

Protective effects of human induced pluripotent stem cell-derived exosomes on high glucose-induced injury in human endothelial cells

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Abstract. Exosomes are a family of extracellular vesicles that are secreted from almost all types of cells and are associated with cell-to-cell communication. The present study was performed to investigate the effects of human induced pluripotent stem cell-derived exosomes (hiPSC-exo) on cell viability, capillary-like structure formation and senescence in endothelial cells exposed to high glucose. Exosomes were isolated from the conditional medium of hiPSCs and confirmed by transmission electron microscopy, nanoparticle tracking analysis and western blot analysis using Alix and cluster of differentiation-63 as markers. hiPSC-exo were labeled with PKH26 for tracking, and it was determined that spherical exosomes, with a typical cup-shape, were absorbed by human umbilical vascular endothelial cells (HUVECs). Cultured HUVECs were treated with high glucose (33 mM) with or without hiPSC-exo (20 μ g/ml) for 48 h, and cell viability, capillary tube formation and senescence were assessed. When exposed to high glucose, viability and tube formation in HUVECs was significantly reduced ($P < 0.0001$), whereas the proportion of senescent cells was higher compared with that in control HUVECs ($P < 0.0001$). Furthermore, hiPSC-exo restored cell viability and capillary-like structure formation, and reduced senescence in HUVECs exposed to high glucose ($P < 0.0001$). However, hiPSC-exo had minimal effects on normal HUVECs. These findings suggest that stem

cell-derived exosomes are able to promote cell proliferation, enhance capillary-like structure formation and reduce senescence in endothelial cells exposed to high glucose.

Introduction

Patients with type 2 diabetes mellitus are at a notably high risk for developing life-threatening cardiovascular complications (1), and there is a clear association between glycemic control and cardiovascular diseases (2). Unlike vascular smooth muscle cells, endothelial cells cannot regulate intrinsic glucose levels, which can result in the accumulation of glucose and its derivatives, leading to an array of metabolic disorders. Furthermore, glucose toxicity can cause reduced cell viability and increased senescence in endothelial cells via multiple signaling pathways. The cardiovascular complications of diabetes have a close association with the function of endothelial cells; however, the underlying mechanism needs further investigation (3-5). Therefore, endothelial cells are potential targets for preventing the cardiovascular complications of diabetes.

Induced pluripotent stem cells (iPSCs) may be used to treat a number of diseases, as they possess the potential for self-renewal and multi-differentiation (6). It has previously been suggested that transplanted iPSCs were able to inhibit vascular apoptosis and fibrosis, thus improving cardiac function in diabetic rats (7). However, little is known about the mechanisms by which iPSCs, or factors released from these cells, inhibit adverse cardiac remodeling (8). Despite their impressive therapeutic ability, teratoma formation has been observed following transplantation of iPSCs (9,10). iPSC-derived lineage cells have avoided this issue; however, the derivatives may still suffer the same difficulties as reported for adult stem cells, particularly in cell survival, retention and coupling in damaged areas (11). Therefore, it is important to exploit the powerful regenerative capacity of pluripotent stem cells while circumventing the problems associated with cell transplantation.

The discovery of cell-free components, including exosomes, may provide a promising alternative for regenerative medicine. Exosomes are small membrane vesicles that contain membrane and cytosolic components, including proteins,

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Abbreviations: hiPSC-exo, human induced pluripotent stem cell-derived exosomes; HUVECs, human umbilical vascular endothelial cells

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lipids and RNAs (12-14). They have an essential role in intercellular communication via transporting this cargo to targeted cells. Furthermore, the role of exosomes depends on their components and cell origin. It has been reported that adipose-derived mesenchymal stromal cells release exosomes that are capable of promoting angiogenesis (15). In addition, embryonic stem cell-derived exosomes have been shown to enhance cell proliferation (16). As one of the most dynamic cells with regenerative potential, iPSCs may also release exosomes, suggesting further potential for disease treatment.

In the present study, human induced pluripotent stem cell-derived exosomes (hiPSC-exo) were isolated and were co-cultured with normal human umbilical vascular endothelial cells (HUVECs) or HUVECs exposed to high glucose. Subsequently, the cell viability, capacity to form capillary-like structures and cell senescence was examined to demonstrate the role for hiPSC-exo in endothelial cell growth.

Materials and methods

Cell culture. HiPSCs were induced at the Cardiovascular Regenerative Engineering Laboratory (Shanghai, China), following a previously described protocol (17) and cultured in mTeSR (Stemcell Technologies, Inc., Vancouver, Canada), which contains mTeSR1 basal medium (400 ml; cat. no. 05851) and mTeSR1 5X supplement (100 ml; cat. no. 05852). The culture medium was replaced every day, and the cells were digested with Accutase (cat. no. A6964; Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) and passaged at a 1:5 dilution every 3 days. Following the culture of passaged cells for 48 h at 37°C in a 5% CO₂ and 95% air gas mixture atmosphere, the media was changed. The media was then collected for exosome isolation following a further incubation with cells at 37°C in a 5% CO₂ and 95% air gas mixture atmosphere for 24 h.

Primary HUVECs were isolated from fresh umbilical cord veins from 3 patients that had undergone normal pregnancy, 24 h following birth, from June 2015 to December of 2015 at the Department of Obstetrics, The Obstetrics and Gynecology Hospital of Fudan University (Shanghai, China). For the collection of HUVECS, informed consent was obtained from all patients and ethical approval was granted by the Experimental Animal and Ethics Committee of the College of Basic Medical Sciences at Fudan University (Shanghai, China). HUVECs were isolated by collagenase digestion as previously described (18) and cultured in endothelial cell medium (ScienCell Research Laboratories, Inc., San Diego, CA, USA) supplemented with 5% fetal bovine serum (cat. no. 0025), 1% endothelial cell growth supplement (cat. no. 1052) and 1% penicillin/streptomycin solution (cat. no. 0503). Furthermore, the culture medium was replaced every 2 days. When the HUVECs became ~80% confluent, cells were sub-cultured with 0.25% trypsin-EDTA and phenol red 1X (cat. no. 25200072; Thermo Fisher Scientific Inc., Waltham, MA, USA), and were incubated at 37°C in an atmosphere containing 5% CO₂.

Exosome isolation. Exosomes from hiPSC were collected and purified using ultracentrifugation. Briefly, timing began when hiPSCs were 70% confluent. After 48 h, 80 ml cell-conditioned medium was collected and centrifuged at 3,000 x g for 30 min

at 4°C and 10,000 x g for 30 min at 4°C to remove dead cells and cell debris. The final supernatant was ultracentrifuged and exosomes were pelleted at 100,000 x g for 70 min at 4°C. The pellet obtained was then washed with PBS at 100,000 x g for 70 min at 4°C to eliminate contaminating proteins. The purified exosome fraction was re-suspended in PBS prior to further use.

HiPSC-exo labeling and uptake assay. Exosomes were labeled using a PKH26 Red Fluorescent Cell Linker kit (PKH26GL-1KT; Sigma-Aldrich; Merck KGaA) according to the manufacturer's protocol. A total of 1 µl PKH26 was mixed with 250 µl Diluent C (provided by the kit). HiPSC-exo with the same volume was then added into the mixture and incubated at room temperature for 5 min. Labeling was stopped by adding 500 µl 0.5% bovine serum albumin followed by an incubation at room temperature for 5 min. The exosomes were then centrifuged at 100,000 x g for 70 min at 4°C and re-suspended in PBS prior to the uptake assay. For the uptake assay, HUVECs were seeded in 96-well plates and left to proliferate for 24 h at 37°C in a 5% CO₂ and 95% air gas mixture. Following 24 h, 20, 50 µg/ml hiPSC-exo or the PBS control were added into the dish and cultured for a further 24 h at 37°C in a 5% CO₂ and 95% air gas mixture. The following steps were applied to the cells prior to observation under an inverted phase contrast microscope at a magnification of x20. The cells were washed with PBS twice, fixed with freshly prepared 4% paraformaldehyde in PBS at room temperature for 15 min, further washed twice with PBS and incubated with 0.1% triton X-100/PBS at room temperature for 15 min. Cells were subsequently washed twice with PBS, stained with DAPI (1:5,000; cat. no. C1002; Beyotime, Shanghai, China) for 5 min at room temperature and further washed with PBS.

Transmission electron microscopy. Transmission electron microscopy images of hiPSC-exo were obtained using an FEI Tecnai G2 Spirit twin transmission electron microscope (FEI; Thermo Fisher Scientific, Inc.) operating at 80 kV. Briefly, 10 µl hiPSC-exo sample was loaded onto a formvar-carbon coated copper grid for 2 min and excess sample was absorbed with filter paper by gently touching the edge of the grid and removed. Next, uranyl acetate was used to stain the grid for 1 min at room temperature and the excess liquid was absorbed using filter paper. The copper grid was then dried under an incandescent lamp.

Nanoparticle tracking analysis. Exosome size and concentration analysis was performed using Nanoparticle Tracking Analyzer (version, 3.1; Build, 3.1.54; Malvern Instruments, Ltd., Worcestershire, UK). For the measurement, 10 µl hiPSC-exo sample was diluted in 1 ml of PBS.

Cell counting assay. A cell counting kit-8 (CCK-8; cat. no. CK04; Dojindo Molecular Technologies, Inc., Kumamoto, Japan) was used to identify the effect of hiPSC-exo on the viability of HUVECs. For this assay, the same initial number (1x10³) of HUVECs was seeded in 96-well plates and allowed to proliferate for 24 h at 37°C, in a 5% CO₂ and 95% air gas mixture. The cells were then treated with different conditions [normal glucose (5 mM); normal glucose + hiPSC-exo (10, 20 or 50 µg/ml);

high glucose (33 mM) and high glucose + hiPSC-exo (10, 20 or 50 $\mu\text{g}/\text{ml}$) for a further 48 h at 37°C in a 5% CO₂ and 95% air gas mixture. The culture media were aspirated and CCK-8 mixed media (CCK-8: Dulbecco's modified Eagle's medium, 1:10) was added to the different groups. After 2 h, the absorbance was measured at 450 nm using a spectrophotometer microplate reader. The assays were performed in sextuplicate and each experiment was repeated five times.

Capillary-like structure formation assay. A capillary-like structure formation assay was performed to identify the functional role of hiPSC-exo on endothelial cells. Briefly, HUVECs (1×10^4 cells) were cultured in 24-well plates at 37°C in a 5% CO₂ and 95% air gas mixture for 24 h and subjected to different treatments (normal glucose (5 mM), normal glucose + hiPSC-exo (20 $\mu\text{g}/\text{ml}$), high glucose (33 mM) and high glucose + hiPSC-exo (20 $\mu\text{g}/\text{ml}$) for another 48 h at 37°C, in a 5% CO₂ and 95% air gas mixture. After 48 h, HUVECs were trypsinized and cultured in 24-well plates (1×10^5 cells/well) coated with 200 μl Matrigel matrix (cat. no. 356234; BD Biosciences, Franklin Lakes, NJ, USA). Tube length was quantified after 6 h by calculating the mean length in five random microscopic fields with an inverted phase contrast microscope at magnification, x10. ImageJ software (version, 1.42q) was then used to analyze the data (National Institutes of Health, Bethesda, MD, USA). Each experiment was repeated three times.

Senescence-associated staining. Senescence-associated staining was applied to demonstrate that hiPSC-exo could inhibit cell senescence in HUVECs. Briefly, HUVECs (1×10^4 cells) were cultured in 24-well plates at 37°C with 5% CO₂ for 24 h and treated with normal glucose (5 mM), normal glucose + hiPSC-exo (20 $\mu\text{g}/\text{ml}$), high glucose (33 mM) and high glucose + hiPSC-exo (20 $\mu\text{g}/\text{ml}$) for a further 48 h at 37°C, in a 5% CO₂ and 95% air gas mixture. Subsequently, senescence-associated β -galactosidase (SA- β -gal) staining was performed using an SA- β -gal staining kit (cat. no. K320-250; BioVision, Inc., Milpitas, CA, USA) according to the manufacturer's instructions. HUVECs were washed three times with PBS and fixed for 15 min at room temperature with fixative solution. After incubation with staining solution overnight at 37°C, senescence was quantified by calculating mean proportion of senescent cells in five random microscopic fields using an inverted phase contrast microscope at magnification, x20 and ImageJ software. Each experiment was repeated three times.

Western blot analysis. hiPSCs and hiPSC-exo were lysed using radioimmunoprecipitation assay buffer (cat. no. 89901; Thermo Fisher Scientific, Inc.) for 30 min on ice, followed by a centrifugation at a speed of 12,000 x g for 20 min at 4°C. The concentration of the protein was determined via bicinchoninic acid assay (Pierce BCA Protein Assay Reagent A; cat. no. 23227; Thermo Fisher Scientific, Inc.). A total of 20 μg protein was mixed with 5X SDS loading buffer and loaded onto 10% SDS-PAGE. The separated protein bands in the gel were transferred onto a polyvinylidene difluoride (PVDF) membrane. The membrane was then blocked with 5% non-fat milk in PBS with 0.1% Tween-20 (PBST) at room temperature for 2 h. PVDF membranes were then incubated with primary

antibodies at 4°C overnight and then washed with PBST. Secondary antibodies were added to each blot, and incubated at room temperature for 2 h; PVDF membranes were then washed with PBST and incubated with Super Signal™ West Pico (cat. no. 34580; Thermo Fisher Scientific, Inc.) and observed using a Western blot visualizer (Tanon 5500; Tanon Science & Technology Co., Ltd., Shanghai, China). Primary antibodies used for exosome identification were goat anti-Alix (N-20; 1:1,000; cat. no. sc-49267; Santa Cruz Biotechnology Inc., Dallas, TX) and anti-cluster of differentiation (CD63; rabbit IgG; 1:1,000; cat. no. EXOAB-CD63A-1; System Biosciences Inc., Palo Alto, CA). Secondary antibodies used were horseradish peroxidase (HRP)-conjugated donkey anti-goat IgG (1:2,000; cat. no. 705-035-003; Jackson ImmunoResearch Laboratories Inc., West Grove, PA) and HRP-conjugated donkey anti-rabbit IgG (1:2,000; cat. no. 711-005-152; Jackson ImmunoResearch Laboratories Inc.). The experiment was repeated three times.

Statistical analysis. Statistical analysis was performed using one-way analysis of variance and a post-hoc Bonferroni test. $P < 0.05$ was considered to indicate a statistically significant difference. Data are presented as the mean \pm standard error of the mean, unless otherwise stated. Furthermore, statistical analysis was performed using GraphPad Prism 6.07 (GraphPad Software, Inc., La Jolla, CA) and SPSS Statistics 17.0 (SPSS Inc., Chicago, IL, USA).

Results

Identification of exosomes derived from hiPSC. A transmission electron microscopic observation of hiPSC-exo showed the presence of spherical vesicles with a typical cup-shape (Fig. 1A). Upon conducting nanoparticle tracking analysis, the concentration/size graph and particle pictorial diagram revealed a homogeneous population of exosomes ranging between 50 and 150 nm (Fig. 1B and C). The mode size of hiPSC-exo was 103.1 nm, and the total concentration of particles with a diameter between 30 and 150 nm was 6×10^{10} particles/ml (Fig. 1D). Additionally, hiPSC-exo expressed the exosomal marker proteins Alix and CD63 (Fig. 1E).

Uptake of hiPSC-exo by HUVECs. In order to function, hiPSC-exo requires the ability to be endocytosed by target cells. Previous research has demonstrated that exosomes express adhesion molecules that may be associated with the adherence of exosomes to cells; however, the cellular and molecular basis for specific targeting to acceptor cells remains to be elucidated. For example, exosomes released by the human intestinal epithelial cell line T84 could be endocytosed by dendritic cells (DCs), but not B or T lymphocytes (19). The present study therefore tested if hiPSC-exo could be taken up by HUVECs. It was demonstrated in Fig. 2A and B that when HUVECs were treated with PKH26 labeled hiPSC-exo (20 and 50 $\mu\text{g}/\text{ml}$) exosomes were endocytosed by cells in a concentration-dependent manner. By contrast, the control group, which was subjected to the same procedure, did not show any intracellular fluorescence as shown in Fig. 2C.

hiPSC-exo reversed high glucose-induced decreased cell viability. Fig. 2 demonstrated that hiPSC-exo may be endocytosed

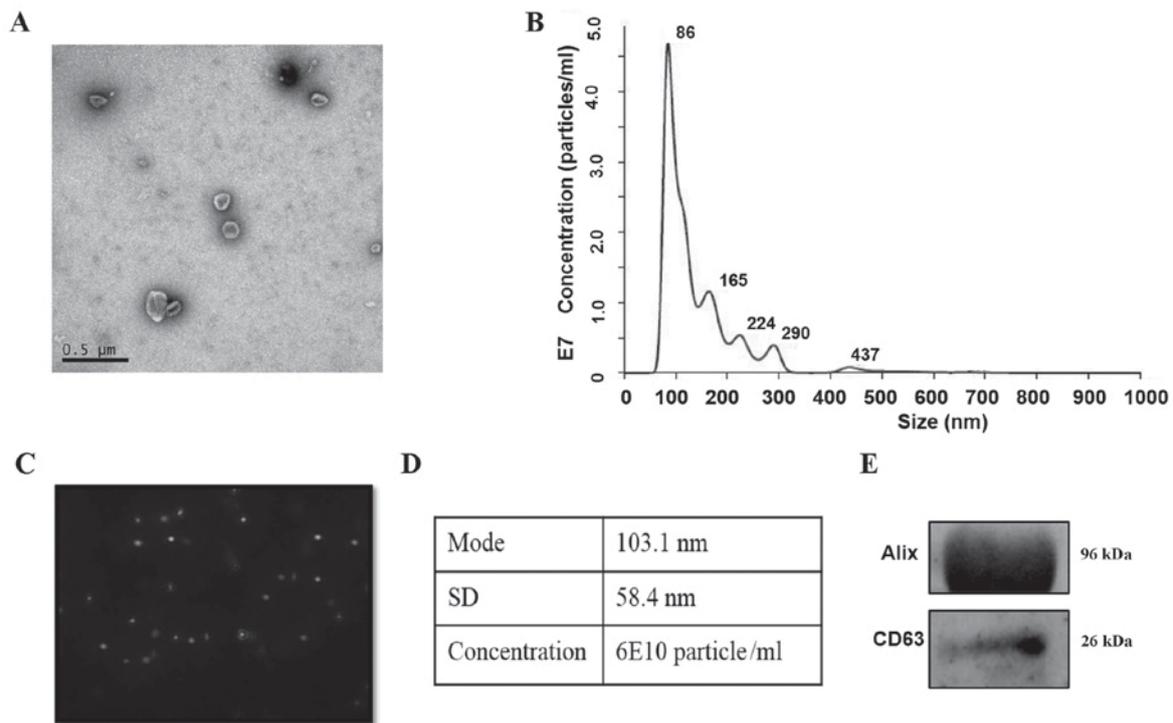


Figure 1. Characterization of hiPSC-exo. (A) Transmission electron microscopy of hiPSC-exo. Scale bar, 0.5 μm . (B-D) NTA of hiPSC-exo fraction. (B) A graph plotting the concentration vs. particle size was obtained from exosome measurements. (C) A particle pictorial diagram. (D) Mode particle size of hiPSC-exo obtained by NTA measurements. (E) Alix and CD63 expression in hiPSC-exo detected by western blot analysis. hiPSC-exo, human induced pluripotent stem cell-derived exosomes; NTA, Nanoparticle tracking analysis; SD, standard deviation; CD, cluster of differentiation.

by HUVECs. It was then investigated if endocytosed hiPSC-exo could influence the fate of HUVECs. In CCK-8 assay the amount of formazan dye in cells is directly proportional to the number of living cells. In the present study, HUVECs were treated with normal (5.5 mM) or high (33 mM) concentrations of glucose, in combination with different concentrations (0, 10, 20 or 50 $\mu\text{g}/\text{ml}$) of hiPSC-exo. After 48 h treatment, the OD value of the eight groups was measured. Compared with the control group [normal concentration of glucose (5.5 mM)], the OD value for high glucose (33 mM) treated HUVECs was significantly decreased (Fig. 3). Furthermore, exosomes had no statistically significant effect on normal HUVECs. HiPSC-exo significantly reversed the harmful effect of high glucose. However, the effects between hiPSC-exo 20 and 50 $\mu\text{g}/\text{ml}$ did not differ significantly (Fig. 3). Thus, a concentration of 20 $\mu\text{g}/\text{ml}$ was selected for use in subsequent experiments.

hiPSC-exo reversed high glucose-induced decreases in capillary-like structure formation. Following treatment with normal glucose, normal glucose + hiPSC-exo, high glucose or high glucose + hiPSC-exo, HUVECs were trypsinized and cultured on Matrigel to assess *in vitro* capillary-like structure formation. The results demonstrated that high glucose significantly decreased capillary-like structure formation in HUVECs, whereas hiPSC-exo reversed this effect. However, hiPSC-exo had a minimal effects on normal HUVECs (Fig. 4).

Anti-senescence effect of hiPSC-exo in high glucose-injured HUVECs. To verify the effects of hiPSC-exo on cell senescence, a senescence detection assay was performed on

HUVECs after being treated with different conditions (normal glucose, normal glucose + hiPSC-exo, high glucose and high glucose + hiPSC-exo). The cells of the control group were transparent and plump whereas the cells of the high glucose group showed a flattened and enlarged morphology (Fig. 5). hiPSC-exo significantly protected HUVECs against cellular senescence induced by high glucose (Fig. 5). Furthermore, there was no statistically significant difference between the normal glucose group and either group treated with hiPSC-exo. Collectively, the results showed that hiPSC-exo are readily absorbed by target cells, wherein they can modulate cell viability and survival.

Discussion

To the best of our knowledge, this is the first report demonstrating that exosomes derived from hiPSCs are able to protect HUVECs from high glucose *in vitro*. In the present study, HUVECs were found to be capable of absorbing hiPSC-exo at a high efficiency. The mechanism of exosome uptake by targeted cells has long been studied. The existing model suggests that exosomes express adhesion molecules to adhere to cells (20); however, the cellular and molecular basis for their specific targeting to acceptor cells remains to be elucidated. Barrès *et al* (21) demonstrated previously that galectin-5 was bound to the surface of rat reticulocyte exosomes and modulated vesicle uptake by macrophages. In the immune system, it has been demonstrated that T cells were able to recruit major histocompatibility complex class II-containing DC exosomes, and that this recruitment was dependent on leukocyte function-associated antigen-1 (22).

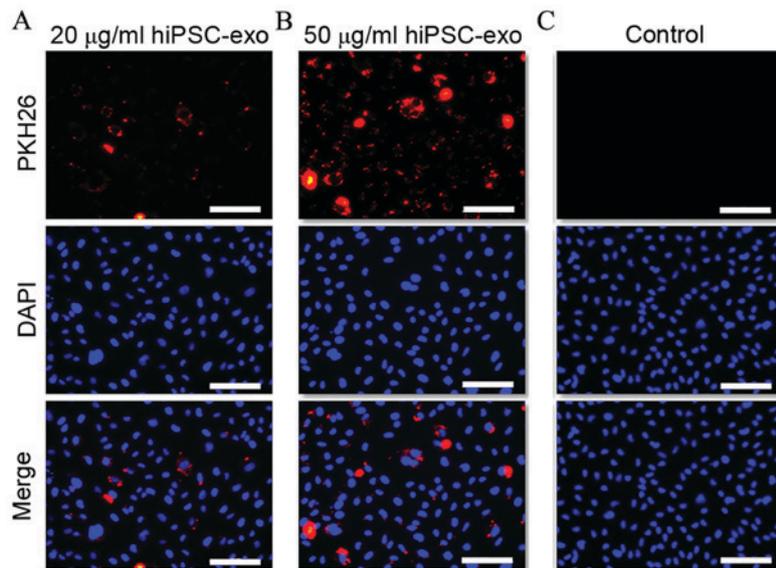


Figure 2. Uptake of hiPSC-exo by HUVECs. Inverted fluorescence microscopy confirmed the location of hiPSC-exo in HUVECs following 24 h treatment with (A) 20 or (B) 50 µg/ml hiPSC-exo. (C) The same methods were used to show the absence of hiPSC-exo or nonspecific localization of labeling dye (control conditions). Red, hiPSC-exo labeled by PKH26; blue, nucleus of HUVECs. Scale bar, 100 µm. hiPSC-exo, human induced pluripotent stem cell-derived exosomes; HUVECs, human umbilical vascular endothelial cells; DAPI, 4', 6-diamidine-2'-phenylindole dihydrochloride.

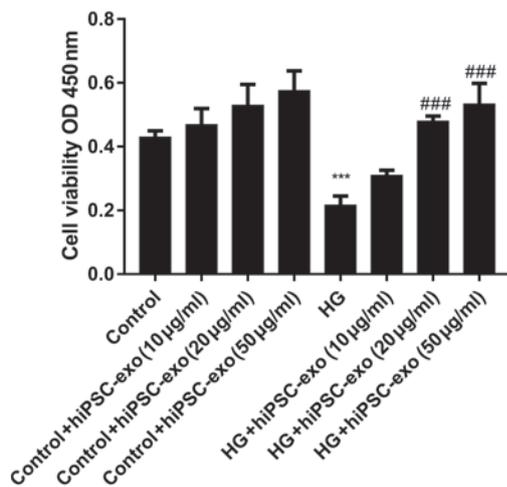


Figure 3. hiPSC-exo ameliorated the high glucose-induced decrease in cell viability in a dose-dependent manner. Cell counting kit-8 assays were performed to evaluate cell viability. HUVECs (1×10^3) were seeded in 96-well plates. The cell viability was assessed by measuring the OD value of HUVECs after 48 h of treatment. Control, normal glucose (5.5 mM); HG, high glucose (33 mM); hiPSC-exo (10, 20 or 50 µg/ml). Control vs. HG, *** $P < 0.0001$, $n = 5$; HG vs. HG + hiPSC-exo (20 µg/ml), *** $P < 0.0001$, $n = 5$; HG vs. HG + hiPSC-exo (50 µg/ml), *** $P < 0.0001$, $n = 5$. hiPSC-exo, human induced pluripotent stem cell-derived exosomes; HUVECs, human umbilical vascular endothelial cells; OD, optical density; HG, high glucose.

The function of exosomes in physiological and pathological conditions depends on their cellular origin and contents. As one of the most dynamic types of cell, hiPSCs have the capacity of self-renewal and multi-differentiation, thus they exert a therapeutic effect when used to treat various diseases, including myocardial infarction (23,24). The role of exosomes derived from hiPSCs was explored to examine their therapeutic effects. The results of the present study demonstrated that hiPSC-exo could promote cell viability and enhance tube formation, and inhibit cell senescence in HUVECs injured by

high glucose. A number of previous studies have demonstrated that exosomes from different types of cells exert different effects. Bang *et al* (25) revealed that cardiac fibroblasts secreted exosomes to mediate cardiomyocyte hypertrophy, suggesting that this is a potential therapeutic target. Intravenous administration of cell-free mesenchymal stromal cells (MSCs)-generated exosomes improved functional recovery and enhanced neurite remodeling, neurogenesis and angiogenesis following stroke in rats (26), suggesting that exosomes may be important in cell therapy. A previous study by Li *et al* (27) demonstrated that exosomes derived from liver nonparenchymal cells mediated the cell-to-cell transmission of interferon type I- α -induced antiviral activity. However, not all types of exosomes mediate cell-protective effects on target cells. Currently, the role of exosomes in tumors is being accepted, particularly with regard to tumor metastasis. The observations of the recent study by Zhang *et al* (28), demonstrated the dynamic and reciprocal cross-talk between tumor cells and the metastatic niche. They found that exosomes prepared the microenvironment of the target organ of metastasis for tumor cell colonization. Another study also demonstrated that tumor exosome integrins determine organotropic metastasis. It also revealed that exosomes from mouse and human lung-, liver- and brain-tropic tumor cells preferentially fuse with resident cells at the predicted destination (29).

In recent years, the incidence of diabetes mellitus has increased significantly, and its chronic vascular complications continue to affect patients' lives (29-31). Although the underlying specific mechanism is unclear, a growing body of research has indicated that the intracellular overproduction of reactive oxygen species (ROS), caused by hyperglycemia, is the common mechanism of vascular complications of diabetes mellitus. In a previous study by Kiritoshi *et al* (30), it was suggested that hyperglycemia increases mitochondrial ROS production, resulting in nuclear factor- κ B activation, cyclooxygenase (COX)-2 mRNA induction, COX-2 protein production

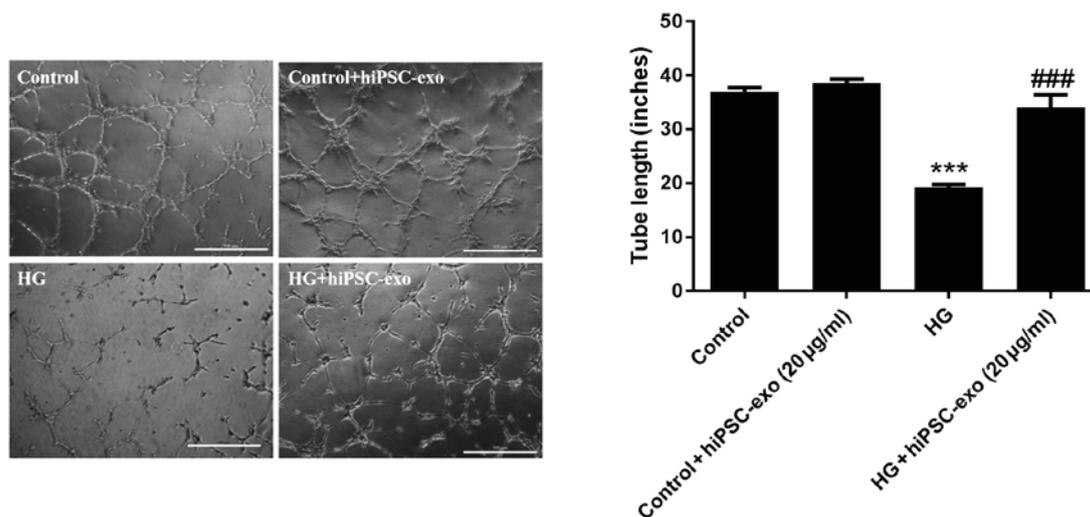


Figure 4. Enhanced capillary-like structure formation in HUVECs treated with hiPSC-exo in the presence of HG. Endothelial capillary-like structure formation was evaluated in HUVECs after being treated for 48 h with different conditions. Control: normal glucose (5.5 mM); HG: high glucose (33 mM); hiPSC-exo: 20 µg/ml. The data were analyzed using one-way analysis of variance and a post-hoc Bonferroni test. Error bars represent the standard error of the mean. Control vs. HG, ^{***}P<0.0001, n=3; HG vs. HG + hiPSC-exo, ^{###}P<0.0001, n=3. Scale bar, 500 µm. HUVECs, human umbilical vascular endothelial cells; hiPSC-exo, human induced pluripotent stem cell-derived exosomes; HG, high glucose.

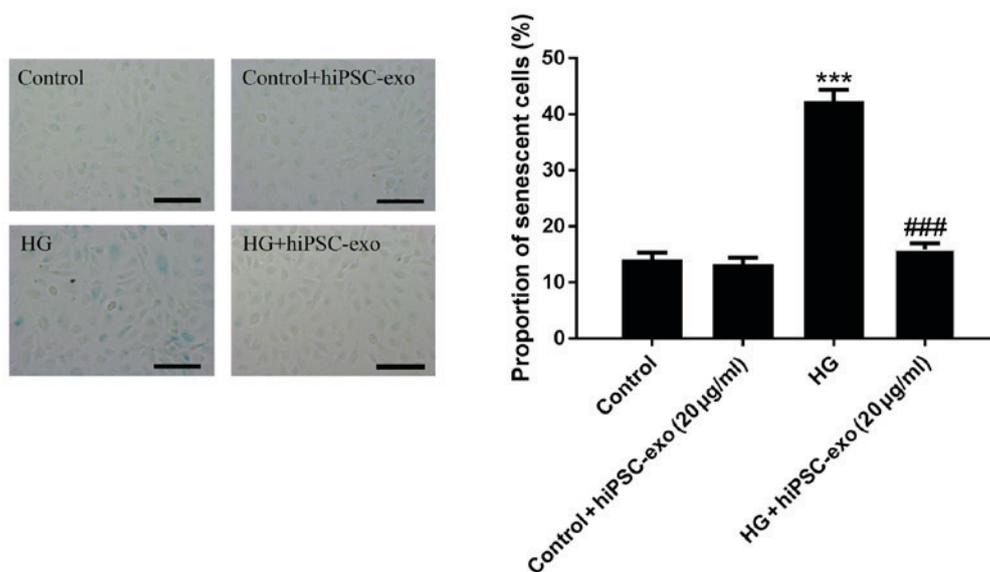


Figure 5. Effects of hiPSC-exo on the proportion of SA-β-gal positive HUVECs with different treatments. Normal glucose (5.5 mM); normal glucose + hiPSC-exo (20 µg/ml); HG (33 mM); HG + hiPSC-exo (20 µg/ml). Representative images of SA-β-gal positive cells (green) are presented in the left panel. Scale bar, 100 µm. Quantification of senescence assay is presented in the right panel. The data were analyzed using one-way analysis of variance and a post-hoc Bonferroni test. Error bars represent the standard error of the mean. Control vs. HG, ^{***}P<0.0001, n=3; HG vs. HG + hiPSC-exo, ^{###}P<0.0001, n=3. hiPSC-exo, human induced pluripotent stem cell-derived exosomes; SA-β-gal, senescence-associated β-galactosidase; HUVECs, human umbilical vascular endothelial cells; HG, high glucose.

and prostaglandin E2 synthesis. Yu *et al* (31) previously determined that a dynamic change in mitochondrial morphology in high glucose conditions contributed to the overproduction of ROS. The present study indicated that mitochondrial fission/fusion machinery may be a previously unrecognized target to control acute and chronic production of ROS in hyperglycemia-associated disorders. Therefore, it is vital to understand the specific mechanisms of ROS overproduction and resulting vascular complications of diabetes mellitus, and to find novel strategies and drugs to treat this problem. This is necessary to prevent or reduce the dysfunction of endothelial

cells, which are important in maintaining vascular function. A number of studies have demonstrated the therapeutic effects of exosomes from different types of cells on normal and injured endothelial cells. Conigliaro *et al* (32) found that exosomes released by CD90+ cancer cells modulated endothelial cells, promoting an angiogenic phenotype and cell-to-cell adhesion. When rats with traumatic brain injury were treated with MSC-derived exosomes, Zhang *et al* (33) found that newly generated endothelial cells in the lesion boundary zone were significantly increased with a parallel reduction in neuroinflammation.

Therefore, the present study may offer a novel strategy for maintaining the normal function of endothelial cells during the vascular complications of diabetes mellitus.

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