

# Curcumin protects renal tubular epithelial cells from high glucose-induced epithelial-to-mesenchymal transition through Nrf2-mediated upregulation of heme oxygenase-1

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**Abstract.** Curcumin has been observed to exhibit an anti-fibrotic effect in the liver, lung and gallbladder. However, the mechanisms underlying the cytoprotective effects of curcumin remain to be elucidated. The epithelial-to-mesenchymal transition (EMT) of mature tubular epithelial cells in the kidney is considered to contribute to the renal accumulation of matrix proteins associated with diabetic nephropathy. The EMT is also closely associated with the progression of renal interstitial fibrosis and oxidative stress. This process may occur through abrogation of high glucose (HG)-induced oxidative stress via activation of nuclear factor (erythroid-derived 2)-like 2 (Nrf2) and heme oxygenase-1 (HO-1) in kidney tubular epithelial cells. In the present study, the effect of curcumin on HG-induced EMT in the NRK-52E normal rat kidney tubular epithelial cell line was investigated, and whether the effect of curcumin was mediated by the induction of Nrf2 and HO-1 expression was examined. The present study revealed that curcumin was able to prevent events associated with EMT, including the downregulation of E-cadherin and the increased expression of  $\alpha$ -smooth muscle

actin. Further analysis revealed that the expression levels of Nrf2 and HO-1 protein were elevated to a greater extent in the curcumin pretreated NRK-52E cells compared with those of the control. Notably, knockdown of Nrf2 with small interfering RNA prevented the curcumin-induced elevation in expression of HO-1 and the associated anti-fibrotic effects. In conclusion, the present findings suggested that curcumin may be significant in cellular antioxidant defense, through the activation of Nrf2 and HO-1, thereby protecting the NRK-52E cells from HG-induced EMT.

## Introduction

Diabetic nephropathy (DN) is the leading cause of chronic kidney failure and end-stage renal disease worldwide, and the prevalence of this disease has progressively increased (1,2). DN is characterized pathologically by the progressive accumulation of extracellular matrix (ECM) proteins in the basement membranes, glomerular mesangium and the peritubular interstitium. DN may eventually lead to kidney scarring and ultimately nephron dropout (3,4). Although glomerulosclerosis is a defining feature of DN, it is the extent of tubulointerstitial injury that fundamentally determines the rate of decline in renal function (5). Data have suggested that tubulointerstitial fibrosis also occurs at an early stage of diabetic renal injury and correlates closely with the decline in renal function observed in certain groups of patients (5-7). Accumulating evidence has implicated the epithelial-to-mesenchymal transition (EMT) of mature tubular epithelial cells in the kidney as a contributing factor to the renal accumulation of matrix proteins associated with DN. In addition, EMT is closely associated with the progression of renal interstitial fibrosis, which is characterized by a loss of the typical features of normal epithelial cells and a gain in the characteristics of ECM-producing myofibroblasts (8-10). Furthermore, blockade of certain stages involved in EMT significantly reduces the formation of fibrotic lesions in specific models of kidney fibrosis, suggesting that EMT may be significant in the development of nephropathy (11-13).

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**Abbreviations:** NRK-52E cells, normal rat kidney tubular epithelial cells; EMT, epithelial-to-mesenchymal transition; HG, high glucose; DMEM, Dulbecco's modified Eagle's medium; SMA, smooth muscle actin; RIPA, radioimmunoprecipitation assay; TGF, transforming growth factor; TBS, Tris-buffered saline; Nrf2, nuclear factor (erythroid-derived 2)-like 2; HO-1, heme oxygenase-1; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide

**Key words:** renal tubular epithelial cells, high glucose, curcumin, epithelial-to-mesenchymal transition, nuclear factor (erythroid-derived 2)-like 2

It has been hypothesized that oxidative stress may contribute to the development of diabetic renal complications, including the EMT of tubular epithelial cells, which are observed in renal tissues even during the early stages of diabetes (14-16). As a key feature of the intracellular antioxidant machinery, nuclear factor (erythroid-derived 2)-like 2 (Nrf2) dissociates from its cytosolic inhibitor Kelch-like erythropore concentrating hormone-associated protein 1 (Keap1), translocates to the nucleus and regulates the coordinated induction of a number of genes, which encode numerous antioxidant and phase II detoxifying enzymes (17-19). One important Nrf2 target gene, heme oxygenase-1 (HO-1), is considered to be significant in the degradation of pro-oxidant heme, which results in the production of anti-inflammatory, antioxidant and anti-apoptotic metabolites (20,21). The essential role of Nrf2 in combating oxidative stress has been demonstrated by investigations revealing the increased sensitivity of Nrf2<sup>-/-</sup> mice to various types of insult, including high glucose(HG)-induced oxidative damage (18,22). HO-1, which is induced by multiple transcription factors, including Nrf2, to protect the kidney from injury may aid in devising a therapeutic approach against the development of DN (23,24). Previous studies have revealed that the increased expression of HO-1 is able to attenuate cytokine- and glucose-mediated cell growth arrest and apoptosis *in vitro* and *in vivo* (25). In addition, HO-1 deficiency has been demonstrated to be associated with increased fibrosis, increased tubular transforming growth factor (TGF)- $\beta$ 1 expression, inflammation and enhanced EMT in obstructive kidney disease (26).

Curcumin (diferuloylmethane) is a commonly used flavoring and coloring agent, and is a major component of the yellow spice, turmeric, derived from the rhizomes of *Curcuma longa* (27). Curcumin exhibits a number of biological effects, including antioxidant, anti-inflammatory and wound-healing properties (28-31). In addition, previous studies have indicated that curcumin has anti-fibrotic effects in the liver and the lungs, providing relief from cystic fibrosis (32,33). In immortalized rat kidney interstitial fibroblasts, curcumin has been observed to attenuate TGF- $\beta$ 1-induced fibrosis through the downregulation of TGF- $\beta$  receptor II (34). In the unilateral ureteral obstruction (UUO) rat kidney fibrosis model, curcumin has been observed to inhibit inflammation and fibrosis of the renal interstitium by inhibition of the NF- $\kappa$ B-dependent signaling pathway (35). Furthermore, the antioxidant properties of curcumin have been observed to be effective in improving renal function in certain diabetic animal models (36,37), as well as in acute kidney failure induced by ischemia-reperfusion (38). However, it remains to be elucidated whether pretreatment with curcumin in tubular epithelial cells leads to an increase in the Nrf2 protein level and alleviates the EMT of tubular epithelial cells. Emerging evidence has suggested that curcumin induces HO-1 mRNA and protein expression in the proximal tubule cells through transcriptional mechanisms and may also involve the NF- $\kappa$ B pathway (39). Notably, Gaedeke *et al* (40) demonstrated that curcumin treatment in nephritic animal models decreased fibrosis by inducing the expression of Nrf2 and HO-1. Therefore, it was hypothesized that administration of curcumin may increase the cellular antioxidant defense capacity via activation of Nrf2 and HO-1 expression, thereby protecting NRK-52E cells from the effects of HG-induced EMT.

## Materials and methods

**Cell culture.** The NRK-52E normal rat kidney tubular epithelial cell line was purchased from the American Type Culture Collection (Manassas, VA, USA). The cells were cultured in a 5% CO<sub>2</sub> atmosphere in complete Dulbecco's modified Eagle's medium (DMEM; low glucose; Gibco Life Technologies, Grand Island, NY, USA), which contained 10% fetal bovine serum (Gibco Life Technologies), 4 mM L-glutamine (Boster Biological Technology Ltd., Wuahn, China) and 1% penicillin/streptomycin (Sigma-Aldrich, St. Louis, MO, USA) at a density of  $5 \times 10^3$  cells/well in six-well culture plates. Once the cells were almost confluent, they were transferred to serum-free DMEM for 24 h at 37°C to arrest and synchronize cell growth. In the control groups, the cells were treated with fresh serum-free DMEM only, which contained 5 mmol/l glucose. In the experimental groups, the cells were subjected to pretreatment with 5, 10 or 20  $\mu$ M curcumin (Santa Cruz Biotechnology, Inc., Dallas, TX, USA) and then cultured for 24 h at 37°C. Subsequently, the medium was changed and the cells were treated for an additional 48 h with 30 mM HG at 37°C (Boster Biological Technology Ltd.). The concentration of glucose was determined as previously described (12) and with reference to preliminary experiments by our group. In some experiments, the cells were treated with 0, 5, 10 or 20  $\mu$ M curcumin for 24 h, or 10  $\mu$ M curcumin for 0, 3, 6, 12, 24 or 48 h, and the expression levels of Nrf2 were detected by western blotting. Cells were also treated with 50  $\mu$ M tin protoporphyrin (SnPP), a known inhibitor of HO, in order to study changes to HO.

**Experimental groups.** The cells were divided into six groups, as follows: Control group, treated with serum-free DMEM; siRNA group, subjected to Nrf2-siRNA transfection; curcumin group, treated with 10  $\mu$ M curcumin for 24 h; HG group, treated with 30 mM HG for 48 h; HG/curcumin group, pretreated with 10  $\mu$ M curcumin for 24 h, followed by 30 mM HG treatment for 48 h; HG/curcumin/Nrf2-siRNA group, 24 h post-transfection the cells were treated with 10  $\mu$ M curcumin for 24 h and 30 mM HG for 48 h.

**Assessment of cell viability.** The cell viability was measured using a 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. Briefly, 10  $\mu$ l MTT (500  $\mu$ g/ml; Sigma-Aldrich) was added to the medium and the sample was incubated for 3 h at 37°C following treatment. Subsequently, the MTT solution was removed and 100  $\mu$ l dimethyl sulfoxide (Sigma-Aldrich) was added to the medium to dissolve the colored formazan crystals. The absorbance of each aliquot at 540 nm was measured using a Sunrise microplate reader (Tecan Group Ltd., Männedorf, Switzerland). The cell viability was determined as the ratio of the signal between the treated and control cultures.

**Western blot analysis.** The NRK-52E cells were pelleted by centrifugation at 125 x g at 4°C for 10 min and then washed once with phosphate-buffered saline. The cells were then lysed using a mixture of radioimmunoprecipitation assay buffer (Sigma-Aldrich) and phenylmethylsulfonyl fluoride (1:100; Sigma-Aldrich), on ice for 30 min with vortexing at intervals.

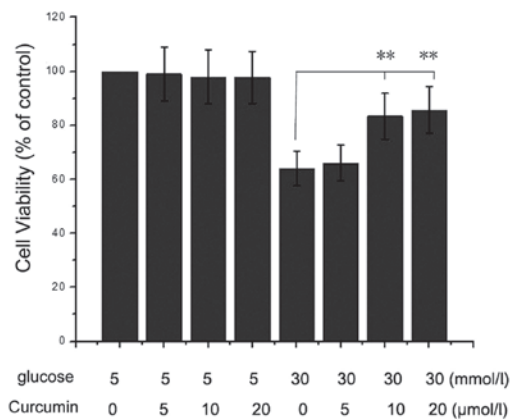


Figure 1. Effects of curcumin on cell viability of NRK-52E cells. Cells were pre-incubated with curcumin (0-20  $\mu$ M) for 24 h. Following curcumin treatment, the medium was changed and cells were treated with 30 mM HG for 48 h. Cell viability was analyzed using an MTT assay. Data are presented as the mean  $\pm$  standard error of the mean (n=5); \*\*P<0.01, vs. HG. HG, high glucose.

The lysates were then centrifuged at 8,000  $\times$  g for 5 min at 4°C. The total protein concentration measurement was performed using the Bradford method (15). The protein samples were boiled for 5 min and 50  $\mu$ g total protein was loaded into the appropriate well for 10% SDS-PAGE (Beyotime Biotechnology, Shanghai, China). The proteins on the gel were then transferred onto a polyvinylidene difluoride membrane (EMD Millipore, Temecula, CA, USA) using Bio-Rad apparatus (A101441, Bio-Rad Laboratories, Inc., Hercules, CA, USA) for 2 h at 4°C and 100 V. The protein-bound membranes were blocked and washed in Tris-buffered saline (TBS)-Tween 20 (20%; Sigma-Aldrich). The membranes were incubated overnight at 4°C with primary antibodies. The primary antibodies used in the present study were as follows: Goat polyclonal anti- $\alpha$ -smooth muscle actin ( $\alpha$ -SMA; 1:400; cat. no. sc-324317), mouse monoclonal anti-E-cadherin (1:400; cat. no. sc-52327), mouse monoclonal anti-Nrf2 (1:400; cat. no. sc-365949), mouse monoclonal anti-HO-1 (1:400; cat. no. sc-136961) and mouse monoclonal anti- $\beta$ -actin (1:400; cat. no. sc-47778) (all Santa Cruz Biotechnology, Inc.). Following extensive washing in TBS-0.1% Tween 20, the membranes were then incubated with horseradish peroxidase-conjugated secondary antibodies, including rabbit anti-goat IgG (1:400; cat. no. sc-2922; Santa Cruz Biotechnology, Inc.) and rabbit anti-mouse IgG (1:400; cat. no. sc-358920; Santa Cruz Biotechnology, Inc.) overnight at 4°C. Subsequently, the membranes were visualized using an enhanced chemiluminescence kit (Walterson Biotechnology Inc., Beijing, China) using the ChemiDoc™ XRS system with Quantity One software version 4.6 (Bio-Rad Laboratories, Inc.) and the G-BOX EF Chemi HR16 gel imaging system (Syngene, Frederick MD, USA). Following development, the band intensities were quantified using Image-Pro Plus 6.0 analysis software (Media Cybernetics, Inc., Rockville, MD, USA). The blots were repeated at least three times for each condition.

**Transient transfection with Nrf2-small interfering RNA (siRNA).** The cells were plated in six-well plates at a density of  $2 \times 10^5$  cells/well in 2 ml DMEM. The cells

were transfected with Nrf2-specific siRNA (sense, 5'-GCACGGUGGAGUUCAUGATT-3' and antisense, 5'-UCAUUGAACUCCACCGUGCCT-3') (Santa Cruz Biotechnology, Inc). The target sequences of the Nrf2 siRNA and control Nrf2 siRNA were aligned against the GenBank database using the Basic Local Alignment Search Tool (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>). Transient transfections were performed according to the manufacturer's instructions using Lipofectamine® 2000 (Invitrogen Life Technologies, Carlsbad, CA, USA) to attenuate Nrf2 expression. All experiments were performed in six-well plates, with cells plated to reach 50-60% confluence on the day of transfection. The cells were incubated in growth medium with 10% fetal bovine serum for 24 h after transfection. The knockdown of Nrf2 was determined using western blot analysis.

**Light microscopy.** The cells (HG group) were cultured for 48 h with 30 mM HG at 37°C. The cells (curcumin/HG group) were cultured with 10  $\mu$ M curcumin pretreatment for 24 h followed by 30 mM HG treatment for 48 h at 37°C. The control cells were cultured for 48 h with 5 mM glucose and 0  $\mu$ M curcumin at 37°C. Subsequently, the cells were observed under a light microscope (Olympus CKX41-A32PH, Olympus, Tokyo, Japan).

**Statistical analysis.** Continuous variables are expressed as the mean  $\pm$  standard error of the mean. One-way analysis of variance was used to analyze the data. Tukey's multiple comparison test was used. P<0.05 was considered to indicate a statistically significant difference.

## Results

**Curcumin rescues HG-induced inhibition of cell viability.** The cell viability of NRK-52E cells under HG (30 mM) and curcumin (0-20  $\mu$ M) conditions were assessed. The results presented in Fig. 1 indicated that the HG condition significantly inhibited NRK-52E cell viability compared with that of the control group (5 mM glucose and 0  $\mu$ M curcumin-treated cells). However, when the cells were treated with 10 or 20  $\mu$ M curcumin and HG, the viability of the cells was increased. Therefore, it was identified that curcumin had a protective effect on NRK-52E cells under HG conditions.

**Curcumin decreases HG-induced EMT in NRK-52E cells.** The HG-induced EMT of NRK-52E cells following curcumin treatment was assessed using light microscopy and western blotting. NRK-52E cells cultured in medium alone for 48 h exhibited typical cobblestone morphology under magnification. As shown in Fig. 2A, a typical epithelial cuboidal shape was observed in the NRK-52E cells cultured in DMEM (5 mmol/l glucose), with the characteristic cobblestone morphology. Following treatment with 30 mM HG, the cell morphology changed to a fibroblast-like shape, with reduced adherence, and the cells lost their apical-to-basal polarity. However, the cellular changes were more noticeable in the cells exposed to 30 mM HG with 20  $\mu$ M curcumin for 48 h. A previous study revealed that HG conditions were able to induce EMT in tubular epithelial cells (8). In fibroblasts,  $\alpha$ -SMA and vimentin proteins



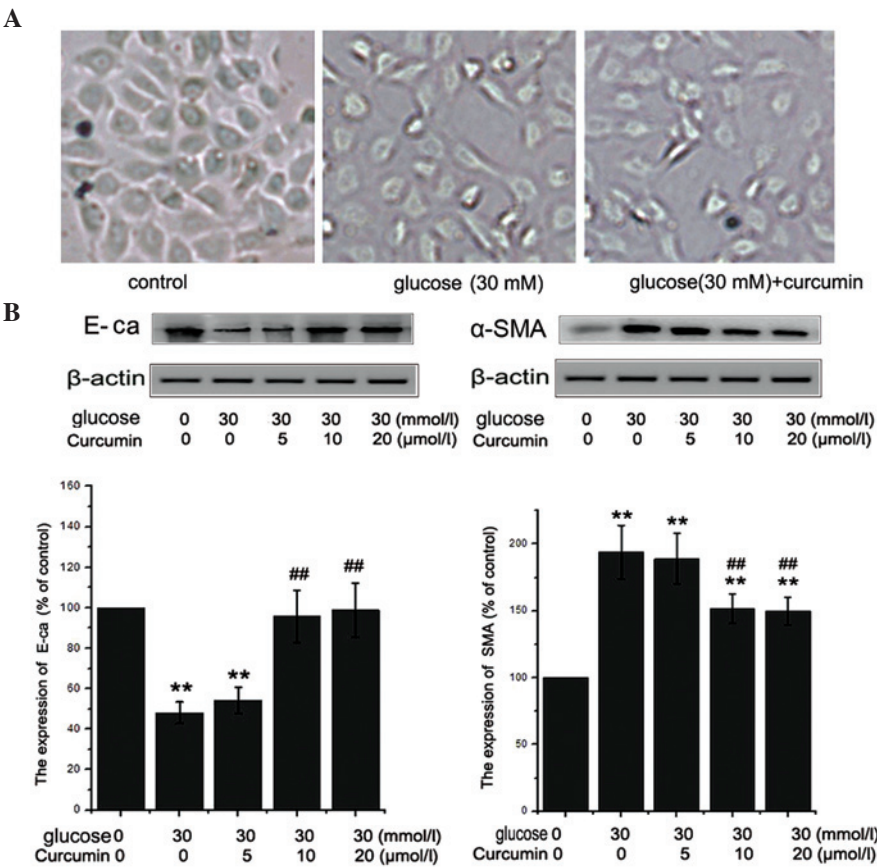


Figure 2. Effects of curcumin on HG-induced epithelial-mesenchymal transition in the NRK-52E cells. (A) NRK-52E cells were treated with 30 mM HG for 48 h, with or without curcumin. Changes in cell morphology were observed using light microscopy (magnification, x200). (B) Representative western blots demonstrated that treatment with 10 or 20  $\mu$ M curcumin prevented HG-induced E-cadherin downregulation and  $\alpha$ -SMA upregulation following 48 h of treatment (n=6). All experiments were repeated three times and results are presented as the mean  $\pm$  standard error of the mean. (\* $P$ <0.01 vs. control, \*\* $P$ <0.01 vs. HG). HG, high glucose;  $\alpha$ -SMA,  $\alpha$ -smooth muscle actin; E-ca, E-cadherin.

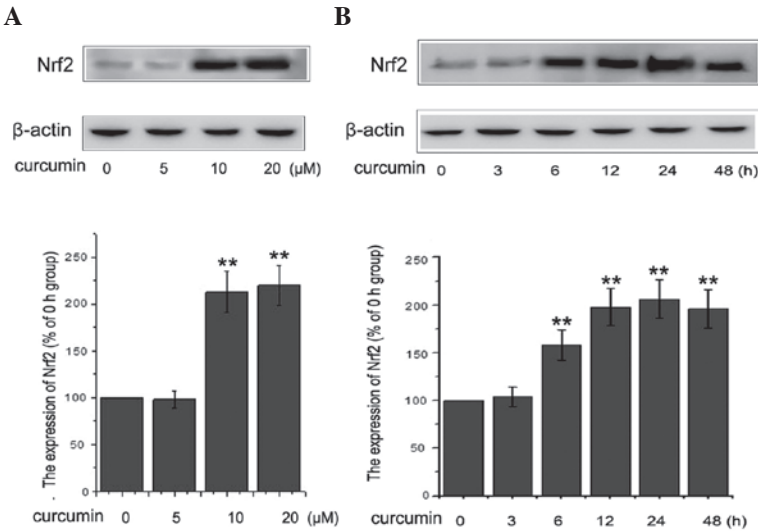


Figure 3. Induction of Nrf2 protein expression in NRK-52E cells by curcumin. Nrf2 expression was analyzed using western blot analysis in cells treated with (A) various concentrations (0, 5, 10, 20  $\mu$ M) of curcumin for 24 h or (B) 10  $\mu$ M curcumin for 0, 3, 6, 12, 24 or 48 h. Results are representative of three independent experiments.  $\beta$ -actin was used as a loading control. (\* $P$ <0.01 vs. control). Nrf2, nuclear factor (erythroid-derived 2)-like 2.

were detected; however, these proteins were not detected in the NRK-52E cells (41). E-cadherin, a  $\text{Ca}^{2+}$ -dependent protein, is crucial in modulating renal epithelial polarity and a decrease in the expression of E-cadherin is considered to indicate

EMT (12). In order to detect HG-induced EMT, the levels of E-cadherin and  $\alpha$ -SMA were assessed using western blotting to analyze samples cultured under HG conditions (30 mM) with or without curcumin pretreatment. A decrease was detected

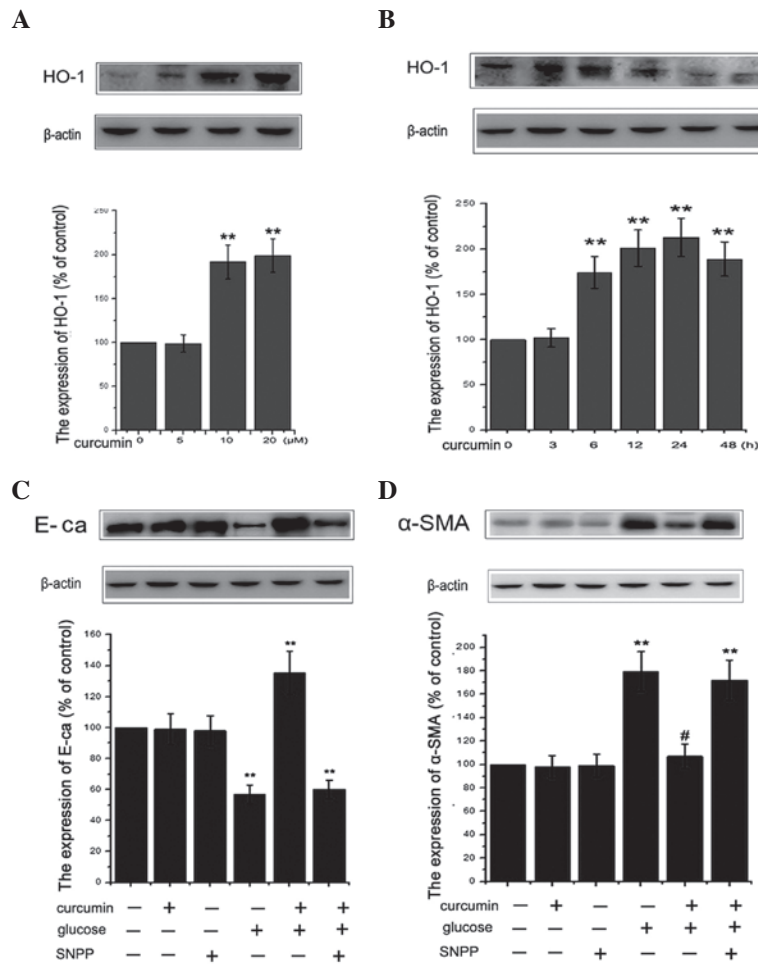


Figure 4. Induction of HO-1 protein expression in NRK-52E cells by curcumin. HO-1 expression was analyzed using western blotting following treatment of cells with (A) various concentrations (0, 5, 10, 20 μM) of curcumin for 24 h or (B) 10 μM curcumin for 0, 3, 6, 12, 24 or 48 h. Results are representative of three independent experiments. β-actin was used as a loading control. (\*\*P<0.01 vs. control). (C and D) Cells were incubated with or without 50 μM SnPP for 12 h and then administered 30 mM glucose for 48 h with or without 10 μM curcumin pretreatment for 24 h. The expression of epithelial-mesenchymal transition proteins, E-cadherin and α-SMA were assessed using western blot analysis. β-actin served as the loading control. Quantitative analysis was performed by measuring the fluorescence intensity relative to the control. Values are expressed as the mean ± standard error of the mean (n=10). All results were obtained from three independent experiments. (\*\*P<0.01 vs. control). α-SMA, α-smooth muscle actin; SnPP, tin protoporphyrin; HO-1, heme oxygenase-1.

in the levels of E-cadherin, accompanied by an increase in α-SMA expression (Fig. 2B), which suggested that these cells had undergone EMT in response to the HG conditions. This reduction in E-cadherin protein expression in response to HG conditions was also accompanied by an increase in α-SMA protein expression in our preliminary experiments, confirming that HG conditions promote EMT in NRK-52E cells. However, this HG-induced EMT was attenuated by pre-treating the NRK-52E cells with 10 or 20 μM curcumin, demonstrated by the reduced upregulation of α-SMA and the ameliorated expression of E-cadherin (Fig. 2B).

**Curcumin increases Nrf2 expression in NRK-52E cells.** Previous studies revealed that Nrf2 was able to regulate cytoprotective genes and cellular antioxidant proteins, as well as allow cells to adapt to stress induced by electrophiles and oxidants (42-44). To analyze the mechanism of action of curcumin on HG-induced EMT in the NRK-52E cells in the present study, the nuclear accumulation of Nrf2 protein in the curcumin-treated NRK-52E cells was examined. The cells were cultured with 0, 5, 10 or 20 μM curcumin for 24 h or 10 μM curcumin for 0, 3,

6, 12, 24 or 48 h and the expression level of Nrf2 was detected using western blot analysis. The results shown in Fig. 3A and B indicated that the nuclear levels of Nrf2 were increased in a concentration and time-dependent manner when cultured with curcumin, compared with those of the control cells. It was therefore concluded that curcumin was capable of effectively inducing the expression of Nrf2 in NRK-52E cells.

**Curcumin promotes expression of HO-1 in NRK-52E cells.** A number of studies have revealed that HO-1 is able to reduce apoptosis by inhibiting cellular oxidative stress (20,45,46). In various types of cell, including glomerular or endothelial cells, the expression of HO-1 was demonstrated to be induced by curcumin in previous studies (47,48). To elucidate the role of curcumin in renal tubular epithelial cells, HO-1 expression was assessed in NRK-52E cells cultured with curcumin. As indicated in Fig. 4A and B, curcumin was observed to upregulate HO-1 protein expression in a dose- and time-dependent manner. Compared with the untreated controls, curcumin treatment led to a significant increase in the level of HO-1 protein expression. In order to determine whether HO-1 exerted a cytoprotective

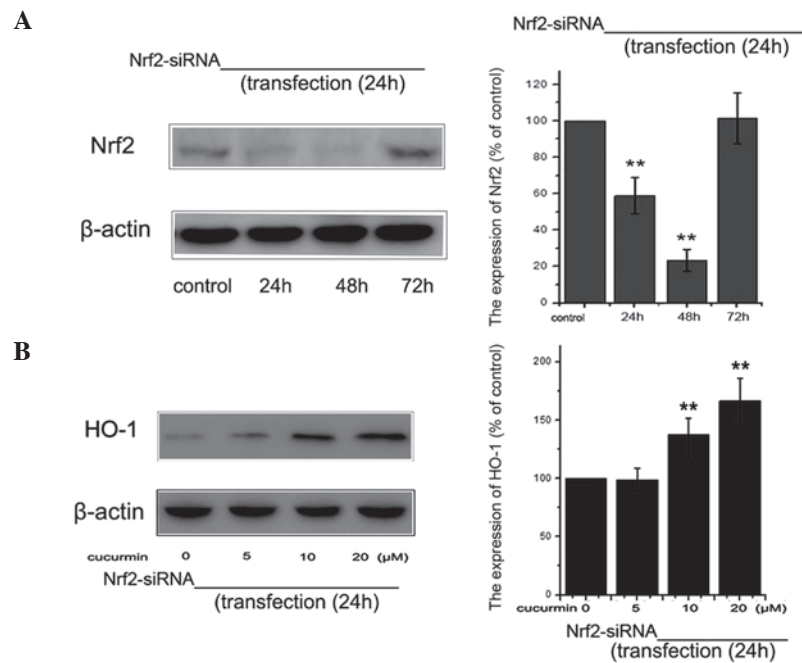


Figure 5. Nrf2 inhibits curcumin-induced HO-1 expression. (A) NRK-52E cells were transfected with Nrf2-siRNA and western blot analysis was performed with an antibody against Nrf2 was performed at various time-points following transfection (24, 48 and 72h). Relative Nrf2 expression levels were calculated and normalized to the loading control. Corresponding protein levels were assessed using densitometry and expressed in relative intensities. All results were obtained from three independent experiments. Values are expressed as the mean  $\pm$  standard error of the mean ( $n=6$ ; \*\* $P<0.01$ , vs. control). (B) Cells were treated with curcumin (0, 5, 10 or 20  $\mu$ M) for an additional 24 h with or without Nrf2-siRNA. Western blotting was performed with an antibody against HO-1. Relative HO-1 expression levels were calculated and normalized to the loading control. Corresponding protein levels were assessed using densitometry and were expressed in relative intensities. All results were obtained from three independent experiments. Values are expressed as the mean  $\pm$  standard error of the mean ( $n=10$ ; \*\* $P<0.01$ , vs. control). HO-1, heme oxygenase-1; Nrf2, nuclear factor (erythroid-derived 2)-like 2; siRNA, small interfering RNA.

effect against the HG-induced EMT, 50  $\mu$ M SnPP, a known inhibitor of HO, was utilized. The concentration of SnPP was determined as previously described (49). As shown in Fig. 4C and D, SnPP treatment attenuated the protective effects of curcumin against HG-induced EMT in renal tubular epithelial cells. Notably, SnPP treatment alone did not affect cell viability in the present study (data not shown). In conclusion, curcumin was demonstrated to have a cytoprotective role, which is mediated through the induction of HO-1 expression.

**siRNA knockdown of Nrf2 abrogates curcumin-induced HO-1 expression.** In order to examine the role of Nrf2 in the upregulation of HO-1 expression, siRNA knockdown of the Nrf2 gene was used. The level of Nrf2 protein was detected using western blot analysis at various time-points following siRNA-Nrf2 transfection (Fig. 5A). siRNA-Nrf2 significantly reduced the HO-1 expression induced by curcumin treatment (Fig. 5B). In conclusion, the present findings supported the hypothesis that curcumin promotes the expression of HO-1 through activation of Nrf2 in NRK-52E cells.

**Anti-fibrotic effects of curcumin are mediated by activation of Nrf2 signaling.** To determine whether curcumin protects cells against HG-induced EMT through the modulation of Nrf2 and HO-1 expression, the role of Nrf2 in EMT was investigated via knockdown of Nrf2. The cells were divided into six groups as follows: i) Control group; ii) siRNA group, subjected to Nrf2-siRNA transfection; iii) curcumin group, subjected to 10  $\mu$ M curcumin treatment for 24 h; iv) HG group, 30 mM HG treatment; v) HG/curcumin group, 10  $\mu$ M

curcumin pretreatment for 24 h followed by 30 mM HG treatment for 48 h; vi) HG/curcumin/Nrf2-siRNA group, following transfection for 24 h, cells were treated with 10  $\mu$ M curcumin for 24 h and 30 mM HG for 48 h. The results revealed that HG-induced EMT was partially prevented by curcumin pretreatment, which resulted in a decrease in the HG-induced increase in  $\alpha$ -SMA expression and an increase in the expression of E-cadherin in NRK-52E cells (Fig. 6). In addition, Nrf2-siRNA alone did not induce EMT in the NRK-52E cells. However, knockdown of Nrf2 with siRNA inhibited the curcumin-induced anti-fibrotic effects (Fig. 6). These results suggested that curcumin protects NRK-52E cells from HG-induced EMT processes through activation of the Nrf2/antioxidant response signaling pathway and subsequent targeting of gene expression.

## Discussion

Multiple studies have demonstrated that curcumin is capable of inhibiting fibrosis in certain chronic inflammatory diseases, including cystic fibrosis, liver fibrosis and myocardial fibrosis (32,33). In the present study, the protective effect of curcumin on HG-induced EMT was analyzed in renal tubular epithelial cells. The results of the present study revealed that curcumin reduced HG-induced EMT in a dose- and time-dependent manner, as indicated by the detected decrease in the upregulation of  $\alpha$ -SMA and the increase in E-cadherin expression. The mechanism underlying this process may involve abrogation of HG-induced oxidative stress via activation of Nrf2 and HO-1 in NRK-52E cells.

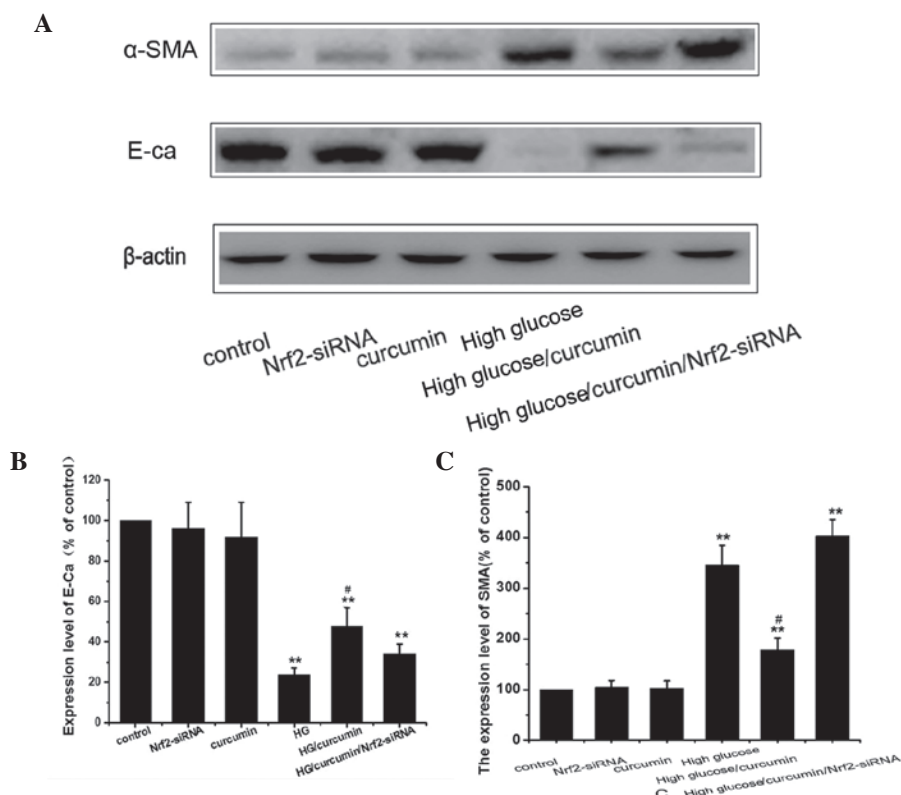


Figure 6. Curcumin-mediated activation of Nrf2 signaling protects cells against HG-induced EMT. Cells were incubated with or without Nrf2-siRNA prior to treatment with 30 mM HG for 48 h with or without 10  $\mu$ M curcumin pretreatment for 24 h. (A) Western blotting was performed and relative expression of EMT proteins was calculated and normalized to the loading control. (B and C) Corresponding protein levels were assessed using densitometry and are expressed as relative intensities. Values are expressed as the mean  $\pm$  standard error of the mean (n=10; \*\*P<0.01, vs. control, \*P<0.01, HG/curcumin vs. HG). siRNA, small interfering RNA; EMT, epithelial-mesenchymal transition; HG, high glucose; Nrf2, nuclear factor (erythroid-derived 2)-like 2.

EMT in mature tubular epithelial cells of the kidney is currently considered to contribute to the renal accumulation of matrix proteins associated with DN and is closely associated with the progression of renal interstitial fibrosis (7,50). In DN, EMT occurs in response to TGF- $\beta$ 1 activation under HG conditions and contributes to the loss of tubular epithelial cells and the accumulation of interstitial fibroblasts, which are associated with a decline in renal excretory function (11,13). Typical epithelial cell alterations, which are associated with EMT, include reorganization of the actin cytoskeleton, *de novo* acquisition of mesenchymal cytoskeletal markers and the downregulation of epithelial adhesion molecules (51,52). To the best of our knowledge, the anti-fibrotic effects of curcumin have previously only been investigated in models of pulmonary or liver fibrosis, and curcumin was found to be associated with a reduction in the expression of inflammatory mediators, decreased expression of the profibrotic cytokine TGF- $\beta$ 1 and a subsequent decrease in the accumulation of collagen (32,33). Curcumin was also found to inhibit renal interstitial inflammation and fibrosis via inhibition of the NF- $\kappa$ B-dependent pathways in a UUO rat model of kidney fibrosis (35). A recent study revealed that curcumin inhibited TGF- $\beta$ 1-induced EMT in renal tubular epithelial cells via the extracellular signal-regulated kinase-dependent and peroxisome proliferator-activated receptor  $\gamma$ -dependent pathways (53). In the present study, it was confirmed that HG-induced changes in EMT markers were more prominent than in the control and were accompanied by a

decrease in the expression of epithelial marker E-cadherin and an increase in  $\alpha$ -SMA. Curcumin pretreatment may provide effective protection against HG-induced EMT, as evidenced by a decrease in the upregulation of  $\alpha$ -SMA and the amelioration of E-cadherin expression, which was associated with the transition from the epithelial to myofibroblastic phenotype in NRK-52E cells.

A number of previous studies have confirmed that EMT in the tubular epithelial cells of patients with DN is generally regarded to be the result of hyperglycemia-induced oxidative stress; notably, antioxidants effectively reverse this induction of EMT in tubular epithelial cells (6,13,16,54). Nrf2-mediated transcriptional responses have been found to be protective in a number of animal models of disease, including those of oxidative lung injury and fibrosis, asthma and brain ischemia-reperfusion (55,56). The induction of kidney ischemia followed by reperfusion in wild-type mice was found to elevate Nrf2 levels and activate downstream target genes (57). By contrast, Nrf2 deficiency was demonstrated to enhance the susceptibility of cells to ischemic and nephrotoxic acute renal injury (58). Additionally, treatment of Nrf2 knockout mice with antioxidants, including N-acetyl-cysteine or glutathione, is able to improve renal function (59). Furthermore, Nrf2 knockout mice with streptozotocin-induced diabetes were found to exhibit progressively increasing levels of nitric oxide metabolites in their urine, eventually developing renal injury (19). Curcumin is able to stimulate the dissociation of Nrf2 from Keap1, a



cytosolic Nrf2 inhibitor, which leads to increased Nrf2 binding to the antioxidant response element in the promoters of target genes (33). Curcumin has also been demonstrated to be a potent inducer of Nrf2-associated antioxidant enzymes and an inhibitor of oxidant-induced NF- $\kappa$ B activation in lung epithelial cells (60). Similarly, in mouse alveolar macrophages *in vitro* and in the lungs *in vivo*, curcumin has been observed to upregulate Nrf2 target antioxidant gene expression (33). In addition, in the present study, it was also confirmed that curcumin induced Nrf2 activation in NRK-52E cells, which may represent the mechanism responsible for the protective effects of curcumin in cells subjected to HG-induced EMT.

To further examine the possible downstream mechanisms through which Nrf2 may elicit protection against HG-induced EMT, the expression of HO-1 was investigated in HG-treated cells transfected with siRNA-Nrf2. HO-1 is a rate-limiting enzyme involved in the degradation of heme to produce equimolar quantities of CO, iron and biliverdin (23,26). Growing evidence indicates that the HO-1 system is a regulator of renal vascular integrity and responses to oxidative stress (23). The induction of HO-1 expression by curcumin has been observed in several cell types, including human renal tubular cells and renal fibroblasts (14). A previous study revealed that curcumin exhibits an anti-fibrotic effect in a model of glomerular fibrosis, and curcumin treatment *in vitro* and *in vivo* may lead to the induction of HO-1 (47). The detailed mechanism through which curcumin induces HO-1 has been investigated in cultured cells (31). In human proximal tubular cells, curcumin-induced HO-1 expression was reduced by co-treatment with an NF- $\kappa$ B inhibitor, implicating this pathway in the modulation of HO-1 in this cell type. Other studies have indicated that, following curcumin treatment, there is an increase in HO-1 protein expression levels in kidney tissue and this mechanism is essential in the prevention of transplant-associated organ injury and rejection (61). Consistent with these results, the present data demonstrated that curcumin induced a marked increase in HO-1 levels and that transfection with siRNA-Nrf2 significantly attenuated this increase. Simultaneously, the HG-induced reduction in E-cadherin and upregulation of  $\alpha$ -SMA were reversed by curcumin, whereas knockdown of Nrf2 with siRNA inhibited the curcumin-induced anti-fibrotic effects. The present results suggested that curcumin-mediated cell protection may occur via the activation of Nrf2 and subsequently the key target gene HO-1, thereby protecting NRK-52E cells from HG-induced EMT processes.

In conclusion, the present study demonstrated that curcumin exhibited inhibition of HG-induced EMT in NRK-52E cells and that this effect was dependent on the activation of Nrf2 and subsequent HO-1 induction. The present data suggested that curcumin is a significant regulator of HG-induced EMT and may be beneficial in the treatment of DN. However, the underlying mechanisms require elucidation through further investigation.

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