ORIGINAL ARTICLE

PSC-MSC-Derived Exosomes Protect against Kidney Fibrosis In Vivo and In Vivo through the SIRT6/ β -Catenin Signaling Pathway

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Background and Objectives: Chronic kidney disease (CKD) has a major impact on the quality of life of patients, and renal fibrosis is a critical pathological change in the disease. It is very important to control the process of renal fibrosis to improve the quality of life of patients with CKD. The pathological mechanism of renal fibrosis is very complicated, and the current treatment strategy also has many flaws.

Methods and Results: To explore a better treatment, we collected exosomes from pluripotent stem cell (PSC)-derived mesenchymal stem cells (MSC) and verified their therapeutic effect on renal fibrosis through *in vivo* and *in vitro* experiments. In this study, we found that PSC-MSC-derived comes could prevent the epithelial differentiation of NRK-52E cells, and with increasing exosome concentrations, the effect was improved. Furthermore, PSC-MSC-derived exosomes could reduce the pathological process of renal fibrosis, reduce inflammatory reactions and improve renal function in UUO mice. Moreover, the protective effect of exosomes against renal fibrosis may be achieved by increasing the expression of SIRT6 and decreasing the expression of β -catenin and its downstream products.

Conclusions: These findings suggest the possibility of PSC-MSC-derived exosomes as a new, effective therapeutic tool for kidney fibrosis.

Keywords: Exosomes, Renal fibrosis, UUO, SIRT6

Introduction

Renal fibrosis (RF) is a common condition and the

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main pathological basis of many chronic kidney diseases, which lead to end-stage renal disease (1, 2). Chronic kidney disease seriously affects the patient quality of life and threatens health (3). Therefore, preventing the progression of renal fibrosis is essential for CKD patients. During renal fibrosis, the sustained injury of endothelial cells and epithelial cells (4), the recruitment of inflammatory cells (5) and the activation of myofibroblasts (6) play important roles. At present, the prevention of renal fibrosis mainly targets the abovementioned mechanism. Although there are various treatment options for renal fibrosis, none of them can achieve satisfactory results (7). A new comprehensive and effective treatment strategy is urgently needed clinically.

Mesenchymal stem cells (MSCs) are a heterogeneous subset of stromal stem cells with multiple surface markers

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(CD13, CD29, CD90, CD71 and CD105, etc.) that can be isolated from many adult tissues. They can differentiate into cells of the mesodermal lineage, such as adipocytes, osteocytes and chondrocytes, as well as cells of other embryonic lineages (8). It has been proved that MSCs can effectively promote tissue regeneration and repair (9-11), and MSCs have also been widely used in the treatment of kidney diseases (12). Studies have shown that exosomes derived from MSCs play important roles in protecting the kidney (13). However, excessive in vitro passage of MSCs leads to cell differentiation, which will seriously affect the composition of MSC-derived exosomes. Tissue repair requires large amounts of exosomes, which places high demands on the number of MSCs, requiring staff to continuously extract MSCs, which undoubtedly increases the workload. Pluripotent stem cell (PSC)-derived mesenchymal stem cells (PSC-MSCs) have been shown to have the characteristics of both MSCs and PSCs (14). However, the function of PSC-MSC remains controversial. For example, some scholars have used PSC-MSC-derived exosomes to treat bone defects (15). While Sfougataki et al. (16) claimed that PSC-MSCs were inefficient in their ability to form hyaline cartilage, which confirmed its limitation. Although multiple studies focused on the MSC-derived exosomes, the effect of PSC-MSC-derived exosomes on the reversal of renal fibrosis and the related mechanisms are still unclear.

Sirtuin 6 (SIRT6) is a conserved nicotinamide adenine dinucleotide-dependent protein deacetylase that has also been suggested to effectively reverse the fibrosis process in many organs, such as myocardial fibers. In kidney disease, Muraoka et al. (17) found that the Nampt-SIRT6 axis plays an active and protective role in the deposition of extracellular matrix in diabetic nephropathy. Liu et al. (18) demonstrated that SIRT6 was downregulated in diabetic nephropathy and adriamycin-induced nephropathy models. Cai et al. (19) found that SIRT6 could block the expression of β -catenin target genes to protect against renal fibrosis.

Exosomes (characterized by surfaced marker CD63, CD73, etc.) are a subset of EVs with an average diameter of ~ 100 nanometers, which are synthesized from endosomes. Through the interactions with other intracellular vesicles and organelles, the final content of the exosomes was determined. The constituents include nucleic acids, proteins, lipids, amino acids, and metabolites. Recently, it has been reported that exosomes play key roles in intercellular communication (20). In the present study, we investigated the antagonistic effects of different concentrations of PSC-MSC-derived exosomes on renal fibrosis. At the same

time, we suggested that PSC-MSC-derived exosomes exert protective effects on the kidney through the SIRT6/ β -catenin signaling pathway.

Materials and Methods

Generation of PSC-MSC-Exosomes

Human-induced PSC-MSCs were purchased from the Fourth Military Medical University and cultured in MGro- 500 chemically defined serum-free MSC medium (StemRD, Burlingame, USA). The cells of $3 \sim 8$ passages were applied in this study. After being cultured for 3 days, the culture medium of PSC-MSCs was collected and centrifuged at 300×g for 10 minutes and at 2,000×g for 10 minutes to remove dead cells and cellular debris. Then, the supernatant was filtered through a 0.22- μ m filter sterilizer (SteritopTM; Millipore, Billerica, USA) to remove the residual cellular debris. Subsequently, the supernatant was centrifuged at 4,000×g and reduced to approximately 200 µ1 by ultrafiltration in a 15 ml Amicon Ultra-15 centrifugal filter unit (Millipore, Billerica, USA). The ultrafiltrated liquid was washed twice with 15 ml of phosphate-buffered saline (PBS) and ultrafiltrated again at 4,000×g to 200 μ l. For exosome purification, the liquid was overlaid on a 30% sucrose-D2O cushion in a sterile Ultra-ClearTM tube (Beckman Coulter, Kraemer Boulevard Brea, USA) and ultracentrifuged at 100,000×g for 2 hours to pellet the small vesicles corresponding to exosomes. The pelleted exosomes were resuspended in 15 ml of PBS and centrifuged at 4,000×g in centrifugal filter units until the final volume was reduced to approximately 200 µ1. All procedures were performed at 4°C. The cell characterization was performed through FACS to determine the expression of CD90, CD45, CD105 and CD19 (all antibodies above were purchased from Abcam, Cambridge, UK). The characterization for exosomes was performed by TEM (transmission electron microscope), Western blot and NTA (Nanoparticle tracking analysis, NanoSight LM10 system, Malvern instruments) according to Théry's guideline (21). In this study, medium after culturing for 3 days as Control in Western blot.

Cell culture and experimental groups

The NRK-52E renal tubular epithelial cell line (ATCC[®] CRL-1730TM) was cultured in Dulbecco's modified Eagle's medium (DMEM, HyClone, USA) supplemented with 5% fetal bovine serum (FBS, BioCytoSci, TX, USA) and 1% antibiotics at 37°C in a humidified 5% CO₂ incubator. Cells were seeded on 6-well plates (Corning, NY, USA) at a density of 1×10^5 cells/ml and incubated with the follow-

ing factors: (1) pure DMEM (Con); (2) medium containing TGF- β 1 (5 ng/ml for 48 h, TGF- β 1); (3) medium containing TGF- β 1 and exosomes (10⁶ particles/ml, Exos-L); (4) medium containing TGF- β 1 and exosomes (10⁷ particles/ml, Exos-M); and (5) medium containing TGF- β 1 and exosomes (10⁸ particles/ml, Exos-H).

Quantitative real-time PCR

NRK-52E cells were cultured under different conditions and collected to evaluate the transcriptional level of related genes on day 2. Briefly, total RNA was extracted according to the instructions for the RNA extraction kit (TRIzol reagent, Invitrogen Life Technologies). The obtained RNA was then used to prepare cDNA by reverse transcription using PrimeScript RT master mix (TaKaRa, Tokyo, Japan), and a Bio-Rad CFX96 PCR manager system (Bio-Rad, Hercules, CA) was used to perform real-time PCR. The primers used in this study are listed in Table 1, and GAPDH was used as a housekeeping gene. The relative gene expression was normalized to that of the housekeeping gene GAPDH through the $\Delta \Delta Ct$ method.

Western blotting

Total protein samples were prepared from kidney tissue and cultured NRK-52E cells using Cell Lytic M (Sigma-Aldrich, St. Louis, MO) according to the instructions. The specimens were stored at -80° C. The primary antibodies used in the study were as follows: anti-E-cadherin (Abcam, Cambridge, UK), anti-Coll-A1 (Abcam, Cambridge, UK), anti- α -SMA (Abcam, Cambridge, UK), anti-SIRT6 (Abcam, Cambridge, UK) and anti- β -catenin (Abcam, Cambridge, UK). β -actin (Sigma-Aldrich) was used to control for protein loading.

Table 1. List of genes and primer sequences

Gene	Primer	Primer (5'-3')	
Col1-A1	Forward	GAGTGAGGCCACGCATGA	
	Reverse	AGCCGGAGGTCCACAAAG	
α -SMA	Forward	CCCCTGAAGAGCATCGGACA	
	Reverse	TGGCGGGGACATTGAAGGT	
Fibro	Forward	TACCAGAGGGGAAAAACGCC	
	Reverse	GCAGCGGTACAGATCCGAATC	
PAI-1	Forward	CACAGGCACTGCAAAAGGTC	
	Reverse	TGTGCCGAACCACAAAGAGA	
Fsp1	Forward	AGCTACTGACCAGGGAGCTG	
	Reverse	CATTGTCCCTGTTGCTGTC	
Axin2	Forward	CGACGCACTGACCGACGATT	
	Reverse	CGAAGGCAGCAGGTTCCACAG	
GAPDH	Forward	CTGATGCCTCCATGTTTGTG	
	Reverse	GGATGCAGGGATGATGTTCT	

Animals and surgical protocol

The animal protocols were approved by the Animal Research Committee of the Xi'an Peihua University (No. IACUC-20190402). Thirty male C57BL/6 mice applied from the Fourth Military Medical University were used for this study and were randomly divided into 4 groups. (1) Mice without any treatment were used as the Con group. (2) After anesthesia, the left abdomen was opened to view the kidney, and the ureter was located inside the lower pole of the left kidney. A 4-0 suture was used to ligate the upper pole of the free ureter near the calyces, and then the ureter was removed (UUO group). (3) After anesthesia, the left abdomen was opened to view the kidney, and then the abdominal cavity was closed (Sham group). (4) PSC-MSC-Exosomes with particles amounts of about 1×10^{11} were injected into UUO mice via the tail vein (Exo-H group).

Creatinine and blood urea nitrogen

Blood was collected from the hearts of mice in each group. The blood samples were coagulated for 2 hours at room temperature (RT) and centrifuged for 20 minutes at 2,000×g. The plasma was collected, and blood urea nitrogen (BUN) and serum creatinine were analyzed. The BUN level in plasma was measured using a BUN-E kit (Asan Pharm, Seoul, Korea), and the serum creatinine level was measured using a QuantiChrom creatinine assay kit (Bioassay Systems, Hayward, CA). Sample analysis was performed according to the manufacturer's recommended protocols.

Immunofluorescence staining

NRK-52E cells were seeded on glass coverslips, washed with PBS and fixed in cold methanol : acetone (1 : 1) for 10 minutes at -20°C. After being washed with phosphatebuffered saline, the fixed cells were blocked in 10% normal goat serum for 30 minutes. Then, the cells were incubated with primary antibodies (SIRT6, Abcam, Cambridge, UK) overnight, followed by incubation with goat anti-rabbit IgG conjugated with Alexa Fluor 594 or 488 (Abcam, Cambridge, UK) for 1 hour at 37°C. The cells were also stained with 4',6-diamidino-2-phenylindole to visualize the nuclei. Eventually, the slides were examined under a ZEISS LSM780 confocal microscope (Oberkochen, Germany).

Histology and immunohistochemical (IHC) staining

The kidney tissues were fixed in 10% formalin at RT. The paraffin-embedded sections were deparaffinized with xylene and dehydrated in gradually decreasing concentrations of ethanol. For immunohistochemical analysis, the dehydrated tissue sections were immersed in 10 mM sodium citrate buffer (pH 6.0) for 5 minutes at 95°C. The last step was repeated using fresh 10 mM sodium citrate solution. The sections were incubated in the same solution while cooling for 20 minutes, and they were then rinsed in phosphate-buffered saline PBS. Next, the sections were incubated with a primary antibody (fibronectin, Abcam, 1 : 100 dilution) for 1 hour at 37°C. After three serial washes with PBS, the sections were processed by an indirect immunoperoxidase technique using a commercial EnVision System kit (DAKO, Carpinteria, CA, USA). Immunohistochemical images were viewed with an Eclipse 80i microscope (Nikon, Tokyo, Japan).

Enzyme-linked immunosorbent assay (ELISA)

Blood was collected from the mice, and the serum was separated. The concentrations of interleukin (IL)-6 and TNF- α in mouse serum were quantified using enzymelinked immunosorbent assay (ELISA) kits (R&D Systems, Hangzhou, China) according to the manufacturer's instructions. The absorbance at 450 nm was measured using an ELISA reader (BMG Labtech, Ortenaukreis, Germany).

Statistical analysis

All data are presented as the means \pm SEM and were subjected to one-way analysis of variance (ANOVA) with Tukey's multiple comparison tests. Statistical significance was examined using GraphPad Prism 5 (GraphPad Software, Inc., San Diego, CA, USA). A value of p<0.05 indicated a significant difference.

Results

Purification, isolation, and characterization of exosomes and PSC-MSCs

The results of characterization for PSC-MSCs were listed in Supplementary Fig. S1 while results of PSC-MSC-Exosomes characterization were listed in Supplementary Fig. S2. The results showed that the expression of CD90 and CD105 was 99.3% and 99.8% respectively, which confirmed that the cells meet the standards of MSCs. PSC-MSCs retained the proliferative ability for at least 40 days (Supplementary Fig. S1). Furthermore, results of nanoparticle tracking analysis (NTA) and transmission electron microscopy (TEM) revealed that the exosomes obtained from PSC-MSCs mainly occupied a diameter of $10 \sim 150$ nm and presented the typical morphological features of exosomes. Results of western blot showed that all exosome specimens were positive for exosome markers (CD63 and CD73), negative for the cellular marker GM130.

Antifibrotic effect of PSC-MSC-Exosomes on TGF- β 1-treated NRK-52E cells

To confirm the preventive effect of PSC-MSC-Exosomes on epithelial dedifferentiation processes, we investigated the protein and gene expression of TGF- β 1-treated NRK-52E cells. In this study, the protein expression of Col1-A1 and α -SMA was increased, and E-cadherin was decreased by TGF- β 1 administration (Fig. 1A). In addition, after the administration of PSC-MSC-Exosomes, we found that Col1-A1 and α -SMA expression was decreased and E-cadherin was increased compared with those of the TGF- β 1 group. Moreover, with increasing exosome con-



Fig. 1. PSC-MSCs-Exosomes inhibit the occurrence of epithelial differentiation in TGF- β 1-treated NRK-52E cells. (A) Western blot analysis of Col1-A1, *a*-SMA and E-cadherin protein expression in NRK-52E cells under different conditions. (B) Real-time PCR analysis of Col1-A1, α-SMA and Fibro gene expression in NRK-52E cells under different conditions. (C) Representative morphology of cells in the different groups. Bar = 100 μ m. n=3 in each group, ${}^{a}p < 0.05$ vs. the Con group, ${}^{b}p < 0.05$ vs. the TGF- β 1 group, ^cp<0.05 vs. the Exo-L group, $^{d}p < 0.05$ vs. the Exo-M group.

centrations, Coll-A1 and α -SMA expression showed a decreasing trend, while E-cadherin showed an increasing trend (Fig. 1A). Subsequently, real-time PCR showed that profibrotic genes, including Coll-A1, α -SMA and Fibro, were significantly increased in the TGF- β 1 group and decreased in the Exo groups (p<0.05), and there was a decreasing trend from the Exo-L to Exo-H group (p<0.05) (Fig. 1B). As shown in Fig. 1C, in the absence of TGF- β 1 exposure, NRK-52E cells presented a typical epithelial morphology and formed a cobblestone monolayer. After treatment with TGF- β 1 for 24 hours, NRK-52E cells exhibited fibroblast-like shuttle-type morphology. We also used Exo-H-treated TGF- β 1-treated NRK-52E cells and found that the morphology of the cells was between the cobblestone monolayer and shuttle type.

PSC-MSC-Exosomes attenuated fibrotic processes in UUO mice

We used IHC staining to examine fibronectin expression in UUO mice. As shown in Fig. 2A, we found that the expression of fibronectin in the control and sham groups was not different but was significantly increased in UUO mice and Exo-treated UUO mice. However, the expression of fibronectin in the Exos-H group was decreased compared with that in the UUO group. Furthermore, we analyzed the fibronectin-positive areas of the different groups. As shown in Fig. 2B, The UUO group $(11.35\% \pm$ 1.79) had the largest area of the 4 groups (p<0.05), and the area in the Exos-H group (5.47%±0.86) was also larger than that of the Con $(1.79\% \pm 0.19)$ and Sham $(1.99\% \pm$ 0.28) groups (p<0.05).

PSC-MSC-Exosomes preserved kidney function and inhibited inflammation in UUO mice

Inflammation plays a crucial role in the development of kidney fibrosis. To confirm the anti-inflammatory effects of PSC-MSC-Exosomes, inflammatory cytokines, including TNF- α and IL-6, were measured by ELISA (Fig. 3A). The secretion of TNF- α and IL-6 was increased in UUO mice (p<0.05). Conversely, treatment with Exosomes



Fig. 2. PSC-MSC-Exosomes inhibited renal fibrosis in UUO mice. (A) Immunohistochemical staining showed that the expression of fibronectin was inhibited by PSC-MSC-Exo administration in UUO-injured mice. (B) Quantitative results of fibronectin expression. Bar=100 μ m. n=3 in each group; ^ap<0.05 vs. the Con group; ^bp<0.05 vs. the UUO group.

Fig. 3. The effect of PSC-MSC-Exosomes on kidney failure and renal inflammation in UUO mice. (A) TNF- α levels in the different mice. (B) IL-6 levels in the different mice. (C) BUN levels in the different mice. (D) Serum creatinine levels in the different mice. n=6 in each group; ^ap< 0.05 vs. the Con group; ^bp<0.05 vs. the UUO group. attenuated the inflammatory response ($p \le 0.05$).

Kidney function decreases in mice with renal fibrosis, and the levels of serum creatinine and BUN are crucial in helping to evaluate kidney function. Fig. 3C and 3D and Table 2 showed that UUO mice developed serious renal dysfunction compared with normal mice, as indicated by significant increases in serum creatinine and BUN levels. However, the level of BUN was significantly decreased in the Exo-H group at day 7 and day 14 (p<0.05) and was similar to the normal level compared with that of the Con and Sham groups (p>0.05). Moreover, the serum creatinine level was also decreased in the Exo-H group compared with the UUO group (p<0.05), although it was higher than that in the Con group at day 14 (p<0.05).

PSC-MSC-Exosomes inhibited renal fibrosis through the SIRT6/ β -catenin signaling pathway

To identify the mechanism by which PSC-MSC-Exosomes inhibited renal fibrosis, this study used Western blotting

to measure the expression of SIRT6 and β -catenin in UUO mice. Fig. 4A shows that obstructive injury significantly decreased the expression of SIRT6, whereas Exo treatment reversed these changes in expression levels. While the results in Supplementary Fig. S3 indicated that the synthesis of β -catenin presented the opposite tendency. We also obtained similar results in vitro (Fig. 4B and 4C). However, the expression of β -catenin, a downstream protein of SIRT6, was downregulated in the Exo-H group, while the expression level in the TGF- β 1 group was between that of the the Con and TGF- β 1 groups (Fig. 4B). These results indicate that Exosomes can inhibit the expression of β -catenin by upregulating SIRT6. Subsequently, we used real-time PCR to measure the expression of genes downstream of β -catenin and found that PAI-1, Fsp1 and Axin2 were all upregulated in the TGF- β 1 group (p<0.05) and significantly decreased in the Exo-H group (p < 0.05).

Table 2. Renal function of mice in the different groups (mean ± SEM)

Group	n -	BUN (mmol/l)		Creatinine (µmol/l)	
		Day 7	Day 14	Day 7	Day 14
Con	6	5.738 ± 0.972	6.247 ± 1.047	25.317 ± 2.558	27.126 ± 3.013
Sham	6	5.213 ± 1.034	6.132 ± 1.217	26.038 ± 2.731	27.996 ± 2.698
UUO	6	11.334 ± 1.361	11.986 ± 1.163	39.442 ± 3.021	44.568 ± 3.139
Exo-H	6	7.439 ± 0.987	8.035 ± 0.972	30.176 ± 2.967	35.698 ± 2.847



Fig. 4. Mechanism by which exosomes inhibit fibrosis. (A) Western blot analysis of SIRT6 protein expression in the different mice. (B) Western blot analysis of SIRT6 protein expression in NRK-52E cells under different conditions. (C) Representative immunofluorescence images showing SIRT6 in NRK-52E cells under different conditions. (D) PAI-1, Fsp1 and Axin2 gene expression in NRK-52E cells under different conditions. Bar = 50 μ m. n=3 in each group; ${}^{a}p < 0.05$ vs. the Con group; ^bp < 0.05 vs. the TGF- β 1 group.

Discussion

Currently, there is still a lack of clinically effective therapeutic drugs for renal fibrosis, leading to a large number of renal fibrosis patients inevitably developing end-stage renal disease. The pathological characteristics of renal fibrosis mainly include excessive formation of extracellular matrix, infiltration of inflammatory cells or the production of inflammatory factors, tubular cell dedifferentiation and excessive activation of myofibroblasts. Therefore, the main reason for the difficulty in treating renal fibrosis is that the mechanism of its development is complex, and interacting signaling pathways easily form a compensation mechanism. There is also a lack of effective indicators to evaluate the effect of drugs on renal fibrosis. Most of the currently used drugs have relatively negative side effects, which seriously affect patient compliance. For example, although ACEI (Angiotensin-Converting Enzyme Inhibitors) can effectively treat renal fibrosis, long-term prognosis is still poor (22). Therefore, when developing a new therapeutic agent, we must not only ensure the long-term safety of the drug but also ensure the effectiveness of the evaluation plan. It is best to clarify the molecular mechanism of the drug to protect the kidney.

Studies have shown that exosomes have therapeutic potential in various diseases, such as the repair of bone defects (23), nerve regeneration (24) and the treatment of myocardial infarction (25), in the absence of the original cells. Exosomes have been proven to be effective in the treatment of kidney diseases (26). Zhou et al. (27) found that exosomes released by human umbilical cord mesenchymal stem cells (huc-MSCs) protect against cisplatin-induced renal oxidative stress and apoptosis, as well as stimulate kidney cell proliferation by the ERK1/2 signaling pathway. Nagaishi et al. (28) used bMSC (bone marrow stem cells)-derived exosomes to treat diabetic nephropathy and found that pathological kidney tissue damage was improved, proinflammatory factors were decreased, and renal tubular EMT was alleviated. Wang et al. (29) verified that MSCs were able to selectively transfer miR-let7c to damaged kidney cells through exosomes, furthermore paving the way for the use of MSCs to delivery miRNA as a targeted therapy for kidney disease. The underlying mechanism of the positive effects is that exosomes contain growth factors, cytokines, chemokines, proteins, mRNAs, and miRNAs. Therefore, the use of exosomes to treat kidney diseases has good prospects. However, exosomes are a kind of cell-derived nanoparticle, and their functional effects and contents are largely restricted by cell type. The

therapeutic effects of MSCs and MSC-Exosomes on kidney disease need not be discussed further. Nevertheless, MSCs will differentiate during the culture process, which affects the function and content of exosomes. Thus, collecting enough exosomes for kidney repair relies on abundant sources of parent cells. Recently, the effectiveness of the application of human-induced pluripotent stem cell-derived mesenchymal stem cells (hiPSC-MSC-Exosomes) in bone defect repair and regeneration has been confirmed. More importantly, hiPSC-MSCs do not differentiate like MSCs but still retain efficacy of MSCs after culturing for 8-9 passages.

TGF- β 1-treated NRK-52E cells are often used as an *in* vitro culture model of renal fibrosis. In this study, we used different concentrations of PSC-MSC-Exosomes to treat TGF- β 1-treated NRK-52E cells and found that PSC-MSC-Exosomes could effectively reverse TGF- β 1-treated cell fibrosis in vitro. We believed this effect was exerted via multiple pathways such as the suppression of inflammation caused by the mRNA and proteins contained in exosomes (30). Moreover, higher concentrations led to better effects. Furthermore, the morphology of TGF- β 1-treated NRK-52E cells treated with PSC-MSC-Exosomes was closer to the morphology of normal NRK-52E cells. To further confirm the antifibrotic effect of Exosomes in kidneys, we used UUO mice for in vivo experiments. Immunohistochemical staining of kidney sections and ELISA analysis showed that exosomes could partially reverse renal fibrosis and reduce inflammation in the kidney, further improving the kidney function of UUO mice.

SIRT6, a protein that has been reported can protect organs from fibrosis in different ways (31). β -catenin is considered to be an important link in the process of renal fibrosis (32). Activated β -catenin can significantly upregulate fibrosis-related proteins, such as fibronectin (FN), matrix metalloproteinase 7 (MMP7), and fibroblast-specific protein 1 (Fsp 1). Recently, Cai et al. (19) found that SIRT6 protects against kidney fibrosis by epigenetically blocking β -catenin target gene expression. Similarly, in our in vitro experiments, SIRT6 was downregulated and β -catenin was upregulated in TGF- β 1-treated NRK-52E cells. Moreover, exosomes derived from PSC-MSCs could increase SIRT6 and reduce β -catenin in TGF- β 1-treated NRK-52E cells. In vivo, SIRT6 expression in Exosomestreated UUO mice was lower than that in control mice, but it was significantly higher than that in unexposed UUO mice. Therefore, we concluded that the protective effect of PSC-MSC-Exosomes on renal fibrosis occurred through upregulating SIRT6, thus downregulating the expression of β -catenin and further reducing the expression of downstream products of β -catenin (Fig. 4, Supplementary Fig. S3).

In summary, this study confirmed that PSC-MSC-Exosomes have a robust protective effect against renal fibrosis, and higher concentrations of Exos were associated with better protective effects. Moreover, this protection was achieved by regulating the SIRT6/ β -catenin signaling axis.

In conclusion, PSC-MSC-Exosomes can effectively inhibit endothelial cell injury, reduce inflammatory reaction, protect renal function and slow down renal fibrosis, and this effect is achieved by up regulation of SIRT6, thereby inhibiting the expression of β -catenin and its downstream products.

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Potential Conflict of Interest

The authors have no conflicting financial interest.

Supplementary Materials

Supplementary data including three figures can be found with this article online at https://doi.org/10.15283/ijsc20184.

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