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REVIEW

- 1035 Stem cells and pain

Silva MDVD, Piva M, Martelossi-Cebinelli G, Stinglin Rosa Ribas M, Hoffmann Salles Bianchini B, K Heintz O, Casagrande R, Verri Jr WA

ORIGINAL ARTICLE

Randomized Controlled Trial

- 1063 Mid-term outcomes of microfragmented adipose tissue plus arthroscopic surgery for knee osteoarthritis: A randomized, active-control, multicenter clinical trial

Wu CZ, Shi ZY, Wu Z, Lin WJ, Chen WB, Jia XW, Xiang SC, Xu HH, Ge QW, Zou KA, Wang X, Chen JL, Wang PE, Yuan WH, Jin HT, Tong PJ

Basic Study

- 1077 ADSC-Exos outperform BMSC-Exos in alleviating hydrostatic pressure-induced injury to retinal ganglion cells by upregulating nerve growth factors

Zheng ZK, Kong L, Dai M, Chen YD, Chen YH

- 1093 Human mesenchymal stem cells exhibit altered mitochondrial dynamics and poor survival in high glucose microenvironment

Abu-El-Rub E, Almahasneh F, Khasawneh RR, Alzu'bi A, Ghorab D, Almazari R, Magableh H, Sanajleh A, Shloul H, Mazari M, Bader NS, Al-Momani J

ABOUT COVER

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Basic Study

Human mesenchymal stem cells exhibit altered mitochondrial dynamics and poor survival in high glucose microenvironment

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Abstract

BACKGROUND

Mesenchymal stem cells (MSCs) are a type of stem cells that possess relevant regenerative abilities and can be used to treat many chronic diseases. Diabetes mellitus (DM) is a frequently diagnosed chronic disease characterized by hyperglycemia which initiates many multisystem complications in the long-run. DM patients can benefit from MSCs transplantation to curb down the pathological consequences associated with hyperglycemia persistence and restore the function of damaged tissues. MSCs therapeutic outcomes are found to last for short period of time and ultimately these regenerative cells are eradicated and died in DM disease model.

AIM

To investigate the impact of high glucose or hyperglycemia on the cellular and molecular characteristics of MSCs.

METHODS

Human adipose tissue-derived MSCs (hAD-MSCs) were seeded in low (5.6 mmol/L of glucose) and high glucose (25 mmol/L of glucose) for 7 d. Cytotoxicity, viability, mitochondrial dynamics, and apoptosis were deployed using specific kits. Western blotting was performed to measure the protein

expression of phosphatidylinositol 3-kinase (PI3K), TSC1, and mammalian target of rapamycin (mTOR) in these cells.

RESULTS

hAD-MSCs cultured in high glucose for 7 d demonstrated marked decrease in their viability, as shown by a significant increase in lactate dehydrogenase ($P < 0.01$) and a significant decrease in Trypan blue ($P < 0.05$) in these cells compared to low glucose control. Mitochondrial membrane potential, indicated by tetramethylrhodamine ethyl ester (TMRE) fluorescence intensity, and nicotinamide adenine dinucleotide (NAD⁺)/NADH ratio were significantly dropped ($P < 0.05$ for TMRE and $P < 0.01$ for NAD⁺/NADH) in high glucose exposed hAD-MSCs, indicating disturbed mitochondrial function. PI3K protein expression significantly decreased in high glucose culture MSCs ($P < 0.05$ compared to low glucose) and it was coupled with significant upregulation in TSC1 ($P < 0.05$) and downregulation in mTOR protein expression ($P < 0.05$). Mitochondrial complexes I, IV, and V were downregulated profoundly in high glucose ($P < 0.05$ compared to low glucose). Apoptosis was induced as a result of mitochondrial impairment and explained the poor survival of MSCs in high glucose.

CONCLUSION

High glucose impaired the mitochondrial dynamics and regulatory proteins in hAD-MSCs ensuing their poor survival and high apoptosis rate in hyperglycemic microenvironment.

Key Words: Mesenchymal stem cells; High glucose; Mitochondrial dynamics; Apoptosis; Poor survival; Phosphatidylinositol 3-kinase/mammalian target of rapamycin pathway

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Core Tip: Mesenchymal stem cells (MSCs) have significant regenerative properties that make them a potential treatment for many chronic diseases. Among these, diabetes mellitus (DM) was found to benefit from MSCs transplantation in which they restored the damaged tissues and prevented hyperglycemia-related complications. However, these therapeutic are short-lived hindering the clinical use of MSCs in the treatment of DM. This study aims to elucidate the mechanisms of hyperglycemia-induced effects on MSCs which will help in improving the therapeutic functions of these cells in this stress environment.

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INTRODUCTION

Mesenchymal stem cells (MSCs) are the most commonly studied forms of stem cells as they possess powerful regenerative and immunomodulatory abilities[1]. MSCs have unique molecular machinery that empowers them to modulate many cellular signaling pathways and restore the functionality of damaged tissues rendering them suitable to treat many acute and chronic diseases[1]. Despite their unprecedented abilities to adapt to many stress conditions, they can still behave unexpectedly under certain stress microenvironments. Many preclinical and clinical studies not only demonstrated the positive outcomes after transplanting MSCs to injured sites[2,3], but also reported the short-lived therapeutic results in hostile microenvironments which need to be figured out[4]. When MSCs are undifferentiated, they rely mainly on glycolysis to support their proliferation, expansion, and immunomodulation[5]. When MSCs are programmed to differentiate, they switch to oxidative phosphorylation (OxPhos)[6], thereby, intact mitochondria are also important to maintain their multi-lineage differentiation capacities[6].

In the last decades, diabetes mellitus (DM) incidence has increased rapidly, and it is deemed among top five chronic diseases[7]. DM is associated with serious multisystem complications that impair the life quality of many patients[8]. DM is characterized mainly by chronic hyperglycemic state that causes many devastating, long-term complications[8]. DM patients can benefit remarkably from MSCs therapy to repair and regenerate damaged tissues, restore their functionality, and reduce the severity of DM-related pathological complications[7]. Several studies reported a significant improvement in glycemic parameters and cardiovascular complications after transplanting MSCs in DM patients[7,9,10]; however, other studies reported that DM can accelerate the decline in MSCs quantity and therapeutic quality, and impair their regenerative capabilities[11]. When MSCs turn into a dysfunctioning cells, they may increase the severity of DM-related complications[11]. Although previous studies had addressed the effects of hyperglycemia on the molecular and cellular characteristics of transplanted MSCs, many aspects still need to be elucidated. Thus, it is important to investigate the effect of high glucose on MSCs biological dynamics, which are essential to understand their fate and performance in

diabetic microenvironment, and provide new avenues on possible molecular targets to modulate their survival and therapeutic outcomes in DM patients.

MATERIALS AND METHODS

Human MSCs

Human adipose tissue-derived MSCs (hAD-MSCs) were commercially purchased from Lonza (Cat# PT5006, Lot# 21TL138912) and expanded using Dulbecco's Modified Eagle's Medium Low Glucose (DMEM-Low Glucose, Euroclone) which contained 5.6 mmol/L glucose and were supplemented with 10% fetal bovine serum (FBS, Gibco)[12], 0.1 mg/mL streptomycin and 100 units/mL penicillin G in standard cell culture incubators (5% CO₂/95% air; 37 °C). Medium was changed every 72 h, and cells were sub-cultured when confluence exceeded 60%[13]. For high glucose conditions, cells were cultured in Dulbecco's Modified Eagle's Medium High Glucose (DMEM-High Glucose, Euroclone) which contained 25 mmol/L glucose[14] and were supplemented with 10% FBS, 0.1 mg/mL streptomycin and 100 units/mL penicillin G for 3, 7, and 14 d in standard cell culture incubators (5% CO₂/95% air; 37 °C). High glucose complete medium was changed every 72 h. MSCs that were cultured in low glucose were considered as our study control. MSCs of passage 5 and 6 were used to perform the experiments. AD-MSCs characterization markers can be found in (Table 1).

Cytotoxicity assays

To measure the level of cytotoxicity in human MSCs after being cultured in low and high glucose, we seeded 5×10^4 cells/well in 96 well plate and we measured the lactate dehydrogenase (LDH) which was released from the damaged MSCs (LDH Assay Kit, Abcam, Cat# ab102526) and the absorbance values were obtained using Cytation 5 (BioTek, United States). The viability of MSCs in low and high glucose were also assessed using 0.4% Trypan blue and the percentage of viable cells as well as representative fluorescent images were obtained using Corning® CytoSmart Cell Counter.

Tetramethylrhodamine ethyl ester (TMRE) measurement

Mitochondrial membrane potential was assessed using tetramethylrhodamine ethyl ester (TMRE)-Mitochondrial Membrane Potential Assay Kit (Abcam, Cat # ab113852). Briefly, human AD-MSCs were placed in 24-well plate at 1×10^4 cells per well and allowed to adhere overnight. After that, cells were cultured either in low glucose or high glucose for 7 d. Then, media were aspirated, and cells were stained using 400 nM TMRE in culture media for 30 min in the incubator, and then media were replaced with 200 μ L phosphate buffered saline (PBS) per well. Fluorescence intensity was detected using Cytation 5 (BioTek, United States). ($\lambda_{ex} = 549$ nm, $\lambda_{em} = 575$ nm), and TMRE fluorescent images were captured at Texas red filter using Cytation 5 (BioTek, United States).

Nicotinamide adenine dinucleotide /NADH assay

Nicotinamide adenine dinucleotide (NAD⁺)/NADH ratio in AD-MSCs that were cultured in low and high glucose media was assessed using NAD/NADH-Glo™ assay kit (Promega, Cat# G9071). Briefly, cells were cultured in low and high glucose media for 7 d. After that, cells were detached and placed in 96 well plate at 5×10^5 in 50 μ L of either low or high glucose media and were allowed to adhere overnight. 50 μ L of NAD/NADH-Glo™ Detection Reagent was added to each well and the plate was gently shaken to mix and lyse the cells and was incubated with the added reagent for 60 min at room temperature. The luminescence values were recorded using Cytation 5 (BioTek, United States) which were corresponding to the total amount of NAD⁺ produced from NADH.

Western blotting

The protein levels for phosphatidylinositol 3-kinase (PI3K) (Santa Cruz Biotechnology Cat # sc-1637), TSC1 (Santa Cruz Biotechnology Cat # sc-377386), and mammalian target of rapamycin (mTOR) (Santa Cruz Biotechnology Cat # sc-293133), NDUFB8 (Abcam, Cat # ab192878), SDHB (Santa Cruz Biotechnology Cat # sc-271548), UQCRC2 (Santa Cruz Biotechnology Cat # sc-390378), COX5a (Santa Cruz Biotechnology Cat # sc-376907), ATP5A (Santa Cruz Biotechnology Cat # sc-136178), and β -actin (Santa Cruz Biotechnology Cat # sc-47778 HRP) were measured by western blot. Briefly, total protein levels were measured using NanoDrop™ Lite Spectrophotometer, and 40 μ g of protein was loaded onto sodium-dodecyl sulfate gel electrophoresis. Following electrophoresis, proteins were transferred to polyvinylidene fluoride membrane and were incubated with appropriate primary and secondary antibodies. The membranes were visualized using VILBER FUSION Gel Documentation System, and bands were quantified using ImageJ for densitometry.

Apoptosis assay

To detect apoptosis in hAD-MSCs after being cultured in low and high glucose, we used RealTime-Glo™ Annexin V Apoptosis live assay (Promega, Cat# JA1011, Lot# 0000400486 following the manufacturer's guidelines. Briefly, cells were seeded 1×10^4 cells/well in 24 well plate and then cultured in low and high glucose for 7 d. The media were aspirated and each well was washed with PBS followed by the addition of 100 μ L of fresh medium having Annexin V-LgBiT, Annexin V-SmBiT, CaCl₂, and Annexin V NanoBiT Substrate to each well. After 1 h incubation, the fluorescent images of green color which represented cells undergoing apoptosis were detected at GFP filter using Cytation 5 (BioTek, United States).

Table 1 Cell surface characterization of human adipose tissue-derived mesenchymal stem cells based on the company analysis report

Positive markers	Negative markers
CD13	CD14
CD29	CD31
CD44	CD45
CD73	HLA-DR
CD90	
CD105	
CD166	

Statistical analysis

Data were reported as mean \pm SD. Comparison of data between multiple groups was performed using one-way analysis of variance (ANOVA) followed by Tukey's post-hoc multiple comparison test, and analysis between two groups was made using Student's *t*-test (two-tailed). Statistical significance is determined as $P < 0.05$. Each figure represents one of at least three independent quantifiable experiments.

RESULTS

Culturing MSCs in high glucose triggers massive cytotoxicity and poor survival

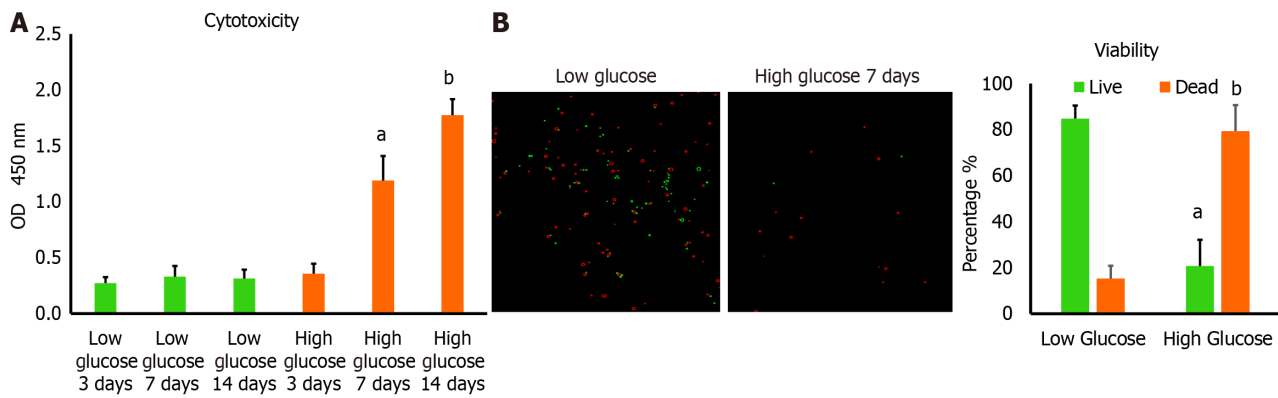
MSCs are known to survive well in low glucose medium which is a physiological requirement to preserve their stemness features. We want to determine the viability of these cells in high glucose microenvironment. hAD-MSCs were cultured in high glucose medium containing 25 mmol/L of glucose for 3, 7, and 14 d. The cytotoxicity was measured by determining the amount of LDH released and was found to be significantly greater ($P < 0.01$) in hAD-MSCs cultured in high glucose for 7 and 14 d compared with hAD-MSCs cultured in normal low glucose medium containing 5.6 mmol/L of glucose (Figure 1A). To confirm these findings, the viability of hAD-MSCs was measured using 0.4% Trypan blue and found to be remarkably low ($P < 0.05$) in hAD-MSCs that were cultured in high glucose for 7 d compared to hAD-MSCs cultured in a low glucose medium (Figure 1B).

The mitochondrial dynamics of MSCs impaired by high glucose

Mitochondria are integral intracellular organelles promoting cells survival and viability. The variance of mitochondrial dynamics depends on the type of microenvironment where cells exist. Previous reports demonstrated that high glucose can induce the fission and fragmentation of mitochondria in glomeruli podocytes[15], but the knowledge surrounding the impact of high glucose on the mitochondrial dynamics of MSCs is still limited. The mitochondrial membrane potential ($\Delta\Psi_m$) generated by proton pumps (complexes I, III and IV) is imperative to ensure proper ATP production by mitochondrial OxPhos. Abnormalities in the mitochondrial membrane potential ($\Delta\Psi_m$) can significantly elicit a decline in cells viability and trigger unwanted signaling pathways. We measured the mitochondrial membrane potential ($\Delta\Psi_m$) in hAD-MSCs after being grown in a high glucose medium for 7 d using TMRE which is sequestered by active mitochondria. Our results illustrated that the fluorescent intensity of TMRE was significantly low ($P < 0.05$) in hAD-MSCs cultured in a high glucose medium compared to hAD-MSCs cultured in a normal low glucose medium indicating an impairment in the mitochondrial membrane potential in these cells (Figures 2A and B). NAD⁺ and its reduced form (NADH) regulate many metabolic pathways, including mitochondrial OxPhos. Maintaining NAD⁺/NADH pool is required to ensure that the influx of electrons is coupled with the translocation of protons to the intermembrane space to generate the required membrane potential ($\Delta\Psi_m$) across the inner mitochondrial membrane to be harnessed by complex V to produce ATP[16]. To investigate if the decrease in the mitochondrial membrane potential ($\Delta\Psi_m$) is correlated with abnormalities in NAD⁺/NADH pool, we measure the NAD⁺/NADH ratio in hAD-MSCs cultured in low and high glucose media (Figure 3). The findings revealed a considerable drop ($P < 0.01$) in NAD⁺/NADH ratio when hAD-MSCs were cultured in high glucose. This may explain the disturbance in the inner mitochondrial potential detected by TMRE assay (Figures 2A and B).

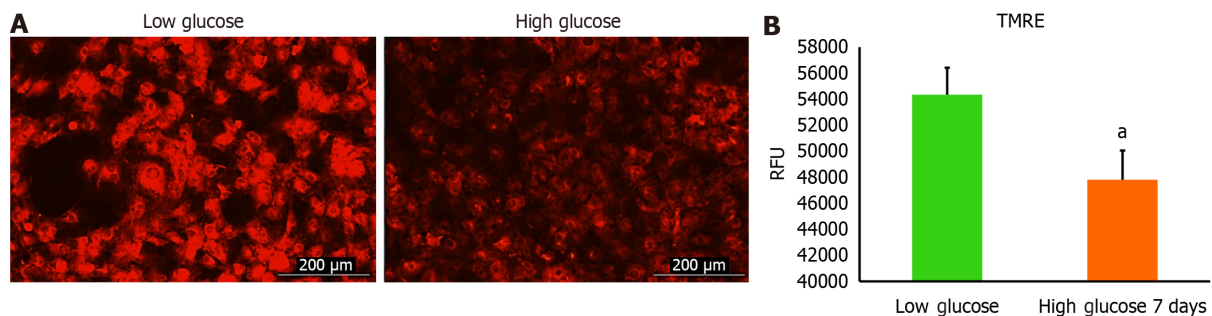
Disturbance of MSCs mitochondria dynamics parameters in high glucose engenders their apoptosis

Apoptosis is a programmed cell death that can be physiological or pathological depending on the triggering factors that enkindle it. Mitochondrial dysfunction is a notable inciting factor which is able to fire up debilitating cellular apoptosis and impact the survival of numerous cells. The noticeable deterioration in the mitochondrial parameters detected in hAD-MSCs after being cultured in high glucose prompted us to examine the level of apoptosis in these cells. To achieve that, we used RealTime-Glo™ Annexin V Apoptosis kit to unveil the presence of apoptosis or not. The luminescence signal which is proportionally linked to apoptosis intensity was considerably higher in hAD-MSCs that were cultured in high glucose (Figure 4). Fluorescent microscopic images showed that apoptosis (marked by green stained cells) was more in



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Figure 1 Culturing human adipose tissue-derived mesenchymal stem cells in high glucose reduced their viability. A: Lactate dehydrogenase cytotoxicity assay revealed higher level of cytotoxicity in human adipose tissue-derived mesenchymal stem cells (hAD-MSCs) cultured in high glucose for 7 and 14 d compared to cells cultured in low glucose. $n = 5$, ^a $P < 0.01$ compared to low glucose 7 d, ^b $P < 0.01$ compared to low glucose 14 d; B: The percentage of cells viability detected by 0.4% Trypan blue showed significant reduction in the number of viable hAD-MSCs after being cultured in high glucose for 7 d. Fluorescent images of live and dead cells were obtained using Corning® CytoSmart Cell Counter. $n = 5$, ^a $P < 0.01$ compared to live cells in low glucose 7 d, ^b $P < 0.01$ compared to dead cells in low glucose 7 d.



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Figure 2 High glucose decreased the mitochondrial membrane potential level of human adipose tissue-derived mesenchymal stem cells. The tetramethylrhodamine ethyl ester (TMRE) fluorescence intensity values of human adipose tissue-derived mesenchymal stem cells (hAD-MSCs) cultured in high glucose for 7 d were substantially declined compared to fluorescent intensity values of cells cultured in low glucose. Live fluorescent images showed a weak signal of TMRE stain in hAD-MSCs after being placed in high glucose for 7 d compared to strong TMRE signal in low glucose cultured cells. $n = 4$, ^a $P < 0.05$ compared to low glucose control. Fluorescent images were captured at Texas Red filter using Cytation 5 (BioTek, United States). TMRE: Tetramethylrhodamine ethyl ester; RFU: Relative fluorescence units.

high glucose cultured hAD-MSCs (Figure 4).

High glucose affects the level of PI3K and mTOR in MSCs

PI3K orchestrates many vital signaling pathways and targets downstream factors that are required to maintain the vivacity of cells. PI3K controls integral proteins that are necessary to preserve mitochondrial dynamics[17,18]. PI3K is a known activator of mTOR which is a central regulator of cell survival and metabolism. It has been reported that mTOR is important to preserve mitochondrial dynamics and generate the required mitochondrial potential to produce ATP. PI3K is required to remove the inhibitory effect of tuberous sclerosis tumor suppressor (TSC1) which binds mTOR and inactivates it[19]. We performed western blotting analysis to measure the expression levels of PI3K, TSC1, and mTOR in high glucose treated hAD-MSCs. Western blotting results demonstrated the loss of PI3K and mTOR in high glucose treated hAD-MSCs ($P < 0.05$ compared to low glucose), while TSC1 is significantly increased in these cells ($P < 0.05$ compared to low glucose) (Figure 5). Those findings indicated impaired PI3K/mTOR axis which may explain the disruption in mitochondrial parameters in high glucose cultured MSCs. Taking together, our findings highlight a new mechanism that can be released by high glucose causing the poor survival of MSCs in diabetic microenvironment.

High glucose disturbs major mitochondrial complexes in MSCs

The reduction in mTOR which is essential for inducing the biogenesis of mitochondrial complexes, most importantly complex I and V[20] prompted us to investigate the expression of mitochondrial complexes in low and high glucose-cultured hAD-MSCs. The results demonstrated that hAD-MSCs that were cultured in high glucose exhibited significant downregulation ($P < 0.05$) in the level of complex I, IV, and V comparing to hAD-MSCs that were cultured in low glucose (Figure 6), while the expression of complex II and III showed no significant changes ($P = 0.85$) in both culturing

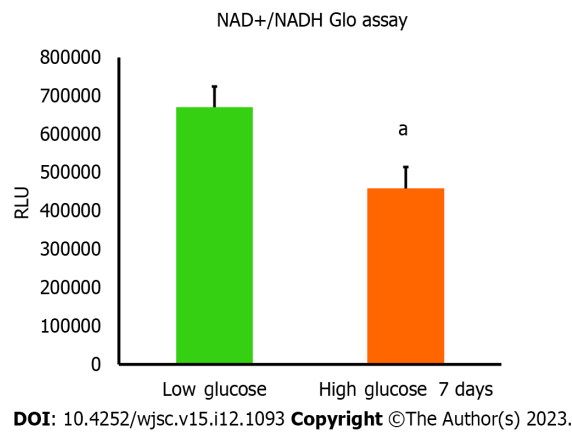


Figure 3 Nicotinamide adenine dinucleotide and NADH amounts were declined in high glucose cultured human adipose tissue-derived mesenchymal stem cells. The amount of nicotinamide adenine dinucleotide (NAD⁺) produced from NADH was measured using NAD⁺/NADH Glo assay. Luminescence intensity was significantly less in human adipose tissue-derived mesenchymal stem cells cultured in high glucose for 7 d compared to cells cultured in low glucose. $n = 4$, $^aP < 0.01$ compared to low glucose control. Luminescence values were measured using Cytation 5 (BioTek, United States). RFU: Relative fluorescence units.

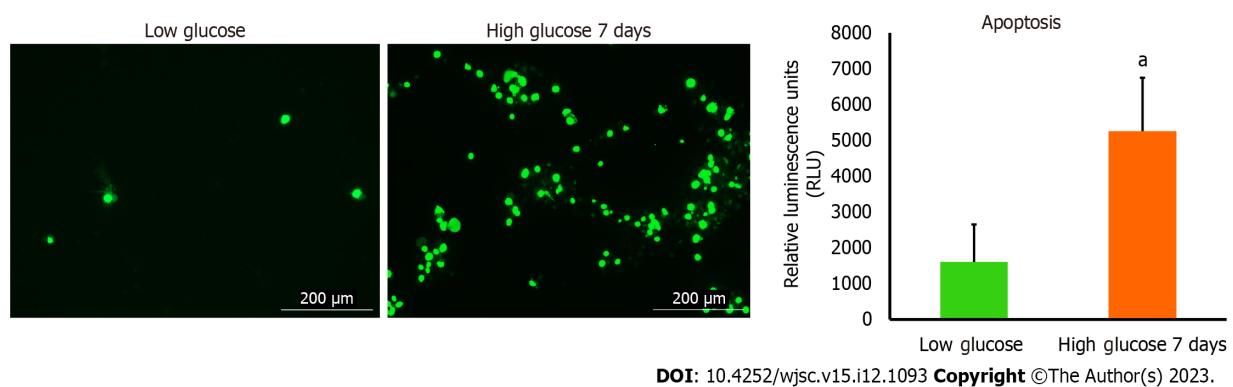
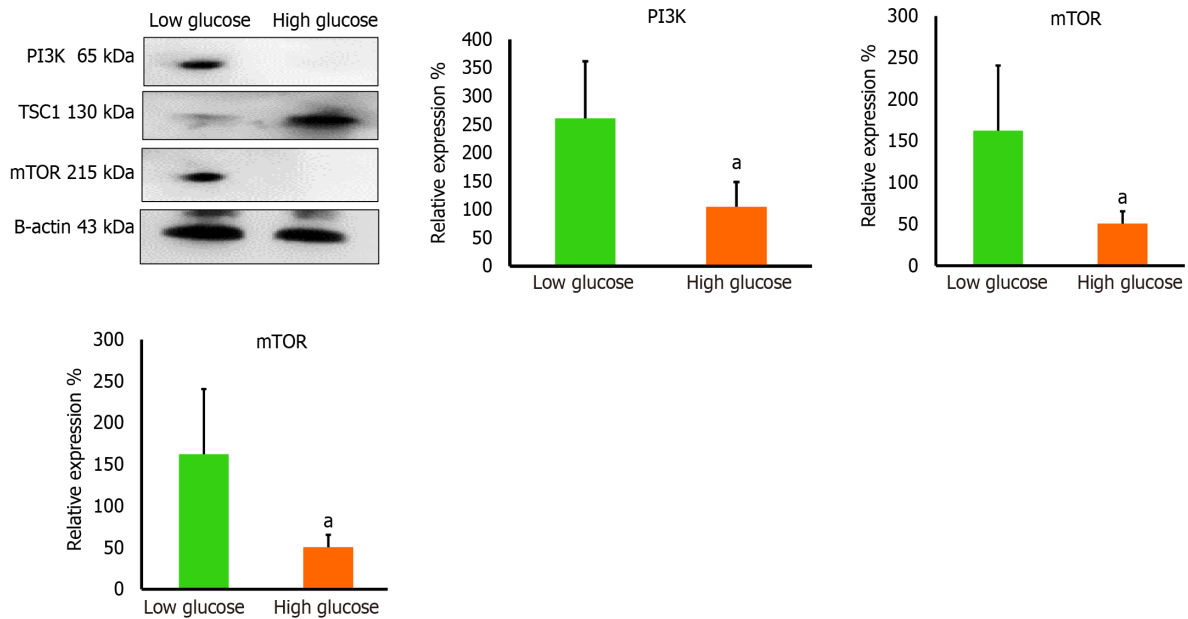


Figure 4 The level of apoptosis was higher in human adipose tissue-derived mesenchymal stem cells cultured in high glucose. RealTime-Glo™ Annexin V Apoptosis live assay was used. The level of apoptosis was remarkably increased in human adipose tissue-derived mesenchymal stem cells cultured in high glucose for 7 d. Green fluorescence intensity is corresponding to the number of cells undergoing apoptosis. Fluorescent images were captured at GFP filter using Cytation 5 (BioTek, United States). $n = 3$, $^aP < 0.05$ compared to low glucose control.

conditions (Figure 6). The disturbance in the level of these major complexes can impact the mitochondrial OxPhos and trigger the mitophagy of mitochondria[21].

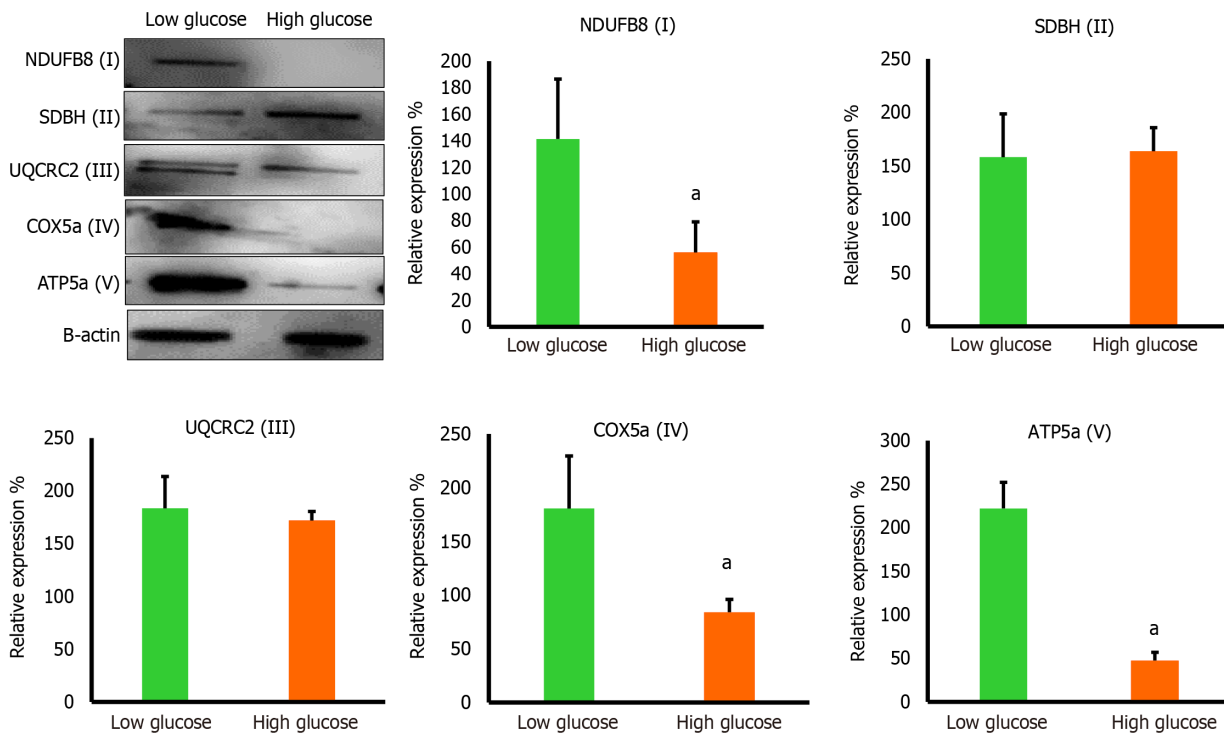
DISCUSSION

MSCs are the most promising type of stem cells for regenerating damaged tissues and restoring their normal functions which can improve the quality of life of many patients who suffer from debilitating disorders[1]. However, the remarkable improvements in pathological markers reported after transplanting MSCs[22,23] were hindered by the short-lived outcomes and poor survival rate, which need extensive investigation[24,25]. Many pathologies are characterized by stress microenvironments, including hypoxia, inflammation, and nutritional imbalances, which can impact the expected outcomes of transplanting MSCs[26-29], by either inhibiting or activating cellular signaling pathways. DM is a frequent chronic disease that leads to significant multi-system complications[8]. The major hallmark of DM is glucotoxicity, which is featured by the presence of high glucose levels and impaired insulin secretion and function[8]. This glucotoxicity produces massive molecular changes in the residing cells and cause the appearance of DM-related complications, such as cardiovascular and neurological complications[8]. Both animal and clinical studies have demonstrated the great therapeutic efficacy of MSCs transplantation in alleviating chronic hyperglycemia by reversing insulin resistance, improving insulin sensitivity, controlling inflammation, relieving metabolic syndrome symptoms, and ameliorating β -cell destructive abilities[10,30]. Nevertheless, studies also revealed that MSCs exhibited short-lived outcomes due to their poor survival after being transplanted in DM disease model[4]. Several studies have asserted the importance of investigating the mechanisms leading to the poor survival of MSCs in hyperglycemia which is the major insult to DM patients [24,31]. Up to date little is known about the molecular and biological changes in MSCs under hyperglycemia. MSCs are



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Figure 5 The protein levels of mitochondrial regulators were reduced in high glucose cultured human adipose tissue-derived mesenchymal stem cells. Western blot protein analysis showed significant reduction in the protein levels of phosphatidylinositol 3-kinase and mammalian target of rapamycin (mTOR) in high glucose cultured cells, while the protein level of mTOR inhibitory protein; TSC1, was elevated in high glucose cultured cells. These proteins are important to maintain proper mitochondrial dynamics. $n = 3$, ^a $P < 0.05$ compared to low glucose control. PI3K: Phosphatidylinositol 3-kinase; mTOR: Mammalian target of rapamycin.



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Figure 6 The protein levels of mitochondrial complexes were reduced in high glucose cultured human adipose tissue-derived mesenchymal stem cells. Western blot protein analysis showed significant reduction in the protein levels of NDUFB8 (complex I), COX5a (complex IV), and ATP5a (complex V) in high glucose cultured human adipose tissue-derived mesenchymal stem cells comparing to low glucose cultured cells, while the protein levels of SDBH (complex II) and UQCRC2 (complex III) were not significantly changed in both culturing conditions. $n = 3$, ^a $P < 0.05$ compared to low glucose control.

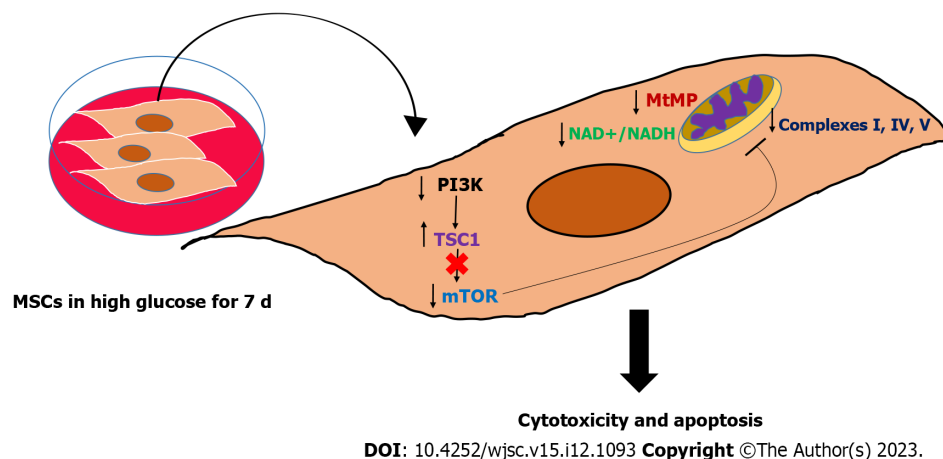


Figure 7 Graphical summary of the study findings. High glucose impaired the mitochondrial function in mesenchymal stem cells (MSCs) by perturbing many mitochondrial regulatory dynamics and factors, particularly mitochondrial membrane potential, nicotinamide adenine dinucleotide (NAD⁺)/NADH pool, and mammalian target of rapamycin protein. This mitochondrial dysfunction is associated with triggering apoptosis in MSCs, which mechanistically explained the poor survival of MSCs in hyperglycemia. MSCs: Mesenchymal stem cells; PI3K: Phosphatidylinositol 3-kinase; mTOR: Mammalian target of rapamycin; NAD⁺: Nicotinamide adenine dinucleotide.

capable of switching their metabolism on to support their regenerative and immunomodulation abilities[32]. MSCs rely primarily on glycolysis when they are undifferentiated as it is important to preserve their stemness and proliferation[33]. Several reports illustrated that glycolysis is required to preserve the immunosuppressive characteristics of MSCs[34]. Upregulation in glycolysis has been found to increase the immunomodulatory abilities of MSCs and promote their capabilities to reprogram immune cells[35]. MSCs use mitochondrial OxPhos when they are induced to differentiate, thus the multi-lineage differentiation capacities of MSCs require intact mitochondria[6]. Maintaining intact mitochondrial dynamics is not only required to support the differentiation capacities of MSCs, but it is also crucial for redox regulation and prevention of excessive production of reactive oxygen species[36]. Recently, it has been proposed by many studies that MSCs are capable of transferring mitochondrial DNA to recipient cells and restoring the normal mitochondrial functions in these cells[37,38]. MSCs-mediated mitochondrial transfer further supports the necessity to conserve normal mitochondrial parameters[38]. Previous reports revealed that the MSCs proliferation decreased in high glucose medium [13,14]. Furthermore, other studies showed that MSCs differentiation capacities might vary when they are cultured in high glucose conditions[39,40]. Our study illustrated that high glucose can reduce the survival rate of MSCs. The reduction in MSCs viability reported in our study is correlated with an impairment in the inner mitochondrial membrane potential ($\Delta\Psi_m$) which is an important parameter and the energy generated from this potential gradient will be utilized by complex V to produce ATP molecules. The maintenance of mitochondrial membrane potential depends on the presence of a plentiful amount of NAD⁺ molecules[41]. NAD⁺ serves as an oxidoreductase cofactor that controls many vital energy production pathways. Maintaining normal NAD⁺/NADH ratio is integral to providing the required electron flux that will be used to generate proper mitochondrial membrane potential ($\Delta\Psi_m$)[41]. High glucose insult causing a substantial drop in NAD⁺/NADH ratio in hAD-MSCs indicates a needed disturbance in the oxidation power to produce NAD⁺ and donating electrons that will keep the functionality of mitochondrial complexes. Another important complex that is required to preserve the mitochondrial dynamics is mTOR complex[42]. mTOR is a known regulator of many cellular processes, including the biogenesis of mitochondrial complexes, most importantly complex I and V and physiological mitophagy[42]. The downregulation in the level of mTOR in MSCs cultured in high glucose was associated with significant disturbances in mitochondrial complexes I, IV, and V which can induce mitochondrial dysfunction, mitophagy and massive oxidative stress. The effect of high glucose on modulating mitochondrial functions by targeting mTOR is barely investigated and with controversial results[43]. While other studies found that hyperglycemia activate mTOR pathway in cardiomyocytes[44,45], other evidence showed that insulin can activate mTOR pathway in DM patients and protect cells from developing mitochondrial dysfunction[46], indicating that high glucose can deactivate mTOR and cause pathological consequences[46]. These discrepancies regarding the effect of high glucose on mTOR level and activity seem to differ depending on the type of cells and the upstream molecular regulators. Based on that, the role of high glucose in modulating the level of mTOR needs further investigation in different cell types. In our study, the upstream regulatory protein PI3K decreased remarkably in hAD-MSCs exposed to high glucose. A plentiful number of reports corroborated that PI3K-AKT pathway is required to activate mTOR by removing the inhibitory effect produced by TSC1 or Hamartin protein. We reported in this study that the reduction in PI3K protein level was coupled with marked elevation in TSC1 and downregulation in mTOR in hAD-MSCs cultured in high glucose. Our findings provide a mechanistic explanation of poor survival of MSCs in hyperglycemic microenvironment and suggests new targets that can be modified to enhance the survival rate of MSCs in DM patients. Moreover, our results highlighted the importance of maintaining the desirable pre-transplantation culturing conditions that preserves the salutary properties of MSCs by controlling the level of glucose in the culture media. Schematic summary of the study findings was illustrated in (Figure 7) which highlighted that high glucose microenvironment reduced the level of PI3K and removed its inhibitory effect on TSC1 and caused its upregulation. The increase in the level of TSC1 caused a reduction in mTOR which

disturbed many mitochondrial dynamics, mainly MtMB, NAD⁺/NADH ratio, and mitochondrial complexes that triggered the apoptosis of hAD-MSCs.

CONCLUSION

MSCs have the potential to regenerate damaged tissues and restore their functions. DM patients can hugely benefit from MSCs transplantation to abrogate the pathological consequences and improve their quality of life. The poor survival and short-lived positive outcomes following MSCs transplantation in DM patients are considered stumbling blocks. High glucose impaired mitochondrial function in MSCs by perturbing mitochondrial regulatory factors, particularly mitochondrial membrane potential, NAD⁺/NADH pool, and mTOR protein. This mitochondrial dysfunction is associated with triggering apoptosis in MSCs. Preserving these factors may help in improving the survival rate of MSCs in diabetic microenvironment and initiate long-lasting therapeutic outcomes. Future studies should focus on providing new strategies to overcome the poor survival of MSCs in high glucose using genetic modification or biomaterials and nanoparticles. Moreover, the impact of other stressors existed in DM patients other than hyperglycemia should be studied.

ARTICLE HIGHLIGHTS

Research background

Mesenchymal stem cells (MSCs) have the ability to cure many chronic diseases, including diabetes mellitus (DM) and its related multisystem complications.

Research motivation

The therapeutic outcomes of MSCs transplantation in DM are short-lived which require thorough investigation.

Research objectives

This study aimed to determine the effects of hyperglycemia microenvironment on various mitochondrial-related parameters in MSCs to better understand their fate in DM patients.

Research methods

Adipose tissue-derived MSCs were exposed to low and high glucose media and mitochondrial dynamics and regulators were measured and analyzed.

Research results

High glucose induces the apoptosis of adipose tissue-derived MSCs by disturbing many mitochondrial parameters, including mitochondrial membrane potential, nicotinamide adenine dinucleotide (NAD⁺)/NADH pool, mammalian target of rapamycin, and mitochondrial complexes I, IV, and V.

Research conclusions

Hyperglycemia decreases the survival of MSCs by triggering mitochondrial dysfunction in these cells causing their short-lived therapeutic outcomes.

Research perspectives

New strategies to improve the survival rate of MSCs in hyperglycemia should be the focus of future studies which can help in increasing the chances of using these cells clinically for the treatment of DM.

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FOOTNOTES

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approve the final manuscript.

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