

Original Paper

TIMP-1 Induces α -Smooth Muscle Actin in Fibroblasts to Promote Urethral Scar Formation

Yinglong Sa^a Chao Li^a Hongbin Li^a Hailin Guo^a^aDepartment of Urology, Shanghai Sixth People's Hospital, Shanghai Jiaotong University, Shanghai, China**Key Words**Tissue inhibitor of metalloproteinases-1 (TIMP-1) • α -smooth muscle actin (α -SMA) • Transforming growth factor β 1 (TGF β 1) • Epithelial-mesenchymal transition (EMT) • Urethral scar**Abstract**

Background/Aims: Tissue inhibitor of metalloproteinases-1 (TIMP-1) has been reported to upregulate in urethral scar. However, the underlying molecular mechanisms remain undefined. **Methods:** Here, we studied levels of TIMP-1 and α -smooth muscle actin (α -SMA) in the fibroblasts isolated from urethral scar tissues, compared to the fibroblasts isolated from normal urethra. Then we either overexpressed TIMP-1, or inhibited TIMP-1 by lentiviruses carrying a transgene or a short hairpin small interfering RNA for TIMP-1 in human fibroblasts. We examined the effects of modulation of TIMP-1 on α -SMA, and on epithelial-mesenchymal transition (EMT)-related genes. We also studied the underlying mechanisms. **Results:** We detected significantly higher levels of TIMP-1 and α -smooth muscle actin (α -SMA) in the fibroblasts isolated from urethral scar tissues, compared to the fibroblasts isolated from normal urethra. Moreover, the levels of TIMP-1 and α -SMA strongly correlated. Moreover, we found that TIMP-1 significantly increased levels of α -SMA, transforming growth factor β 1 (TGF β 1), Collagen I and some other key factors related to an enhanced EMT, suggesting that TIMP-1 may induce transformation of fibroblasts into myofibroblasts to promote tissue EMT to enhance the formation of urethral scar. Moreover, increases in TIMP-1 also induced an increase in fibroblast cell growth and cell invasion, in an ERK/MAPK-signaling-dependent manner. **Conclusion:** Our study thus highlights a pivotal role of TIMP-1 in urethral scar formation.

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Introduction

Urethral stricture is a fibrotic process in incompliant scarred tissue, resulting in constricted urethral lumen [1-4]. The common causes for developing urethral stricture are indirect or direct trauma-like side effects, such as surgical correction of urinary incontinence [1-4]. Although urethrotomy under direct vision could be an effective therapy for minor urethral stricture in the penile segment, the recurrence rate is pretty high [1-4]. Scar tissue for larger strictures is usually treated with excision by anastomotic urethroplasty, free buccal mucosal transplants or foreskin pedicled grafts [1-4].

The molecular basis of urethral stricture involves activation of tissue fibroblasts, their transformation into myofibroblasts, following outgrowth and increased deposition of collagens, and consequent changes in extracellular matrix components [5-8]. The matrix metalloproteinase (MMP) family members are involved in the breakdown of extracellular matrix during embryonic development, tissue remodeling and disease processes [9-13].

Tissue inhibitor of metalloproteinases 1 (TIMP-1) is a glycoprotein ubiquitously expressed in various human cells and tissues [14-16]. TIMP-1 controls the activity of MMPs and appears to an important regulator of extracellular matrix turnover [14-16]. Moreover, TIMP-1 also regulates cellular processes including cell growth, apoptosis, and differentiation that are independent of its metalloproteinase inhibitory activity [14-16]. Recently, TIMP-1 has been reported to upregulate in urethral scar tissues, which has been suggested to be responsible for decreased collagenase activities [17-20]. However, little is known about the effects of TIMP-1 on fibroblasts and the exact downstream mechanisms of TIMP-1-mediated cell signaling.

Here, we found significantly higher levels of TIMP-1 and α -smooth muscle actin (α -SMA) in the fibroblasts isolated from urethral scar tissues, compared to the fibroblasts isolated from normal urethra. Moreover, the levels of TIMP-1 and α -SMA strongly correlated. Then we either overexpressed TIMP-1, or inhibited TIMP-1 by lentiviruses carrying a transgene or a short hairpin small interfering RNA for TIMP-1 in human fibroblasts. We found that TIMP-1 significantly increased levels of α -SMA, transforming growth factor β 1 (TGF β 1), Collagen I and some other key factors related to an enhanced epithelial-mesenchymal transition (EMT), suggesting that TIMP-1 may induce transformation of fibroblasts into myofibroblasts to promote tissue EMT to enhance the formation of urethral scar. Moreover, increases in TIMP-1 also induced an increase in fibroblast cell growth and cell invasion, in an ERK/MAPK-signaling-dependent manner.

Materials and Methods

Patient tissue specimens

A total of 21 resected urethral scar tissue specimens from patients were collected for this study. The normal urethral tissues from the same patient were used as controls. All specimens had been histologically and clinically diagnosed at the Department of Urology, Shanghai Sixth People's Hospital of Shanghai Jiaotong University from 2010 to 2013. For the use of these clinical materials for research purposes, prior patient's consents and approval were collected, and the approvals by the Institutional Research Ethics Committee were obtained.

Isolation of fibroblasts from urethral scar tissues

A six-well plate was coated with a 0.1% gelatin solution and then incubated for 20 min. The dissected tissue parts from urethral scar tissue or control normal urethral tissue were put in separated large petri dishes and a scalpel was used to cut pieces as small as possible. Hereafter, the pieces were carried over to a new 10 ml tube, and 2.5 ml 30mg/ml collagenase (Sigma-Aldrich, St. Louis, MO, USA) was added to it to incubate for 60 min. at 37 °C. And 5 ml 0.25% trypsin (Sigma-Aldrich) was added in the last 10 min. The tubes were centrifuged for 5 min. at 300 g. The gelatin was removed from the dish, and the volume was distributed equally over 2 coated wells and placed in the incubator for 2 days. Passaging of the cells

was done when the culture plate grown confluent. A small plane of 0.25% trypsin was added to the cells after removing the medium and placed in the incubator at 37 °C for 10 minutes. Now the loosen cells were put in a tube. The culture plate was washed with PBS and added to the cells. After centrifugation for 5 minutes at 1000 RPM, cells were resuspended and seeded at 1:3 dilution on a new coated plate which surface is now approximately 3 times larger. New culture medium was added and the cells were placed in the incubator. The culture media was Dulbecco's modified Eagle's media (DMEM, Invitrogen, Carlsbad, CA, USA) supplemented with 15% fetal bovine serum (Invitrogen). PD98059, LY294002 and IWP-2 were both purchased from Sigma-Aldrich.

Transduction of human fibroblasts

The coding sequence of TIMP-1 was amplified using human cDNA as a template, and cloned into pLVX-ZsGreen1-C1 vector (Clontech, Mountain View, CA, USA). Short hairpin interfering RNAs (shRNAs) targeting TIMP-1 used the published sequences [21]. The shRNA was also cloned into pLVX-ZsGreen1-C1 vector, with pLVX-ZsGreen1-C1 itself as the mock control. To generate TIMP-1 or shTIMP-1 lentiviral particles, NIH HEK293T cells were seeded in a 100 mm dish at 50,000 cells/cm² and co-transfected with 10 µg of recombinant DNA plasmids and 5 µg each of packaging plasmids (REV, pMDL and VSV-G) using Lipofectamine-2000 (Invitrogen). The supernatant containing lentiviral particles was collected 48 hours after transfection and filtered through a 0.45 µm syringe filter. The fibroblasts were seeded in 100 mm plates at 15,000 cells/cm² one day prior to lentiviral infection. The lentiviral particles were added along with 10 µg/ml polybrene (Sigma-Aldrich) to the cell culture for 24 hours. Then the cells were washed twice with complete media and purified for the transduced cells, based on expression of GFP.

Transwell migration assay

The transwell migration assay was performed using a Fluorometric Cell Migration Assay kit with polycarbonate membrane inserts (5-µm pore size; Cell Biolabs, San Diego, CA, USA). Cells (4×10^4) were serum-starved overnight in DMEM prior to initiation of the experiment, and added to the upper chamber. Cells were then incubated at 37 °C for 24 hours to allow cell migration through the membrane to the lower chamber. The cells inside the upper chamber were then removed with cotton swabs. Migratory and invasive cells on the lower membrane surface were fixed, stained with hematoxylin, and counted for 10 random 100X fields per well. Cell counts are expressed as the mean number of cells per field of view. Five independent experiments were performed and the data are presented as mean ± standard deviation (SD).

Scratch wound healing assay

Scratch wound healing assay was performed as has been described previously [22]. Cells were seeded in 24-well plates at a density of 10^4 cells/well in complete DMEM and cultured to confluence. The cell monolayer was serum starved overnight in DMEM prior to initiating of the experiment. Confluent cell monolayer were then scraped with a yellow pipette tip to generate scratch wounds and washed twice with media to remove cell debris. Cells were incubated at 37 °C for 24 hours. Time lapse images were captured after 12 hours using a Nikon Eclipse TE2000-5 microscope. Images were captured from five randomly selected fields in each sample, and the wound areas are calculated by NIH ImageJ software.

Cell proliferation assay

For assay of cell proliferation, the cells were seeded into 96 well-plate at 4000 cells per well and subjected to a Cell Proliferation Kit (MTT, Roche, USA), according to the manufacturer's instruction.

RNA extraction, reverse transcription and quantitative RT-PCR

Total RNA was extracted from the cultured cells, or tissue, or patient specimens using Trizol (Invitrogen), according to the manufacturer's instruction. For mRNA analysis, complementary DNA (cDNA) was randomly primed from 2.0µg of total RNA using the Omniscript reverse transcription kit (Qiagen, Hilden, Germany). Quantitative Real-time PCR (RT-qPCR) was subsequently performed in duplicate with a 1:4 dilution of cDNA using the Quantitect SyBr green PCR system (Qiagen) on a Rotorgene 6000 series PCR machine. Data were collected and analyzed using the Rotorgene software accompanying the PCR machine. Relative expression levels were determined using the comparative quantification feature of the Rotorgene software. Levels of gene transcripts were normalized to α -tubulin, and then compared to controls.

Western blot and ELISA

Protein was extracted from the cultured cells or conditioned media by RIPA buffer (Sigma-Aldrich) for Western Blot. The supernatants were collected after centrifugation at 12000×g at 4°C for 20min. Protein concentration was determined using BCA protein assay, and whole lysates were mixed with 4×SDS loading buffer (125 mmol/l Tris-HCl, 4% SDS, 20% glycerol, 100 mmol/l DTT, and 0.2% bromophenol blue) at a ratio of 1:3. Samples were heated at 100 °C for 5 min and were separated on SDS-polyacrylamide gels. The separated proteins were then transferred to a PVDF membrane. The membrane blots were first probed with a primary antibody. After incubation with horseradish peroxidase-conjugated second antibody, autoradiograms were prepared using the enhanced chemiluminescent system to visualize the protein antigen. The signals were recorded using X-ray film. Primary antibodies for Western Blot are anti-TIMP-1, anti- α -SMA, anti-Collagen I and anti- α -tubulin (all purchased from Cell Signaling, San Jose, CA, USA). Secondary antibody is HRP-conjugated anti-rabbit (Jackson Labs, Bar Harbor, ME, USA). Images shown in the figure were representative from 5 repeats. Densitometry of Western blots was quantified with NIH ImageJ software. ELISA for TIMP-1 in the conditioned media was performed using a commercial ELISA kit (R&D Systems, Minneapolis, MN, USA), according to the manufacturer's instruction.

Statistical analysis

All statistical analyses were carried out using the SPSS 19.0 statistical software package. All data were statistically analyzed using one-way ANOVA with a Bonferoni Correction. Bivariate correlation was calculated by Spearman's Rank Correlation Coefficients. All values are depicted as mean \pm standard deviation from 5 individuals and are considered significant if $p < 0.05$. The figures were generated in Prism 6.0 (GraphPad Software Inc, USA).

Results

Correlated increases in TIMP-1 and α -SMA levels in urethral scar tissues

In order to figure out whether TIMP-1 may induce the transformation of fibroblasts into myofibroblasts to promote EMT, we examined the transcript levels of TIMP-1 and α -SMA, a marker for myofibroblasts and the levels of EMT in tissue, in the resected urethral scar tissue specimens (US), compared to the normal urethral tissues (NU) from the same patient. We also analyzed the correlation between TIMP-1 and α -SMA levels in individual patients. We detected significantly higher levels of TIMP-1 (about 8 fold increase, Fig. 1A), and significantly higher levels of α -SMA (about 6 fold increase, Fig. 1A) in US, compared to NU. In addition, a strong correlation between TIMP-1 and α -SMA levels was detected (Fig.

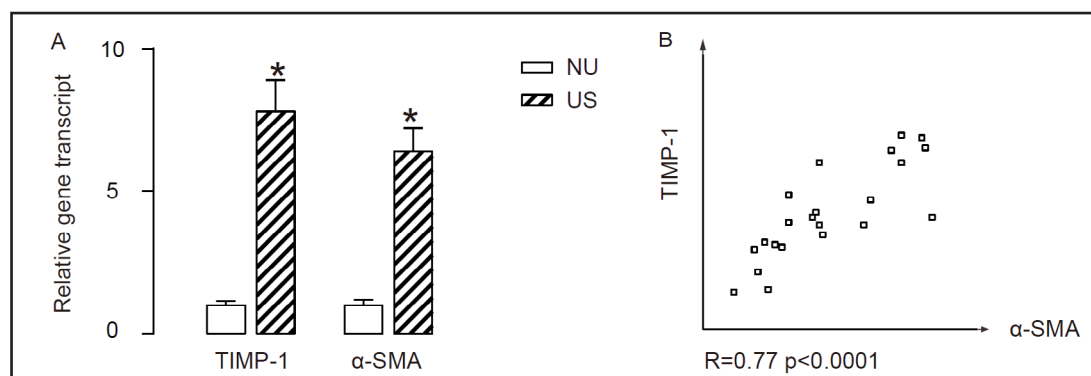


Fig. 1. Correlated increases in TIMP-1 and α -SMA levels in urethral scar tissues. The transcript levels of TIMP-1 and α -SMA in the resected urethral scar tissue specimens (US) were compared to the normal urethral tissues (NU) from the same patient. (A) Significantly higher levels of TIMP-1 (about 8 fold increase) and α -SMA (about 6 fold increase) were detected in US, compared to NU. (B) A strong correlation between TIMP-1 and α -SMA levels was detected ($R=0.77$, $p<0.0001$). * $p<0.05$. $n=21$. Statistics: one-way ANOVA with a Bonferoni Correction. Bivariate correlation was calculated by Spearman's Rank Correlation Coefficients.

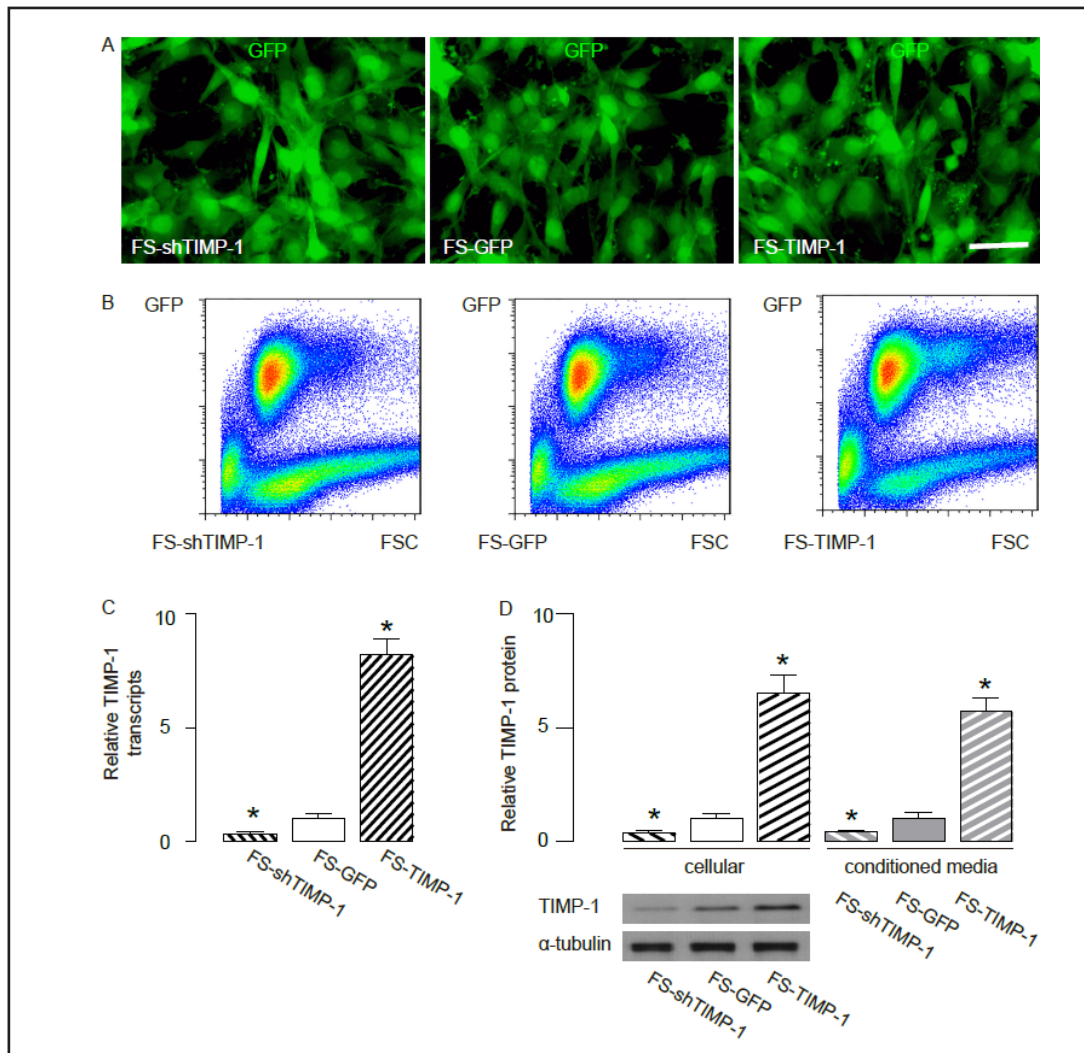


Fig. 2. Preparation of TIMP-1-modified human fibroblasts from urethral scar tissues. (A-B) Human fibroblasts (FS) were isolated from urethral scar tissues and transduced with either a TIMP-1 expressing lentivirus (FS-TIMP-1), or a lentivirus carrying shRNA for TIMP-1 (FS-shTIMP-1), or a control lentivirus (FS-GFP). All viruses contained GFP as a reporter, to allow transduced cells to be visualized in culture (A), and to be purified by flow cytometry (B). (C-D) The modifications of TIMP-1 levels in these cells were confirmed by RT-qPCR on TIMP-1 transcripts (C), by Western blot on cell extracts and by ELISA on conditioned media (D). FSC: forward scatter. * $p < 0.05$. $N = 5$. Representative images are randomly selected from 5 repeats. Statistics: one-way ANOVA with a Bonferroni Correction. Scale bar is $20\mu\text{m}$.

1B, $r = 0.77$, $p < 0.0001$). These data suggest a causal link between TIMP-1 and α -SMA in the formation of urethral scar.

Preparation of Timp-1-modified human fibroblasts from urethral scar tissues

Then we isolated human fibroblasts (FS) from urethral scar tissues. To examine whether TIMP-1 directly regulates α -SMA and other events of EMT, we transduced FS with either a TIMP-1 expressing lentivirus, or a lentivirus carrying small short hairpin interfering RNA for TIMP-1, or a control lentivirus. All viruses contained GFP as a reporter, to allow transduced cells to be visualized in culture (Fig. 2A), and to be purified by flow cytometry (Fig. 2B). The purified cells were termed as FS-TIMP-1, FS-GFP and FS-shTIMP-1, respectively. The modifications of TIMP-1 levels in these cells were confirmed by RT-qPCR (Fig. 2C), and by Western blot on the cell extracts (Fig. 2D), and by ELISA on the conditioned media (Fig. 2D).

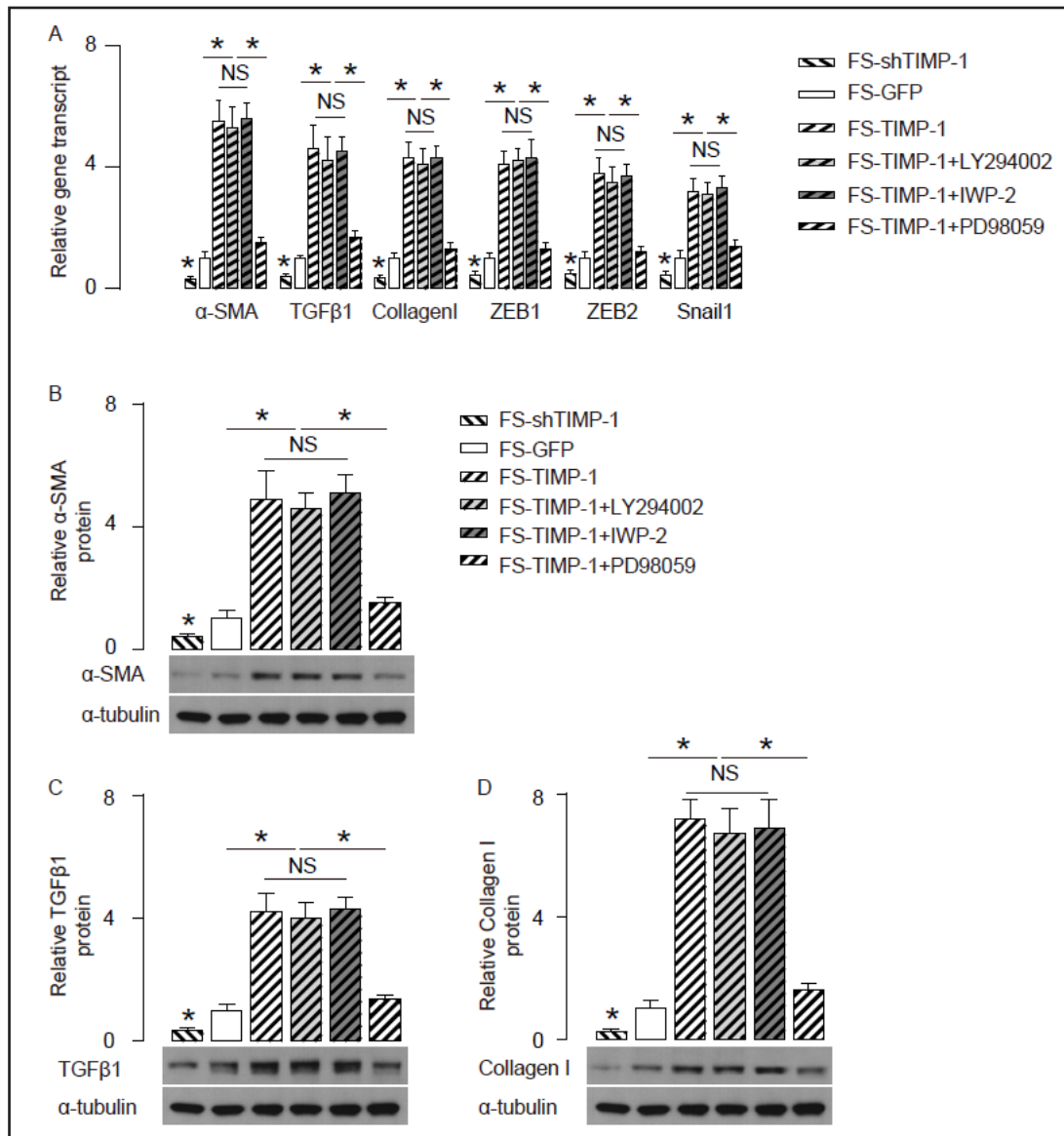


Fig. 3. TIMP-1 induced transformation of fibroblasts into myfibroblasts. (A) RT-qPCR for α -SMA, TGF β 1, Collagen I, ZEB1, ZEB2 and Snail1 in Timp-1-modified fibroblasts (cell extracts). (B-D) Western blot on cell extracts for α -SMA (B), TGF β 1 (C) and Collagen I (D) in TIMP-1-modified fibroblasts. LY294002 (20 μ mol/l), IWP-2 (1 μ mol/l), and PD98059 (10 μ mol/l) were added to cell culture to inhibit respective signaling pathways. * p <0.05. NS: non-significant. N=5. Representative images are randomly selected from 5 repeats. Statistics: one-way ANOVA with a Bonferoni Correction.

TIMP-1 induced transformation of fibroblasts into myfibroblasts

We found that overexpression of TIMP-1 in FS significantly increased α -SMA and TGF β 1 levels, while inhibition of TIMP-1 in FS significantly decreased α -SMA and TGF β 1 levels, by RT-qPCR (Fig. 3A), and by Western blot (Fig. 3B-C). Since both α -SMA and TGF β 1 are markers for myfibroblasts that trigger EMT, we then examined the levels of Collagen I, a key player in urethral scar formation, and the levels of some EMT-related genes. We found that overexpression of TIMP-1 in FS significantly increased Collagen I levels, while inhibition of TIMP-1 in FS significantly decreased Collagen I levels, by RT-qPCR (Fig. 3A), and by Western blot (Fig. 3D). Moreover, overexpression of TIMP-1 in FS also significantly increased EMT-related genes ZEB1, ZEB2 and Snail1, while inhibition of TIMP-1 in FS significantly decreased

Fig. 4. TIMP-1 increased fibroblast growth. Cell growth was examined in a MTT assay, and the values on O.D. at 570nm were shown. LY294002 (20 μ mol/l), IWP-2 (1 μ mol/l), and PD98059 (10 μ mol/l) were added to cell culture to inhibit respective signaling pathways. * p <0.05. NS: non-significant. N=5. Statistics: one-way ANOVA with a Bonferoni Correction.

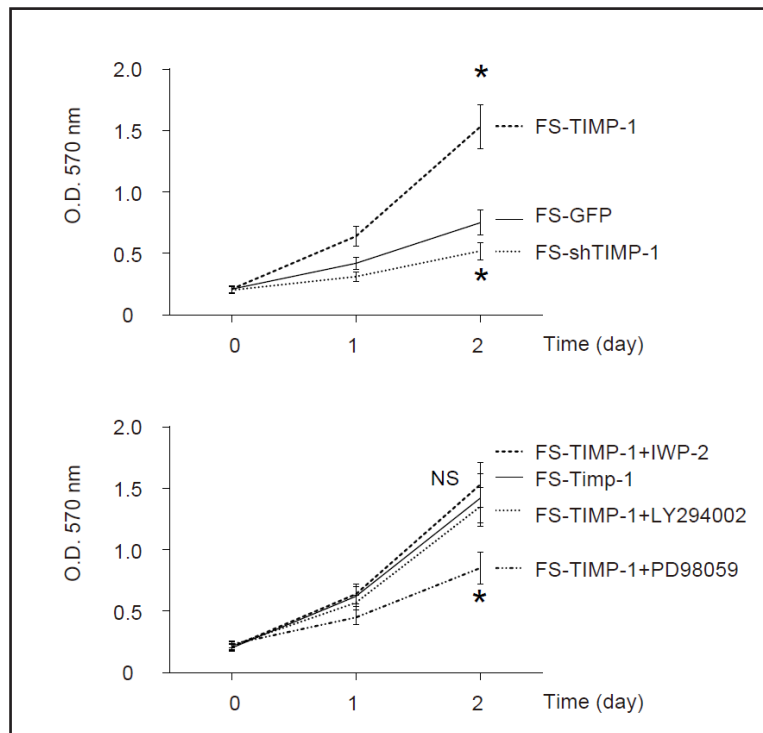
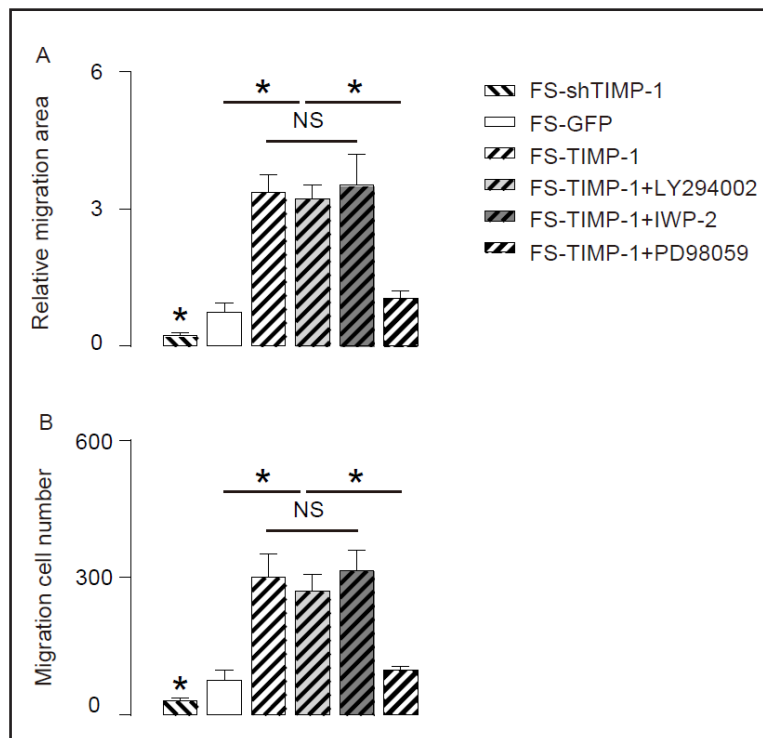


Fig. 5. TIMP-1 increased fibroblast invasiveness. (A-B) Overexpression of TIMP-1 in FS significantly increased cell migration and invasion, while inhibition of TIMP-1 in FS significantly decreased cell migration and invasion, in either a scratch wound healing assay (A), or a transwell migration assay (B). LY294002 (20 μ mol/l), IWP-2 (1 μ mol/l), and PD98059 (10 μ mol/l) were added to cell culture to inhibit respective signaling pathways. * p <0.05. NS: non-significant. N=5. Statistics: one-way ANOVA with a Bonferoni Correction.



these genes (Fig. 3A). Together, these data suggest that TIMP-1 may induce transformation of fibroblasts into myofibroblasts to promote processes of EMT.

TIMP-1 increased fibroblast growth

Moreover, we found that overexpression of TIMP-1 in FS significantly increased cell growth, while inhibition of TIMP-1 in FS significantly decreased cell growth, in a MTT assay

(Fig. 4, upper panel), suggesting that TIMP-1-induced transformation of fibroblasts may also promote fibroblast outgrowth.

TIMP-1 increased fibroblast invasiveness

In either a scratch wound healing assay (Fig. 5A), or a transwell migration assay (Fig. 5B), we found that overexpression of TIMP-1 in FS significantly increased cell migration and invasion, while inhibition of TIMP-1 in FS significantly decreased cell migration and invasion. These data suggest that TIMP-1-induced transformation of fibroblasts may also increase fibroblast invasiveness, which may eventually contribute to the formation of urethral scar.

TIMP-1 stimulated myofibroblast differentiation through ERK/MAPK signaling

Since PI3k/Akt, Wnt/ β -catenin and ERK/MAPK pathways may be activated by TIMP-1 to transduce downstream intracellular signaling, we used either 20 μ mol/l LY294002, a specific inhibitor to Akt, to inhibit PI3k/Akt signaling cascades in FS-TIMP-1 cells, or 1 μ mol/l IWP-2, a specific inhibitor to β -catenin, to inhibit Wnt/ β -catenin signaling cascades in FS-TIMP-1 cells, or 10 μ mol/l PD98059, a specific inhibitor to ERK1/2, to inhibit ERK/MAPK signaling cascades in FS-TIMP-1 cells. We found that only application of PD98059 significantly abolished the TIMP-1-mediated induction of EMT-related factors (Fig. 3A-D), the increases in cell growth (Fig. 4, lower panel), and the increases in cell invasiveness (Fig. 5A-B). These data suggest that TIMP-1 stimulates myofibroblast differentiation through ERK/MAPK signaling.

Discussion

Urethral stricture is a disease of reduced lumen in a section of the urethra, often concomitant with decreased tissue flexibility due to severe fibrosis and collagen deposition, resulting in decreased urine flow or even retention [1-4]. Surgery is not sufficient to provide satisfactory treatments for urethral stricture [1-4]. Thus efficient therapeutic interventions, e.g. inhibition of fibrosis, or decrease of collagen deposition, or increase of collagenolysis, are highly needed [5-8].

Although TIMP-1 has been recently recognized to play a role in the urethral scar formation, it is mainly thought to inhibit collagenase activity [5-8, 23]. The effect of TIMP-1 on fibroblast transformation, however, has not been studied.

Here, we addressed to this important question. We found that the TIMP-1 directly regulated transformation of fibroblasts into myofibroblasts, evident from the examination on α -SMA and TGF β 1 levels. α -SMA is specific marker for myofibroblasts, and is not expressed by normal fibroblast [24-27]. The EMT is a process by which epithelial cells acquire a migratory, mesenchymal phenotype, as a result of its repression of E-cadherin [28-30]. Several signaling pathways (TGF β , FGF, EGF, HGF, Wnt/ β -catenin and Notch) regulate EMT, among which TGF β receptor signaling triggered by TGF β 1 has been shown to be the most important one and extensively studied [31-42]. Loss of E-cadherin is considered to be a fundamental event in EMT [28-30]. Transcription factors Snail1, Slug1, ZEB1 and ZEB2 directly bind to E-cadherin promoter and repress its transcription [28-30]. Here, TIMP-1-induced TGF β 1 may bind to TGF β receptor to activate downstream SMAD proteins, and then subsequently activate EMT-related factors ZEB1, ZEB2 and Snail1, as has been previously reported [33, 43, 44]. Of note, we detected activation of all these factors by TIMP-1 overexpression, except for Snail2, which may be due to a cell/tissue-specific manner.

The exact causal relationship between fibroblast transformation and EMT was not further examined in the current study, since it is a well-established model [39, 45-51]. However, further proof of this regulatory axis is needed in follow-up studies. Together, our data highly suggest that in the current model, the EMT and fibrosis should be resulted from the transformation of fibroblasts into myofibroblasts. A set of loss-of-function and gain-of-function approaches, with the help of gene-knockout and gene-overexpression, should be taken to elucidate the exact signal transduction that regulates the whole process.

Our study illustrates a novel molecular mechanism that underlies the regulation of fibrosis and scar formation for urethral stricture, and suggests that besides affecting collagenase activity, TIMP-1 may also promote transformation of fibroblasts into myofibroblasts to enhance an EMT-mediated fibrosis and scar formation. These findings may provide novel insights upon the therapeutic approaches for patients with urethral stricture.

Disclosure Statement

The authors have declared that no competing interests exist.

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