ORIGINAL ARTICLE

Smad3 signalling affects high glucose-induced podocyte injury via regulation of the cytoskeletal protein transgelin

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Funding information

National Natural Science Foundation of China. Grant/Award Number: 81400718; Research Foundation of Beijing Friendship Hospital, Capital Medical University, Grant/Award Number: yyqdkt2018-21

Abstract

Aim: The aim of the present study was to characterize the role of Smad3 signalling on high glucose-induced podocyte injury.

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Methods: Synchronized conditionally immortalized mouse podocyte cell line (MPC5) cells were treated with either D-glucose alone or D-glucose plus the Smad3 inhibitor SIS3. The distribution of F-actin and transgelin in a high glucose-induced model of podocyte injury were examined by immunofluorescence. Levels of transgelin and Smad3 signalling proteins in MPC5 cells were determined by Western blot.

Results: A disordered distribution of F-actin, as well as co-localization of F-actin and transgelin, was observed in podocytes exposed to high glucose. Increased levels of transgelin were first observed 10 minutes after treatment with glucose, suggesting that this protein is sensitive to hyperglycaemic injury. Levels of phosphorylated Smad3 and cleaved caspase 3 increased significantly with glucose stimulation. Moreover, expression of the downstream protein c-Myc, but not JAK1/STAT3, was induced in conditions of high glucose. The Smad3-specific inhibitor SIS3 prevented the effects of high glucose on Smad3 phosphorylation, expression of transgelin and c-Myc, caspase 3 cleavage and cytoskeletal organization. Expression of the tumour suppressor protein p15^{INK4B} increased after podocyte injury but was unaffected by Smad3 inhibition, suggesting that Smad3 regulation of high glucose-induced podocyte injury occurs through a p15^{INK4B}-independent mechanism.

Conclusion: Smad3 signalling plays a critical role in the modulation of hyperglycaemic injury. Targeted inhibition of the Smad3 pathway may offer a novel route for treatment of podocyte damage, especially in cases of diabetic nephropathy.

KEYWORDS

high glucose, podocyte injury, Smad3, transgelin

Kidney injury is often accompanied by de-differentiation and apoptosis of podocytes, differentially terminated cells that are involved in kidney diseases such as diabetic nephropathy.¹ Podocyte cytoskeletal proteins, actin-binding proteins and polysaccharide proteins in the apical region

of the foot process function together to maintain the integrity of slit diaphragms.² Podocyte loss, deteriorated microstructure and foot process effacement are frequently observed in diabetic nephropathy.^{3,4} Loss of podocyte function is also a key feature of diabetic nephropathy.

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However, the precise mechanism underlying hyperglycaemic podocyte injury remains unclear. Therefore, understanding the mechanism of high glucose-induced podocyte injury will contribute to therapeutic strategies for treatment of diabetic nephropathy.

Morphological changes of podocytes are typically accompanied by disruption of the actin cytoskeleton. It has been reported that high glucose induces actin remodelling in a murine podocyte cell line⁵ as well as in a mouse model.⁶ In addition, transgelin has also been identified as a marker of cytoskeleton remodelling.⁷ Transgelin is an actin-binding protein that is absent in normal rat glomeruli, but highly expressed in podocytes during nephrotoxicity, diabetic nephropathy, IgA nephropathy and membrane kidney diseases. Transgelin is known to mediate morphological changes of podocyte foot processes in a rat model of puromycin aminonucleosideinduced kidney injury.⁸ However, the role and location of transgelin during high glucose-induced podocyte cells injury have yet to be described.

In a model of high fat diet-induced diabetic nephropathy in rats, the protective effects of Taxus chinensis occurred in part via downregulation of TGF-β1 expression and Smad2/3 phosphorylation.⁹ In another study, betulinic acid alleviated high glucose-induced cardiac fibroblasts mainly through inhibition of the TGF-^β1-Smad signalling pathway.¹⁰ The damage to cardiomyocytes via reperfusion injury, aggravated by high glucose, was also reported to be mediated by TGF-β1-Smad signalling.¹¹ During high glucose-induced epithelial-tomesenchymal transition in peritoneal mesothelial cells, Smad3 protein was significantly more phosphorylated than in low-glucose conditions.¹² In addition to its involvement in high glucose stimulation, the Smad signalling pathway is also activated during podocyte injury.¹³ Podocyte injury was alleviated in obese Smad3-deficient mice.¹⁴ and Smad3 phosphorylation was induced by puromycin aminonucleosidestimulated podocyte injury.¹⁵ In a mouse podocyte cell line, treatment with gremlin aggravates high glucose-induced cell injury and also enhances activation of TGF-β1-Smad signalling.¹⁶ Interestingly, TGF- β 1 was also able to disrupt the cytoskeleton and induce podocyte apoptosis by regulating the interaction of Smad2/3 with other proteins.¹⁷ The precise mechanism by which TGF- β 1 signalling regulates high glucose-induced podocyte injury is still unclear.

In this study, changes in the distribution of F-actin and transgelin during high glucose-induced kidney injury were examined, and the expression of transgelin and related Smad3 signalling pathways were analysed. These results provide an experimental basis for clarifying the pathogenesis of kidney injury under conditions of high glucose.

1 | MATERIALS AND METHODS

1.1 | Podocyte cultures and treatments

The conditionally immortalized podocyte mouse cell line (MPC5) (gifted by Professor Peter Mundel, Mount Sinai Medical School, New York) was cultured at 33° C with recombinant murine γ -interferon (PEPRO Tech), and then was incubated at 37° C for

SUMMARY AT A GLANCE

Podocyte damage is characteristic of diabetic kidney disease. This study showed that inhibition of Smad-3 signalling using the specific inhibitor SIS3 in vitro, could alleviate podocyte abnormalities induced by high glucose including actin cytoskeletal rearrangement and activation of cell death pathways.



F-actin; Transgelin; Nucleus

FIGURE 1 Distribution of cytoskeletal proteins upon treatment of podocytes with high glucose. A, Time-dependent disruption of Factin in MPC5 cells treated with high glucose. B, High glucose induces co-localization of F-actin (red) and transgelin (green) in MPC5 cells. Scale bar = $50 \mu m$

1 week to promote differentiation.¹⁸ Mature podocytes were seeded onto 60-mm Petri dishes coated with 10 μ g/mL type-I collagen (Sigma), and cultured in the RPMI-1640 medium (Gibco) in 37°C. When mature podocytes reached 60% confluency they were synchronized first in the RPMI-1640 media with 0.5% fetal bovine serum (Gibco) for 12 hours, then treated with 35 or 5 mmol/L D-glucose or 5 mmol/L D-glucose (Sinopharm) with 30 mmol/L mannitol for 2, 5, 10, 15 and 30 minutes. Total cell lysates were extracted at different time points to be used for immunoblot assay.

To detect the effect of the Smad3 inhibitor SIS3 (Calbiochem) on high glucose-induced podocyte injury, MPC5 cells were grouped to receive one of the following five treatments: (a) blank control, (b) osmotic pressure control, (c) SIS3 treatment, (d) high glucose

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FIGURE 2 Expression of transgelin and signalling proteins upon treatment of podocytes with high glucose. Protein levels of transgelin, cleaved caspase 3, Smad3 and phospho-Smad3 protein were assessed by Western blot. β -Actin was used as an internal control. A, Representative blot, B, Average levels of transgelin, C, cleaved caspase 3, D, phospho-smad3 and E, total smad3 were normalized with respect to β -Actin. Data are representative of three independent experiments, and were analysed using one-way analysis of variance. **P* < .05, ***P* < .01 compared with control or mannitol group

treatment and (e) SIS3 with high glucose treatment. This experiment was repeated at least three times.

1.2 | Immunofluorescence staining

When mature podocytes reached 80% confluency, they were synchronized first in the RPMI-1640 media with 0.5% serum for 6 hours, and then treated with 35 mmol/L D-glucose for 0, 24, 48 and 72 hours. To examine the effect of SIS3 on cytoskeletal morphology after treatment with high glucose, the synchronized MPC5 cells were incubated with or without 5 μ mol/L SIS3 for 30 minutes, followed by 35 mmol/L Dglucose for 12 hours. These cells were subjected to immunofluorescence staining as previously described.¹⁹ Briefly, cells were fixed with 4% paraformaldehyde for a quarter of an hour at room temperature, and then washed several times with phosphate buffer (PBS). After permeabilizing with Triton X-100 for 15 minutes, podocytes were blocked with 5% bovine serum albumin for half an hour at room temperature. Cells were incubated with primary antibodies (mouse anti-Factin, 1:100, Ab205, Abcam; rabbit anti-transgelin, 1:100, Ab155272, Abcam) overnight at 4°C, and then washed three times with PBS and incubated with secondary antibodies (donkey anti-mouse IgG (H + L) Alexa Fluor 546-conjugated, cat. No. A10036, ThermoFisher Scientific; goat anti-rabbit IgG (H + L) Alexa Fluor 488-conjugated, cat. No. A11034, ThermoFisher Scientific) for 1 hour at room temperature, then washed three times with PBS. These cells were then mounted with the Prolong Gold antifade mountant with DAPI (cat. No. P36931, ThermoFisher Scientific). Cellular morphology was observed using laser scanning confocal microscopy (TE300, Nikon), and analysed with the ImageQuant software v5.3. The distribution of F-actin and transgelin were observed in 10 randomly selected high-power fields of podocytes (original magnification ×600).



FIGURE 3 Expression of transgelin and signalling proteins upon treatment of podocytes with high glucose and SIS3. A, Representative Western blot used to quantify protein levels of B, transgelin, C, phospho-smad3, D, total smad3, E, cleaved caspase 3, F, phospho-JAK, G, phospho-STAT3, H, c-Myc and I, p15^{INK4B}. Protein levels were normalized with respect to β -Actin. Data are representative of three independent experiments, and were analysed using two-way analysis of variance. *P < .05. **P < .01

1.3 | Western blot

MPC5 cells were treated with D-glucose, mannitol or SIS3 as described above. Cell lysates were extracted with lysis buffer (P001, UKzybiotech, China) and proteinase inhibitor cocktail (04693116001, Roche). Protein levels were quantified using the BCA Protein Assay Kit (cat. No. 23225, ThermoFisher Scientific). Protein extracts were run through 12% of SDS-polyacrylamide gel electrophoresis gels, then transferred to polyvinylidene fluoride membranes (0.45 μ m pore size, IPVH00010, Millipore). Membranes were first washed with PBS and then blocked with 5% skim milk for 1 hour. Primary antibodies used were as follows: rabbit anti-Smad3 (1:750, 9523T), rabbit anti-phospho-Smad3 (1:1000, 9520T), rabbit anti-phospho-JAK (1:1000, 3331S), rabbit anti-phospho-STAT3 (1:1000, 9145T), rabbit anti-caspase 3 (1:1000, 9661T) and rabbit anti-c-myc (1:1000, 9402S) antibodies were purchased from the cell signalling technology; rabbit anti-transgelin (1:1000, Ab155272) and rabbit anti-p15^{INK4B} (1:1000, Ab53034) antibodies from Abcam; and mouse anti-beta-Actin (1:1000, TA-09) from Zsbio. Peroxidase-

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FIGURE 4 Distribution of F-actin upon inhibition of Smad3. A, control or SIS3 group (top left): untreated control podocytes display well-organized F-actin; high glucose group (top right): disordered cytoskeleton was observed in podocytes treated with high glucose; SIS3 with high glucose group (bottom left): the Smad3-specific inhibitor had no effect on F-actin organization, and (bottom right) appeared to block the disruptive effects of high glucose. Scale bar = 50 μ m. B, Percentage ratios of shrunken cells to total cells for each group



F-actin; Nucleus

AffiniPure goat anti-mouse (1:4000, 115-035-003) and anti-rabbit (1:4000, 111-035-003) secondary antibodies were purchased from Jackson Laboratory. Immunoblot signals were detected with the Super Signal West Pico-enhanced chemiluminescent substrate (WBKLS0500, Millipore) and then analysed with the Alphalmager mini gel imaging system.

1.4 | Statistical analysis

The grey value of the immunoblotting bands of at least three independent experiments was quantified with the ImageJ software. The data were then compared by one-way and two-way analysis of variance (ANOVA) with the least significant difference test (equal variance) or Dunnett's T3 test (unequal variance) where appropriate. *P* value less than .05 was considered significant.

2 | RESULTS

2.1 | Transgelin is induced and co-localized with Factin in a model of high glucose-induced podocyte injury

Control cells displayed intact cytoskeleton, and evenly distributed Factin in cytoplasm as well as in podocyte foot processes (Figure 1A). Cells that were treated with high glucose displayed cytoskeletal disruption that increased with the duration of glucose treatment; the most profound foot process retraction and cytoskeletal damage were observed at the 72-hour time point.

Immunofluorescence localization of transgelin showed low levels of the cytoskeletal marker protein in control cells, but high levels after treatment with high glucose for 12 hours (Figure 1B). Levels of F-actin were high in both groups, although mild disarrangement of cytoskeleton was observed after glucose treatment as described above. Finally, it was observed that F-actin and transgelin were mostly co-localized in cells exposed to high glucose.

2.2 | The time course of transgelin expression upon high glucose stimulation

As assayed by Western blot, levels of transgelin and the cleaved (activated) form of the apoptosis marker caspase 3 varied significantly after exposure to high glucose. Specifically, transgelin levels increased as early as 10 minutes after exposure to high glucose, and remained at a high level 30 minutes after exposure (Figure 2). In contrast, cells treated with mannitol or low glucose displayed low levels of transgelin at all time points. Levels of the cleaved (activated) apoptosis marker caspase 3 were also notably higher 10 minutes after treatment with high glucose, implying that apoptosis of podocytes had been induced.

It is well known that TGF- β signalling is important for the apoptosis of podocyte cells,^{20,21} so we next examined whether and when Smad3 was phosphorylated (activated) during high glucose treatment. Smad3 phosphorylation was first observed 10 minutes after treatment with high glucose, and levels continued to be high throughout the observation period. This suggests that activation of TGF- β signalling may be important for high glucose-induced apoptosis.

2.3 | Inhibition of Smad3 decreases high glucoseinduced podocyte injury and transgelin expression

To further elucidate the relationship between Smad3 signalling and high glucose-induced apoptosis, the SIS3 was applied to MPC5 cells. As described above, levels of transgelin, phospho-Smad3 and cleaved caspase 3 increased upon treatment with high glucose; this effect was significantly inhibited by SIS3 based on the treatment with high glucose, suggesting that SIS3 may be able to block apoptosis in podocytes (Figure 3). Proteins downstream of Smad3, c-myc and p15^{INK4B}, followed a similar trend, but SIS3 had no effect on the glucose-induced expression of p15^{INK4B}. Levels of phosphorylated JAK and STAT3 in cells exposed to high glucose did not differ from those of the low glucose or mannitol groups, indicating that the JAK/STAT signalling pathway was not affected.

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2.4 | Smad3 inhibition rescues the morphological changes induced by high glucose stimulation

Complete cytoskeletal filaments were readily observed in the control and SIS3 groups. In the high glucose group, however, the cytoskeleton appeared disordered and there were decreased numbers of podocyte microfilaments (Figure 4A). Cells treated with both high glucose and SIS3 resembled controls, with organized F-actin and increased numbers of podocyte filaments (Figure 4B).

3 DISCUSSION T

In the present study, we investigated changes to the location and expression of newly identified cytoskeleton protein transgelin during high glucose-induced podocyte injury. High glucose induced a striking disarrangement of cytoskeletal F-actin and enhanced the expression of transgelin; moreover, F-actin and transgelin were observed to mostly co-localize. Concomitant with these changes to F-actin and transgelin was a significant up-regulation of Smad3 phosphorylation and its downstream proteins c-Mvc and p15^{INK4B}. Inhibition of Smad3 via SIS3 was protective against cytoskeletal disarrangement and transgelin up-regulation. These results begin to define the mechanism by which Smad3 signalling affects high glucose-induced podocyte injury, and provides an experimental basis for novel treatments for diabetic nephrology.

The cytoskeletal protein transgelin was identified in our previous study using a rat model of kidney injury induced by puromycin aminonucleoside.⁷ Others have reported that transgelin expression can be induced in proteinuric human diseases, including diabetic nephropathy.⁸ In human skeletal cells, transgelin was reported to be important for differentiation and cytoskeletal integrity.²² In mice with crescentic glomerulonephritis, more podocytes were observed in transgelin knockout mice than in wild-type controls, suggesting that transgelin may aggravate glomerular cell injury. In the present study, transgelin expression was also increased upon high glucose treatment. This induction occurred as early as 10 minutes after treatment, suggesting that transgelin can be induced quickly during high glucoseinduced podocyte injury.

Localization of transgelin was reported to be different from that of the myofibroblast marker protein aSMA in nephritic rat model.²³ The induced transgelin was preferentially located in podocytes, whereas aSMA was mainly expressed in mesangial cells, indicating that transgelin specifically functions in podocytes in this kidney injury model.

In the present study, transgelin protein levels increased after treatment with high glucose, an effect that was blocked by Smad3 inhibition, implying that Smad3 acts as a regulator of transgelin under high glucose conditions. Smad3 may directly regulate expression of transgelin by binding to the transgelin promoter. It has been reported that the promoter region of transgelin is activated by the phosphorylated Smad.²⁴ There is also a TGF-^β control element within the transgelin promoter,²⁵ suggesting that TGF- β /Smad signalling has a direct role in the regulation and control of transgelin expression. Smad1/3/4 proteins can be activated by TGF- β to form heterotrimers, which then translocate to cell nuclei to induce transcription of downstream genes. During this process, Smad3 plays a critical role in inducing transgelin transcription, due to the Smad binding site in its promoter region.²⁶

In addition to TGF- β 1/Smad3 signalling, expression of transgelin is also influenced by pro-inflammatory signals such as interleukin-1beta (IL-1 β) and TGF β 2²⁴ and JNK and p38 MAPK signalling²⁷ in smooth muscle cells. Smad3 can be regulated by TGFβ2 at the mRNA level in chondrocytes²⁸ and in primary human trabecular meshwork cells.²⁹ In addition, reduced levels of Smad3 expression were reported during IL-1_B-induced chondrocyte injury.³⁰ Together, these data suggest that TGF-\beta1/Smad3 signalling may also indirectly regulate the expression of transgelin via cross talk with other signalling pathways such as IL-1 β or TGF β 2.

In the present study, addition of a Smad3 inhibitor attenuated podocyte injury upon treatment with high glucose. However, in a mouse model of diabetic glomerulosclerosis disease, proteinuria and foot process fusion still occur even after Smad3 knockout,³¹ indicating that there may be other signalling pathways involved.

Smad3 and c-Myc signalling are known to interact. During liver fibrosis, activated Smad3 interacts with BRD4 to promote its direct binding to the promoter region of c-Myc.³² Smad3/c-Myc signalling was also reported to be important for oncogenesis of hepatocellular carcinoma.³³ It is also known that TGF- β can inhibit cell proliferation by down-regulating expression of c-Myc and up-regulating expression of the cyclin-dependent kinases inhibitor p15^{INK4B.34} In the present study, c-Myc was identified to be a downstream target of Smad3 during high glucose-induced podocyte injury. However, p15^{INK4B} expression was induced by podocyte injury, but unaffected by Smad3 inhibition, suggesting that the involvement of Smad3 in regulating high glucoseinduced podocyte injury was not through p15^{INK4B}. Although JAK/STAT3 signalling has been shown to be activated in other cell types exposed to high glucose conditions,³⁵ we did not observe any such involvement here. Therefore, JAK/STAT3 activation by high glucose may be dependent on cell type. There are likely additional, as-yetunidentified, signalling pathways involved as well. Finally, all cited experiments were conducted at the molecular or cellular level. Further in vivo experiments should be done to confirm these results at a wholeorganism level.

In conclusion, disordered cytoskeleton, foot process retraction and Smad3 activation were observed during high glucose-induced podocyte injury. In addition, transgelin protein was observed to increase and to co-localize with the cytoskeletal protein F-actin. The expression of downstream protein c-Myc, but not JAK1/STAT3 signalling proteins, was increased by high glucose condition. Addition of SIS3 blocked the effects of high glucose on Smad3 phosphorylation, expression of transgelin and c-Myc, cleavage of caspase 3 and cytoskeletal structure, implying a critical role for Smad3 signalling in modulating hyperglycaemic injury. Targeted inhibition of the Smad3 pathway may be a novel treatment route for alleviating podocyte damage, especially in cases of diabetic nephropathy.

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ACKNOWLEDGEMENTS

This study was supported by the National Natural Science Foundation of China (Grant No. 81400718) and Research Foundation of Beijing Friendship Hospital, Capital Medical University (Grant No. yyqdkt2018-21).

CONFLICT OF INTEREST

The authors declare no conflicts of interest.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of this article.

How to cite this article: Jiang L, Cui H, Ding J. Smad3 signalling affects high glucose-induced podocyte injury via regulation of the cytoskeletal protein transgelin. *Nephrology*. 2020;25:659–666. https://doi.org/10.1111/nep.13701