

Differentiation of Multipotent Stem Cells to Insulin-Producing Cells for Treatment of Diabetes Mellitus: Bone Marrow- and Adipose Tissue-Derived Cells Comparison

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Abstract

Background: Mesenchymal stem cells (MSCs) from human adipose tissue and bone marrow have a great potential for use in cell therapy due to their ease of isolation, expansion, and differentiation. Our intention was to isolate and promote in vitro expansion and differentiation of MSCs from human adipose and bone marrow tissue into cells with a pancreatic endocrine phenotype and to compare the potency of these cells together.

Methods and Results: MSCs were pre-induced with nicotinamide, mercaptoethanol, B-27 and b-FGF in L-DMEM for 2 days and re-induced again in supplemented H-DMEM for another 3 days. Expression of five genes in differentiated beta cells was evaluated by Real-time PCR and western blotting and the potency of insulin release in response to glucose stimulation was evaluated by insulin and C-peptide ELISA kit.

Quantitative RT-PCR results showed up-regulation of four genes in differentiated beta-islet cells (Insulin, Ngn-3, Pax-4 and Pdx-1) compared with the control. Western blot analysis showed that MSCs cells mainly produced proinsulin and insulin after differentiation but nestin was more expressed in pre-differentiated stem cells. Glucose and insulin secretion assay showed that insulin levels and C-peptide secretion were significantly increased in response to 10 mM glucose.

Conclusions: Our study showed that both adipose and bone marrow stem cells could differentiate into functional beta-islet cells but it seems that adipose stem cells could be a better choice for treatment of diabetes mellitus according to their more safety and potency.

Introduction:

Insulin dependent diabetes mellitus (IDDM) is one of the most common chronic childhood diseases that caused usually by destruction of insulin-secreting pancreatic β -islet cells due to self-immune attack of T-lymphocytes [1]. Although daily insulin injections as main treatment of IDDM can control blood glucose level, but exogenous insulin does not completely imply the physiology of natural insulin secretion, therefor it may have complications such as severe hypoglycemia and coma [2–4]. The replacement of beta cells by pancreatic transplant to the patient is another method which associated with major problems, including limitation in the pancreatic donor and immunologic rejection of the transplant [5, 6]. Stem cell therapy has provided many hopes for the curable treatment of many autoimmune diseases, including diabetes [7]. There are several sources of stem cells with different potencies. Pluripotent cells are embryonic stem cells derived from the inner cell mass of the embryo and induced pluripotent stem cells (iPSCs) are formed following reprogramming of somatic cells. Pluripotent cells can differentiate into tissue from all 3 germ layers (endoderm, mesoderm, and ectoderm). Multipotent stem cells may differentiate into tissue derived from a single germ layer such as mesenchymal stem cells (MSCs) which form adipose tissue, bone, and cartilage. Tissue-resident stem cells are oligopotent since they can form terminally differentiated cells of a specific tissue. In fact, over the years, various types of stem cells, including embryonic stem cells (ESCs) [8, 9] and mesenchymal stem cells (MSCs) [10], have the ability to

differentiate into insulin producing cells. According to some difficulties such as destroying human embryo in its early stages of development, using embryonic stem cells (ESCs) involved ethical issues. MSCs are multipotent stem cells that have been introduced as new sources of insulin-producing cells and do not have ethical problems related to ESCs [11, 12]. MSCs have ability to homing in damaged tissues [13], and naturally produce cytokines and growth factors, such as IL-6, M-CSF, IL-10, HGF, TGF- β and PGE2 that preventing apoptosis of beta-pancreatic cells and increasing their reengineering and angiogenesis [14–16]. Furthermore, through various immunomodulatory pathways, in a non-MHC restricted manner these cells can suppress active immune reactions and have emerged as an interesting therapeutic option [17, 18]. A wide range of CD antigens/Cluster of differentiation markers have been reported to be a useful tool for characterization of stem cells, especially for human species. The CD antigens can be used to exclude undifferentiated cells from other populations, and are most commonly used markers to identify mesenchymal stem cells. The most common surface marker molecules used to describe MSCs are positive surface markers such as CD29, CD44, CD54, CD105 whereas negative antigens are CD11, CD18, CD19, and CD45. These surface markers are different and naturally expressed on stem cell precursors, endothelial or epithelial cells, natural killer cells and etc. MSCs can be obtained from variety of sources including bone marrow, adipose tissue, umbilical cord, endometrium, peripheral blood monocytes, liver cells and dental pulp [19]. But the adipose tissue and bone marrow have preferable advantage in terms of availability and abundance. The clinical applicability of bone marrow stem cells (BMSCs) may be limited due to the invasive procedure required for sample collection. On the other hand, for adipose stem cells (ADSCs) liposuction aspirates are widely available and many studies reported that adipose tissue contain a significantly greater number of mesenchymal stem cells than bone marrow per unit weight and should not be wasted. Collectively, the properties of ADSCs render them well suited for applications in regenerative medicine and can provide a viable alternative for BMSCs. In this study we aimed to compare the potency of human ADSCs and BMSCs to differentiate to functional beta-islet pancreatic cells for in vitro insulin production that it may offer a potentially effective therapeutic approach in cell therapy.

Materials And Methods:

1. Isolation and culture of human adipose and bone marrow stem cells:

Human adipose tissue is collected by needle biopsy or liposuction aspiration at surgery ward of Shariati Hospital, Tehran, Iran. The protocol was performed in accordance with the Declaration of Helsinki and approved by the Cellular Transplantation Ethics Committee of Iranian public hospitals and also Zabol University of Medical Sciences ethics committee. All participants informed about the proposed intervention and we emphasized their roles in our decision-making process and also all of the participants informed about the risks of proposed intervention. Adipose tissue was washed extensively with sterile phosphate-buffered saline (PBS) and 1% penicillin-streptomycin to remove red blood cells. Washed tissue fragments placed in a sterile tissue culture plate with 0.075% Collagenase Type I (Sigma-Aldrich, USA) prepared in PBS containing 2% penicillin-streptomycin for tissue digestion. After incubation

for 30 min at 37 °C and 5% CO₂, collagenase activity was neutralized by adding 5 ml of α-MEM containing 20% heat inactivated fetal bovine serum (Gibco, Thermo Fisher Scientific, USA) and centrifuged in 4000 g for 10 minutes. To lyse remaining red blood cells, pellet was re-suspended in 160 mM NH₄Cl at room temperature for 10 minutes and then centrifuged at 1,200×g for 10 min to pellet the MSC-rich dense cell fraction. Afterward, the collected cells were filtered through 100-µm cell strainer (Grainer, USA) and incubated for 48 h at 37 °C in 5 % CO₂ in culture medium with serum and antibiotics. Bone marrow was isolated from one patient in hematology ward of Shariati hospital, Tehran, Iran under aseptic condition and dispersed into single cell suspension, L-DMEM cells were cultured in a density of 1.5×10^5 /cm² in alpha-MEM (Sigma, USA), 10 % FBS, 1% penicillin-streptomycin and incubated at 37 °C and 5 % CO₂ for 3 days and later the medium was refreshed every 48 h till confluence was reached.

2. Characterization of cultured adipose and bone marrow stem cells

Flow cytometry analysis was used to determine positive (CD29 and CD105) and negative (CD18 and CD34) markers. The cells were washed with PBS containing 2 % FBS. FITC conjugated anti human CD18, CD29, CD34 and CD105 monoclonal antibodies (Cell Signaling Technology, USA). The fluorescence of the cells was immediately determined by a FACS-Calibur flow cytometry (Becton Dickinson, San Jose, USA). The FACS flow cytometry data were analyzed using freely available WinMDI software ver.2.9 (<http://facs.scripps.edu/software.html>).

3. Differentiation of MSCs into functional islet beta cells

At passage 3, adipose and bone marrow derived stem cells were induced to differentiate into functional pancreatic cells. According to previous reports about beta cell differentiation, we used design of experiment (DOE) to find cause-and-effect relationships and to determine the relationship between factors affecting our differentiation process. Among our designed experiments we decided to choose our best quality method. Therefore, cells were plated in L-DMEM medium supplemented with 20% FBS and 1% penicillin-streptomycin for 10 days at 37°C and 5% CO₂. Second, the cells were cultured in differentiation medium containing L-DMEM supplemented with 10% FBS, 2% B27, b-FGF, and 10 mM nicotinamide (all purchased from Sigma-Aldrich) for 7 days and finally, the cells were cultured in induction medium containing H-DMEM supplemented with 2% B27, b-FGF, 10 ng/ml activin A, 10 mM nicotinamide, and 1 mM 2-mercaptoethanol (Sigma-Aldrich) for another 6 days. Cells cultured in L-DMEM medium without an inducer were used as controls.

4. Real-time PCR analysis

Total RNA was isolated utilizing the high pure RNA isolation kit (Roche, Mannheim, Germany), according to the manufacturer's instructions. For synthesis of first strand cDNA, total RNA (0.2 µg) from each sample, before and after differentiation was used in a random primed reverse transcription (RT) reaction utilizing the RevertAid™ First Strand cDNA Synthesis Kit (Fermentas, Lithuania). Quantitative real-time PCR using custom made gene specific primers (Invitrogen, Carlsbad, CA) was performed to evaluate mRNA expression before and after differentiation. Target genes and the primer sequences used for qRT-PCR were described in Table 1. Quantitative real-time PCR (qPCR) reactions were carried out using the HOT MOLPol EvaGreen qPCR Mix Plus (Solis BioDyne, Estonia) in a total volume of 20 µl on ABI Prism 7900 Sequence Detector System (Applied Biosystems) according to the manufacturer's instructions. The following thermal profile was used: One cycle at 95 °C for 15 min and 40 cycles of denaturation at 95 °C for 15 s, annealing at 60–65 °C for 20 s and elongation at 72 °C for 20 s. All qRT-PCR results were normalized to GAPDH gene as reference gene. The results were expressed as the target/reference ratio of differentiated cells divided by the target/reference ratio of the calibrators (undifferentiated control samples) [28].

Table 1

Forward and reverse primers used for beta-cell differentiation in real-time PCR

5. Western blot analysis

Protein expression of nestin and insulin as two markers of ADSCs and BMSCs before and after differentiation to beta-islet cells was evaluated by western blot analysis. Cells before and after differentiation were homogenized in a lysis buffer containing 50 mM Tris (pH 7.4), 2 mM EDTA, 2 mM EGTA, 2 mM NaF, 1 mM Na₃VO₄, 10 mM B-glycerophosphate, 10 mM 2-mercaptoethanol, sodium desoxycholate, and 1 µL protease and phosphatase inhibitor cocktail and 1 mM phenylmethanesulfonyl. Proteins were separated on a 12% SDS-PAGE under reducing conditions at a constant voltage (120 v) for 1:45 h. The separated proteins were electro-transferred to the polyvinylidene difluoride (PVDF, Bio Rad) membrane. To block non-specific binding sites, membranes were incubated in 5% non-fat dry milk in TBST buffer (20 mM Tris–HCl pH 7.5, 137 mM NaCl, 0.5% Tween 20) for 1 h at room temperature. After blocking, the membranes were incubated overnight at 4°C with primary antibodies against nestin (1/1000), insulin (1/1000) and β-actin (1/1000). The blots were then incubated for 1 h at room temperature with a HRP conjugated secondary antibody (1:3000) (All antibodies were purchased from Abcam, USA). Protein bands were visualized using the enhanced chemiluminescence method (ECL detection kit, Pierce Biotech, Inc.) and Alliance 4.7 Gel Doc (UVTEC, Cambridge). Band intensities were quantified using UV BAND image analysis software (UVTEC, Cambridge).

6. Insulin and C-peptide secretion assay

Gene	Sequence	Annealing temperature (°C)	To determine whether the
<i>Insulin</i>	Forward: 5'- AGCCTTTGTGAACCAACACC-3' Reverse: 5'- GCTGGTAGAGGGAGCAGATG-3'	54	
<i>Nestin</i>	Forward: 5'- GCGGGGCGGTGCGTGACTAC-3' Reverse: 5'- AGGCAAGGGGGAAGAGAAGGATGT-3'	56	
<i>NGN-3</i>	Forward: 5'- AGGCAGTCTGGCTTTCTCAG-3' Reverse: 5'- GAACAAGTGCTTTTGAGGGC-3'	53	
<i>PAX-4</i>	Forward: 5'- TGGAGAAAGAGTTCCAGCGTG-3' Reverse: 5'- TGGAAAACCAGACCCTCACC-3'	56	
<i>PDX-1</i>	Forward: 5'- TTCAACAAGTACATCTCACGGC-3' Reverse: 5'- TTGTCCTCCTCCTTTTTCCA-3'	54	
<i>NeuroD1</i>	Forward: 5'-GCGGCCCCAAAAGAAGAAG-3' Reverse: 5'-TCCGACAGAGCCCAGATGTA-3'	57	
<i>NKX6.1</i>	Forward: 5'- GGGCTCGTTTGGCCTATTCGTT-3' Reverse: 5'- CCACTTGGTCCGGCGGTTCT-3'	55	
<i>PC1/3</i>	Forward: 5'- CGCTGACCTGCACAATGACT-3' Reverse: 5'- CAGACAACCAGGTGCTGCAT-3'	56	
<i>IAPP</i>	Forward: 5'- TGCTCTCTGTTGCATTGAACC- 3' Reverse: 5'- GTTGCATTTCCGCTTTTCCA-3'	54	
<i>GAPDH</i>	Forward: 5'- TGGTATCGTGGAAGGACTCATGA-3' Reverse: 5'- ATGCCAGTGAGCTTCCCGTTCAGC-3'	56	

differentiated ADSCs and BMSCs were responsive to glucose stimulation, insulin release was measured after exposure to high glucose level using human insulin ELISA kit (Abcam, Cambridge, UK). After differentiation of adipose and bone marrow stem cells, they were washed twice with PBS and incubated in low glucose DMEM culture media for 3 h and then the medium was collected and stored at -20°C. In next step, cells were washed twice with PBS and incubated for 3 h in high glucose DMEM culture media and again, the medium was collected and stored at -20°C. C-peptide content of culture medium was determined by using C-peptide ELISA kits (Abcam, Cambridge, UK) according to the manufacturer's instructions. Briefly, 50 µl of samples was added to 100 µl C-peptide HRP and biotin conjugate in a 96-well plate and incubated for 2h at room temperature. After washing three times with 300 µl washing

solution, 100 μ l tetramethylbenzidine (TMB) was added and incubated for 15 minutes at room temperature in the dark. Then, 100 μ l stop solution was added to the wells and mixed for 10 seconds on a shaker. Finally, the absorbance of the sample at 450 nm within 5 minutes of addition of the stop solution against a reference wavelength of 620-630 nm was measured.

7. Statistical analysis

The results obtained from the experiment were presented as means \pm SD. Each experiment will be repeated for 3 times. Student's t-test was used to compare the means of two groups. P value of less than, or equal to 0.05 was considered to be statistically significant. The statistical analysis was performed using the SPSS 16.0 software program (Statistics Package for Social Sciences, SPSS Inc. Chicago, Illinois, USA).

Results:

1. Characterization and differentiation of adipose and bone marrow stem cells

After ADSC and BMSC isolation and cell culture process, they rapidly proliferated and were confluent after 4-5 days of stimulation. Flow cytometry method was used to analyze the expression pattern of cell surface markers and the results showed that these cells displayed CD29 and CD105 as a positive surface markers but they did not express CD18 and CD34 as negative surface markers (Fig. 1). During differentiation, morphological changes of ADSCs and BMSCs to beta-cell-like cells were investigated by an invert microscope. Before differentiation, ADSCs and BMSCs exhibited a fibroblast and spindle-like morphology but after differentiation they formed clusters with a round shape morphology.

2. Gene expression analysis of genes involved during ADSCs and BMSCs differentiation

To determine whether the adipose and bone marrow stem cells had differentiated to beta-cell-like cells, the expression of nine genes related to the pancreatic endocrine development and function evaluated by real-time PCR analysis. Pdx-1 (pancreas duodenum homeobox-1) a pancreatic master gene; Pax-4 and Ngn-3, two genes important in the early pancreatic development; NeuroD1, a gene that regulates expression of the insulin gene and contributes to the regulation of several cell differentiation pathways like endocrine islet cells of the pancreas; Insulin-1 gene encodes insulin specially after differentiation of MSCs to beta-cell-like cells and finally, Nestin is a gene required for survival, renewal and mitogen-stimulated proliferation of neural progenitor cells but also important as a pre-marker for islet cell

differentiation. NKX6.1 gene is necessary to confer beta cell identity to differentiating endocrine precursors in the embryo and raising the possibility to maintain the differentiated state of mature beta cells. The prohormone convertase PC1/3 is endopeptidase found in islet cells that convert the prohormones to mature hormone. Islet cells express other secreted peptides such as islet amyloid polypeptide (IAPP). In adult islet, IAPP is produced by β -cells and some δ -cells and is expressed in the developing endocrine pancreas. According to Figure 2a that is related to gene expression after differentiation, except for Pdx-1 and Nestin, a significant difference (For Ngn-3; $P < 0.001$, For Insulin; $P < 0.01$ and for Pax-4, NKX6.1, PC1/3, IAPP and NeuroD1; $P < 0.05$) in gene expression between adipose and bone marrow stem cells was obviously cleared. For example Ngn-3 was 10.9 times more expressed in adipose stem cells compare to bone marrow stem cells. Figure 2b Shows that except Nestin gene that was downregulated after differentiation, all of the other genes were upregulated in both type of stem cells. According to Figure 3, gene expression of all genes mentioned previously evaluated before and after differentiation in adipose and bone marrow stem cells. Results showed that Nestin gene expression was significantly ($P < 0.001$) decreased after differentiation, on the other hand the expression of genes involved in differentiated state of adult and mature beta cells (NKX6.1, PC1/3 and IAPP) was significantly increased ($P < 0.01$)

3. Western blot analysis for ADSCs and BMSCs

Western blot analysis was performed to test insulin or nestin protein expressions in MSCs. It was expected that after differentiation process, stem cells produce insulin but in different quantity in ADSC and BMSCs. After differentiation culture, the cells mainly produced proinsulin and insulin with detectable bands corresponding to 7 and 5.5 kDa, respectively. However, MSCs cultured in media alone (before differentiation), showed no (pro)-insulin reactivity. Nestin as an important pre-marker for islet cell differentiation, and its expression was tested. Western-blotting using anti-nestin showed a single band at 220 kDa in pre-differentiated stem cells while nestin expression in differentiated islet-like cells was decreased significantly. Our results showed that insulin and nestin proteins were more expressed in adipose stem cells compared to bone-marrow stem cells (Figure 4).

4. Glucose-induced insulin and C-peptide secretion analysis

ADSCs and BMSCs after differentiation were checked because of their viability. The viability of ADSCs and BMSCs at the end of differentiation were 95 and 93%, respectively. Insulin and C-peptide secretion in response to different concentration of glucose was assessed in differentiated cells. Amount of insulin and C-peptide secretion in adipose and bone marrow stem cells, before and after exposure of these cells with different concentration of glucose (5mM, 10 mM and 20 mM) were evaluated and results had been shown in table 2,. According to the results (Table 2), insulin levels and C-peptide secretion were significantly increased in response to 10 mM glucose but increase of glucose concentration to 20 mM was not significantly elevated insulin and C-peptide secretion.

Table 2

Amount of insulin and C-peptide secretion of ADSCs and BMSCs in response to different glucose concentration

Glucose concentration (mM)	Insulin secretion (ng/10 ⁶ cells)		C-peptide secretion (µg/10 ⁶ cells)	
	ADSCs	BMSCs	ADSCs	BMSCs
0	0.64 ± 0.04	0.59 ± 0.1	0.16 ± 0.01	0.18 ± 0.02
5	12.4 ± 1.3	12.1 ± 0.9	0.71 ± 0.07	0.63 ± 0.06
10	20.5 ± 1.9	17.5 ± 0.9	1.25 ± 0.06	1.16 ± 0.1
20	20.9 ± 1.3	17.1 ± 0.4	1.26 ± 0.04	1.15 ± 0.1

Discussion:

Treatment of diabetes mellitus by pancreatic islet transplantation combined with severe shortage of donor pancreas, have forced scientists to find a new prospect to generate insulin-secreting beta cells from various progenitor populations. Because of self-renewal and multipotent characteristics, multipotent stem cells (MSCs) have attracted a great deal of attention as potential candidates for transplantation [20]. MSCs have been isolated from different adult tissues, such as bone marrow, adipose, liver, lung, etc. Adipose and bone marrow derived MSCs have a good proliferative capacity, as well as the advantage of being autologous sources. The clinical applicability of BMSCs may be limited due to the possible pain and morbidity associated with the bone marrow aspiration procedure. On the other hand, liposuction aspirates are widely available and attractive features, including availability, abundance, 500-fold higher frequency of mesenchymal stem cells (MSCs) and higher immunomodulatory capacity when compared to bone marrow-derived counterparts [21]. In this study we used nicotinamide, beta mercaptoethanol, B27 and b-FGF in DMEM media to differentiate MSCs to beta-cell-like cells. Nicotinamide could increase the mitotic indices of beta cells and also is a potent inducer of endocrine differentiation and could play an important role in terminal endocrine differentiation. It was previously reported that using low glucose medium (5 mM) with nicotinamide supplementation together, increased insulin content approximately 20-fold. It seems that β-mercaptoethanol could increase the potency of nicotinamide [22, 23]. B27 supplement is a mix of hormones and other molecules that were empirically added to media to promote growth and proliferation of cells. Proliferation of pancreatic progenitors can be induced by addition of mitogenic signals such as basic fibroblast growth factors (bFGFs). During pancreatic islet development, activating A is required for the specialization of pancreatic precursors from the gut endoderm during midgestation [24]. According to results of our study, after differentiation of MSCs to beta-cell-like cells, the cells lost *nestin* expression but on the other hand, they turned on mRNA expression of transcription factors such as *Pax-4*, *Pdx-1*, *NeuroD1* and Insulin. These genes play an important role in the development and formation of pancreatic beta-cell-like cells. In this study, we observed that except *Pdx-1*

and *nestin*, there was significant differences between expression level of other genes in adipose and bone marrow stem cells ($P < 0.05$ for *Pax-4* and *NeuroD1*; $P < 0.01$ for *Insulin* and $P < 0.001$ for *Ngn3*). Our results clarified that adipose stem cells expressed significantly more levels of *Pax-4*, *NeuroD1*, *Insulin* and *Ngn-3* compared to bone marrow stem cells but the expression of *NKX6.1*, *PC1/3* and *IAPP* genes was not significantly different. Many reports have shown that exendin-4 upregulated *Pdx-1* during β -cell regeneration. Therefore, low expression of *Pdx-1* gene in our study in comparison with other genes could be result of not using of exendin-4 in our media [25]. Nestin was regarded as an important pre-marker for islet cell differentiation and this marker transiently expressed in early stages in many tissues, including pancreas [26]. So, down regulation of nestin after ADSCs and BMSCs differentiation to beta-cell-like cells is a logical event. The results of western blot analysis for insulin and nestin expression in our experiment validated the results of qPCR. Considering nestin expression in pre-differentiated MSCs, MSCs might differentiate into pancreatic islet-like cells through intermediate neurocyte stage. *NKX6.1*, *PC1/3* and *IAPP* genes involved in developing of endocrine pancreas and their gene expression investigated to confirm the ability of functional beta islet-like cells to produce insulin. Results of our study showed a significant increase ($P < 0.01$) in expression of these related genes after differentiation to beta-like cells. Results of insulin and C-peptide secretion analysis showed that they could respond to different glucose concentrations, among which 20 mM glucose stimulated a significant increase in release of insulin and C-peptide in both ADSCs and BMSCs. However, there is no significant differences between insulin and C-peptide secretion in response to 10 mM and 20 mM glucose concentration. Several studies reported that beta cells replicates more when they induced by increasing glucose levels but at extremely hyperglycemic conditions, beta cell replication was reduced or stopped [27].

In conclusion, our study showed that both adipose and bone marrow stem cells had advantage of being autologous sources and also they could differentiate into functional beta-cell-like cells but according to the results of this study, by comparing these two kind of cells, it seems that adipose stem cells could be a good choice for treatment of diabetes mellitus according to their more safety and potency. Further researches is required in terms of in vivo evaluation of glucose levels in diabetic model as well as their immune reaction. To proof of functionality of stem cell-derived beta-cell-like cells, it's required to transplant these differentiated cells into streptozocin-mediated diabetic mice. Such analyses would confirm the nature of the differentiated insulin⁺ cells.

Declarations

Acknowledgment:

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Author Disclosure:

There is no conflict of interest.

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Figures

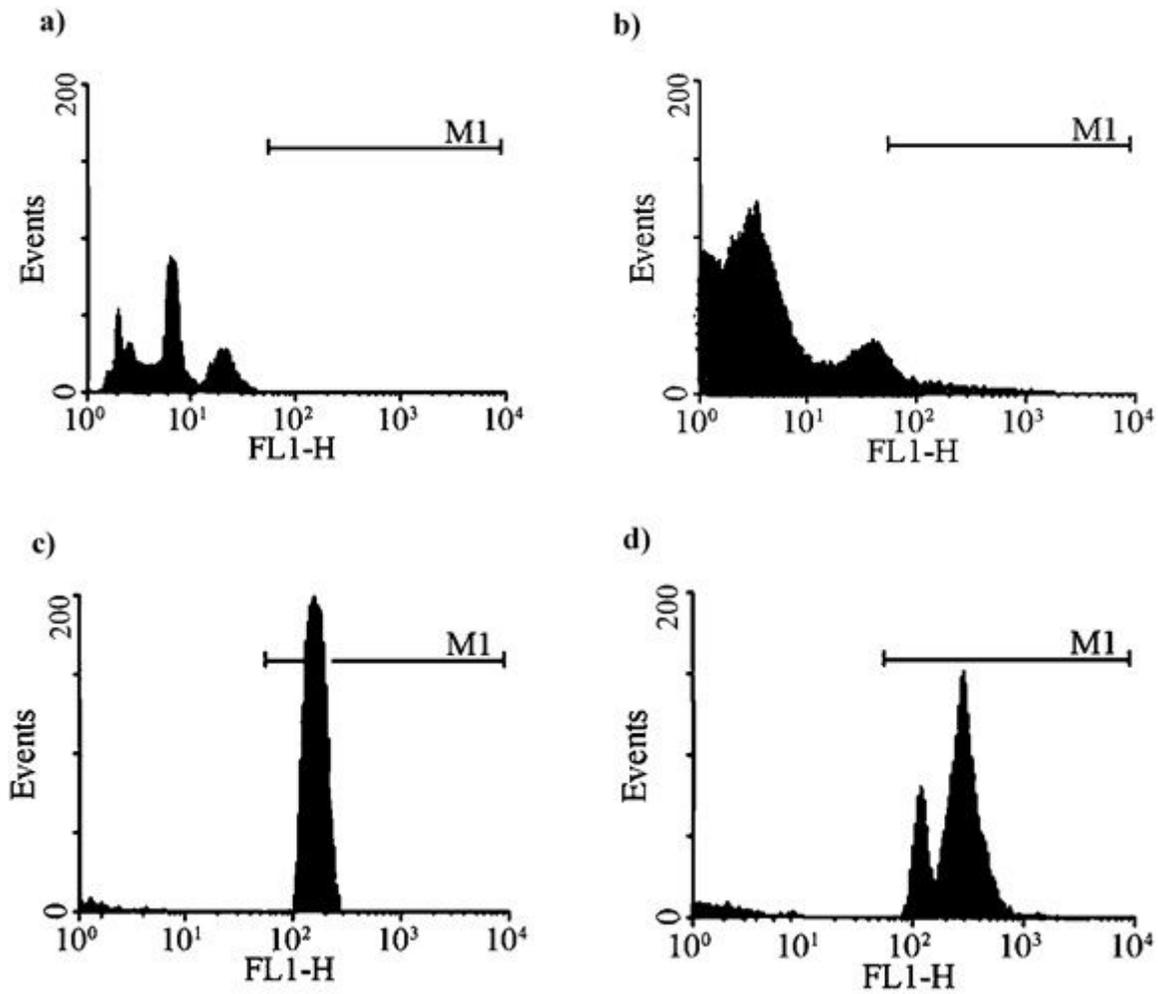


Figure 1

Characterization of isolated stem cells by detection of surface markers by flow cytometry method showed that these stem cells did not express (a) CD18 and (b) CD34 but they are positive in (c) CD29 and (d) CD105.

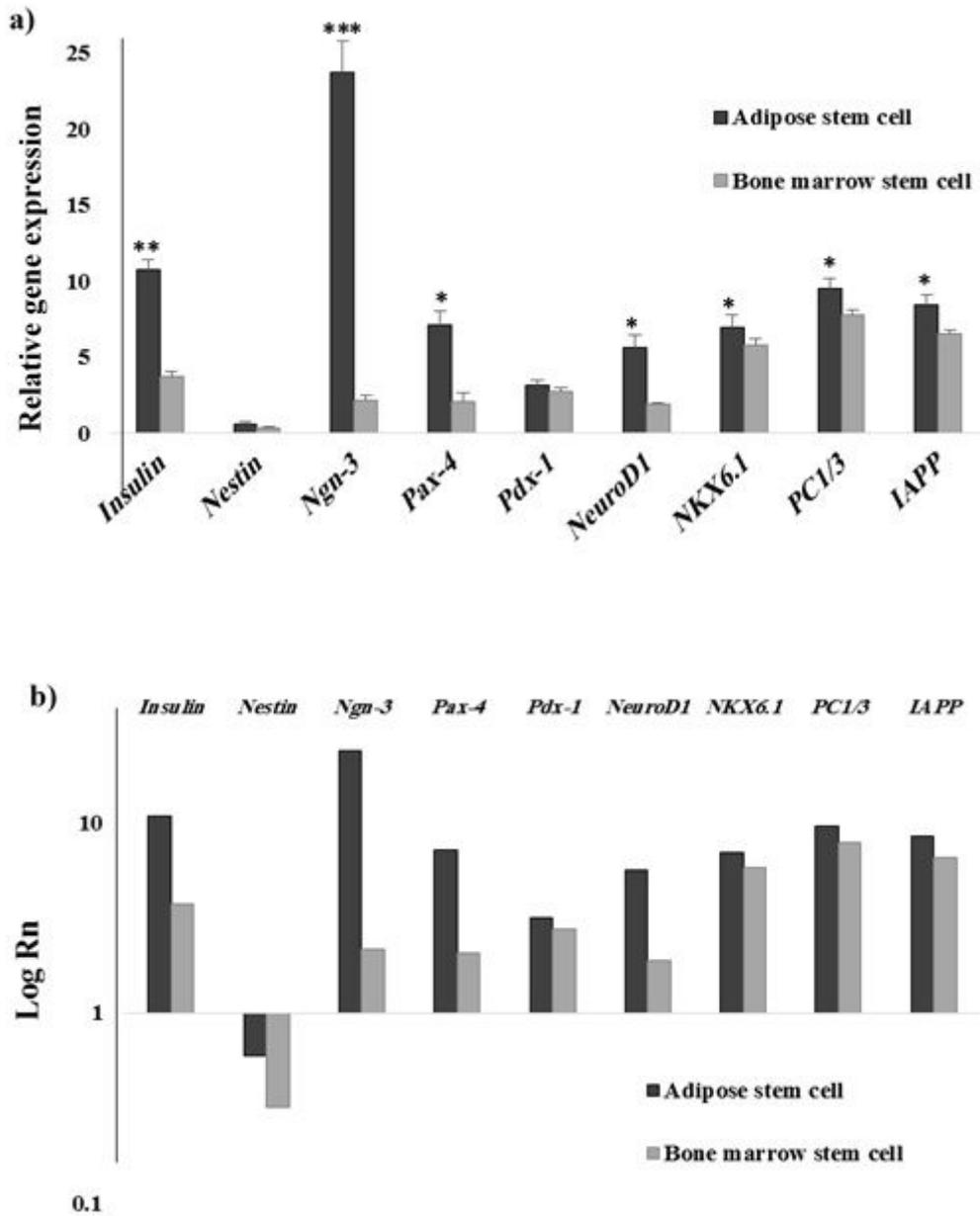


Figure 2

QPCR results of nine genes related to the pancreatic endocrine development after differentiation to beta cells compared to GAPDH gene as control in a) linear and b) logarithmic gene expression plot. * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$.

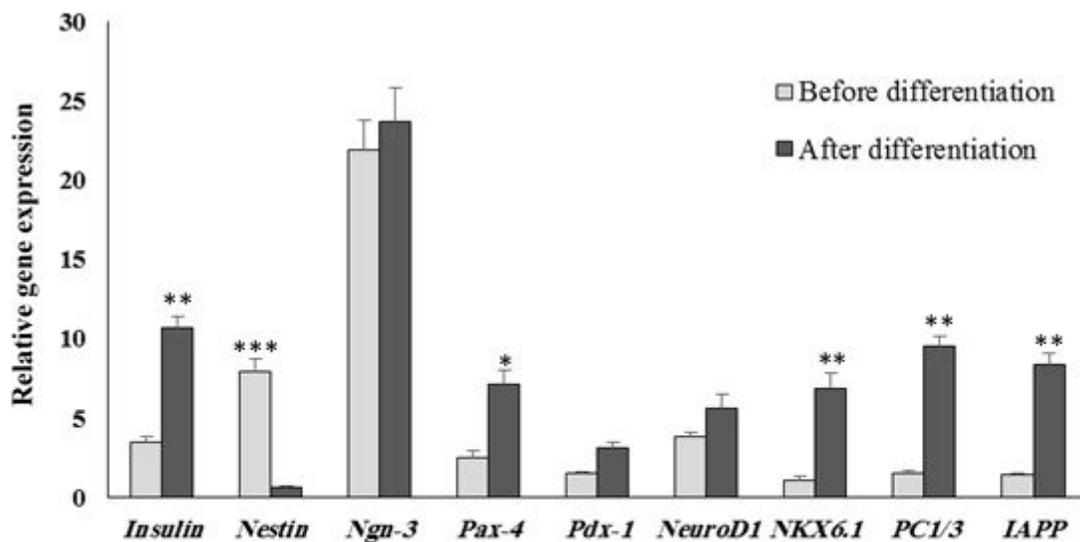


Figure 4. QPCR results of nine genes related to adipose derived stem cells before and after differentiation to beta-cell-like cells. * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$.

Figure 3

QPCR results of nine genes related to adipose derived stem cells before and after differentiation to beta-cell-like cells. * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$.

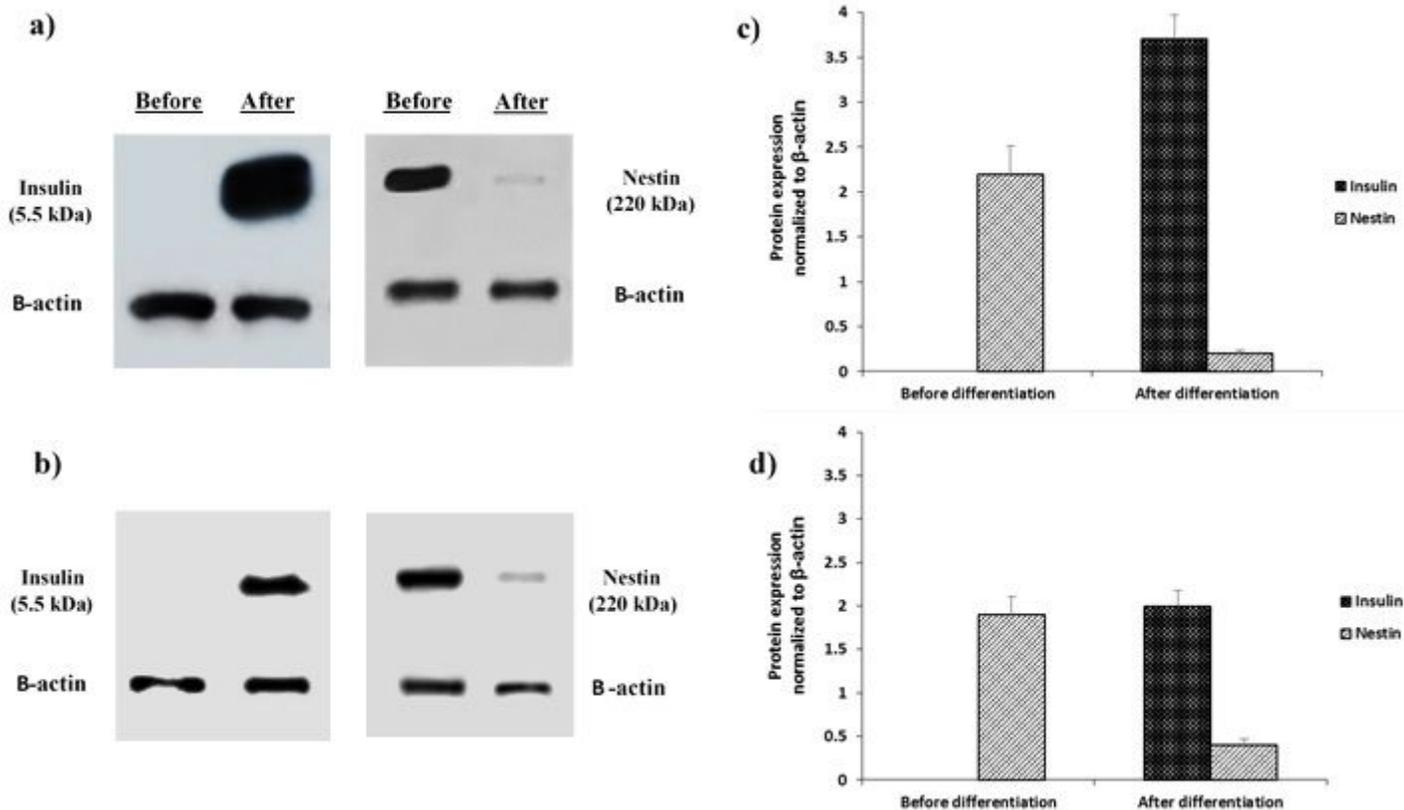


Figure 4

a) Western blot analysis of insulin and nestin proteins in adipose stem cells and b) in bone-marrow stem cells before and after differentiation to beta-cells. Insulin and nestin were quantified with densitometry of immune-reactive bands and these proteins were expressed as average density to β -actin in c) adipose stem cells and d) in bone-marrow stem cells.