



Melatonin Protects Chronic Kidney Disease Mesenchymal Stem/Stromal Cells against Accumulation of Methylglyoxal via Modulation of Hexokinase-2 Expression

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Abstract

Treatment options for patients with chronic kidney disease (CKD) are currently limited; therefore, there has been significant interest in applying mesenchymal stem/stromal cell (MSC)-based therapy to treat CKD. However, MSCs harvested from CKD patients tend to show diminished viability and proliferation due to sustained exposure to uremic toxins in the CKD environment, which limits their utility for cell therapy. The application of melatonin has been demonstrated to improve the therapeutic efficacy of MSCs derived from and engrafted to tissues in patients suffering from CKD, although the underlying biological mechanism has not been elucidated. In this study, we observed overexpression of hexokinase-2 (HK2) in serum samples of CKD patients and MSCs harvested from an adenine-fed CKD mouse model (CKD-mMSCs). HK2 upregulation led to increased production levels of methylglyoxal (MG), a toxic metabolic intermediate of abnormal glycolytic processes. The overabundance of HK2 and MG was associated with impaired mitochondrial function and low cell proliferation in CKD-mMSCs. Melatonin treatment inhibited the increases in HK2 and MG levels, and further improved mitochondrial function, glycolytic metabolism, and cell proliferation. Our findings suggest that identifying and characterizing metabolic regulators such as HK2 in CKD may improve the efficacy of MSCs for treating CKD and other kidney disorders.

Key Words: Melatonin, Hexokinase, Methylglyoxal, Mesenchymal stem/stromal cells, Mitochondria, Glycolysis

INTRODUCTION

Chronic kidney disease (CKD) remains a highly prevalent condition that affects all ages and has a high mortality rate (Lv and Zhang, 2019), constituting a significant portion of the global disease burden (Romagnani *et al.*, 2017). Despite the great efforts made by both clinical and basic science researchers to develop an effective therapy, the burden of CKD has not declined over the past few decades (GBD Chronic Kidney Disease Collaboration, 2020). Progression of kidney injury in CKD leads to ultimate loss of kidney function, which causes the accumulation of uremic toxins in the kidney tissues (Nigam and Bush, 2019), significantly increasing the mortality and risk of morbidity (Lee *et al.*, 2018). Mitochondrial damage and dysfunction play a critical role in the pathophysiology of kidney

injury. Despite ongoing efforts to elucidate the specific regulatory mechanisms of mitochondrial dysregulation in CKD, currently available knowledge is insufficient to design clinically successful interventions without adverse side effects (Breyer and Susztak, 2016).

Mesenchymal stem/stromal cells (MSCs) are attractive candidates for developing cell-based therapies for CKD, as they can contribute to tissue repair or regeneration by releasing appropriate factors related to cell proliferation, survival, and differentiation (Yun and Lee, 2019). Several reports have suggested that the angiogenic and regenerative properties of MSCs can improve various pathological conditions *in vivo* (Hu and Li, 2018; Keshtkar *et al.*, 2018; Thalakiriyawa *et al.*, 2021), supporting the idea that MSC-based therapy may be highly efficacious for treating CKD and other renal diseases. However,

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CKD patient-derived human MSCs (CKD-hMSCs) show decreased potency for autologous cell-based therapy because of their exposure to uremic toxins circulating in the body during their growth and after their administration to CKD patients (Hickson *et al.*, 2016). Therefore, more work is needed to find an improved method for increasing and protecting MSCs in the context of CKD to realize their therapeutic potential.

Melatonin is a peptide hormone released by the pineal gland to modulate several physiological functions, including sleep, but there has been growing interest in its broader therapeutic potential. Accumulating evidence suggests that melatonin not only ameliorates myocardial infarction (Fu *et al.*, 2020), pulmonary ischemia reperfusion injury (Bhatia *et al.*, 2019), and kidney injury (Ohashi *et al.*, 2019) when administered directly but also improves the viability, motility, and proliferation of MSCs (Hu and Li, 2019), thereby enhancing the therapeutic potential of MSC-based therapy (Yoon *et al.*, 2020c). Our previous work demonstrated that melatonin treatment can improve the stress tolerance of mouse MSCs (mMSCs) by improving mitochondrial function (Han *et al.*, 2019). In a different study, we found that prion protein-rich exosomes from melatonin-treated normal hMSCs can modulate microRNA-4516 in CKD-hMSCs to improve their therapeutic potential (Yoon *et al.*, 2020c). Although melatonin-induced improvements in mitochondrial ATP synthesis have been confirmed *in vitro* and *in vivo*, the mechanism by which melatonin engages the broader metabolic network in the context of its protective effect against CKD has not been well investigated.

Hexokinase-2 (HK2) catalyzes the ATP-mediated phosphorylation of glucose to glucose-6-phosphate (G6P), which is the rate-limiting step of glucose metabolism. Overexpression of HK2 has been linked to diabetic nephropathy (Rabbani and Thornalley, 2019), or diabetic kidney disease, which is another form of CKD induced by hyperglycemia (Kato and Natarajan, 2019). Elevation of HK2 has also been speculated to be related to ischemic CKD (Rabbani and Thornalley, 2019). Although HK2 isoforms have distinct roles in non-pathologic cellular processes, HK2 overexpression can occur in some pathogenic contexts such as in mediating tumorigenic activity (Liu *et al.*, 2017) or the profibrotic actions of transforming growth factor- β (Yin *et al.*, 2019). A recent report suggested that HK2 is usually bound to the voltage-dependent anion channel (VDAC) on the outer mitochondrial membrane, but abnormally high cytosolic glucose concentrations displace and stabilize HK2 without upregulating other enzymes involved in glycolysis. This causes the accumulation of methylglyoxal (MG) and MG-derived advanced glycation end-products, all of which contribute to dysregulation of metabolism, leading to mitochondrial dysfunction and oxidative stress.

In the current study, we investigated whether HK2 overexpression is related to non-diabetic CKD and MSCs from the CKD condition. Our data showed that HK2 expression was markedly elevated in the serum of CKD patients, as well as in MSCs derived from a CKD mouse model (CKD-mMSCs). Thus, we hypothesized that melatonin may influence the accumulation of abnormal glycolytic intermediates, or glycolytic overload, to improve mitochondrial function and protect MSCs against CKD. Melatonin-treated CKD-mMSCs demonstrated lower HK2 expression, a reduced MG concentration, improved mitochondrial function, and normal glycolysis. Collectively, our data show that melatonin downregulates HK2 and MG to improve the therapeutic potential of MSCs for CKD.

MATERIALS AND METHODS

Serum collection of CKD patients and healthy controls

The local ethics committee approved this study, and informed consent was obtained from all individuals participating in the study. Explanted sera (n=30) were obtained from patients with CKD at Gyeongsang National University, Jinju, Korea (IRB: SCHUH 2018-04-035-002). Upon fulfilling the transplantation criteria, the control samples were obtained from healthy controls (n=30) at the Korea Institute of Radiological & Medical Sciences (IRB: SCHUH 2018-04-035-002). CKD diagnoses were made based on abnormal kidney function with an estimated glomerular filtration rate <25 mL \cdot min $^{-1}$ \cdot 1.73 m $^{-2}$ for over 3 months (stages 3b-5).

Establishment of the CKD mouse model and harvesting bone marrow-derived MSCs

We prepared a CKD mouse model following a previously established protocol (Yoon *et al.*, 2020a, 2020b). In brief, 8-week-old male BALB/c mice were fed an adenine-containing diet (0.75% adenine in diet) for 1-2 weeks. Mouse body weights were measured weekly. Bone marrow-derived MSCs were harvested from the hind limb long bones. Isolated bone marrow was centrifuged and suspended in primary culture medium, and bone marrow-derived MSCs were cultured in a humidified incubator at 37°C with 5% CO $_2$. Healthy- and CKD-MSCs were separated from each group with five mice per group. All MSCs were characterized in each cell line.

Detection of MG in serum and cellular lysates

The concentrations of MG in the human healthy control group or CKD patient group, or in whole cell lysates of healthy mMSCs, CKD-mMSCs, and CKD-mMSCs treated with melatonin were determined using a commercially available enzyme-linked immunosorbent assay (ELISA) kit (Lifespan Biosciences, Seattle, WA, USA). A total of 100 μ L of the sample from each group was used for the experiments. The levels of MG were quantified by measuring the absorbance at 450 nm using a microplate reader (BMG Labtech, Ortenberg, Germany).

Melatonin treatment to mMSCs

Healthy- and CKD-mMSCs were washed twice with phosphate-buffered saline (PBS), and then cultured in fresh alpha-minimal essential medium (HyClone, Logan, UT, USA) supplemented with 10% fetal bovine serum (GIBCO, Grand Island, NY, USA). To examine the protective effect of melatonin (Sigma-Aldrich, St. Louis, MO, USA), mMSCs were incubated with melatonin (1 μ M) at 37°C for 24 h and then subjected to various experimental assays for respective purposes. For luzindole (Sigma-Aldrich) treatment, the mMSCs were pretreated with luzindole (1 μ M for 48 h) before melatonin treatment.

Preparation of the mitochondrial fraction

The harvested healthy-mMSCs or CKD-mMSCs were lysed in mitochondria lysis buffer (Thermo Fisher Scientific, Waltham, MA, USA) and were subsequently incubated for 10 min on ice. The mixtures were centrifuged at 5,000 g at 4°C for 20 min. The residual pellet containing the mitochondrial fraction was lysed with RIPA lysis buffer (Thermo Fisher Scientific) and then used for subsequent analysis.

Western blot analysis

Whole cell lysates, cytosol fraction lysates, or mitochondrial lysates from healthy-mMSCs or CKD-mMSCs (30 μ g protein) were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis on an 8%-12% gel, and the proteins were transferred to a nitrocellulose membrane. After the blots were washed with TBST (10 mM Tris-HCl [pH 7.6], 150 mM NaCl, 0.05% Tween 20), the membranes were blocked with 5% skim milk for 1 h at room temperature and incubated with the appropriate primary antibodies against HK1 (#2024S, Cell Signaling Technology, Beverly, MA, USA), HK2 (#2867S, Cell Signaling Technology), VDAC1 (NB100-695, Novus), CDK2 (sc-6248, Santa Cruz Biotechnology, Santa Cruz, CA, USA), CDK4 (SC-56277, Santa Cruz Biotechnology), cyclin D1 (SC-20044, Santa Cruz Biotechnology), cyclin E (SC-377100, Santa Cruz Biotechnology), and β -actin (sc-47778, Santa Cruz Biotechnology). The membranes were then washed, and the primary antibodies were detected using goat anti-rabbit IgG or goat anti-mouse IgG secondary antibodies (Santa Cruz Biotechnology). The bands were detected by enhanced chemiluminescence (Amersham Pharmacia Biotech, Little Chalfont, UK). Densitometric quantification was performed using ImageJ software 1.48v (National Institutes of Health, Bethesda, MD, USA).

Measurement of mitochondrial superoxide and mitochondrial membrane potential

The production of mitochondrial superoxide in mMSCs was measured using MitoSOX (Thermo Fisher Scientific). To measure mitochondrial membrane potential, tetramethylrhodamine, methyl ester, and perchlorate (TMRE; Thermo Fisher Scientific) dye was used. The mMSCs in each group were trypsinized and centrifuged at 1,200 rpm for 3 min. After washing with PBS twice, the cells were treated with either 10 μ M MitoSOX in PBS or 200 nM TMRE in PBS at 37°C for 15 min. MitoSOX-positive or TMRE-positive cells were analyzed using flow cytometry (Sysmex, Kobe, Japan). Flow cytometry data were analyzed using FCS Express 5 Flow research software (DeNovo Software, Los Angeles, CA, USA).

Measurement of oxygen consumption rate (OCR) and extracellular acidification rate (ECAR) with a Seahorse assay

OCR and ECAR were measured in mMSCs using an XF96 extracellular flux analyzer (Seahorse Bioscience, MA, USA). In brief, mMSCs were seeded at 45,000 cells per well in XF96 cell culture multi-well plates in Dulbecco's modified Eagle medium, and incubated for 24 h at 37°C and 5% CO₂. The XF96 cartridges were then incubated overnight in an XF calibrant at 37°C in a non-CO₂ incubator. After the growth medium of mMSCs was changed with XF medium, the plates were incubated at 37°C in a non-CO₂ incubator for 1 h. Inhibitors were diluted to appropriate concentrations in XF medium and loaded into corresponding microwells in the XF96 cartridge plate. Following the equilibration of the sensor cartridges, the XF96 cell culture plate was loaded into the XF96 Extracellular Flux analyzer at 37°C. OCR and ECAR were measured after cycles of mixing and data acquisition (basal), or inhibitor injection, mixing, and data acquisition using Seahorse Wave Desktop software (Agilent Technologies, Santa Clara, CA, USA).

Ethics statement

All animal care procedures and experiments were approved by the Institutional Animal Care and Use Committee of Soonchunhyang University Seoul Hospital (Seoul, Korea) (IA-CUC2013-5) and performed in accordance with the National Research Council (NRC) Guidelines for the Care and Use of Laboratory Animals. All animal experiments were performed using 8-week-old male BALB/c mice (Biogenomics, Seoul, Korea). The animals were maintained on a 12 h light/dark cycle at 25°C, in accordance with the regulations of Soonchunhyang University Seoul Hospital.

Statistical analysis

Values are expressed as the mean \pm SEM and were evaluated by Student's t-test, or one- or two-way analysis of variance to compute the significance of differences between groups. Dunnett's test was used for comparisons of three or more groups. Data were considered significantly different at $p < 0.05$.

RESULTS

Patients with CKD show an increased level of HK2, and CKD mice-derived MSCs show impaired mitochondrial functionality

To analyze the relationship between HK2 expression and CKD, we measured HK2 expression levels in the serum of patients with CKD. ELISA revealed a high level of HK2 in the serum of CKD patients compared with that of the healthy control group (Fig. 1A). To confirm that HK2 expression leads to metabolic dysregulation, we examined the serum concentration of MG, a cytotoxic glycolytic intermediate. MG levels were also significantly elevated in CKD patients compared with those in the healthy control group (Fig. 1B). These results are consistent with previous reports suggesting that HK2 overexpression may play a significant role in kidney injury (Smith *et al.*, 2014; Lan *et al.*, 2016).

To examine the relationship between HK2 and MSCs in the context of CKD, MSCs were isolated from both healthy and CKD mice. Immunoblotting revealed that CKD mice-derived MSCs exhibited elevated HK2 levels compared with those of their healthy counterparts, whereas HK1 expression remained unaffected (Fig. 1C). In addition, CKD-mMSCs had increased levels of mitochondrial reactive oxygen species (mtROS) and significantly decreased mitochondrial membrane potential, as assessed using the MitoSOX assay and TMRE staining, respectively (Fig. 1D, 1E). Taken together, these results indicate that in patients with CKD, increased HK2 expression promotes toxic glycolytic intermediates such as MG, and the effect of these intermediates might carry over to MSCs, undermining the functionality of mitochondria in harvested MSCs and their therapeutic potential.

Melatonin improves oxidative stress and mitochondrial membrane potential by lowering mitochondrial HK2 expression in CKD-mMSCs

Our data verifying the high HK2 levels in patients with CKD as well as in those with CKD-mMSCs suggested that CKD-mMSCs may suffer from impaired mitochondrial function related to the role of HK2 in metabolism. Hence, we hypothesized that decreasing the expression of HK2 may increase the

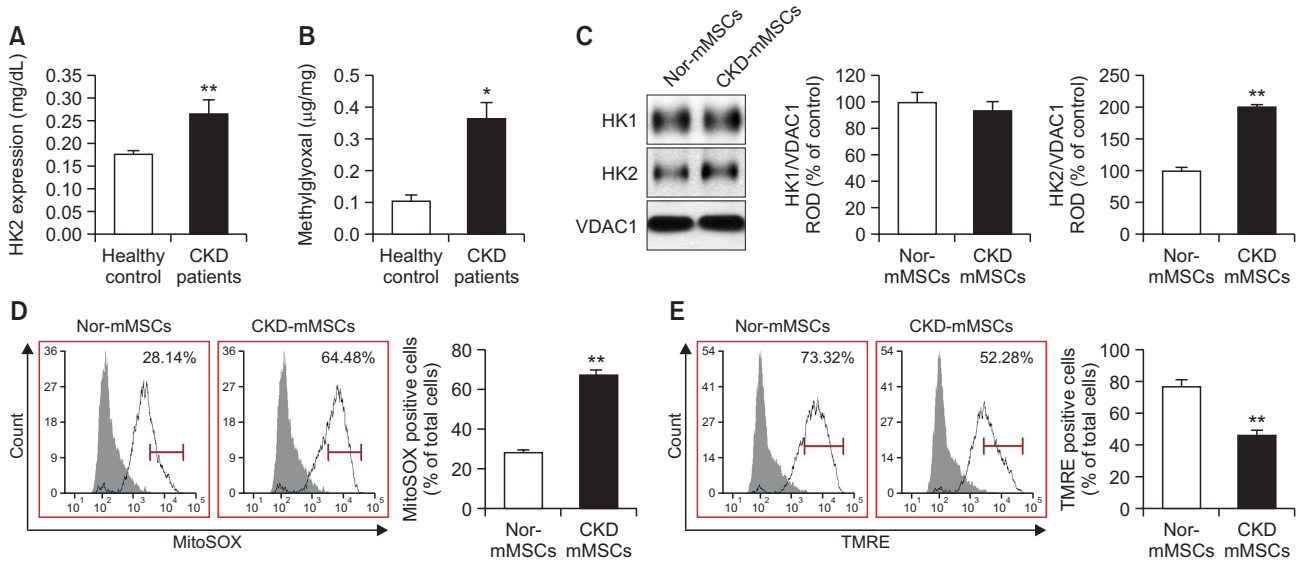


Fig. 1. Patients with chronic kidney disease (CKD) show an increased level of HK2, and CKD mice-derived mesenchymal stem/stromal cells (MSCs) show impaired mitochondrial functionality. (A) Serum samples from patients with CKD (n=30) and healthy controls (n=30) were subjected to enzyme-linked immunosorbent assay (ELISA) for HK2. (B) Methylglyoxal (MG) content in human serum samples from the same experimental groups (CKD patients vs. healthy control group). (C) Expression of HK1 and HK2 in mitochondrial fractions of MSCs derived from the CKD mouse model (CKD-mMSCs) and normal control mice (Nor-mMSCs), as detected by immunoblotting (n=5). Expression levels of HK1 and HK2 were normalized relative to levels of the mitochondrial marker VDAC1. (D) Normal-mMSCs and CKD-mMSCs were assayed using MitoSOX to assess mitochondrial oxidative stress. MitoSOX signals were quantified using flow cytometry (n=5). (E) Normal-mMSCs and CKD-mMSCs were assayed using TMRE to assess the mitochondrial membrane potential. TMRE signals were quantified using flow cytometry (n=5). Values represent the mean \pm standard error of the mean (SEM). * $p < 0.05$ or ** $p < 0.01$ vs. control.

functionality of MSCs by positively influencing mitochondrial activity. On a different note, we also questioned whether melatonin can modulate glycolytic overload via HK2. We previously reported that melatonin treatment provides protective effects against CKD, and improves mitochondrial functions in renal proximal tubular cells (Han *et al.*, 2020) and CKD-MSCs (Yoon *et al.*, 2020c). Others have proposed that melatonin can act as a glycolytic agent, which inhibits Warburg effect-dependent cancer cells by shifting the metabolic focus from anaerobic glycolysis to normal mitochondrial oxidative phosphorylation (Reiter *et al.*, 2020). In light of these previous findings, we expected melatonin to downregulate the overabundant HK2 and its glycolytic side product MG in CKD-MSCs. Immunoblotting against HK2 showed increased HK2 expression levels in CKD-mMSCs, which was effectively reduced with melatonin treatment (Fig. 2A). Evaluation of mitochondrial oxidative stress and membrane potential demonstrated reduced mtROS and increased active mitochondria in melatonin-treated CKD-mMSCs compared with those of the no-treatment group (Fig. 2B, 2C). We also determined the effect of melatonin on the glycolytic process in TH1 cells, which are human renal proximal tubule epithelial cells. Treatment of TH1 cells with p-cresol increased the expression of HK2. However, melatonin treatment of p-cresol-treated TH1 cells reduced the expression of HK2, similar to the results in CKD-mMSCs (Supplementary Fig. 1). Treatment of TH1 cells with p-cresol also increased HK1 expression levels, which was reversed by treatment with luzindole, a melatonin inhibitor, confirming that the observed results were indeed dependent on melatonin. Taken together, these results indicate that melatonin can modulate HK2 expression to improve the functionality of mitochondria in CKD-

mMSCs.

Melatonin decreases the production of HK2-related toxic metabolites and improves the metabolic activity of CKD-mMSCs

After confirming the melatonin-mediated reduction of HK2 expression in CKD-mMSCs, we questioned whether the functional improvements observed with mitochondria in melatonin-treated CKD-mMSCs were related to the role of HK2 in glycolytic processes. Consistent with the differences in HK2 expression levels, the MG concentrations were elevated in CKD-mMSCs compared with those of mMSCs harvested from healthy mice (Nor-mMSCs), and this abnormal elevation of MG concentrations was inhibited by melatonin treatment (Fig. 3A).

To determine whether this change in MG concentration is reflective of the larger shift in the bioenergetic profile of MSCs in response to melatonin treatment, we employed a seahorse assay to measure the OCR and ECAR. Plotting the OCR vs. ECAR for CKD-mMSCs, melatonin-treated CKD-mMSCs, and Nor-mMSCs (Fig. 3B) showed a consistently positive correlation with similar slopes, which suggests a similar metabolic profile across different conditions, and that the increase in OCR and ECAR values corresponded to an increase in the metabolic activity of mMSCs. Melatonin-treated CKD-mMSCs showed improved metabolic activity compared with that of the Nor-mMSC group, whereas CKD-mMSCs, which have no efficacy, and melatonin inhibitors were grouped together given their low metabolic activity. Specific measurements of ECAR (Fig. 3C) and OCR (Fig. 3D) demonstrated improved glycolysis and mitochondrial respiration, respectively. Overall,

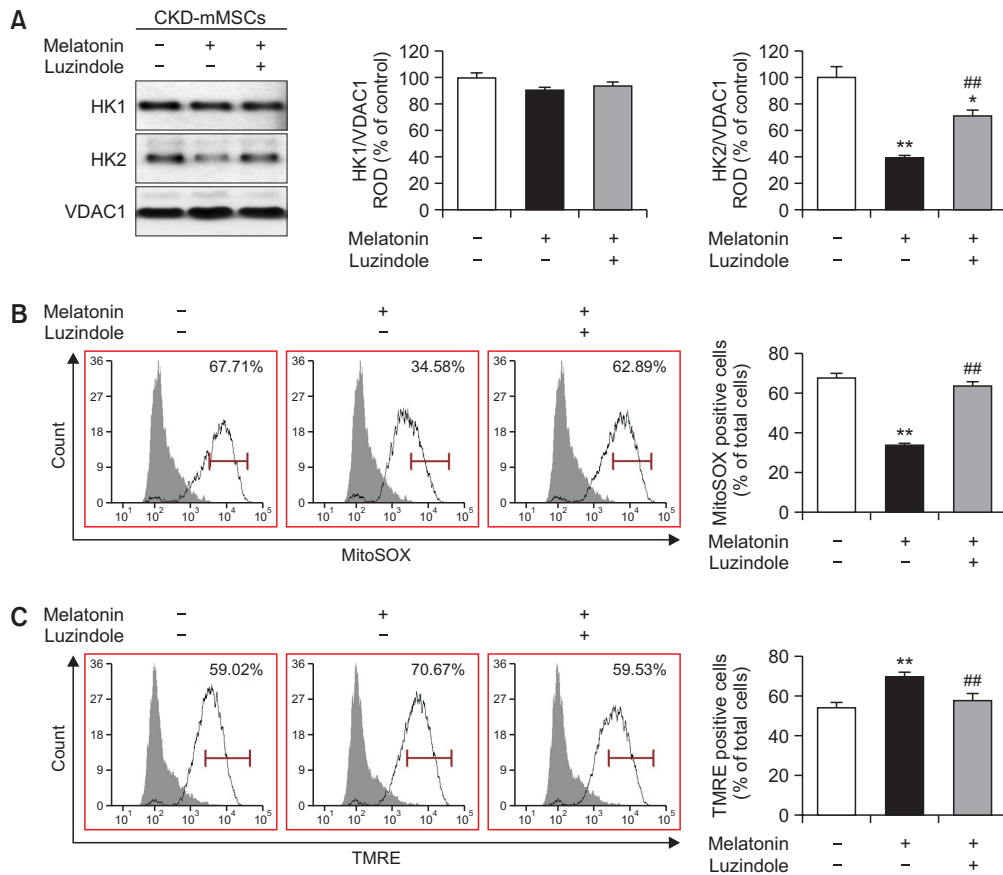


Fig. 2. Melatonin improved oxidative stress and mitochondrial membrane potential by lowering mitochondrial HK2 expression in chronic kidney disease mouse model-derived mesenchymal stem cells (CKD-mMSCs). (A) Immunoblotting results showing the expression of HK1 and HK2 in each experimental mMSC group: untreated CKD-mMSCs, CKD-mMSCs treated with melatonin, and melatonin-treated CKD-mMSCs pre-treated with luzindole. Expression levels of HK1 and HK2 were normalized relative to VDAC1 levels. Values represent the mean \pm SEM (n=3). * p <0.05 or ** p <0.01 vs. control; ## p <0.01 vs. melatonin-treated CKD-mMSCs. (B) CKD-mMSCs from each experimental group assayed using MitoSOX to assess mitochondrial oxidative stress. MitoSOX signals were quantified using flow cytometry. Values represent the mean \pm SEM (n=5). ** p <0.01 vs. control, ## p <0.01 vs. melatonin-treated CKD-mMSCs. (C) CKD-mMSCs from each experimental group assayed using TMRE to assess mitochondrial membrane potential. TMRE signals were quantified using flow cytometry. Values represent the mean \pm SEM (n=5-6). ** p <0.01 vs. control, ## p <0.01 vs. melatonin-treated CKD-mMSCs.

experimental mMSC groups with high MG levels were metabolically less active with worse performance in glycolysis and mitochondrial respiration, and melatonin treatment was able to enhance the metabolic performance of CKD-mMSCs closer to that of Nor-mMSCs through regulation of MG by influencing HK2 expression.

Melatonin restores the cell cycle and expression patterns of cell cycle-associated proteins in CKD-mMSCs

Since melatonin was found to be capable of regulating HK2 expression and MG concentration in CKD-mMSCs to restore their mitochondrial function and metabolic activity, we next analyzed whether these improvements translated into an enhanced proliferative capacity of MSCs, which is an important feature for autologous MSC treatment. To assess the improvement in cell proliferation induced by melatonin in CKD-mMSCs, we determined the cell cycle of CKD-mMSCs using flow cytometry. CKD-mMSCs showed a decrease in the cell population in the S phase compared with that of Nor-mMSCs. However, melatonin treatment effectively restored the cell

population in the S phase (Fig. 4A). We further determined the expression of cell cycle-associated proteins in CKD-mMSCs. Western blotting data showed marked downregulation in the expression of CDK2, cyclin E, CDK4, and cyclin D1, which are all important for the G1-phase to S-phase transition, whereas melatonin treatment successfully restored the expression of these proteins in CKD-mMSCs (Fig. 4B). Densitometric analysis showed that pre-treatment with luzindole almost completely negated the protective effect of melatonin, indicating that all improvements observed with CKD-mMSCs were attributable to the melatonin treatment. These results suggest that the ability of melatonin to regulate HK2 and its downstream metabolite MG may have implications for restoring cell proliferation in CKD-mMSCs.

Melatonin improves the therapeutic efficacy of CKD-mMSCs in the CKD mouse model

After confirming that melatonin treatment improved the mitochondrial function and cell proliferation of CKD-mMSCs, we tested whether melatonin could improve the therapeutic

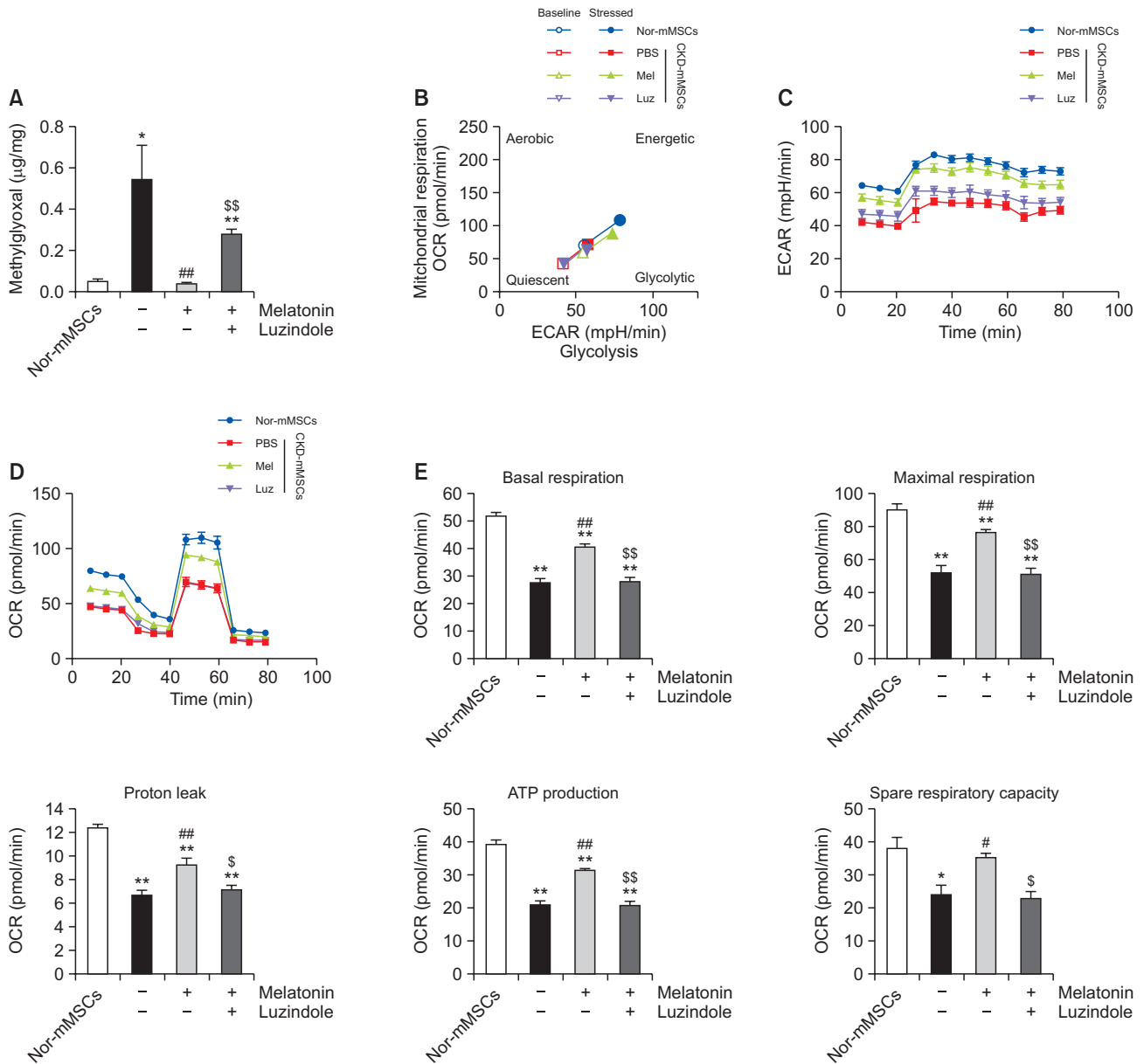


Fig. 3. Melatonin decreases the levels of HK2-related toxic metabolites and improves the metabolic activity of chronic kidney disease mouse model-derived mesenchymal stem cells (CKD-mMSCs). (A) Measurement of methylglyoxal (MG) in mMSCs derived from normal healthy mice (Nor-mMSCs) or CKD mice (CKD-mMSCs) with enzyme-linked immunosorbent assay (ELISA). CKD-mMSCs were either treated with melatonin alone or pre-treated with luzindole before melatonin treatment. Values represent the mean \pm SEM (n=5). * p <0.05 or ** p <0.01 vs. Nor-mMSCs; ### p <0.01 vs. CKD-mMSCs; \$\$\$ p <0.01 vs. melatonin-treated CKD-mMSCs. (B) Mitochondrial respiration rate (OCR) vs. extracellular acidification rate (ECAR) equated to the glycolytic rate of Nor-mMSCs, CKD-mMSCs, and CKD-mMSCs with melatonin treatment (Mel), or melatonin-treated CKD-mMSCs pre-treated with luzindole (Luz). Assessment of the baseline metabolic profile of each mMSC group was followed by a stress test to measure the metabolic shift toward the energetic state. (C) Seahorse assay for determining the ECAR of Nor-mMSCs and CKD-mMSCs under different experimental conditions. (D) Mitochondrial OCR of each group over time (min). (E) Histogram representing the bioenergetics parameters are based on the OCR data by administering oligomycin (20 min, 1 μ M), FCCP (40 min, 0.75 μ M), and antimycin A (60 min, 1 μ M). Values represent the mean \pm SEM (n=5). * p <0.05 or ** p <0.01 vs. Nor-mMSCs; # p <0.05 or ## p <0.01 vs. CKD-mMSCs; \$ p <0.05 or \$\$\$ p <0.01 vs. melatonin-treated CKD-mMSCs.

efficacy of CKD-mMSCs in the CKD mouse model. We examined the renal function recovery in CKD mice after the injection of melatonin-treated CKD-mMSCs by measuring serum blood urea nitrogen (BUN) and creatinine levels. When Nor-mMSCs were injected into CKD mice, BUN and creati-

nine levels decreased, whereas the injection of CKD-mMSCs did not decrease BUN and creatinine levels. However, when melatonin-treated CKD-mMSCs were injected into CKD mice, BUN and creatinine levels significantly increased (Fig. 5A, 5B). Further, treatment with luzindole reversed these effects,

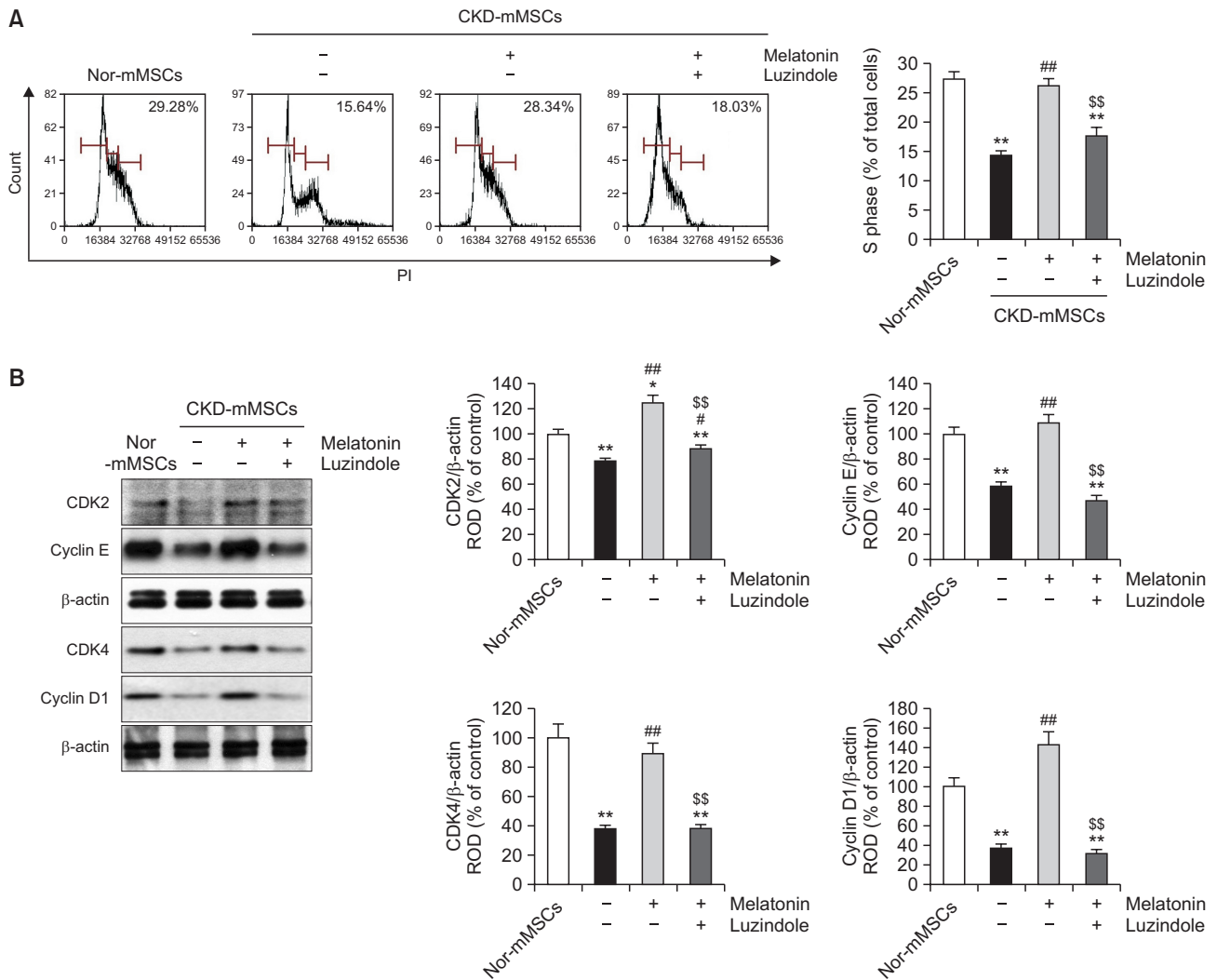


Fig. 4. Melatonin restores the cell cycle and expression patterns of cell cycle-associated proteins in chronic kidney disease mouse-derived mesenchymal stem cells (CKD-mMSCs). (A) Cell cycle analysis of control mouse MSCs (Nor-mMSCs) and CKD-mMSCs using flow cytometry. (B) Immunoblotting data showing the expression of cell cycle-associated proteins, CDK2, cyclin E, CDK4, and cyclin D1, in Nor-mMSCs, CKD-mMSCs, CKD-mMSCs with melatonin treatment, or melatonin-treated CKD-mMSCs pre-treated with luzindole. Band signals from western blotting were quantified by densitometry and were normalized to β -actin levels (n=3). Values represent the mean \pm SEM. * p <0.05 or ** p <0.01 vs. Nor-mMSCs; # p <0.05 or ## p <0.01 vs. CKD-mMSCs; \$\$ p <0.01 vs. melatonin treated CKD-mMSCs.

confirming that the observed effects were due to melatonin treatment (Fig. 5A, 5B). These results suggest that melatonin enhances the mitochondrial function and cell proliferation of CKD-mMSCs, and improves the therapeutic efficacy of CKD-mMSCs in a CKD mouse model.

DISCUSSION

Sustained damage to the kidney tissues leads to CKD and progressive loss of kidney function. The compromised glomerular filtration, excretion, and reabsorption causes the increased circulation of uremic toxins. These effects can be detrimental to the viability and normal functions of MSCs and limit their application in regenerative therapy for CKD patients. Although previous studies to improve the efficacy of MSC-

based therapy for CKD have demonstrated that melatonin treatment provides a broad range of protective effects (Han *et al.*, 2019; Hu and Li, 2019; Yoon *et al.*, 2020c), the complex biological mechanisms underlying this phenomenon have not been well elucidated. In this study, we demonstrated that treatment of MSCs with melatonin modulates the glycolytic process in these cells by influencing HK2 expression and MG concentration, which improved mitochondrial function and cell proliferation in CKD-mMSCs (Fig. 6).

Although recent studies on the mechanisms underlying the therapeutic effect of melatonin against kidney diseases have focused on its effect on mitochondrial function (Yoon *et al.*, 2020a), homeostasis (Qi and Wang, 2020), and autophagy (Yoon *et al.*, 2019), whether melatonin modulates the glycolytic pathways in CKD remains a relatively under-researched topic. Several review articles in the field of cancer research

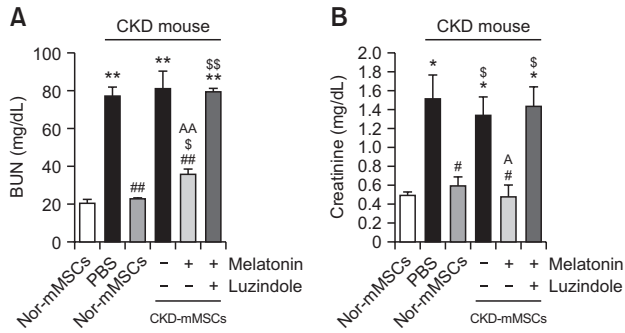


Fig. 5. Melatonin treatment improves the therapeutic efficacy of chronic kidney disease mouse-derived mesenchymal stem cells (CKD-mMSCs). Measurement of blood urea nitrogen (A) and creatinine (B) levels in the serum of CKD mice. Each CKD mouse was injected with phosphate-buffered saline (PBS) or with control mouse MSCs (Nor-mMSCs), CKD-mMSCs, melatonin-treated CKD-mMSCs, or melatonin-treated CKD-mMSCs pretreated with luzindole. Values represent the mean \pm SEM. * p <0.05 or ** p <0.01 vs. healthy mouse; # p <0.05 or ## p <0.01 vs. PBS; \$ p <0.05 or \$\$ p <0.01 vs. Nor-mMSCs; ^ p <0.05, ^^ p <0.01 vs. CKD-mMSCs.

have elucidated the metabolism-altering anti-cancer effects of melatonin (Li *et al.*, 2017; Reiter *et al.*, 2017; Talib, 2018; Wang *et al.*, 2018). For example, melatonin has been found to inhibit tumor growth in Ewing sarcoma (Sanchez-Sanchez *et al.*, 2015), leiomyosarcoma (Mao *et al.*, 2016), breast cancer (Hill *et al.*, 2015), prostate cancer (Hevia *et al.*, 2017), and hepatocellular carcinoma (Feng *et al.*, 2020) by suppressing excessive aerobic glycolysis (Warburg effect). Furthermore, a recent review article suggested that the modulation of glycolytic functions by melatonin is also associated with inhibition of various diseases (Reiter *et al.*, 2021). On a different note, HK2-mediated glycolytic dysregulation has been implicated in fibrotic processes (Xie *et al.*, 2015; Yin *et al.*, 2019; Gopu *et al.*, 2020), which is an important part of the CKD pathophysiology. With the above findings providing a hint for the link between melatonin and glycolysis in CKD, our present study provides more tangible evidence for the melatonin-mediated modulation of glycolysis in the CKD environment, although our experiments were more focused on MSCs.

hMSCs are among the most commonly used adult stem cells in experimental cell therapies (Pittenger *et al.*, 2019). Despite the initial focus on the multi-potency of MSCs as the rationale behind applying MSC-based cell therapy, more recent studies have led to the novel perspective that the more significant aspect of MSCs for clinical applications comes from

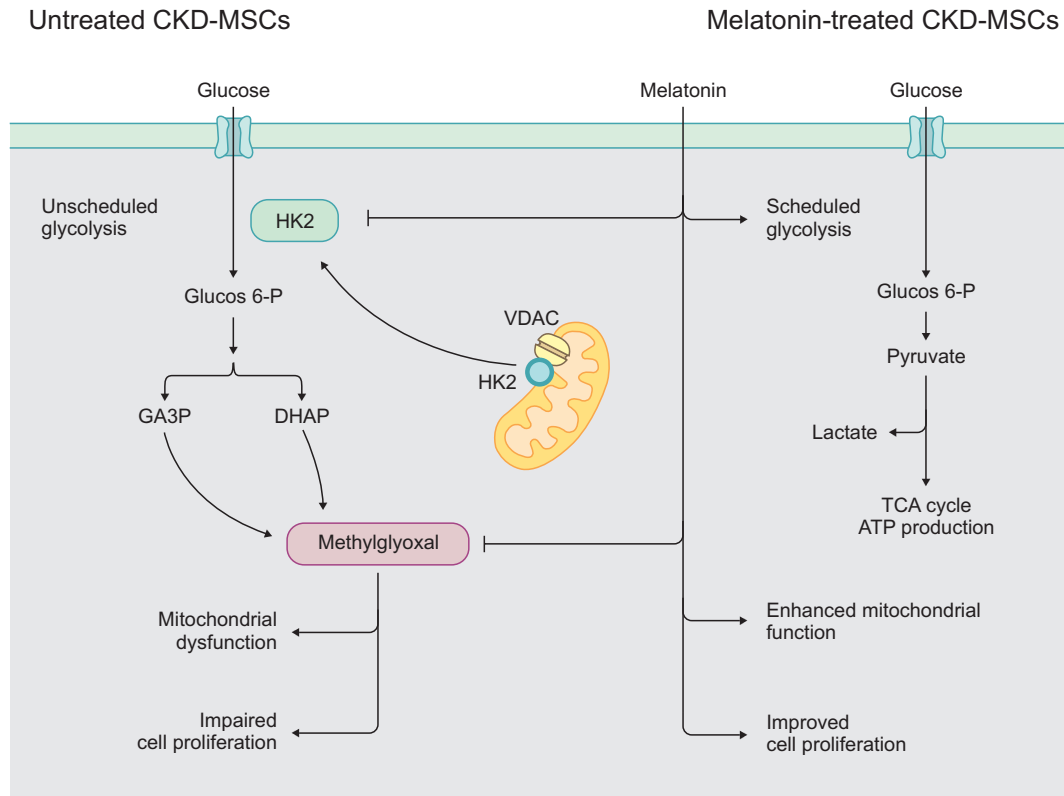


Fig. 6. Schematic representation of the proposed protective effect of melatonin treatment for mouse mesenchymal stem/stromal cells in chronic kidney disease (CKD-mMSCs) through modulating hexokinase-2 (HK2), thereby reducing the damage caused by the accumulation of methylglyoxal (MG) during abnormal or unscheduled glycolysis. In CKD-mMSCs, HK2 dissociates from the mitochondrial voltage-dependent anion channel (VDAC) and induces excessive production of a toxic glycolytic intermediate product (MG). This leads to mitochondrial dysfunction and impaired cell proliferation. Melatonin treatment of CKD-mMSCs enhances mitochondrial functions and rescues the normal expression of cell cycle-related proteins.

their ability to secrete immunoregulatory and pro-regenerative factors (Yuan *et al.*, 2019). There is growing interest in using allogeneic and culture-expanded cells for MSC-based cell therapy because their characteristics are well documented and they have potential for mass production. Nonetheless, this approach also lends itself to other barriers for the therapeutic application of MSCs, as the specific morbidities of MSC donors or extensive cell culturing can alter the metabolism of MSCs and compromise their therapeutic potential (Mastrolia *et al.*, 2019). Therefore, understanding the underlying regulatory mechanisms of MSC metabolism and functions, and finding ways to control or bioengineer these properties of MSCs may help to drastically improve the efficacy of MSC-based therapy for specific target diseases. In this light, our successful attempt to improve the glycolytic profile of CKD-MSCs with melatonin treatment may help build the foundation for additional research efforts to develop an effective cure for CKD.

Although melatonin is primarily known for its role in sleep regulation, the broad impact of melatonin on regenerative processes in various tissues provides an excellent entry point for exploring different regulatory mechanisms for therapeutic purposes (Tordjman *et al.*, 2017). For example, our previous work with CKD-mMSCs revealed that melatonin enhances the viability, motility, stemness, and proliferation of MSCs (Han *et al.*, 2019; Lee *et al.*, 2020). We also reported that melatonin-treated MSCs can better withstand the pathophysiological conditions of the CKD environment when used for CKD cell therapy in a mouse model (Han *et al.*, 2019; Lee *et al.*, 2019). In addition, we established specific mechanisms for the protective effects of melatonin against CKD in MSCs involving antioxidant enzymes (Lee *et al.*, 2020), proangiogenic cytokines (Lee *et al.*, 2020), and cellular prion proteins (Han *et al.*, 2019). Our current research revealed that the glycolytic enzyme HK2 and its intermediate metabolite MG can be a target of modulation in melatonin treatment for CKD, providing another potential option for treating CKD-MSCs prior to their use in further cell-based CKD therapies.

In this study, we demonstrated that melatonin restores mitochondrial function and promotes cell proliferation by inhibiting the expression of HK2 and the accumulation of MG. There are conflicting reports on the correlation between mitochondrial function and cell proliferation. In cancer cells with active cell division, mitochondrial function is decreased and glycolysis is increased (Galli *et al.*, 2003; Antico Arciuch *et al.*, 2012). Recent studies have shown that mitochondrial function restoration can promote cell proliferation (Yan *et al.*, 2017; Yao *et al.*, 2019). Therefore, in this study, melatonin may have promoted cell proliferation by enhancing mitochondrial function. Moreover, there is a possibility that melatonin directly promotes cell proliferation through other mechanisms.

Given that the clinical treatment of CKD patients with hMSC-based cell therapy requires substantial improvements in efficacy, we examined the efficacy of the melatonin-mediated inhibition of HK2 expression in CKD-mMSCs derived from a CKD mouse model. Not only were abnormal elevations of HK2 and its downstream metabolite MG inhibited by melatonin treatment, but also mitochondrial function as assessed by the respiratory OCR, and cell proliferation as measured by the expression of pro-proliferative proteins, were also improved. This experimental outcome may reflect the significant role of HK2 in the cellular activity and metabolism of MSCs. It is noteworthy that the broad impact of melatonin on CKD-

mMSCs influences other metabolic processes in addition to hexokinase-induced glycolysis. Therefore, identification and characterization of the various intertwined metabolic regulators under the influence of melatonin with respect to MSCs in a CKD condition may build a foundation for developing more effective therapeutic strategies, cell-based or otherwise, for treating CKD and other kidney disorders.

CONFLICT OF INTEREST

The authors have no conflict of interest to declare.

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