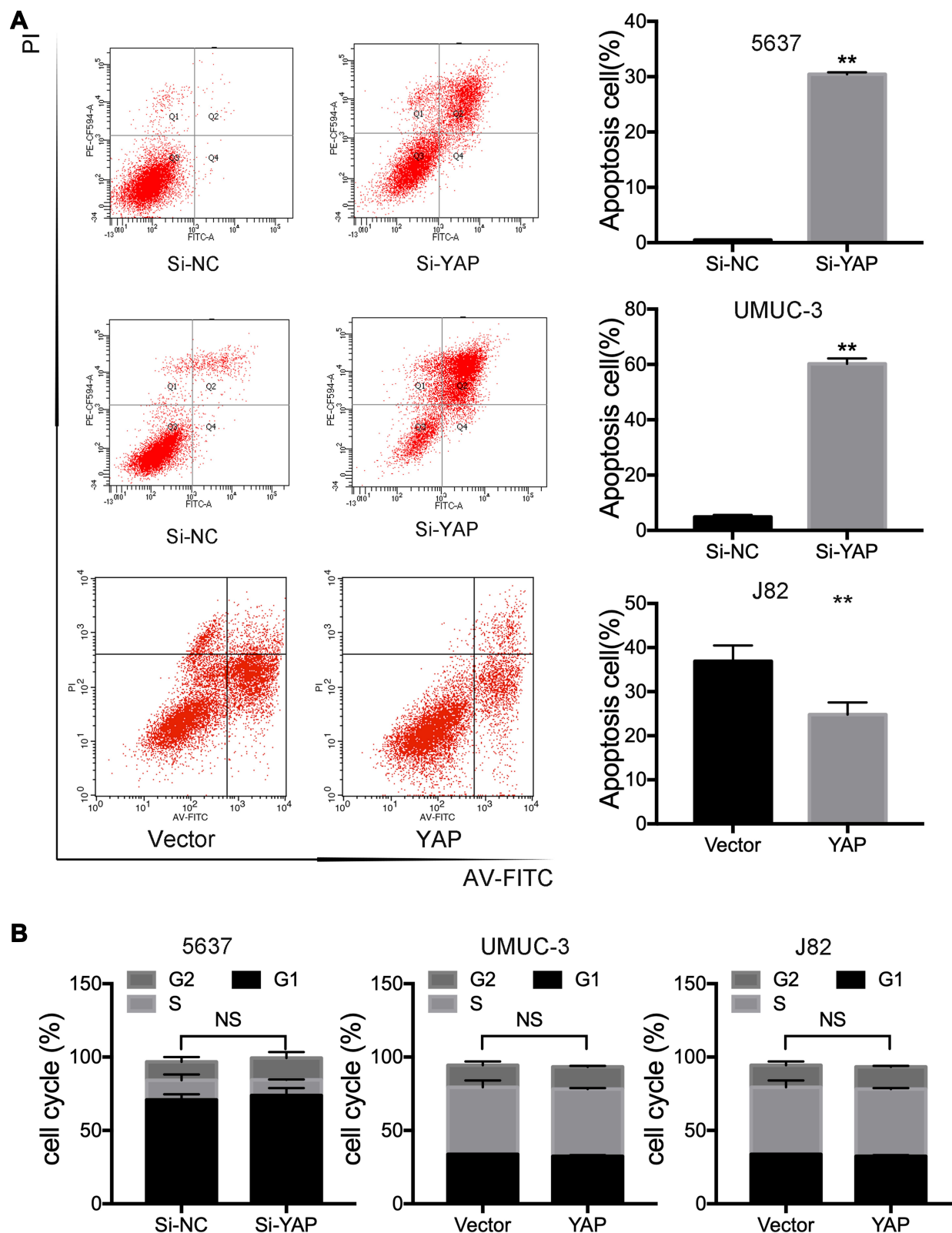


Figure 2 YAP promotes bladder cancer cell proliferation. **(A)** Knockdown YAP expression by siRNAs (si-YAP#1, si-YAP#2, and si-YAP#3) in 5637 and UMUC-3 cells. Overexpression of YAP by plasmid in J82. Protein level was detected by Western blot. **(B–D)** Analysis of YAP expression by RT-PCR in 5637, UMUC-3 and J82 cells. Si-NC or control as a control. **(E and F)** Inhibition of YAP could inhibit 5637 and UMUC-3 cells proliferation. Si-NC was chosen as a control. **(G)** Growth curve of J82 cells after transfection for 24 h with YAP plasmid was determined by CCK8 assays. Vector was chosen as a control. Values were presented as mean \pm SD from three independent experiments. ** $P < 0.01$.



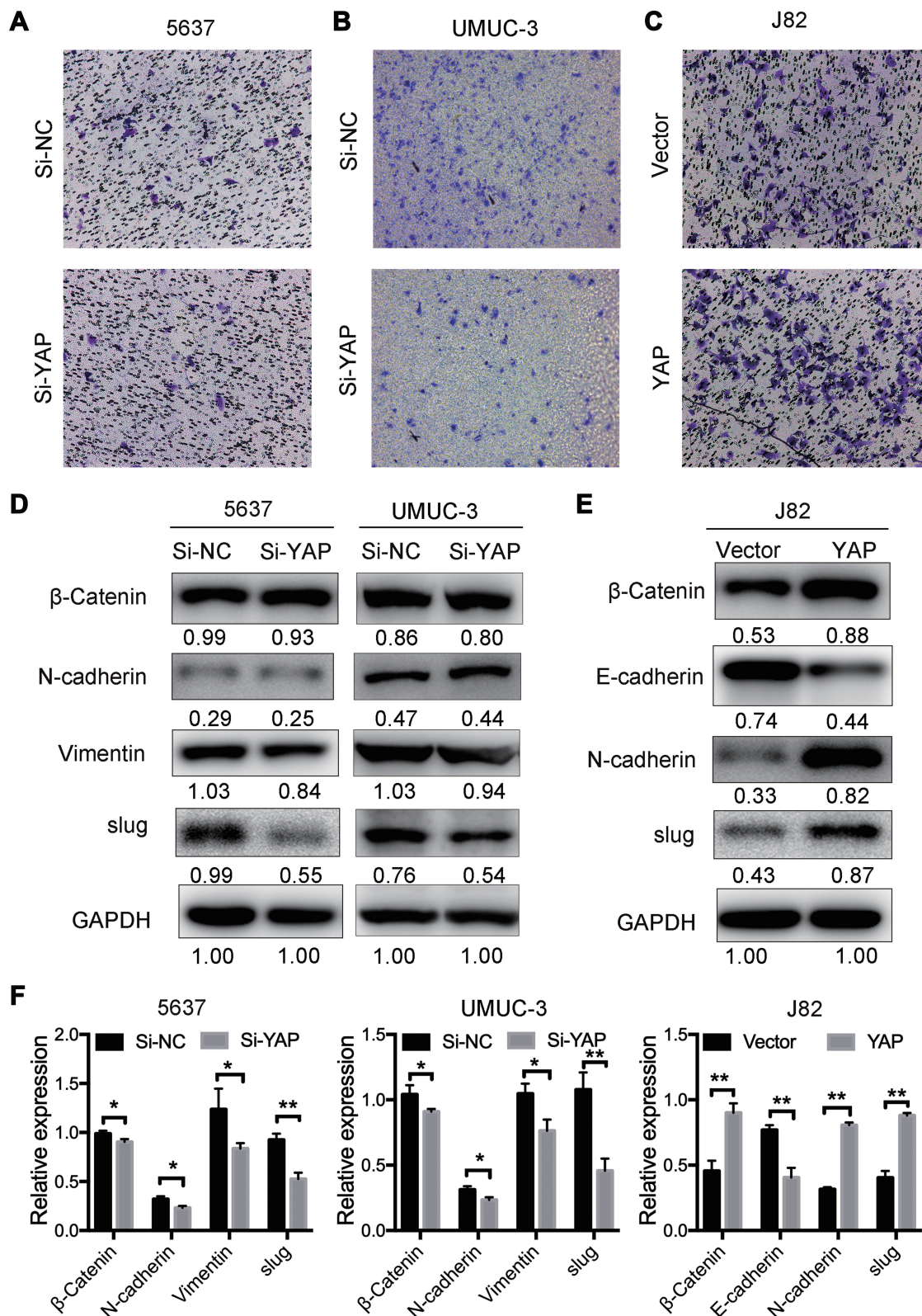


Figure 4 YAP induces bladder cancer cell migration. (A and B) Knocked down YAP expression by siRNAs in 5637 and UMUC-3 cells could reduced cell migration which was detected by Transwell assay. (C) Overexpression of YAP in J82 by plasmid after transfection for 24 h using Transwell assay. (D) Western blot analysis of the relative expression of EMT-related markers in 5637 and UMUC-3 cells 72 h after transfection. (E) YAP could increase β -Catenin, N-cadherin and slug and decrease E-cadherin detected by Western blot in J82 cells. (F) The expression was quantified and normalized to GAPDH. The error bars correspond to the means \pm SD. The values are presented as the means \pm SD. * $P < 0.05$; ** $P < 0.01$.

Abbreviation: NS, not significant.

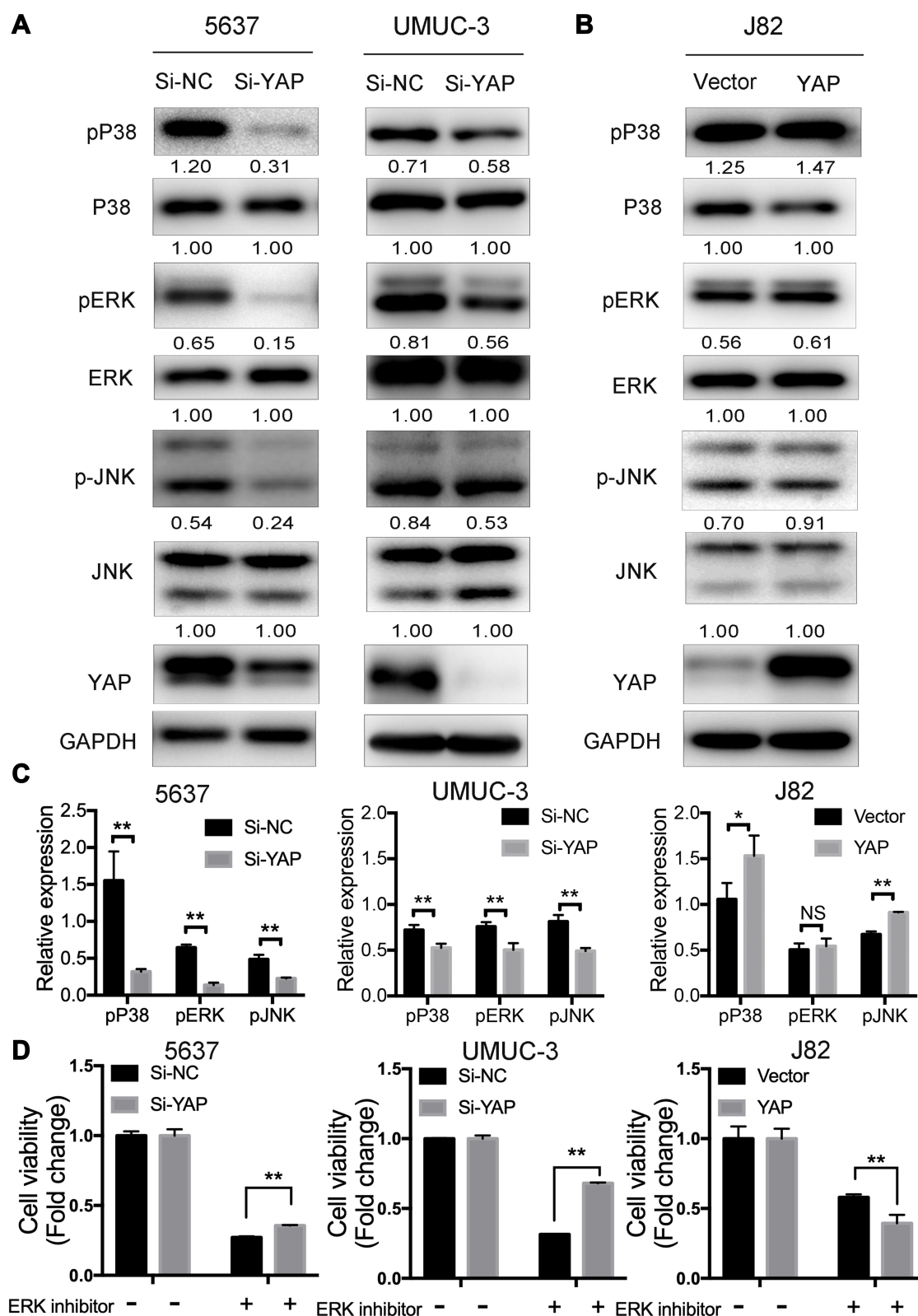


Figure 5 YAP regulates bladder cancer progression through MAPK pathway. (A and B) MAPK pathway-related markers, including ERK, JNK, P38, phosphorylated ERK, phosphorylated p38 (pP38), and phosphorylated JNK (pJNK) protein levels, were detected in 5637, UMUC-3 and J82 cells after transfection with YAP siRNAs or plasmid using Western blot. The relative expression was quantified and normalized to GAPDH. (C) The expression was quantified and normalized to ERK, P38 and JNK. (D) Cells were transfected 48h and subsequently cultured without or with SCH772984 10 μ m for 48h. The error bars correspond to the means \pm SD. The values are presented as the means \pm SD. **P < 0.01.

Abbreviation: NS, not significant.

immune cells.²⁴ YAP1-NMU expression can regulate pancreatic cancer metastasis and is related to poor prognosis.²⁵ YAP is associated with bladder cancer progression, ALDH1A1 expression, and bladder cancer stem cell properties and thus is a promising target for the prevention and treatment of bladder cancer. YAP/Nrf2 crosstalk is critical for bladder cancer chemoresistance; however, the mechanism underlying YAP regulation in bladder cancer remains unclear.

These results provide direct evidence for the protumorigenic role of YAP in bladder cancer. YAP expression in bladder cancer tissues is higher than that in normal tissues and is related to poor survival. For confirmation, cell proliferation assay and Transwell assay were performed, and the results showed that YAP enhanced tumor growth and induced cell migration in vitro as indicated by the increase in the proliferation capacity and migration of tumor cells. On the contrary, YAP knockdown significantly inhibited these abilities. These results revealed that YAP is involved in bladder cancer progression. The mitogen-activated protein kinase family, which includes ERK, p38, and JNK, modulates tumor progression by upregulating downstream kinases and transcription factors.^{3,15,16,26} ERK and p38 MAPK activate cell growth and invasive ability in bladder cancer cells. Further assays were performed to determine whether YAP can regulate the MAPK pathway, and the results revealed that the protein levels of MMP-9, XIAP, VEGF, and Cyclin-D1 were significantly reduced by PD98059 or SB203580 treatment. In addition, the protein levels of pERK, pP38, and pJNK were diminished by YAP silencing but promoted by its overexpression.

In summary, YAP is an oncogene that could actively promote cell viability and proliferation by acting as an upstream regulator and activating the MAPK pathway in bladder cancer. The YAP/MAPK pathway is a novel axis in bladder cancer tumorigenesis. This work provides novel insights into the molecular pathogenesis of bladder cancer.

Data Sharing Statement

All data analyzed during this study are included in this published article.

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Author Contributions

Dandan Qiu, Yan Zhu and Zhicheng Cong designed the study and wrote the manuscript. Dandan and Yan Zhu performed all of the experiments. Dandan Qiu and Zhicheng Cong conducted the statistical analysis. All authors read and approved the final manuscript. All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; agreed to submit to the current journal; gave final approval of the version to be published; and agree to be accountable for all aspects of the work.

Disclosure

The authors declare no competing financial interests.

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