Regulation of transforming growth factor β-mediated epithelial-mesenchymal transition of lens epithelial cells by c-Src kinase under high glucose conditions

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Abstract. Recent studies have reported that high glucose (HG) conditions may contribute to the acceleration of renal cell apoptosis and renal fibrosis by inducing epithelial-mesenchymal transition (EMT) of tubular epithelial cells, in which c-Src kinase and transforming growth factor (TGF)-β are key modulators. In the present study, the roles of c-Src kinase and TGF-β in EMT of lens epithelial cells (LECs) under HG conditions were investigated. Results indicated human lens epithelial B3 (HLE-B3) cells under HG conditions exhibited significantly increased protein expression levels of phosphorylated c-Src (p-Src⁴⁴⁶) (P<0.05) and secreted a significantly increased amount of TGF-β compared with HLE-B3 cells under normal glucose conditions (P<0.05). Notably the c-Src inhibitor PP1 and the activin receptor-like kinase 5 (ALK5) inhibitor SB431542 suppressed EMT of HLE-B3 cells. Results indicated that PP1 significantly inhibited the activities of c-Src and ALK5 and the secretion of TGF-β, whereas SB431542 only significantly downregulated the protein expression levels and secretion of TGF-β (P<0.05). Following c-Src knockdown, the protein expression levels of p-Src⁴⁴⁶, ALK5 and TGF-β were significantly decreased, the secretion of TGF-β was significantly suppressed (both P<0.05) and EMT was decreased in HLE-B3 cells. These results suggest that c-Src and TGF-β may promote EMT of LECs under HG conditions, with c-Src as the upstream regulatory molecule. Thus, the signal axis of c-Src/TGF-β in EMT of LECs may be a potential novel therapeutic target for the prevention of diabetic subcapsular cataract.

Introduction

Patients with diabetes mellitus typically develop irreversible renal and ocular tissue impairments due to the high levels of blood glucose (1,2). Diabetic cataract is one of the major ocular complications of diabetes mellitus and subcapsular cataract is considered the distinctive phenotype (3). Notably, a previous study demonstrated that human lens epithelial cells (LECs) are involved in modifications of proliferation and apoptosis in diabetic subcapsular cataract in an aldose reductase 2 (ALR2) transgenic mouse model (4). It has been reported that a number of aberrant nucleated cells assemble underneath the posterior subcapsular region in experimentally diabetic mice (5). However, the exact mechanism in diabetic cataract formation remains unclear.

Under physiological conditions, LECs are located in the less proliferative center zone or the mitotically active germinative zone of the lens. These LECs regulate the majority of the homeostatic functions of the lens, including its growth and differentiation, repair of damage and maintenance of lens transparency (6). However, under pathological conditions, LECs may promote the emergence of various types of cataracts, including posterior capsule opacity (PCO), anterior subcapsular cataract (ASC) and diabetic cataract (7).

Transforming growth factor (TGF)-β is a pleiotropic growth factor that controls cell growth, proliferation and migration, morphological plasticity and epithelial-mesenchymal transition (EMT) in tumor cells and specific epithelial cells (8-11). Furthermore, TGF-β exerts its biological effects through two membrane-bound receptors, designated type II and activin receptor-like kinase 5 (ALK5) (6). In the TGF-β family, TGF-β1 and TGF-β2 are expressed abundantly in the human lens and ocular media (12), and are considered key factors that regulate LEC proliferation, migration and EMT in ASC and PCO (13,14). Pathological studies regarding posterior subcapsular cataract (PSC) have indicated that various abnormal nucleated cells and the extracellular matrix accumulate underneath the capsule, resulting in transformation of the epithelial cell phenotype into fibroblastic cells (13-15). These studies suggested EMT of LECs may be an important initial step in the development of PCO and ASC, in which TGF-β is a major inducing factor. Du et al (16) studied the mechanism of high glucose (HG)-induced lens opacity accompanied by lens fibrosis in mice. The study reported that the TGF-β2/phosphoinositide 3-kinase/protein kinase B axis served an important role in HG-induced EMT of ex vivo human LECs and the inhibition of TGF-β delayed the EMT of LECs. Kim et al (17) identified that aldose reductase (AR) and
TGF-β exerted vital actions in a rat diabetic cataract model. The study indicated that the inhibition of AR resulted in the decrease of mRNA expression levels of TGF-β and fibronectin (FN) in HG-induced ex vivo human LECs. In another study, Kim et al (18) reported that the expression levels of TGF-β and FN were increased in HG-induced ex vivo human LECs and the expression levels of TGF-β2, Smad2/3, p38 mitogen-activated protein kinase (MAPK) and FN were decreased when the inhibitor KIOM-79 was applied. The results suggest that TGF-β2/p38MAPK and TGF-β2/Smad2/3 signaling pathways may serve key roles in EMT of LECs.

Lu et al (19) reported that rat lenses cultured in HG conditions exhibited increased expression levels of TGF-β, E-cadherin and α-smooth muscle actin (SMA). These results suggest that TGF-β may be a key molecule in EMT of HG-induced LECs and diabetic cataract. However, it remains unclear whether EMT of LECs is involved in PSC. c-Src kinase, converting into p-Src418 when activated, is a non-receptor intracellular tyrosine kinase that mediates a variety of cellular responses (20). c-Src may be activated by oxidative stress, osmotic stress, ultraviolet radiation and inflammation, which are associated with cataract formation (21-23). In various types of tumor cells, activated c-Src serves a principal role in regulating EMT properties, cell adhesion, proliferation and migration (24-27) and tumor metastasis (28,29). Notably, in the ocular media, activated c-Src is considered a predominant factor in the formation of cortical cataracts (30,31). Previously, it was revealed that activated c-Src mediated the EMT of LECs in normal culture medium, resulting in the occurrence of POC in vitro (32).

In the present study, HLE-B3 cells were employed in vitro to investigate the roles of c-Src and TGF-β in the EMT of LECs. Furthermore, the association between c-Src and TGF-β was assessed.

Materials and methods

Cell culture, reagents and treatments. Immortalized human lens epithelial cells (HLE-B3) cells were obtained from Dr Zhang Wei at the Beijing Institute of Ophthalmology (Beijing, China). HLE-B3 cells were maintained in Dulbecco's modified Eagle medium (DMEM) supplemented with 10% fetal bovine serum (HyClone; GE Healthcare Life Sciences, Logan, Utah, USA), penicillin (100 units/ml) and streptomycin (100 µg/ml). All cells were maintained at 37˚C in a humidified incubator containing 5% CO2. Cells were treated with 0.25% trypsin-0.02% EDTA to detach cells and passaged at 75‑80% confluence.

Stock solutions of PPI, a c-Src inhibitor, and SB431542, an ALK5 inhibitor, (Enzo Life Sciences, Inc., Farmingdale, NY, USA) were prepared in dimethyl sulfoxide and diluted with cell culture media to obtain a final concentration of 10 µmol/l. Antibodies against c-Src (cat. no. 2123), E-cadherin (cat. no. 3195), β-tubulin (cat. no. 2128) and β-actin (cat. no. 4970) were obtained from Cell Signaling Technology, Inc. (Danvers, MA, USA). Antibodies against TGF-β1 (cat. no. ab922486), p-Src418 (cat. no. ab4816), α-SMA (cat. no. ab5694), ALK5 (cat. no. ab31013) and GAPDH (cat. no. ab181602) were obtained from Abcam (Cambridge, MA, USA). Anti-TGF-β2 antibody (cat. no. ABE586) was obtained from EMD Millipore (Bedford, MA, USA).

A preliminary study was performed to optimize the time and glucose concentration for subsequent experiments. HLE-B3 cells were subsequently divided into three groups: HG DMEM (glucose, 35.5 mM), HG DMEM with PPI (10 µmol/l) and HG DMEM with SB431542 (10 µmol/l). The glucose concentration for the normal glucose (NG) group was 5.5 mM. The time-dependent effects were measured at 0, 3, 6 and 12 h for further experiments. The HLE-B3 cells in exponential growth phase were seeded at a cell density of 1x10^5 cells/ml, cultured at 37˚C, 5% CO2 and then assessed at 0, 3 and 6 h.

Short hairpin (sh)RNA knockdown assay. shRNA for human c-Src (GenBank no. NM: 005417) and random control lentiviral plasmid (pLKO.1-puro Control Vector) were obtained from GenePharma (GenePharma Co. Ltd, Beijing, China). The human HLE-B3 cells were cultured in the 60 mm diameter culture dish under 37˚C, 5% CO2 condition. When cells growth covered 50-60% of the dish, the cells were transfected with 5 µg shRNA-c-Src or 5 µg shRNA-vector using Lipofectamine 2000 transfection reagent (Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA) according to the manufacturer's recommendations. Following 72 h of incubation, transfection efficiency was measured by determining the green fluorescent protein expression ratio via fluorescence microscopy (Olympus BX53; Olympus Corporation, Tokyo, Japan; data not shown).

In c-Src knockdown experiments, 5x10^5 cells/ml HLE-B3 cells were divided into two groups: shRNA-vector and shRNA-c-Src. Cells were cultured in HG DMEM (35.5 mM) for a total of 12 h, and the time-dependent effects were monitored at 0, 3, 6, 9 and 12 h. The knockdown of c-Src was confirmed via western blot analysis.

Western blot analysis. Total protein in HLE-B3 cells was extracted using radioimmunoprecipitation assay lysis buffer with protease inhibitor cocktail set 1 (cat. no. 539131) and phosphatase inhibitor cocktail set V (cat. no. 524629; all EMD Millipore). Protein concentration was determined using BCA phosphatase inhibitor cocktail set V (cat. no. 524629; all EMD Millipore). Protein concentration was determined using BCA protein assay reagent (Beyotime Institute of Biotechnology, Shanghai, China). Proteins (30 µg/lane) were separated using 10% SDS-PAGE and electrophoretically transferred onto nitrocellulose filter membranes (EMD Millipore). Membranes were blocked in 4˚C overnight with 5% non-fat milk and incubated with various primary antibodies, including rabbit anti-c-Src monoclonal antibody (1:10,000), rabbit anti-p-Src418 monoclonal antibody (1:1,000), rabbit anti-TGF-β1 polyclonal antibody (1:100), rabbit anti-TGF-β2 polyclonal antibody (1:1,000), rabbit anti-E-cadherin polyclonal antibody (1:1,000), rabbit anti-α-SMA polyclonal antibody (1:1,000), rabbit anti-ALK5 polyclonal antibody (1:200), rabbit anti-β-actin monoclonal antibody (1:1,000), rabbit anti-β-tubulin monoclonal antibody (1:1,000) and rabbit anti-GAPDH monoclonal antibody (1:10,000) in 4˚C for 18 h. Following 2 h incubation at room temperature with horseradish peroxidase-conjugated secondary antibody (1:1,000; cat. no. 7074; Cell Signaling Technology, Inc.), the protein bands were quantified using a Bio-Rad chemiluminescent gel imaging system and standardized to β-actin, β-tubulin or GAPDH protein expression using Image Lab™ software (version 6.0; both Bio-Rad Laboratories, Inc., Hercules, CA, USA).
Determination of TGF-β1 and TGF-β2 secretion using ELISA. An ELISA kit (eBioscience; Thermo Fisher Scientific, Inc.) was employed to evaluate the secretion of TGF-β1 (cat. no. BMS249-4) and TGF-β2 (cat. no. BMS254) according to the manufacturer's protocol. Three samples of each group were examined.

Statistical analysis. All data were analyzed using SPSS statistical analysis software, version 16.0 (SPSS, Inc., Chicago, IL, USA). Data were expressed as the mean ± standard deviation. For multiple group comparisons at different time points, statistical analysis was performed using repeated measures analysis of variance followed by Fisher's Least Significant Difference post hoc test. For groups at the same time point, the Student's t-test was used. P<0.05 was considered to indicate a statistically significant difference.

Results

HG conditions induce EMT of LECs. To investigate the effect of HG conditions on LECs, the protein expression levels of EMT marker protein E-cadherin and interstitial cell marker protein α-SMA in HLE-B3 cells under HG and NG conditions were studied using western blot analysis. Results indicated that the protein expression levels of E-cadherin were marginally decreased whereas α-SMA protein expression levels were marginally increased under normal glucose (NG) conditions between 0 to 12 h. However, a time-dependent decrease in E-cadherin expression levels and a time-dependent increase in α-SMA expression levels was indicated when HLE-B3 cells were exposed to HG conditions. The differences were significant at 6 and 12 h compared with 0 h under the same conditions (P<0.05). Additionally, results indicated that the protein expression levels of E-cadherin were significantly decreased whereas α-SMA protein expression levels were significantly increased at 6 and 12 h in HLE-B3 cells under HG conditions compared with that in HLE-B3 cells under NG conditions at the same time points (P<0.05; Fig. 1). These results suggested that LECs may slowly trans-differentiate into mesenchymal cells under HG conditions.

HG conditions activate c-Src. To further investigate the influence of HG conditions on the activity of c-Src, the protein expression levels of p-Src418 were determined using western blot analysis under NG and HG conditions. The results demonstrated that there was little p-Src418 expression observed following 12 h incubation under NG conditions. By contrast, the protein expression levels of p-Src418 were elevated and reached a peak at 6 h of culture in HG conditions, which was followed by a decrease in p-Src418 expression at 12 h (Fig. 2). Statistical analysis indicated that the protein expression levels of p-Src418 were significantly increased under HG conditions at 3, 6 and 12 h compared with at 0 h under HG conditions (P<0.05). Furthermore, the p-Src418 protein expression levels of HLE-B3 cells at the corresponding time points in the HG conditions were significantly higher than those indicated under NG conditions (P<0.05), which suggested that the activation of c-Src was stimulated by HG conditions.

HG conditions activate TGF-β. The influence of HG conditions on the activation of TGF-β was observed using western blot analysis. There were no notable changes in the protein expression levels of TGF-β1 and TGF-β2 under NG conditions. However, under HG conditions the expression levels of TGF-β1 and TGF-β2 were marginally increased over 12 h (Fig. 3A and B). These findings suggested that HG conditions exerted little impact on the protein expression levels of TGF-β1 and TGF-β2.

The secretion of TGF-β1 and TGF-β2 was also measured under NG and HG conditions using ELISA. Results revealed that the secretion of TGF-β1 and TGF-β2 under NG conditions
was not significantly altered over 12 h. However, a significant increase in the secretion of TGF-β1 at 6 and 12 h and TGF-β2 at 3, 6 and 12 h was observed under HG conditions compared with 0 h under the same conditions (P<0.05; Fig. 3C and D). Furthermore, significantly increased secretion of TGF-β1 at 6 to 12 h and TGF-β2 at 3, 6 to 12 h was indicated under HG conditions compared with those indicated under NG conditions at the same time points (P<0.05).

Effect of c-Src and TGF-β on EMT of LECs. To further investigate the regulatory effects of c-Src and TGF-β in EMT of LECs, SB431542 and PP1 were added to the HG media. PP1 was used to investigate the regulatory effect of c-Src on TGF-β, while SB431542 was used as the positive control. The protein expression levels of c-Src and TGF-β were measured using western blot analysis and TGF-β secretion was analyzed using ELISA. As indicated in Fig. 4, the protein expression levels of p-Srcα18, TGF-β1 and TGF-β2 in HLE-B3 cells under HG conditions and PP1 were significantly decreased compared with those indicated in HLE-B3 cells under HG conditions alone (P<0.05). Notably, the secretion of TGF-β1 and TGF-β2 was significantly decreased in HLE-B3 cells under HG conditions and PP1 compared with cells under HG conditions alone at 6 h (P<0.05; Fig. 5). These findings suggest that active c-Src promotes the activation of TGF-β. Under HG conditions, SB431542, which is an inhibitor of TGF-β, significantly downregulated the protein expression levels and secretion of TGF-β1 and TGF-β2 (P<0.05) but exerted no significant effect on p-Srcα18 expression levels compared with that in HLE-B3 cells under HG conditions alone (Figs. 4A-C and 5).

Effect of c-Src on the biological function of TGF-β in EMT of LECs. The influence of c-Src on the biological function of TGF-β was assessed by determining the activity of ALK5 using western blot analysis. The results revealed that the protein expression levels of ALK5 in HLE-B3 cells cultured under HG conditions with PP1 or SB431542 were significantly decreased compared with those in HLE-B3 cells under HG conditions alone at 6 h (P<0.05, Fig. 4D). This was consistent with the results obtained regarding TGF-β1 and TGF-β2 secretion in HLE-B3 cells cultured under HG conditions with SB431542. The downregulation of ALK5 suggested that c-Src may influence the biological function of TGF-β.

c-Src and TGF-β mediate EMT of LECs. The roles of c-Src and TGF-β in EMT of HLE-B3 cells induced by HG conditions were determined by measuring the alterations of EMT protein markers E-cadherin and α-SMA in HLE-B3 cells in the presence of PP1 or SB431542 using western blot analysis. The protein expression levels of E-cadherin in HLE-B3 cells under HG conditions were decreased in a time-dependent manner. However, HG conditions and PP1 or SB431542 increased the protein expression levels of E-cadherin in HLE-B3 cells compared with that in HLE-B3 cells cultured in HG conditions alone. This difference was statistically significant at 6 h (P<0.05; Fig. 4E). Notably, the protein expression levels of α-SMA in HLE-B3 cells under HG conditions were increased in a time-dependent manner. However, the protein expression levels of α-SMA were significantly decreased in HLE-B3 cells treated with PP1 or SB431542 under HG conditions compared with HLE-B3 cells under HG conditions alone at 6 h (P<0.05, Fig. 4F). These findings suggested that PP1 or SB431542 hindered the decrease of E-cadherin and the increase of α-SMA, which suggested the role of c-Src and TGF-β in promoting EMT of LECs.

Influence of c-Src knockdown on TGF-β and EMT of LECs. HLE-B3 cells transfected with shRNA-c-Src or shRNA-vector were exposed to HG conditions (35.5 mM). As indicated in Fig. 6, the expression levels of p-Srcα18 protein increased in a time-dependent manner up until 9 h, after which the levels were decreased in the shRNA-vector group. In the shRNA-c-Src group, the p-Srcα18 protein expression levels were significantly decreased compared with those indicated in the shRNA-vector group at the same time points (P<0.05; Fig. 6A). The decreased expression level of p-Srcα18 protein, instead of completely silenced p-Srcα18, suggested that the knockdown of c-Src may not be completely successful.

The protein expression levels of TGF-β1 and TGF-β2 in shRNA-c-Src or shRNA-vector groups stimulated by HG conditions were observed using western blot analysis. In the shRNA-c-Src group, the expression levels of TGF-β1 protein were increased in a time-dependent manner up until
9 h, after which the expression levels were decreased. In the shRNA-c-Src group, TGF-β1 protein expression levels significantly decrease compared with the shRNA-Vector group \( (P<0.05; \text{Fig. 6B}) \). Notably, TGF-β2 protein expression levels in the shRNA-c-Src and shRNA-vector groups were increased in a time-dependent manner up until 9 h after which the expression levels were decreased (Fig. 6A). However, the TGF-β2 protein expression levels were significantly increased in the shRNA-vector group compared with the shRNA-c-Src group at all time points \( (P<0.05) \). These findings suggested that suppression of c-Src significantly reduced the protein expression levels of TGF-β.

The secretion of HG-stimulated TGF-β1 and TGF-β2 proteins, were measured using ELISA in the shRNA-c-Src and shRNA-vector groups. In shRNA-c-Src and shRNA-vector groups, the secretion of TGF-β1 and TGF-β2 increased in a time-dependent manner without significant differences. Furthermore, the secretion of TGF-β1 and TGF-β2 was significantly decreased in the shRNA-c-Src group compared with that in the shRNA-vector group at all time points \( (P<0.05; \text{Fig. 6F and G}) \), which suggested that the suppression of c-Src effectively reduced the secretion of TGF-β.

**Suppression of c-Src inhibits the biological function of TGF-β in the EMT of LECs.** To evaluate whether c-Src activity may influence the biological function of TGF-β, the expression levels of ALK5 stimulated by HG conditions in transfected HLE-B3 cells were determined using western blot analysis. In the shRNA-vector and shRNA-c-Src groups, the protein expression levels of ALK5 were increased in a time-dependent manner. However, the protein expression levels of ALK5 were significantly decreased in the shRNA-c-Src group compared with the shRNA-vector group at 0 to 12 h \( (P<0.05; \text{Fig. 6C}) \). These results suggested that c-Src silencing inhibited the activity of ALK5, which is the downstream mediator of the TGF-β signaling pathway.

**Suppression of c-Src inhibits EMT of LECs.** The influence of c-Src knockdown on EMT of LECs induced by HG conditions was investigated by evaluating the expression of EMT protein markers using western blot analysis. In the shRNA-vector and shRNA-c-Src groups, the protein expression levels of E-cadherin were decreased in a time-dependent manner. However, the expression levels of E-cadherin in the shRNA-vector group were significantly decreased compared with that in the shRNA-c-Src group at all time points \( (P<0.05; \text{Fig. 6D}) \). Notably, the protein expression levels of α-SMA were increased in the shRNA-vector and shRNA-c-Src groups in a time-dependent manner. Additionally, the protein expression levels of α-SMA were significantly decreased in the shRNA-c-Src group compared with that in the shRNA-vector group at 0 to 12 h \( (P<0.05; \text{Fig. 6E}) \). These results indicated that knockdown of c-Src may effectively inhibit EMT of LECs induced by HG conditions.
EMT is a process by which epithelial cells lose their original characteristics and gain mesenchymal cell properties during specific physiological and pathological stages, including organ fibrosis (33,34). Previous studies on the EMT of renal cells cultured in HG conditions (30 or 60 mM) demonstrated significant increases of α-SMA and decreases of E-cadherin (33,34). It was also reported that EMT is the primary factor in the pathogenesis of renal fibrosis, in which TGF-β and c-Src are key inducing factors (35-38). Subcapsular cataract is a common type of diabetic cataract (39). Studies have revealed that a number of aberrant nucleated cells assemble underneath the posterior capsule in animal models of subcapsular cataract (13-15). However, at present, the exact mechanism and key trigger factors of diabetic subcapsular cataract remains unclear. In Rap39 transgenic animal models of diabetic cataract, the presence of α-SMA was confirmed in.

Discussion

EMT is a process by which epithelial cells lose their original characteristics and gain mesenchymal cell properties during specific physiological and pathological stages, including organ fibrosis (33,34). Previous studies on the EMT of renal cells cultured in HG conditions (30 or 60 mM) demonstrated significant increases of α-SMA and decreases of E-cadherin (33,34). It was also reported that EMT is the primary factor in the pathogenesis of renal fibrosis, in which TGF-β and c-Src are key inducing factors (35-38). Subcapsular cataract is a common type of diabetic cataract (39). Studies have revealed that a number of aberrant nucleated cells assemble underneath the posterior capsule in animal models of subcapsular cataract (13-15). However, at present, the exact mechanism and key trigger factors of diabetic subcapsular cataract remains unclear. In Rap39 transgenic animal models of diabetic cataract, the presence of α-SMA was confirmed in.
the muddy plaque underneath the anterior capsule of the lens in the early stage of cataract (4), suggesting that HG-induced EMT of LECs may be a key mechanism in animal models of diabetic cataract. The data obtained in the present study suggested a similar process in EMT of LECs under HG conditions (35.5 mM) as the protein expression levels of E-cadherin were decreased and the protein expression levels of α-SMA were increased over time.

The present study indicated that HG conditions contributed to the increased activity of TGF-β and promoted the phosphorylation of c-Src. Glomerular mesangial cells incubated with HG (30 mM) concentrations also demonstrated similar effects on the levels of p-Src (40, 41). Furthermore, it has been reported that exposure of A549 cells to HG (25 mM) concentrations for 48 h resulted in increased activity of TGF-β (42). This is in agreement with the present findings, which indicated that c-Src and TGF-β were activated during EMT of LECs under HG conditions.

As c-Src and TGF-β were concurrently activated in EMT of LECs, the roles of c-Src and TGF-β in the EMT were further investigated in the present study. The results revealed that PP1 and SB431542 delayed the EMT progression in LECs induced by HG stimulation. Previous studies also suggested that PP1 and SB431542 suppressed the activities of c-Src and TGF-β following HG treatment in different cell phenotypes (30, 31, 43, 44). In addition, it was reported that activated c-Src promoted EMT and c-Src-specific inhibitors blocked EMT in renal tubular epithelial cells (45). Furthermore, various studies have reported that TGF-β is a central mediator of EMT and the inhibition of the TGF-β signaling pathway with SB431542 prevented the progression of EMT (43, 44, 46). These studies support the present findings, which indicated that c-Src and TGF-β mediate the EMT of LECs.

The present study indicated that PP1 directly inhibited the activity of c-Src and TGF-β in HLE-B3 cells following HG stimulation. However, although SB431542 also significantly inhibited the activity of TGF-β, SB431542 had no significant effect on activity of c-Src. It is likely that PP1 may induce the indirect inhibition on TGF-β by the PP1-induced inhibition of c-Src activity. Previous studies have reported that PP1 and PP2 inhibited TGF-β-mediated cell invasion and migration in pancreatic ductal adenocarcinoma cells and non-small-cell lung carcinoma cells via directly targeting ALK5, suggesting that PP1 and PP2 may be dual inhibitors of TGF-β and c-Src in vitro (47, 48). This is consistent with the speculations from the present study. Furthermore, incubation with SB431542 resulted in a significant decrease in the protein expression level of TGF-β at 6 h, which suggests that SB431542 may affect the cell growth and the cell cycle and promote cell apoptosis of HLE-B3 cells (49, 50). The present study has demonstrated
that PP1 not only restrained the c-Src-induced EMT of LECs, but also downregulated TGF-β-mediated EMT of LECs. However, it is not known whether c-Src participates in the TGF-β-mediated EMT of LECs.

In order to further assess the association between c-Src and TGF-β and to confirm the inhibiting effect of PP1 on the intra-cellular function of TGF-β, shRNA was used to knockdown the c-Src gene in HLE-B3 cells. The results demonstrated that, when phosphorylation of c-Src was inhibited by c-Src gene knockdown, the activities of TGF-β and ALK5 were significantly inhibited compared with the shRNA-vector group. This suggests that c-Src may be an upstream molecule that regulates the bioactivity of TGF-β. In addition, the EMT of HLE-B3 cells was blocked following c-Src knockdown, which indicates that active c-Src may promote the EMT of LECs. It has been reported that the activation of c-Src promoted TGF-β-mediated EMT of glomerular endothelial cells, in which c-Src is the key regulatory molecule (51). However, studies with regards to c-Src and TGF-β have provided various results, including the suggestion that TGF-β may regulate the activity of c-Src in vascular smooth muscle cells and non-small-cell lung cancer A549 cells (52,53). Maeda et al (46) indicated that Src activation was not involved in the processing of TGF-β-mediated EMT in mammary epithelial cells and pancreatic ductal adenocarcinoma cells. These different results regarding c-Src and TGF-β in the regulation of EMT of various cells may be due to the differences in cell type and stimuli.

In conclusion, the findings of the present study suggest that c-Src may be an upstream molecule that regulates the TGF-β-mediated EMT of LECs under HG conditions. Therefore, the c-Src/TGF-β signal axis, particularly c-Src, may be a potential target for preventing and treating complications associated with diabetes mellitus, including diabetic subcapsular cataract. However, further in vitro investigations on the exact mechanism and signaling pathway of c-Src and TGF-β are required.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

JZ designed the experiments. ZHH and FW performed the experiments. FLW and QL analyzed and interpreted the results of the experiments. QL was a major contributor in writing the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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