

Transplantation of Human Menstrual Blood Progenitor Cells Improves Hyperglycemia by Promoting Endogenous Progenitor Differentiation in Type 1 Diabetic Mice

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Recently, a unique population of progenitor cells was isolated from human menstrual blood. The human menstrual blood progenitor cells (MBPCs) possess many advantages, such as the noninvasive acquisition procedure, broad multipotency, a higher proliferative rate, and low immunogenicity, and have attracted extensive attention in regenerative medicine. Preclinical studies to test the safety and efficacy of MBPCs have been underway in several animal models. However, relevant studies in type 1 diabetes mellitus (T1DM) have not yet been proceeded. Herein, we studied the therapeutic effect of MBPCs and the mechanism of β -cell regeneration after MBPC transplantation in the T1DM model. Intravenous injection of MBPCs can reverse hyperglycemia and weight loss, prolong lifespan, and increase insulin production in diabetic mice. Histological and immunohistochemistry analyses indicated that T1DM mice with MBPC transplantation recovered islet structures and increased the β -cell number. We further analyzed *in vivo* distribution of MBPCs and discovered that a majority of MBPCs migrated into damaged pancreas and located at the islet, duct, and exocrine tissue. MBPCs did not differentiate into insulin-producing cells, but enhanced neurogenin3 (*ngn3*) expression, which represented endocrine progenitors that were activated. *Ngn3*⁺ cells were not only in the ductal epithelium, but also in the islet and exocrine tissue. We analyzed a series of genes associated with the embryonic mode of β -cell development by real-time polymerase chain reaction and the results showed that the levels of those gene expressions all increased after cell transplantation. According to the results, we concluded that MBPCs stimulated β -cell regeneration through promoting differentiation of endogenous progenitor cells.

Introduction

TYPE 1 DIABETES MELLITUS (T1DM), which can lead to hyperglycemia and severe complications [1,2], is an insulin-dependent metabolic disorder characterized by auto-immune destruction of pancreatic islet β cell and inadequate insulin production. Human islet transplantation is an effective therapy by controlling blood glucose and appropriate preventing hyperglycemia without exogenous insulin administration. However, a lack of pancreas donors and the need for long-term immunosuppression limit the widespread use of this treatment [3,4]. Recently, the use of human bone marrow-derived mesenchymal stem cells (BM-MSCs) or human umbilical cord blood (HUCB) cells to treat experimental diabetes had some positive results [5–8]. Transplantation of BM-MSCs or HUCB cells could reduce blood

glucose levels [5,6] or improve pancreatic insulinitis [7]. Thus, BM-MSCs and HUCB cells may be potential sources for β -cell replacement therapy. However, BM-MSCs and HUCB cells are limited in more widely use for invasiveness of extraction, restricted differentiation potential, or in some cases, a limited proliferative capacity [9–11].

Recently, a novel population of progenitor cells is isolated from human menstrual blood, which can be easily obtained without invasive procedures [12–15]. The human menstrual blood progenitor cells (MBPCs) have showed highly proliferative capabilities and broad multipotency. In absence of induction stimuli, MBPCs are able to expand at least 18 passages without chromosome abnormalities [13]. Meng et al. *in vitro* induced MBPCs to differentiate into all three germ lineages, including cardiomyocytic, respiratory epithelial, neurocytic, myocytic, endothelial, pancreatic,

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hepatic, adipocytic, and osteogenic [13]. A clinical trial and an *in vitro* immunologic test demonstrated that MBPCs possessed low immunogenicity properties and immunomodulatory effects [16,17]. Animal experiments showed that MBPCs had tissue repair effects in some diseases such as Duchenne muscular dystrophy (DMD), myocardial infarction (MI), critical limb ischemia (CLI), and stroke [12,18–21]. Based on the advantages in characteristics and repair effects in diseases, the therapeutic potential and mechanism of MBPCs in diabetes should be notable and investigated.

Thus, in this study, we have two purposes: one is to investigate the therapeutic effect of MBPCs to T1DM mice and the other is to study involved repair mechanism. Using a mouse model of streptozotocin (STZ)-induced type 1 diabetes, we show that transplantation of MBPCs reverses hyperglycemia, recovers islet structures, and stimulates endogenous β -cell regeneration. MBPCs migrate to the pancreatic duct, exocrine tissues, and islet and promote endogenous pancreatic progenitor differentiation.

Materials and Methods

Experimental animals

Six- to eight-week-old male BALB/c mice were purchased from the SLAC Laboratory Animal Corporation (Shanghai, China). Mice were fed *ad libitum* and housed in a 12-h light and 12-h dark cycle under specified pathogen-free conditions. Eight-week-old male BALB/c mice, weighing 32–36 g, were chosen in animal experiments. All animal experiments were according to the institutional animal welfare guidelines and approved by the Animal Care and Use Committees of Zhejiang University, China.

Isolation and culture of MBPCs

The MBPCs were isolated from female donors according to the protocol previously reported [13] with slight modification. The entire procedures were with consent of the donors and approved by the Ethics Committee of The First Affiliated Hospital, College of Medicine, Zhejiang University, China. The menstrual blood samples were collected with a Divacup (Kitchener, ON) from healthy women ($n=9$) during a menstrual cycle. The cells were transferred into phosphate-buffered saline (PBS) with mixed antibiotics, including amphotericin B, gentamycin sulfate, kanamycin sulfate, cephalexin, vancomycin hydrochloride, and heparin at 4°C for 24 h. Then, the samples were centrifuged at 1,600 g for 10 min at 4°C, and the supernatants were used for microbiological testing. Mononuclear cells were separated by a density gradient centrifugation with FicolI-Paque (Fisher Scientific, Portsmouth NH). The interlayer cells were collected and cultured with the Chang Medium [14] in a tissue culture flask (Corning, Corning, NY) to obtain adherent cells. The media were changed every 2–3 days until adherent cells grew to 80%–90% confluency, and then the cells were subcultured using 0.25% trypsin (Invitrogen, Carlsbad, CA) and cultured on flasks or plates seeded at 8,000–10,000 cells/cm². The cells used in the experiments were at fourth to eighth passage.

Identification of MBPCs by flow cytometry

Isolated MBPCs were utilized for flow cytometry analysis. About 10⁵ cells were resuspended in 100 μ L PBS and

incubated with primary antibodies (PE-conjugated anti-human CD13, CD29, CD34, CD44, CD45, CD73, CD90, CD105, CD117, CD166, SSEA4, or FITC-conjugated anti-human HLA-DR; Becton Dickinson, Franklin Lakes, NJ) (1:100) at 4°C for 1 h. Then, the cells were washed twice with PBS. The corresponding isotype antibodies (Becton Dickinson) were set as negative controls. Cells were analyzed by a Flow Cytometer (FC500MCL; Beckman Coulter, Pasadena, CA). The results were analyzed by FlowJo software.

Induced differentiation of MBPCs to osteoblasts, adipocytes, and chondroblasts

For osteogenic differentiation, the MBPCs were plated at 3×10^3 cells/cm² and treated with the human MSC osteogenic differentiation medium (Cyagen, Guangzhou, China). The osteogenic medium consisted of 100 nM dexamethasone, 10 mM β -glycerophosphate, 0.2 mM ascorbate, 1 mM glutamine, 1% penicillin–streptomycin, and 10% fetal bovine serum (FBS) and was completely changed every 3 days for up to 21 days. Cells were then fixed by 4% formaldehyde and stained with Alizarin red for 5 min.

For adipogenic differentiation, cells were plated at 2×10^4 cells/cm² in the human MSC adipogenic differentiation medium A (ADMA; Cyagen); ADMA consisted of 1 mM dexamethasone, 0.5 mM 3-isobutyl-1-methyl-xanthine (IBMX), 10 μ g/mL recombinant human insulin, 100 mM rosiglitazone, and 10% FBS. Cells were cultured in the ADMA for 3–5 days, in the adipogenic differentiation medium B (ADMB; 1% penicillin/streptomycin + 10% FBS + 10 μ M insulin + 1 mM glutamine) for 24 h, and then interchanged from ADMB to ADMA of three to five cycles of induction/maintenance. The cells were cultured in the ADMB for an additional 7 days by replacing the medium every 3 days. For Oil Red O stain analysis, cells were fixed in 4% formaldehyde and then stained with Oil Red O for 30 min.

For chondrogenic differentiation, cells were plated onto 12-well tissue culture plates at 2.5×10^5 cells/cm² in a complete chondrogenic medium. The chondrogenic medium consisted of 1 mM dexamethasone, 0.2 mM ascorbate, 1% ITS, 1 mM sodium pyruvate, 1 mM proline, and 20 ng/mL TGF- β 3 (Cyagen). The cells were cultured for 14–28 days by completely replacing the medium every 2 days. Cells were formalin fixed and stained with Alcian blue for 30 min.

Karyotype analysis

The 10th and 20th passages of cultured MBPCs were used to prepare for chromosome samples according to [12]. Karyotype analysis was performed by standard cytogenetic protocol with G staining. For cells, 50 metaphases were prepared for G-banding analysis.

Preparation of mouse T1DM model

The BALB/c mice were injected intraperitoneally (i.p.) with 180 mg/kg STZ (Sigma-Aldrich, St. Louis, MO) at a single injection [22]. STZ was dissolved in 0.1 M of sodium citrate buffer, pH 4.5 [23], and injected within 15 min after preparation. The blood glucose level was assayed with tail vein blood by a standard blood glucose meter (Accu-check,

Performa). Hyperglycemia developed within 3–5 days after STZ treatment. Mice with blood glucose levels over 13.9 mM [24] at two random measurements were considered diabetic and used in the subsequent transplantation experiments.

Cell transplantation

Cell transplantation assay was performed to evaluate whether MBPCs had effect to ameliorate diabetic symptoms in the T1DM model. About 3×10^5 cells were injected through a tail vein into each diabetic mouse, 5–7 days after STZ injection. T1DM mice without cell transplantation were set as control. Untreated normal mice were used as normal control. Body weight and fasting blood glucose were monitored overnight, before and after cell administration. Additionally, the lifespan of cell-treated mice and untreated diabetic mice after cell transplantation was recorded. MBPCs at passage 5 were used in the transplantation model.

Cell labeling and imaging

For cell tracking, MBPCs were labeled by the fluorescent lipophilic tracer 1,1'-dioctadecyl-3,3',3',3'-tetramethylindocarbocyanine perchlorate (DiI; Molecular Probes, Eugene, OR). For labeling, the cells were resuspended at 1×10^6 cells/mL in the α MEM, and added DiI in the cell suspension to obtain a final concentration of 10 μ g/mL. After incubation for 20 min at 37°C with 5% humidified CO₂, the cells were centrifuged at 400 g for 5 min, washed twice with PBS [25,26], and then resuspended in PBS for 2×10^6 cells/mL. For transplantation, MBPCs were infused in diabetic mice and normal mice. After 24 h, 3, 7, and 14 days, the mice were sacrificed. The heart, liver, lungs, pancreas, spleen, and kidney were immobilized and prepared for tissue imaging with a Night Owl LB 981 low-light luminograph (Berthold Technologies, Bad Wildbad, Germany) equipped with a cooled CCD camera. Digital images were analyzed with WinLight³² Software. At each time point, the diabetic group and normal group both had three parallel mice. Each mouse received 5×10^5 cells.

Intraperitoneal glucose tolerance test

To further evaluate the functional recovery effect of islet β cell after MBPC transplantation, an intraperitoneal glucose tolerance (IPGT) test was performed on day 14 after transplantation followed by the protocol from the Animal Models of Diabetic Complications Consortium (AMDCC). Briefly, mice were fasted for 6 h by removal to a clean cage without food at the end of their dark (feeding) cycle and administered i.p. with glucose (1 mg/g body weight; Sigma-Aldrich). Blood samples were obtained from a small tail clip and analyzed by a Roche Accu-check glucose meter. Blood glucose values were obtained at 0, 15, 30, 45, 60, 75, 90, 105, 120, and 135 min.

ELISA

To evaluate the recovering effect of MBPCs, the levels of insulin and C-peptide in serum of T1DM mice, cell-treated mice, and control mice were comparatively analyzed by ELISA kits (Merckodia, Uppsala, Sweden). For this purpose,

a human insulin-specific ELISA kit (Ultrasensitive Insulin ELISA), a mouse insulin-specific ELISA kit (Ultrasensitive Mouse Insulin ELISA), and a human C-peptide-specific ELISA kit (Ultrasensitive C-peptide ELISA) were used, respectively. Fasting blood was collected on day 28 after cell administration. Serum samples were obtained from tail vein and allowed to clot for 2 h at room temperature. The clotted material was discarded by centrifugation at 3,000 rpm for 15 min. Serum samples were stored at -80°C for test as above.

Histology, immunohistochemistry, and immunofluorescence

For histological analysis, pancreas samples were fixed in 10% formalin overnight at 4°C and embedded in paraffin. Two micrometer pancreatic sections were cut serially. For histopathology analysis, sections were stained with hematoxylin and eosin (H&E; Sigma-Aldrich). The slides were visualized using a microscope (Zeiss, Jena, Germany).

For the immunohistochemistry test, sections were peroxidase blocked in 3% H₂O₂ for 10 min, and then they were pretreated using heat-mediated antigen retrieval with the sodium citrate buffer (pH6.0). Tissue sections were blocked with 10% FBS for 20 min at room temperature and then incubated with a rabbit polyclonal antibody against mouse insulin (1:500 dilution; Abcam, Cambridge, United Kingdom) for 2 h at room temperature. Slides were washed three times with PBS, 5 min at every time. Then, they were further incubated with a secondary antibody (HRP-conjugated goat anti-rabbit IgG; Chemicon, Billerica, MA). In addition, to detect whether MBPCs migrated into the injured pancreas, we used a mouse monoclonal anti-human nuclei antibody (Millipore, Darmstadt, Germany) to stain MBPCs. The tissues were incubated with this primary antibody (1:50 dilution) for 1 h at room temperature, followed by incubation with a secondary antibody (HRP-conjugated goat anti-mouse IgG; Chemicon) for 1 h at room temperature. 3,3'-diaminobenzidine (DAB) was used as the reaction substrate.

For the immunofluorescence test, sections were costained with a rabbit polyclonal anti-human insulin antibody and a mouse monoclonal anti-human nuclei antibody (1:50 dilution; Millipore) for 90 min at room temperature. Then, they were incubated with secondary antibodies (DyLight 405-labeled goat anti-rabbit IgG, Cy3-labeled goat anti-mouse IgG) for 1 h at room temperature. Additionally, tissues were immunostained with a rabbit polyclonal anti-mouse neurogenin3 (ngn3) antibody (1:500 dilution; Millipore) or costained with a rabbit polyclonal anti-mouse ngn3 antibody and a mouse monoclonal anti-human nuclei antibody for 2 h at room temperature. They were further incubated with secondary antibodies (Alexa Fluor 555-labeled goat anti-rabbit IgG, FITC-labeled goat anti-mouse IgG) for 1 h at room temperature. Immunostaining analysis was performed in a Zeiss fluorescence microscope and images were captured and digitalized with the Axiovision 4.6 software.

Reverse transcriptase–polymerase chain reaction and real-time PCR

Total RNA of pancreas was extracted on day 3, 7, 10, 14, 21, and 28 after cell transplantation by using the TRIzol

Reagent (Invitrogen). The samples were reverse transcribed with PrimeScript Reverse Transcriptase (TaKaRa, Shiga, Japan) for 30 min at 42°C in the presence of an oligo-dT primer. Real-time polymerase chain reaction (PCR) analysis was performed by the Eppendorf Mastercycler Realplex detection system using the SYBR Green Premix Ex Taq (Perfect real-time; TaKaRa). The PCR reaction consisted of 5 µL of SYBR Green PCR Ex Taq Mix, 1 µL of 10 mM forward and reverse primers, 3.5 µL of water, and 0.5 µL of template cDNA in a total volume of 10 µL. Cycling was performed using the default conditions of Realplex Software 1.5: 95°C for 2 min, followed by 40 cycles of 95°C for 15 s and 58°C for 20 s. The relative expression of each gene was normalized to endogenous β-actin. The primers of assayed genes, including *ngn3*, *foxa2*, *pax4*, *mafa*, *mafB*, and *nkx6.1*, were designed specifically for mouse. The primers of insulin, *glut2*, and *pdx1* are referenced in [23]. The primer sequences are shown in Table 1. The related gene expression level was calculated using the $2^{-\Delta\Delta CT}$ method. Each sample was run in three parallel reactions.

Statistical analyses

Statistical data were analyzed by one-way analysis of variance (ANOVA) with SPSS 16.0 software. Results are expressed as mean ± standard deviation; a *P*-value < 0.05 or < 0.01 was considered significant. Survival curves were plotted using the Kaplan–Meier method.

Results

Characterization of MBPCs

Flow cytometry results revealed that MBPCs exhibited phenotypes of MSCs. They expressed CD13, CD29, CD44, CD73, CD90, CD105, and CD166, but did not express HLA-DR, CD34, CD45, CD117, and SSEA4 (Fig. 1a).

MBPCs had the capacity for trilineage mesenchymal differentiation. MBPCs were induced to chondroblasts, adipocytes, and osteoblasts (Fig. 1b) using the commercial differentiating medium. Cell staining showed multipotent characteristics such as sulfated proteoglycans formed in cartilage induction, fat vacuoles detected in adipocyte differentiation, and calcium deposits in bone formation, whereas the controls were negative of trilineage staining (Fig. 1b). Isolated MBPCs were plastic adherent, fibroblast like, and spindle shaped (Fig. 1c). One doubling time of MBPCs at passage 5 was near 20 h (Fig. 1d). MBPCs expressed the multipotent marker Oct-4 at passage 10 (Fig. 1e). MBPCs maintained diploid karyotype without chromosomal aberrations at passage 10 and 20 (Fig. 1f).

Therapeutic effect of MBPCs in diabetic mice

To analyze the potential therapeutic effect of MBPCs to T1DM, we transplanted MBPCs into T1DM mice at the onset of hyperglycemia by tail vein infusion and recorded body mass, blood glucose levels, and survival rates for 42 days. T1DM mice experienced a significant mean weight loss from 32.6 ± 1.3 to 23.7 ± 1 g, whereas mice transplanted with MBPCs had a relative stable weight at 32.3 ± 1.6 g (Fig. 2a). The mean body mass of cell-transplanted mice was significantly greater compared with T1DM mice. Glycemic monitoring showed that blood glucose increased in T1DM mice at all time points (from 21 ± 1.6 to >33 mM) (Fig. 2b), but it dramatically reduced in cell-transplanted mice on day 7 post-transplantation (from 21.2 ± 1.6 to 15 ± 1.5 mM). The mean blood glucose level from day 10 to 42 was 14.7 ± 1.5 mM in cell-transplanted mice and was 7.7 ± 1 mM in normal mice (Fig. 2b). Cell-transplanted mice had a survival rate of 95%, in contrast to T1DM mice with a rate of 30% (100% morbidity) (Fig. 2c). Furthermore, polyuria was significantly improved compared with untreated diabetic mice (data not shown).

TABLE 1. PRIMERS USED FOR PCR AND REAL-TIME PCR

Primer name	Sequence (5'-3')	Species
Insulin1	F: TAGTGACCAGCTATAATCAGAG	Mouse
Insulin1	R: ACGCCAAGGTCTGAAGGTCC	
Pdx1	F: TGTAGGCAGTACGGGTCCTC	Mouse
Pdx1	R: CCACCCAGTTTACAAGCTC	
Glut2	F: CCACCCAGTTTACAAGCTC	Mouse
Glut2	R: TGTAGGCAGTACGGGTCCTC	
Foxa2	F: GAACTCCATCCGCCACTCT	Mouse
Foxa2	R: GGTCTTCTTGCCCTCCGCTA	
Pax4	F: GGATGCGACCCTGTGAC	Mouse
Pax4	R: TCCTGAAGTGCCCCGAAG	
Ngn3	F: CGGATGACGCCAAACTTA	Mouse
Ngn3	R: GCCTCCACTACCTCCACTC	
Nkx6.1	F: GGGGACTTCGGAGAATGAG	Mouse
Nkx6.1	R: GGCGAGCAGCCAGGATA	
MafA	F: ATGGCCCGCGGAGCTGGCGATGGGCG	Mouse
MafA	R: CGAAGAGGGCACCGAGGAGCAGGGC	
MafB	F: CTGCGCCCCTAGCCCTGGACT	Mouse
MafB	R: GGCGCCCCTGGCACTCAAA	
β-Actin	F: ATGGATGACGATATCGCTG	Mouse
β-Actin	R: ATGAGGTAGTCTGTGAGGT	

PCR, polymerase chain reaction.

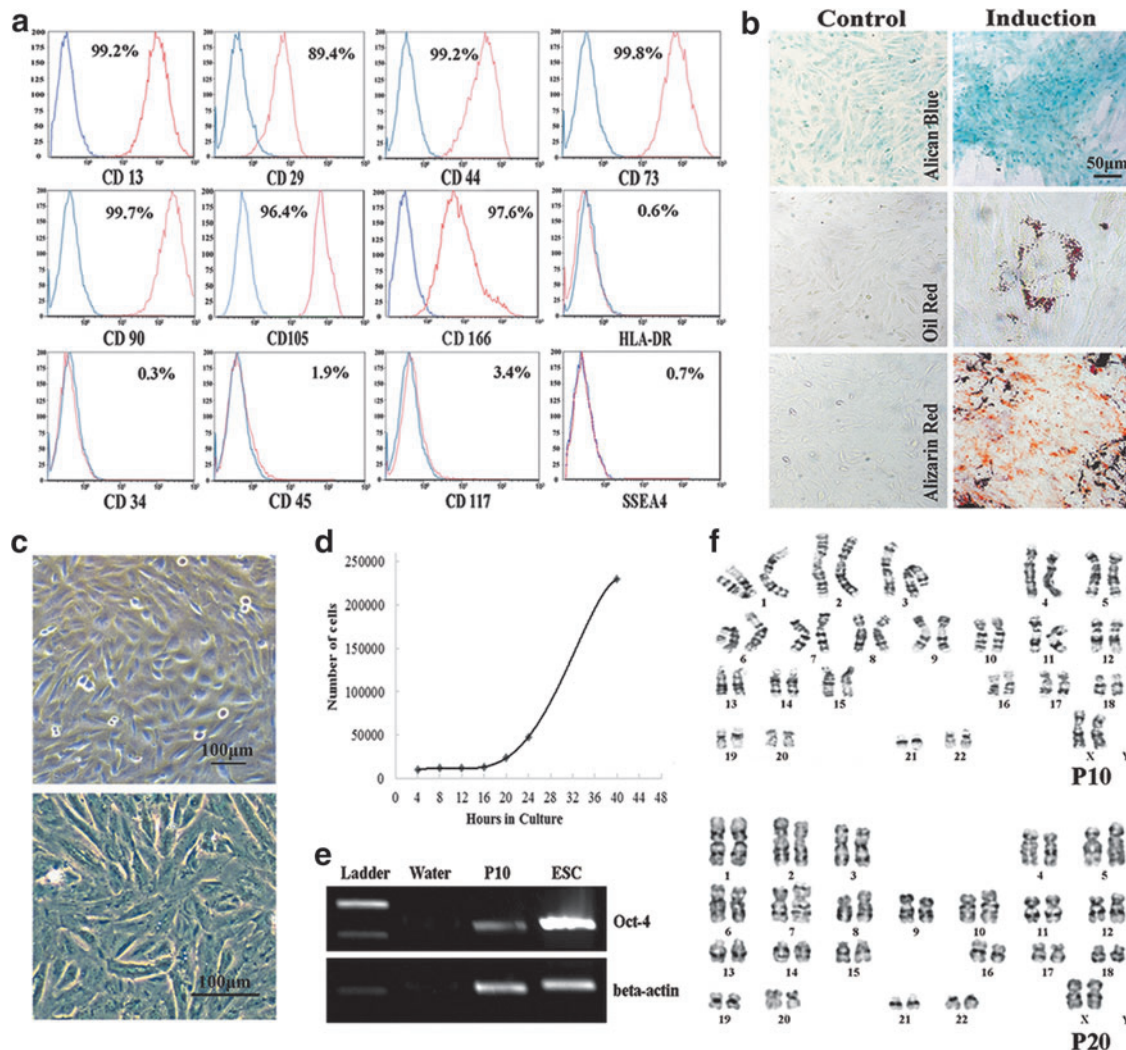


FIG. 1. Identification of human menstrual blood progenitor cells (MBPCs). **(a)** Representative mesenchymal stem cell (MSC) phenotypes of MBPCs (four to six passages) by flow cytometry. MBPCs are positive for MSC markers such as CD13, CD29, CD44, CD73, CD90, CD105, and CD166 and negative for HLA-DR, CD34, CD45, CD117, and SSEA4. **(b)** Representative MBPCs having the potential to differentiate into chondroblasts, adipocytes, and osteoblasts of mesoderm tissues under conditioned culture. **(c)** Representative photomicrographs showing adherent MBPCs with spindle shapes on a plastic tissue culture flask. **(d)** Growth curve of MBPCs at passage 5 by MTT assay. **(e)** MBPCs expressing the multipotent marker Oct-4 at the RNA level at passage 10 by reverse transcriptase–polymerase chain reaction (RT-PCR). **(f)** Representative normal chromosome morphology of MBPCs by karyotype analysis at passage 10 and 20.

Islet function and structure repair by MBPC engraftment

To test the repair effect of MBPCs to impaired islet, we estimated glucose control in cell-transplanted mice using an IPGT test and found a similar rate of hypoglycemic activity as normal control, but T1DM mice showed an impaired glucose control (Fig. 3a). The serum insulin levels in cell-transplanted mice ($0.81 \pm 0.08 \mu\text{g/L}$) were ~ 10 -fold higher than in T1DM mice ($0.08 \pm 0.005 \mu\text{g/L}$), but were lower than in normal control ($1.12 \pm 0.24 \mu\text{g/L}$) (Fig. 3b).

Pathological and histological results (Fig. 3c) indicated that the islet structure and morphology were restored by MBPC administration. Following transplantation of MBPCs, mice exhibited normal islet structure and size in the pancreas, whereas in T1DM mice, the islet changed mor-

phologically and appeared atrophied. In the inner part of the islet, some endocrine cells of T1DM mice formed pseudo-ducts, while in the normal mice and cell-treated mice, there were just normal vessels in the islet. In addition, in T1DM mice, some acinar cells of the exocrine gland were swollen and trapped in the islet, whereas in the cell-transplanted mice, there were very few acinar cells around islet displaying intumescence. From the results of H&E staining, it displayed that cell-transplanted mice did not experience apparent pathological changes compared with normal control (Fig. 3c).

Immunohistochemical results (Fig. 3d) showed that the islet size and β -cell number in cell-transplanted mice were higher than in T1DM mice. We carried out a statistical analysis for islet size and β -cell number in normal control mice, T1DM mice, and cell-transplanted mice (Fig. 3e, f).

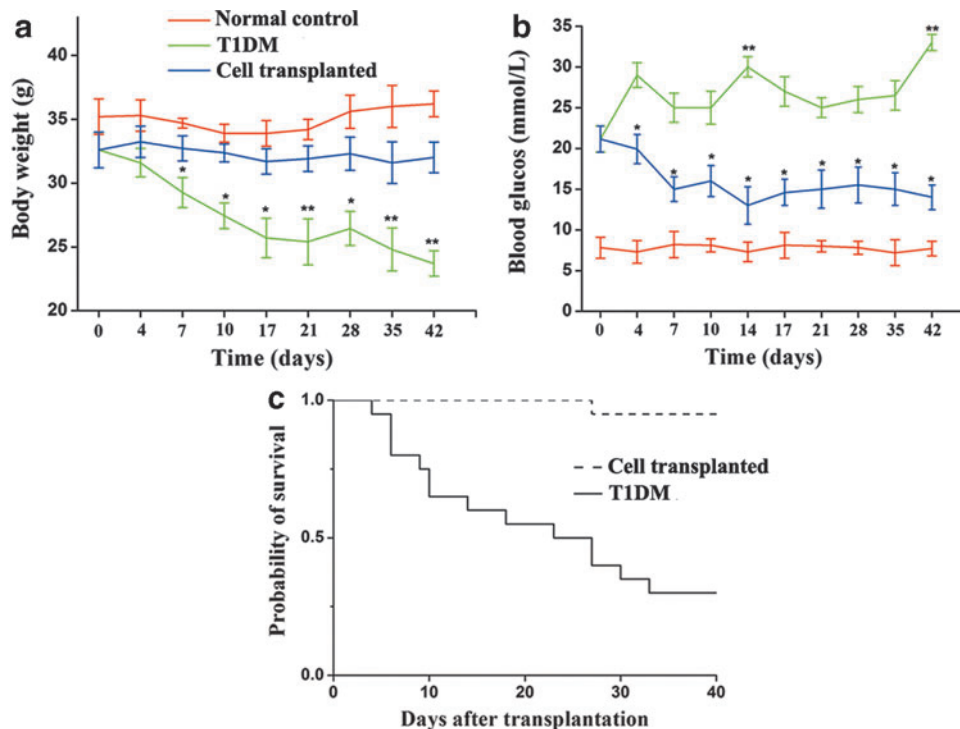


FIG. 2. Transplantation of MBPCs ameliorates diabetic symptoms in streptozotocin (STZ)-induced mice. Each STZ-treated (1 injection at 180 mg/kg) hyperglycemic mouse received 3×10^5 MBPCs by tail vein transplantation. STZ, streptozotocin. **(a)** Compared with type 1 diabetes mellitus (T1DM) mice (green line, $n=8$), cell-transplanted mice (blue line, $n=10$) maintained their body weight and had no significant difference from normal controls (red line, $n=6$). **(b)** After transplantation, the mean level of blood glucose from day 10 to 42 reduced to 14.7 ± 1.5 mM in MBPC-treated mice (blue line, $n=10$). The mean level of blood glucose in T1DM mice (green line, $n=8$) was at 27.4 ± 2.8 mM. **(c)** Compared with T1DM mice (full line, $n=20$), mice transplanted with MBPCs (dotted line, $n=18$) showed a higher survival rate (95% vs. 30%). Data are presented as mean \pm standard deviation (SD). * $P < 0.05$, ** $P < 0.01$.

According to the statistical result, compared with T1DM mice, islet size (Fig. 3e) and the quantity of β cell in visible islet (Fig. 3f) were significantly increased ($P < 0.05$) in cell-transplanted mice, but were not significantly different from normal control ($P > 0.05$). Even though there was a slight reduction in the β -cell number in cell-transplanted mice, islet size and morphology still remained normal relative to normal control mice (Fig. 3d–f). Compared with normal control, the total number of islets in T1DM mice also had a significant decrease (data not shown).

MBPC distribution in vivo

Tissues were dissected under the CCD camera for tracking distribution of DiI-marked MBPCs in the diabetic mice and normal mice. In the diabetic mice, MBPCs mainly located in the lungs, pancreas, and liver on day 3 after cell implantation, and relatively more cells resided in diabetic pancreas on day 3 and 14 (Fig. 4a). In the normal mice, on day 14 after transplantation, MBPCs mainly distributed in the lungs, liver, and kidney and most of them located in the lungs (Fig. 4a,b); the quantity of MBPCs significantly reduced compared with cell-transplanted diabetic mice, and MBPCs were almost undetectable in normal pancreas (Fig. 4b). However, on day 14 after transplantation, many MBPCs still survived in diabetic mice, and most of them existed in the injured pancreas (Fig. 4a,b). During the first 24 h after cell infusion, part of the MBPCs migrated into diabetic

pancreas and the number of MBPCs increased on day 3, then had a significant reduction on day 7 and 14 (Fig. 4c). The DiI signal increased during the first 3 days (Fig. 4c), indicating the acceptance of the grafted cells and proliferation of the cells. However, after 3 days, the DiI signal dropped step by step (Fig. 4c), indicating cell death in long-term implantation. Additionally, we also evaluated MBPC distribution on day 28 and 42. The quantity of MBPCs in pancreas gradually decreased on day 28 and 42, but there were still a few MBPCs in pancreas. Although the infused cell number had an obviously decrease in course of time, the number of MBPCs in diabetic pancreas was still higher than in diabetic other tissues (data not shown).

Recruitment and differentiation of MBPCs in damaged islet

To confirm whether MBPCs migrated to the diabetic pancreas, we immunohistochemically stained human specialized nuclei antigen (Fig. 5a). The results showed that infused MBPCs migrated to the injured pancreatic tissue and located at islet and exocrine tissue near the pancreatic duct and pancreatic vessel (Fig. 5a).

To test whether MBPCs promoted β -cell regeneration via self-differentiation into insulin-producing cells (IPCs) under the injured microenvironment, ELISA for human insulin and human C-peptide were performed by using serum samples from cell-transplanted mice. The results showed that no human

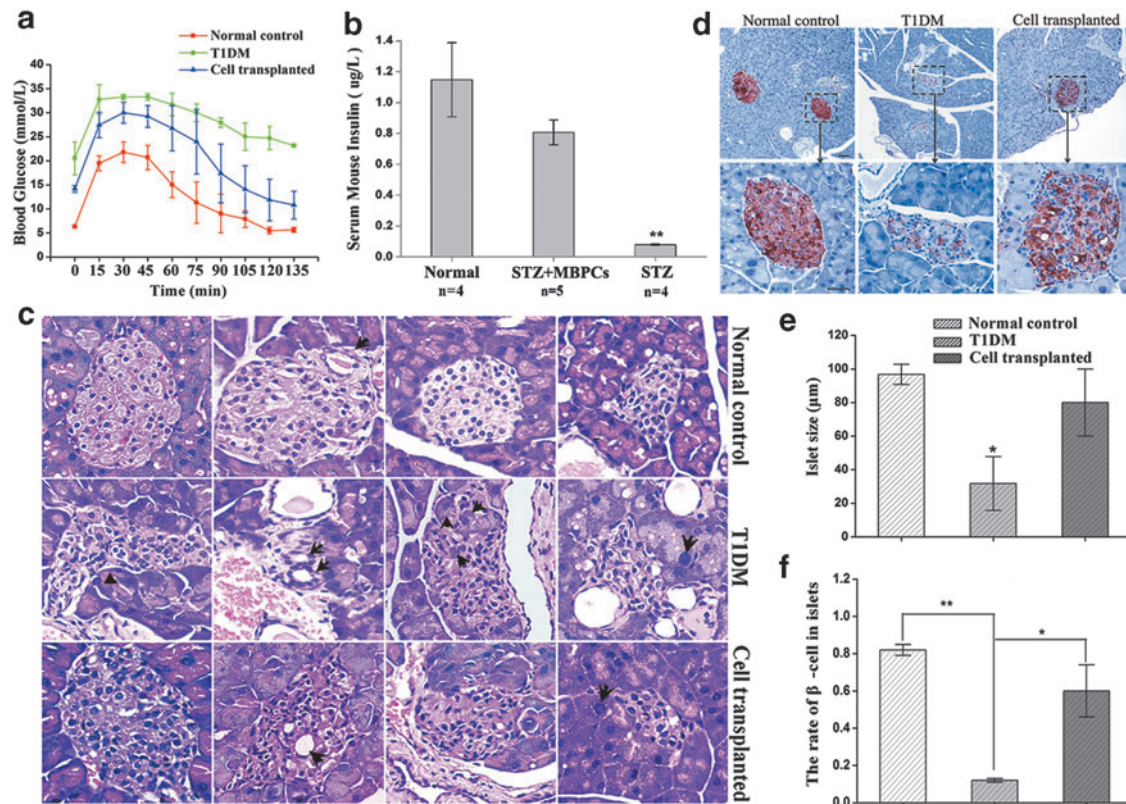


FIG. 3. Transplantation of MBPCs improves the function and structure of damaged islets in STZ-induced mice. **(a)** An intraperitoneal glucose tolerance test (IPGT) was performed on day 14 after transplantation. Compared with normal control (red line, $n=5$), cell-transplanted mice (blue line, $n=6$) showed a similar rate of lowering blood glucose, whereas T1DM mice had uncontrolled blood glucose levels during 135 min (green line, $n=5$). **(b)** The serum samples of mice were collected on day 28 after cell administration for ELISA. Compared with T1DM mice (right bar, $n=4$), mice transplanted with MBPCs (middle bar, $n=5$) showed a significantly higher level of serum insulin ($P<0.01$), but lower than normal control (left bar, $n=4$) with no significant difference. **(c)** Representative pathological variances in the pancreas on day 21 after cell infusion. Islet in T1DM mice was shrinking and deformed (arrow on the left), and exocrine acinar cells swelled (arrow on the right), and exocrine cells inserted in islet (arrows on third line), and pseudoductal structures appeared in the islet (arrows on the second line). However, islets in cell-transplanted mice had no obvious pathological changes compared with normal control. The four to six sliders of each group were used to analyze. Original magnification, $400\times$. **(d)** Representative immunohistochemistry results showed that insulin production in islet tissue increased in the cell-transplanted mice relative to T1DM mice. Mice were sacrificed on day 28 after cell transplantation. The three to five sliders of each group were used to analyze. Scale bar = $10\ \mu\text{m}$. **(e, f)** These two figures are the statistical analysis of figure 3d. The islet size of T1DM mice decreased significantly compared with cell-transplanted mice. The islet size of cell-transplanted mice had no significant difference from normal control. The number of β cells in cell-transplanted mice significantly increased compared with T1DM mice and had no significant difference from normal control. The average 20 islets of each group were used to analyze. Data are mean \pm SD ($n=3$). * $P<0.05$, ** $P<0.01$.

insulin or human C-peptide was detected (data not shown). To further confirm the *in vivo* differentiation of MBPCs, we costained the human-specific nuclei antibody and the human insulin antibody on day 28 after cell transplantation by immunofluorescence. The results displayed a lack of insulin expression in MBPCs (Fig. 5b). Based on the specificity of the human ELISA kit and antibodies, we concluded that MBPCs did not differentiate into IPCs in injured pancreas.

MBPC-promoted endogenous progenitor cell differentiation

To confirm whether ngn3^+ cells differentiated from MBPCs, we costained ngn3 and the human nuclei antibody (Fig. 6a, b) on day 28 after transplantation. As a result, in cell-transplanted mice, ngn3^+ cells were detected in islet,

but they were negative for the human nuclei antigen (Fig. 6b). Compared with cell-transplanted mice (Fig. 6b), fewer ngn3^+ cells were observed in islet in T1DM mice (Fig. 6a). To verify whether MBPCs stimulate the differentiation of pancreatic progenitor cells associated with the duct epithelium, we analyzed ngn3 expression in pancreatic structures, including the duct and exocrine tissue (Fig. 6c, d). We observed that in cell-transplanted mice, on day 14 after transplantation, a few of the ngn3^+ cells were detected not only in the ductal epithelium, but also in exocrine tissue near the duct (Fig. 6d), while in T1DM mice, no ngn3^+ cells were detected in the exocrine tissue and duct epithelium (Fig. 6c). On day 28 after cell transplantation, a small cluster of insulin-positive cells were detected in exocrine tissue near the duct (Fig. 6f), where ngn3^+ cells also appeared on day 14 in cell-transplanted mice (Fig. 6d). The results

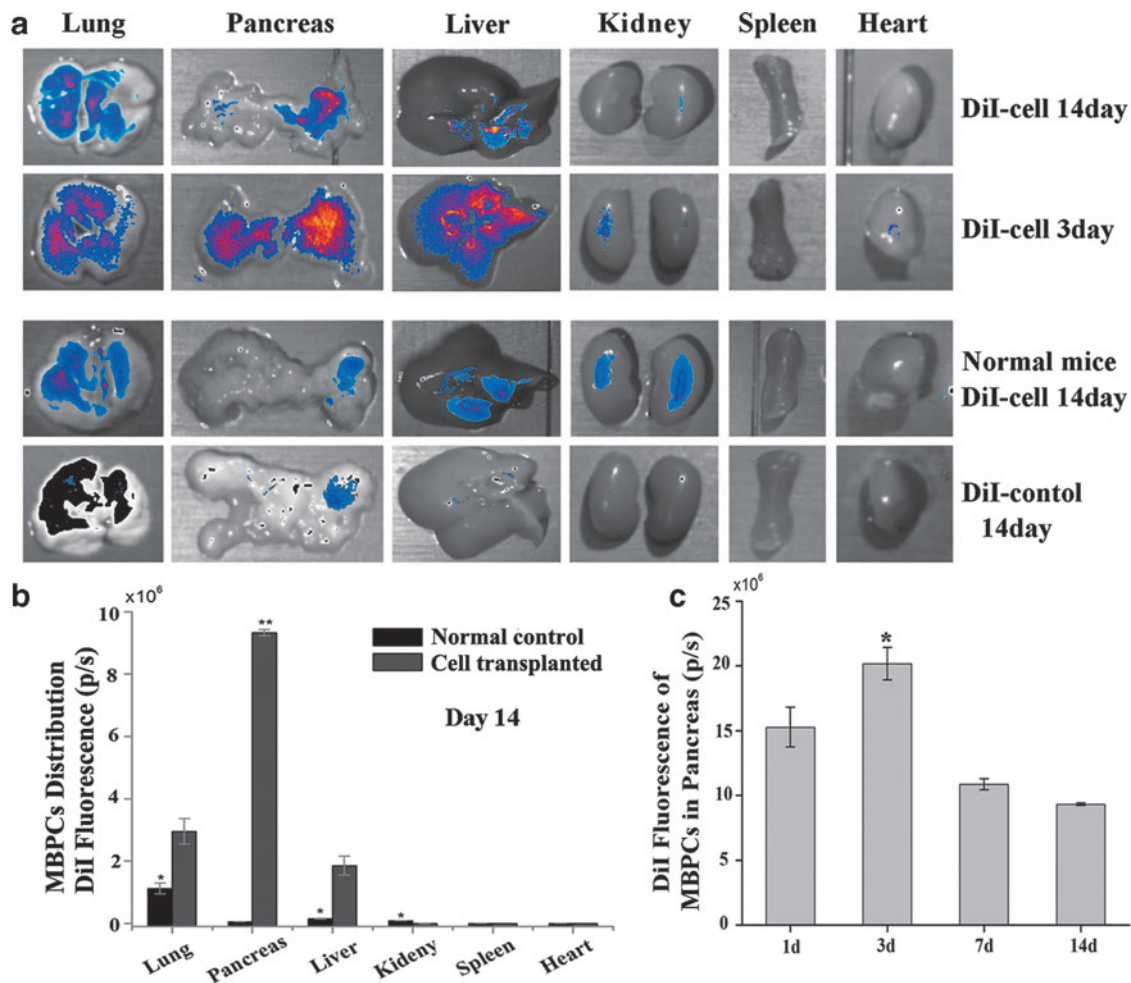


FIG. 4. The distribution of MBPCs in vivo. **(a)** 1,1'-dioctadecyl-3,3,3',3'-tetramethylindocarbocyanine perchlorate (DiI) marked MBPC distribution in vivo of diabetic mice and normal mice on day 3 and/or 14 after transplantation. Each mouse had an intravenous infusion with 5×10^5 MBPCs. In diabetic mice, MBPCs mainly distributed in the pancreas, lungs, and liver (*upper two rows*). And diabetic pancreatic tissue acquired more cells. In the normal mice, MBPCs disseminated in the lungs, liver, and kidney, and were almost undetectable in the pancreas on day 14. **(b)** Representative quantity of MBPCs migrated in different tissues after infusion in T1DM mice and normal mice. The majority of MBPCs resided in injured pancreas in T1DM mice, while most of cells resided in the lungs in normal mice on day 14. **(c)** Representative quantity of MBPCs in diabetic pancreas at different time points. Data are mean \pm SD ($n=3$). * $P < 0.05$, ** $P < 0.01$.

demonstrated that MBPCs could activate more endocrine progenitor cells, which resided not only in the duct but also in islet and exocrine tissue.

To further analyze whether endogenous progenitor cell differentiation was involved in β -cell regeneration, we assayed a series of genes associated with embryonic β -cell development (Fig. 7a) by real-time PCR. In cell-transplanted mice, the average expression levels of *foxa2* and *pdx1* genes (associated with early stages of pancreatic differentiation) were much higher than that in the other two groups. Compared with T1DM mice and normal control, the expression of *ngn3* significantly increased throughout 28 days after MBPC treatment. The expression of *nkx6.1* (associated with mid-stage of β -cell development and maturation) was significantly upregulated in cell-transplanted mice at all time points. The expression of *pax4* (associated with immature and mature stages of β cell) also significantly increased in cell-transplanted mice compared with T1DM mice and normal mice at all time points. In cell-transplanted mice, the gene *mafb* only

expressed in immature β cells had significantly increased on day 3 and 7, and was lower on day 10, 14, 21, and 28. It indicated a transition from immature β cells to mature β cells. The gene *mafa* only expressed in mature β cells had a higher expression in the cell-transplanted mice than in the T1DM mice, and there was no difference in the levels of *mafa* between the cell-transplanted mice and the normal control on day 10, 14, 21, and 28. Moreover, insulin and *glut2*, which were expressed in functional β cells, were significantly increased in cell-transplanted mice compared with T1DM mice.

In T1DM mice, those genes are of irregular expression. On day 3 and 7, gene *foxa2* and *pdx1* significantly increased compared with normal control and *ngn3* increased on day 3, 7, 21, and 28. On day 3, 7, 10, and 28, *pax4* comparatively increased relative to normal control and *nkx6.1* was only upregulated on day 14. Levels of *mafa*, insulin, and *glut2* were much lower than cell-transplanted mice and normal control at all time points. In the normal mice, the above genes, except *mafa*, insulin, and *glut2*, were lower than in

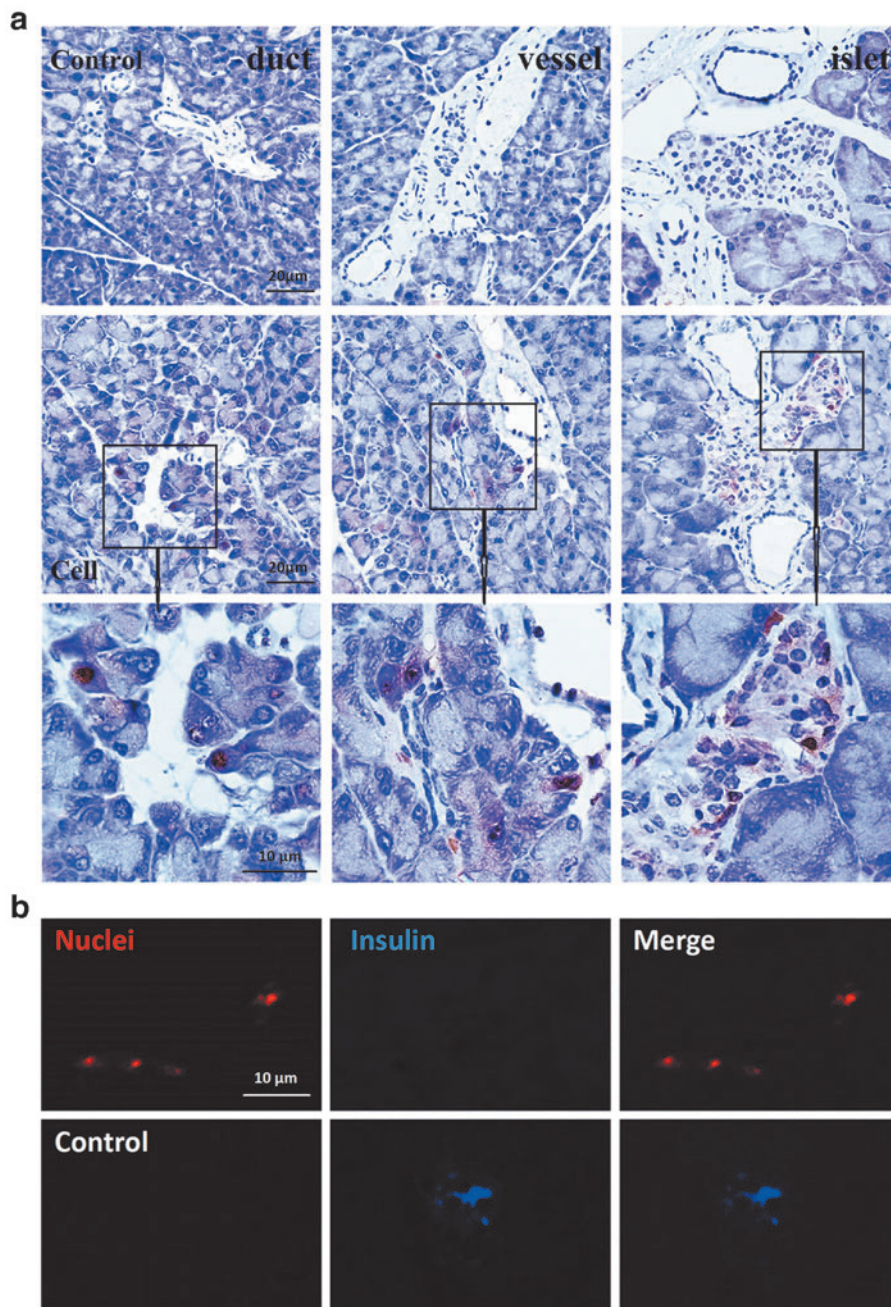


FIG. 5. Immunostaining analysis of location and differentiation of MBPCs in pancreatic tissue. To detect MBPC migration and differentiation in pancreas, mice were sacrificed on day 28 after cell transplantation. **(a)** Immunohistochemical results showed that MBPCs migrated to the damaged pancreas and located at the exocrine regions near the pancreatic duct and vessel, and islet. T1DM mice without cell transplantation were as control. **(b)** Fluorography indicated that MBPCs in pancreas did not differentiate into insulin-producing cells. *Red* fluorescent staining was representative of specific human nuclei, whereas *blue* fluorescence was representative of human insulin.

the other two groups (Fig. 7b), which indicated their homeostasis in the β -cell mass.

Discussion

The characteristics of MBPCs

In this study, we isolated and identified a population of progenitor cells from human menstrual blood (MBPCs). Although MBPCs exhibit some characteristics of MSCs, they also have some distinct properties. First, MBPCs do not express SSEA4 and CD117 (Fig. 1a), which are common markers of BM-MSCs [27–31], whereas they express an embryonic stem cell maker Oct-4 (Fig. 1e). Second, MBPCs are much more proliferative. Under in vitro culture conditions, they can stably expand more than 20 passages without

karyotype abnormalities (Fig. 1f), and their one doubling time is near 20 h (Fig. 1d). MSCs isolated from BM or UCB have a longer doubling time [27,32] and their proliferation capacity will slow down in long-term culture. Additionally, compared with BM-MSCs, MBPCs seem to possess broader multipotent plasticity. MBPCs are able to generate cells of all three germ lines [13]. Thus, these characteristics make MBPCs a unique population different from MSCs obtained from other tissues such as BM and UCB.

The therapeutic effect of MBPCs to T1DM mice

Because of the unique characteristics of MBPCs, their therapeutic potential in diseases deserves to be studied. In our study, we first investigated the therapeutic effect of MBPCs in STZ-induced mice. The results of animal experiments

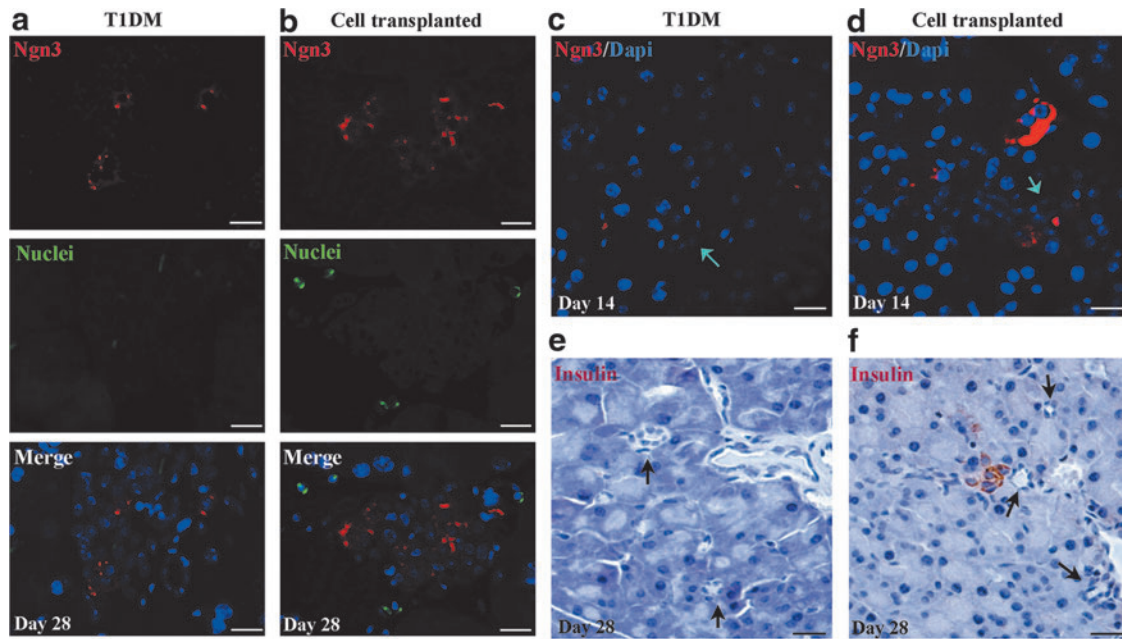


FIG. 6. MBPCs promote endogenous pancreatic progenitor differentiation to compensate for β -cell loss. **(a, b)** Fluorescence photomicrographs showed that neurogenin3 (*ngn3*)-positive cells, which were representative of endocrine progenitor cells, appeared in islet of cell-transplanted mice on day 28 after MBPC transplantation, and there were a few of MBPCs around islet. Less of *ngn3*⁺ cells were observed in islet of T1DM mice. **(c)** *Ngn3* did not express in the exocrine tissue close to the ductal epithelium in T1DM mice. However, **(d)** *ngn3*⁺ cells were detected in exocrine tissue and ductal epithelium in MBPC-transplanted mice. **(c, d)** *Arrows* are representative of pancreatic ducts. **(e, f)** *Arrows* are representative of pancreatic ducts.

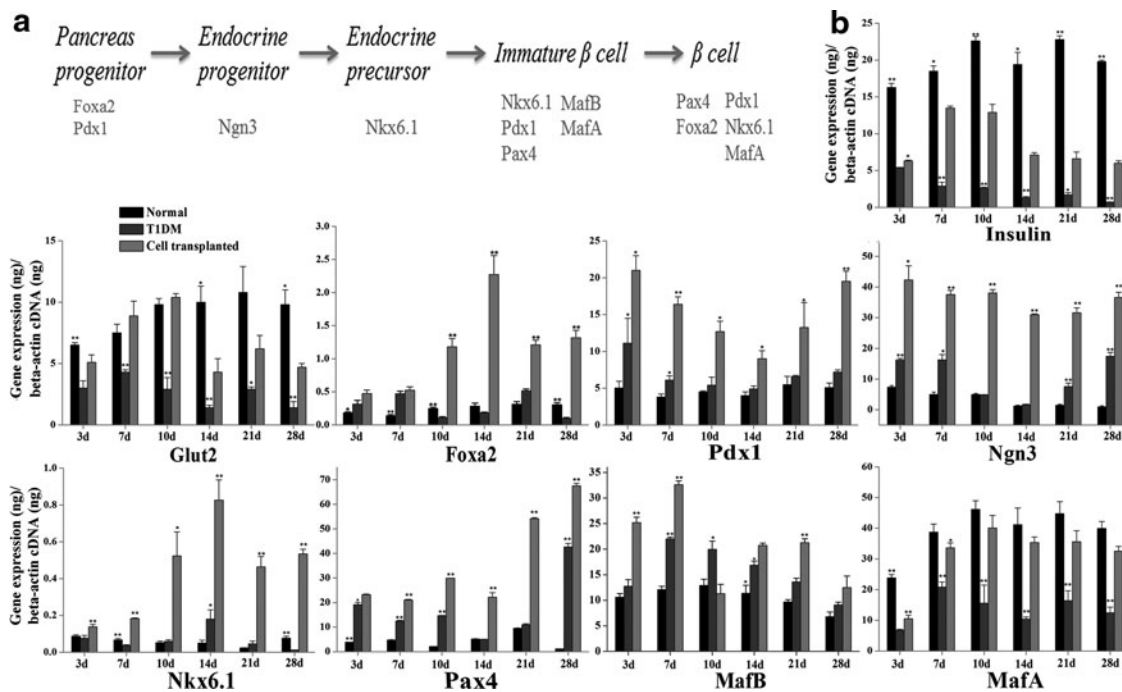


FIG. 7. MBPCs activated the expression of genes associated with embryonic β -cell development and promoted differentiation of pancreatic progenitor to β cell. **(a)** Representative of important genes in the development of islet β cell, from pancreatic progenitor to mature β cell. **(b)** Representative of real-time PCR analysis of genes relating to islet β -cell development at time point of day 3, 7, 10, 14, 21, and 28 after cell transplantation. In functional β cells, the insulin and *glut2* gene levels significantly increased in cell-transplanted mice compared with T1DM control. Genes associated with embryonic β -cell development (*foxa2*, *pdx1*, *ngn3*, *nkx6.1*, and *pax4*) were time dependent and significantly increased in cell-transplanted mice compared with normal mice and T1DM mice. Compared with T1DM mice and normal control, gene *mafb* representative of immature β cell was significantly upregulated in cell-transplanted mice on day 3, 7, 14, and 21. *Mafa* associated with β -cell maturation was significantly upregulated in cell-transplanted mice compared with T1DM mice at all time points. Values are expressed as mean \pm SD ($n=3$).

demonstrated that transplantation of MBPCs was an effective treatment for T1DM mice by ameliorating diabetic symptoms, such as reversing hyperglycemia, maintaining the body weight, improving the survival rate, insulin levels, and glucose tolerance. The levels of blood glucose could be reduced and stabilized at about 15 mM after an infusion of 3×10^5 MBPCs for each mouse. However, to obtain a similar hypoglycemic effect with BM-MSc treatment, the cells need to be purified and sorted with high-aldehyde dehydrogenase activity (ALDHhi), and the dose of cells must achieve to at least 2×10^5 for each mouse [33]. A 10-fold or more of transplanted UCB-MSCs still could not improve hyperglycemia in NOD mice [34]. So, MBPCs reduced blood glucose more efficiently than BM-MSCs and UCB-MSCs.

Recovery effect of MBPCs to diabetic islet

Pathological and histological analyses showed that diabetic mice without MBPCs had severe histological changes in islet and exocrine tissue. However, in cell-transplanted mice, islet structures and morphology remained normal. This showed that MBPCs had a recovery effect in islet. It has reported that STZ selectively destroyed pancreatic β cells through the generation of reactive oxygen species (ROS) and alkylation of DNA [35,36]. In addition, the oxidative stress (OS) caused by ROS contributed to β -cell death or dysfunction in type 1 diabetes [37–39]. Some studies showed that MSCs are able to resist OS under in vitro culture and ionizing radiation circumstances [40,41]. Valle-Prieto et al. found that human MSCs had a high resistance to OS-induced death by expression of enzymes required to manage OS and basal glutathione (GSx) [42]. The study of Cho et al. proved that MSCs could restore liver injury by promoting an antioxidant response [43]. In view of these results together, the fact that MBPCs have a recovery effect on the pathologic structure of diabetic pancreas possibly has a link with OS.

Endogenous β -cell regeneration stimulated by MBPCs

β -cell regeneration stimulated by MBPCs was another important observation in our study. According to the immunohistochemistry analyses, the islet size and β -cell number were recovered to normal levels. Previous studies have indicated that exogenous human progenitor cells migrated into impaired pancreas and then differentiated into IPCs [34,44,45]. To investigate whether MBPCs differentiate into IPCs, we detected the in vivo distribution and differentiation of MBPCs. The results demonstrated two points: (i) the injury signal can recruit a majority of MBPCs to the diabetic pancreas and make them locate at the ductal, exocrine, and islet structures. In normal mice, although part of cells still survive in the lungs and liver tissues on day 14 after transplantation, their survival time is much shorter than cells in diabetic pancreas. The different survival time of MBPCs between injured pancreas and other normal tissues indicates a sealed dependence relationship between injured tissues and progenitor cells; and (ii) MBPCs do not differentiate into IPCs under this diabetic environment. That MBPCs do not differentiate into IPCs implies an endogenous β -cell regeneration mechanism stimulated by MBPCs.

Endogenous pancreatic progenitors have been identified in adult mice pancreas, which revealed that the adult pan-

creas retained the potential to reactivate β -cell development. During embryonic islet development, *ngn3* is considered as an essential master switch for differentiation of endocrine progenitors [46–48]. To investigate whether the differentiation of pancreatic progenitor cells contributed to endogenous β -cell regeneration, we detected *ngn3* and insulin expression by immunostaining in our models. In T1DM mice, *ngn3* was activated, but insulin was reduced (Figs. 3d and 6a). This indicated that injury signals could stimulate endogenous progenitor cell differentiation, but have no effect on functional β -cell formation [49]. This result could be confirmed by quantitative analysis. In T1DM mice, although those genes associated with β -cell development (*pdx1*, *foxa2*, *ngn3*, *nkx6.1*, *pax4*, and *mafb*) are not all upregulated at all time points, the expression levels of them were higher than in normal mice in some time points, but almost much lower than in cell-transplanted mice. Gene *mafa*, insulin, and *glut2* (associated with mature β cell) are much less than in cell-transplanted mice and normal mice. Thus, a damage signal of STZ induction can stimulate pancreatic progenitor differentiation, but cannot repair β -cell damage.

In cell-transplanted mice, *ngn3* was also activated and was higher than in T1DM mice. Insulin was increased as well. Both higher expression of *ngn3* and insulin in cell-transplanted mice indicate that MBPCs promote pancreatic progenitor cell differentiating into β cell. In addition, the results of quantitative analysis further confirm this conclusion. Compared with T1DM mice and normal mice, average levels of the key genes associated with embryonic β -cell development are all the more significantly upregulated in cell-transplanted mice during 28 days after transplantation. Also, the gene levels of insulin, *glut2*, and *mafa* (associated with mature β cells) in cell-transplanted mice also significantly increased compared with T1DM mice. Thus, MBPC-promoted pancreatic progenitor differentiation is a possible mechanism of β -cell regeneration in our models. Interestingly, Bell et al. [33] did not detect any *ngn3*⁺ cells in diabetic mice and cell-transplanted mice in their study. They demonstrated that another putative neogenic mechanism may contribute to β -cell regeneration. It is noteworthy that we use different cell sources and mouse models. The unique characteristics of MBPCs may stimulate different pathways of β -cell regeneration.

Additionally, we also discovered that in cell-transplanted mice, *ngn3*⁺ cells not only resided in the ductal epithelium but also appeared in exocrine tissue and islet, which was in contrast to the viewpoint that *ngn3*⁺ progenitor cells associated with the ductal epithelium [49–51]. The result indicates that MBPCs activated both duct progenitors and other population of pancreatic progenitor cells, which are distinct from ductal progenitor cells [52].

Conclusions

Our study reveals a novel promising cell source for cell replacement and regeneration therapy for diabetes mellitus. MBPCs can be easily isolated under a noninvasive manner. After being transplanted in vivo, MBPCs are efficient to reverse hyperglycemia and restore islet structures. We have also discussed the in vivo repair mechanism about MBPC-stimulated β -cell regeneration. Our data demonstrate that MBPCs promote endogenous progenitor cells differentiating into β cells. The β -cell regeneration may be due to the

paracrine mechanism of MBPCs [22,53]. Clarifying which factors are involved in MBPC-induced β -cell regeneration will aid the future use of MBPCs to treat diabetes.

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Author Disclosure Statement

No competing financial interests exist.

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