Research Article

Islet Stellate Cells Isolated from Fibrotic Islet of Goto-Kakizaki Rats Affect Biological Behavior of Beta-Cell

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We previously isolated islet stellate cells (ISCs) from healthy Wistar rat islets. In the present study, we isolated “already primed by diabetic environment” ISCs from islets of Goto-Kakizaki rats, determined the gene profile of these cells, and assessed the effects of these ISCs on beta-cell function and survival. We detected gene expression of ISCs by digital gene expression. INS-1 cell proliferation, apoptosis, and insulin production were measured after being treated with ISCs supernatant (SN). We observed the similar expression pattern of ISCs and PSCs, but 1067 differentially expressed genes. Insulin production in INS-1 cells cultured with ISC-SN was significantly reduced. The 5-ethynyl-2'-deoxyuridine-positive INS-1 cells treated with ISC-SN were decreased. Propidium iodide-(PI-) positive INS-1 cells were 2.6-fold higher than those in control groups. Caspase-3 activity was increased. In conclusion, ISCs presented in fibrotic islet of GK rats might be special PSCs, which impaired beta-cell function and proliferation and increased beta-cell apoptosis.

1. Introduction

Type 2 diabetes mellitus (T2DM) has reached pandemic proportions, and current predictions show that this trend will continue [1, 2]. Therefore, achieving a better understanding of this complex disease is imperative. Islet fibrosis in T2DM has received increasing scientific attention [3–10]. Studies showed that pancreatic stellate cells (PSCs) have important functions in islet fibrogenesis in both rodent animal models and human patients with T2DM [6, 7, 11, 12]. In addition, we observed that high glucose aggravates the detrimental effects of pancreatic stellate cells on beta-cell function [13]. To elucidate the underlined mechanisms responsible for islet fibrosis in the late stage of T2DM, we find that endocrine pancreatic islets contain cells resembling PSCs and suggest these may contribute to islet fibrosis in T2DM [14].

In the normal pancreas, PSCs are quiescent and are found in low abundance [15, 16]. Upon pancreatic injury or pancreatic inflammation, PSCs lose their vitamin A stores and transform from “activated” into myofibroblast-like phenotypes, which highly proliferate, migrate, synthesize, and secrete excessive amounts of the extracellular matrix (ECM) proteins, resulting in tissue fibrosis [15–23].

The function of PSCs in islet fibrosis has been the subject of several studies for years. In vivo studies have shown that PSCs are present in rat islets and are involved in islet fibrogenesis in several animal models of T2DM [10, 11, 24–27]. In humans, PSCs are also present in islets of T2DM patients and possibly have a function in the progression of islet fibrosis [11]. Fibrosis is one of the major factors leading to progressive pancreatic beta-cell loss and dysfunction [6, 11, 28–33]. Efforts have been made for developing antifibrotic strategies to ameliorate islet fibrosis and the progression of T2DM [6, 12, 34–36]. In vivo studies showed that the attenuation of PSC activation reduces islet fibrosis [6, 12, 26, 36] or increases insulin content [26].
Kikuta et al. recently reported that indirect coculture of RIN-5F cells with PSCs results in decreased insulin production and increased cell apoptosis [37], and very recently published data showed that activated PSCs can impair pancreatic islet function in mice [38]. We observed that PSCs transplantation exacerbated the impaired β-cell function in GK rats [13] and demonstrated that PSCs resemble cells within endocrine pancreatic islets [14]. However, in T2DM, the PSCs that presented intra-/peri-islets were exposed to the islet niche, such as marked hyperglycemia [28, 39, 40], oxidative stress [41–43], and inflammation [4, 28, 39]. These factors could affect PSCs activation and proliferation and subsequently stimulate the production of endogenous inflammatory mediators in PSCs [17]. Thus, PSCs were already “activated” by the environmental conditions of T2DM in vivo. Therefore, we isolated stellate cell from fibrotic islets of Goto-Kakizaki (GK) rats, and our previous data pushed us to question whether these population cells were a special type of PSCs and which effect is on biological behavior of beta-cells.

2. Methods

2.1. Animals and Ethics Statement. Rats were housed in cages (three rats per cage) under a 12 h/12 h light/dark cycle. Rats were given free access to food and water ad libitum. Animal experiments were approved by the Southeast University Animal Care and Use Committee according to institutional guidelines and national animal welfare.

2.2. Isolation of PSCs. PSCs were isolated from 8-week-old Wistar rats as described previously [16]. PSCs were cultured in Dulbecco’s modified Eagle’s medium (DMEM)/Ham’s F12 (1:1 v/v) containing 10% fetal bovine serum (FBS) (Invitrogen, Carlsbad, USA). Cell purity was assessed by immunostaining for vimentin (100%), α-SMA (>95%), and desmin (20–50%).

2.3. Isolation and Culture of ISC and Preparation of ISC-SN. Islets were isolated from four-month-old male GK rat pancreas as previously described [4]. In brief, pancreas tissues were digested with collagenase V (1 mg/mL, w/v) (Sigma, St. Louis, MO, USA) at 37°C for 15 min to 18 min. Islets were purified by handpicking twice under a stereomicroscope. Then, islets were precultured in RPMI-1640 supplemented with L-glutamine containing 10% (v/v) fetal bovine serum (FBS) (Invitrogen, Carlsbad, CA, USA) overnight followed by handpicking.

After 48 h in culture, ISC-SN began to grow out of GK islets. After 5 d, cells were subcultured in DMEM/Ham’s F12 (1:1, v/v) containing 10% FBS. Cells at passages 3 to 8 were used for experiments.

To prepare ISC-SN, cells grown near confluence were cultured with DMEM/F12 serum-free medium plus 0.2% BSA for another 48 h. The culture medium was collected, centrifuged, filtered, and stored at −80°C until use.

2.4. Immuno-Fluorescence Staining. After shortly being washed with cold phosphate buffer saline (PBS), cells were fixed with 4% paraformaldehyde at room temperature for 20 min and subsequently stained with primary antibodies at the following dilutions: α-SMA, (1: 100, DAKO, Hamburg, Germany), vimentin (1: 200, DAKO), and desmin (1: 50, DAKO), followed by fluorescent secondary antibodies (DAKO). The nuclei were then counterstained with bisbenzimide. Photos of eight different areas in each well were taken using a microscope with 100x magnification. A minimum of 500 cells in each experimental group was analyzed (n = 3).

2.5. Digital Gene Expression (DGE) Profile of ISC and PSC. The passage 3 ISC and PSCs were used for DGE analysis. DGE was performed by the BGI Tech (Shenzhen, China).

2.6. Treatment of INS-1 Cells with ISC-SN. Insulin-producing β-cell line (INS-1 832/13) cells were seeded into RPMI-1640 medium with L-glutamine containing 10% FBS. Upon reaching confluence, INS-1 cells were treated with or without 35% ISC-SN for up to 48 h. After exposure to ISC-SN, INS-1 cells were washed in RPMI 1640, and the following experiments were performed, representing n = 3 to 4.

2.6.1. 5-Ethynyl-2'-deoxyuridine (EdU) Incorporation Assay. EdU incorporation assay was performed as previously described [44]. In brief, EdU (Molecular Probes, Eugene, OR, USA) was added 4 h before the experiments ended and stained. Photos of eight different areas in each well were taken using a microscope with 200x magnification. The average percentage of EdU+/DAPI+ was calculated.

2.6.2. Propidium Iodide (PI) Staining. After removing the medium, INS-1 cells were incubated with PI (20 μg/mL) and Hoechst (2 μg/mL) (Sigma, St. Louis, MO, USA) for 10 min in the dark. Photos of eight different areas in each well were taken using a microscope with 200x magnification. The average PI staining rate was expressed as the percentage of PI-positive nuclei compared with the total cell count.

2.6.3. Caspase-3 Fluorometric Assay (CFA). CFA was performed according to the manufacturer’s instructions (R&D, Minneapolis, MN, USA). CFA was performed in a 96-well flat-bottom microplate. After incubation with the substrate for 1.5 h, the plate was read on a fluorescent microplate reader equipped with filters set at 400 nm/505 nm (excitation/emission wavelengths). Data were normalized with total protein content in each well and expressed as OD/μg/μL.

2.6.4. Quantitative RT-PCR (qRT-PCR). Total RNA of INS-1 cells was extracted using TRIzol reagent (Invitrogen). The primer sequences used for amplification of genes encoding insulin and GAPDH are listed in Table 1. qRT-PCR analyses were performed using a standard SYBR-Green PCR kit protocol on a Step One Plus system (Applied Biosystems, Foster City, CA, USA) according to the manufacturer’s instructions. The relative level of insulin transcripts was calculated and normalized to GAPDH, with at least four repeats per experimental group.
2.6.5. Potassium-Stimulated Insulin Secretion (KSIS) Assay. Measurements of potassium-stimulated insulin secretion (KSIS) by INS-1 cells were performed as described [13, 45]. Briefly, after preincubation in glucose-free Krebs-Ringer bicarbonate buffer for 1 h, cells were treated with low glucose (2.8 mmol/L) and high potassium (50 mmol/L) for 1 h. Insulin secretion after stimulation and insulin content in the cell lysate were measured using an insulin radioimmunoassay kit (Beijing Technology Company, Beijing, China). Data were normalized to the cellular insulin content and expressed as a percentage.

2.6.6. Measurement of Cytokines in ISC-SN. To determine the cytokines present in ISC-SN, the RayBio Biotin Label-Based Rat Antibody Array 1 (RayBiotech, Norcross, GA, USA) was performed according to the manufacturer's recommendations.

2.7. Statistical Analysis. Data are presented as the mean ± SE. Statistical significance was determined by unpaired Student's t-test or ANOVA, followed by the Bonferroni–Dunn post hoc test. \( P < 0.05 \) was considered statistically significant. All statistical analyses were performed using the Statistical Product and Services Solutions (SPSS) package (Version 11.5, SPSS Science, Chicago, IL, USA).

3. Results

3.1. Isolation and Characterization of ISCs. After 48 h of culture, cells with triangular shapes and large nuclei began to grow out of GK rat islets (Figure 1(a)). When the culture period was extended, these cells migrated away from the islet (Figure 1(b)). In addition, these stellate cells had a shorter doubling time (28 h). To determine these cells markers, immunofluorescence staining was performed. The ISC was positive for \( \alpha \)-SMA (Figure 2(a)), vimentin (Figure 2(b)), and desmin (Figure 2(c)), thereby resembling the pattern of PSCs (Figures 2(d), 2(e), and 2(f)).

3.2. The DGE Profile of ISC Compared with PSC. Taking advantage of the DGE performed by the BGI Tech, we observed that the relative mRNA expression levels of ISCs were similar to PSCs mRNA expression pattern in intermediate filaments (\( \alpha \)-SMA, vimentin, and desmin), ECM (Procollagen, Pro-\( \alpha \)-1 collagen type, Collagen \( \alpha \)-1 type IX, Collagen \( \alpha \)-1 type XI, Collagen \( \alpha \)-1 type III, Fibronectin, Laminin, Tenasin, TIMP1, TIMP2, MMP1, MMP2, MMP4, RECK, and TUBA6), cytokines (IL-1\( \beta \), IL-6, IL-7, IL-15, IL-18, TGF-\( \alpha \), TGF-\( \beta \)1, TGF-\( \beta \)2, TGF-\( \beta \) binding protein 2, TGF-\( \beta \) binding protein 3, HGF, PDGF-A, PDGF-B, CTGF, RANTES, MCP-1, ET-1, and VEGF), signal transduction (Smad1, Smad2, Smad7, Smad8, ERK1, ERK3, and ERK5), integrins (integrin \( \alpha \)-1, integrin \( \alpha \)-E2, integrin \( \alpha \)-M, integrin \( \alpha \)-v, integrin \( \beta \)-3, integrin \( \beta \)-5, and integrin associated protein), cytokine receptors (TGF-\( \beta \)R type 1, PDGFR\( \beta \), PGFR\( \beta \), FGFR1, IL-3R\( \beta \), and ActivinR), and PPAR\( \gamma \). However, there were 1657 genes differentially expressed (600 genes upregulated and 1057 genes downregulated) of ISCs compared with that of PSCs, which enriched GO terms in metabolic pathways (196, 13.75%), pathways in cancer (63, 4.42%), MAPK signaling pathway (48, 3.37%), focal adhesion (46, 3.23%), regulation of actin cytoskeleton (44, 3.09%), biosynthesis of secondary metabolites (39, 2.74%), Alzheimer's disease (38, 2.67%), cell cycle (29, 2.04%), tight junction (37, 2.6%), vascular smooth muscle contraction (34, 2.39%), microbial metabolism in diverse environments (34, 2.39%), and chemokine signaling pathway (33, 2.32%), as well as Wnt signaling pathway (32, 2.5%) (for more information please see Supplementary Table 1 in the Supplementary Material available online at http://dx.doi.org/10.1155/2016/6924593).

3.3. ISC-SN Impaired INS-1 Cell Survival. Using EdU incorporation assay, the effects of ISC-SN on INS-1 cell proliferation were assessed. The proliferative capacity of INS-1 cells...
Figure 2: Stellate shape-like cell expression of (a) α-SMA, (b) vimentin, and (c) desmin and PSC expression of (d) α-SMA, (e) vimentin, and (f) desmin. Representative images for α-SMA and desmin staining were red, and those for vimentin were green at 100x magnification.

3.4. ISC-SN Reduced Insulin Secretion of INS-1 Cells. To investigate the effects of ISC-SN on beta-cell function, insulin mRNA expression and secretion in INS-1 cells incubated with ISC-SN for 24 h were evaluated. Figure 5(a) showed that the insulin mRNA expression in INS-1 cells treated with ISC-SN (0.21 ± 0.10) was more significantly decreased than that in the control group (P < 0.01). In response to potassium challenge, insulin secretion in INS-1 cells incubated with ISC-SN (1.03% ± 0.23%) was more significantly reduced than that in control groups (1.71% ± 0.17% μIU/mL, P < 0.01) (Figure 5(b)).

3.5. Cytokine Profile of ISC-SN. To determine the cytokines that were highly expressed in ISC-SN, RayBio Biotin Label-Based Rat Antibody Array 1 was performed. Cytokines, such as integrin αMβ2, IFN-γ, MCP-1, MMP-2, PDGF, TNF-α, TLR4, and TROY, were highly expressed in both ISC-SN and PSCs-SN (Table 2), whereas CD106, IL-1β, CXCR4, IL-3, IL-5, and MIF were highly expressed in ISCs-SN (Table 3).

3.6. ISC-SN Reduced Beta-cell Function. To observe the effects of ISC-SN on INS-1 cell survival, apoptosis in INS-1 cells incubated with ISC-SN was detected by PI staining and CFA. As shown in Figure 4(a), a higher number of PI-positive INS-1 cells were observed in ISC-SN group. Quantitative analysis of PI-positive index (Figure 4(b)) showed that the dead cells in the ISC-SN group (0.18 ± 0.03) were 2.6-fold higher than those in the control (0.07 ± 0.02) group. Caspase-3 activity in the ISC-SN group (338.09 ± 62.51) was significantly higher than that in control groups (80.40 ± 16.59, P < 0.01) (Figure 4(c)).

4. Discussion
In this study, we suggested an effective method for isolating ISCs from fibrotic islets of four-month-old GK rats. We observed a significant overlap and differentially expressed genes of ISCs and PSCs. In addition, ISCs exerted deleterious effects on beta-cells.

Previous studies have proven the existence of islet fibrosis in T2DM patients and rodent animal models, including GK
Table 3: Cytokines were highly expressed in ISC-SN than in PSCs-SN. MFI of ISC-SN ≥ 1.5-fold of PSCs-SN was considered as highly expressed.

|                | ISC-SN (MFI) | PSCs-SN (MFI) | Fold |
|----------------|--------------|---------------|------|
| Negative control | 5828.5       | 6319.8        |      |
| CD106          | 47241.0      | 10725.0       | 4.40 |
| IL-1β          | 22605.5      | 10431.0       | 2.24 |
| CXCR4          | 23640.0      | 10119.5       | 2.34 |
| ICAM-1/CD54    | 41550.0      | 12306.5       | 3.38 |
| IL-3           | 23586.0      | 1006.5        | 2.14 |
| IL-5           | 12685.0      | 5296.0        | 2.40 |
| MIF            | 10394.0      | 6536.5        | 1.59 |
|                | 37598.0      | 17121.5       | 2.20 |

Table 3: Cytokines were highly expressed in ISC-SN than in PSCs-SN. MFI of ISC-SN ≥ 1.5-fold of PSCs-SN was considered as highly expressed.

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Figure 3: Effect of ISC-SN on INS-1 cell proliferation. (a) Cell proliferation was performed using EdU incorporation assay after treatment of INS-1 cells with ISC conditional medium for 48 h. Representative images for EdU staining (green) and nuclei labeled by DAPI (blue) at 200x magnification. (b) Data were expressed as mean ± SE (n = 3), **P < 0.01.

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The interesting results in this study are the deleterious effects of ISC on INS-1 cell function and survival. Previous studies have clearly shown that PSCs reduced insulin expression and induced apoptosis in pancreatic beta-cell [37], and high glucose aggravates the detrimental effects of PSCs on beta-cell function [13]. In this study, our data indicated as circulating fibrocytes. To determine which cell types are present in fibrotic islets of T2DM, islets from four-month-old GK rats were isolated and cultured. Although most islets in 16-week-old GK rats were significantly deformed with massive fibrosis, 90 ± 16 islets were harvested per pancreas using handpicking after mild digestion with collagenase V for 15 min to 18 min. As shown in Figure 1, stellite-like cells began to grow out of the islets after 48 h of culture, and these cells often exhibited a triangular shape and a large nucleus. Immunofluorescence staining demonstrated that these cells were positive for α-SMA, vimentin, and desmin, thereby resembling the protein expression profile of PSCs [15, 16]. Given that these cells were isolated from fibrotic islets of GK rats, which were presumably already influenced by the surrounding islet microenvironment in T2DM, were highly similar to activated PSCs. The isolation of ISCs from fibrotic islets in GK rats may provide a tool to investigate the interactions between ISCs and pancreatic beta-cell in vitro.
Figure 4: Effect of ISC-SN on INS-1 cell apoptosis. (a) Cell apoptosis was determined by PI staining (red), (b) PI-staining index, and (c) caspase-3 activity in INS-1 cells cultured with ISC-SN for 48 h. Data are expressed as mean ± SE (n = 3), **P < 0.01.

Figure 5: Effects of ISC-SN on insulin mRNA transcript and INS-1 cell function. (a) qRT-PCR analysis of insulin mRNA expression and (b) KSIS assay on INS-1 cells cultured with ISC-SN for 24 h. Insulin mRNA data were normalized by GAPDH gene, and insulin secretion data were normalized by cellular insulin content. Data are expressed as mean ± SE (n = 4 to 5), **P < 0.01.
that ISC also exhibited deleterious effects on INS-1 beta-cells by reducing insulin mRNA transcription and inhibiting insulin secretion. In addition, ISCs significantly decreased cell proliferation and increased cell apoptosis. To determine the factors that possibly contribute to the effect of ISCs on INS-1 cells, cytokine profiles in ISCs were assessed. ISCs-SN was shown to contain high levels of cytokines, such as IFN-γ and TNF-α, which are well-known factors mediating or promoting cell death [50–55]. Our data also show that ISCs-SN contained higher levels of cytokines than PSCs-SN, such as IL-1β, which can impair beta-cell function and induce cell death at higher concentrations [56].

5. Conclusion

In conclusion, our in vitro study shows that fibrotic islets of GK rats contain a population of stellate cells resembling but not identical to PSCs, which exerted deleterious effects on beta-cells by inducing beta-cell death and suppressing insulin production and cell proliferation.

Conflict of Interests

The authors declare that there is no conflict of interests regarding the publication of this paper.

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