

Melatonin attenuates high glucose-induced endothelial cell pyroptosis by activating the Nrf2 pathway to inhibit NLRP3 inflammasome activation

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Abstract. Endothelial injury induced by hyperglycemia is the most critical initial step in the development of diabetic vasculopathy. The aim of this present study was to explore the prevention and treatment strategies and elucidate the specific mechanism of diabetes-induced vascular endothelial injury. Melatonin, a hormone secreted by the pineal gland to regulate biological rhythm, serves an important role in maintaining human physiological function. Pyroptosis is a type of newly discovered inflammatory cell death. The current study first found by western blotting that melatonin could activate nuclear factor erythroid 2-related factor 2 (Nrf2) pathway in human umbilical vein endothelial cells (HUVECs) under high glucose (HG) condition. Second, it found that pretreatment with Luzindole, a specific inhibitor of melatonin receptor (MT1/MT2), significantly reduced the activation of Nrf2 pathway by melatonin in HUVECs. It also found that pretreatment with melatonin or a specific NOD-like receptor family, pyrin domain-containing 3 (NLRP3) inhibitor (MCC950) pretreatment reduced HG-induced endothelial cell pyroptosis. Finally, it was found

that the protective effect of melatonin against reactive oxygen species/NLRP3 inflammasome pathway activation induced by HG in HUVECs was decreased after Nrf2 knockdown. In conclusion, the present study showed that melatonin may serve a protective role in HG-induced vascular endothelial cell pyroptosis by activating the Nrf2 pathway to inhibit NLRP3 inflammasome activation. In addition, it was further found that melatonin attenuated HG-induced vascular endothelial cell injury by interacting with its receptors (MT1/MT2) to promote activation of Nrf2 pathway.

Introduction

Vascular disease is the core of diabetes mellitus (DM), which endangers life and health. Severe vascular disease can cause insufficient blood supply and function loss in the heart, brain, kidney, eyes, lower limbs and other important tissues and organs, which seriously threatens human life and health (1). Endothelial injury induced by hyperglycemia is the most critical initial step in the development of diabetic vasculopathy (2). Damaged endothelial cells can secrete a large number of proinflammatory and procoagulant cytokines, which can induce platelet aggregation, thrombosis, adhesion and infiltration of mononuclear macrophages and release of inflammatory factors. Long-term chronic vascular wall inflammation can stimulate the proliferation and migration of vascular smooth muscle cells and the deposition of extracellular matrix, eventually leading to vascular stenosis and occlusion (3). Therefore, it is of great clinical value to explore the prevention and treatment strategies and elucidate the specific mechanism of diabetes-induced vascular endothelial injury.

Melatonin, a hormone secreted by the pineal gland to regulate biological rhythm, serves an important role in maintaining human physiological function (4). Studies have found that melatonin has anti-inflammatory and antioxidant protective effects in diabetes-induced vascular endothelial cell injury (5,6), but the specific mechanism remains to be elucidated.

Pyroptosis, a type of newly discovered inflammatory cell death, depends on the activation of caspase-1 (7). The NOD-like receptor family pyrin domain containing 3 (NLRP3)

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inflammasome, one of the most extensively studied pyroptotic inflammasomes, is composed of NOD-like receptor family, pyrin domain-containing 3 (NLRP3), the adaptor ASC and pro-caspase-1 (8). Activation of the NLRP3 inflammasome involves two sequential steps (priming and assembly) triggered by two signals respectively. The priming step triggered by the first signal (e.g., pathogen-associated molecular pattern molecules such as lipopolysaccharide) activates NF- κ B signaling and induces the transcription of pre-IL-1 β and NLRP3. The second signal (e.g., damage-associated molecular patterns such as ATP) triggers several signaling pathways, including potassium efflux, reactive oxygen species (ROS) production, and lysosomal damage, all of which stimulate the assembly of the NLRP3 inflammasome. Activation of the NLRP3 inflammasome induces the transformation of caspase-1 into its activated form, cleaved-caspase-1. For one thing, this enzyme cleaves gasdermin D (GSDMD) into gasdermin-N domain (GSDMD-N), which oligomerizes on the cell membrane to form cell membrane pores. For another, cleaved-caspase-1 promotes the maturation of the pro-inflammatory cytokines IL-1 β and IL-18, which are released into the extracellular space through GSDMD cell membrane pores, to induce a strong inflammatory response in cells and finally pyroptosis (9). Previous studies have found that melatonin can inhibit NLRP3 inflammasome activation and pyroptosis (10,11). However, whether melatonin can inhibit endothelial pyroptosis and serve a protective role in diabetes-induced vascular endothelial injury is rarely reported.

Nuclear factor erythroid 2-related factor 2 (Nrf2) is an activator of antioxidant response elements (ARE) and a key target when reducing oxidative stress. Nrf2 binds to kelch-like ECH-associated protein 1 (Keap1) in the cytoplasm under physiological conditions (12). When exposed to oxidative stress, extracellular signal-regulated protein kinases, protein kinase C and phosphatidylinositol 3-kinase signaling pathways promote the dissociation of Keap1-Nrf2 and the activation of Nrf2 through phosphorylation. Activated phosphorylated-Nrf2 is translocated into the nucleus and binds to the cis-acting AREs, initiating the transcription of a series of antioxidant enzyme genes, including heme oxygenase-1 (HO-1), NADPH: Quinone Oxidoreductase 1 (NQO1), superoxide dismutase (SOD) and glutathione peroxidase (GSH-PX), to maintain the stability of the intracellular environment (13). Previous studies have found that melatonin serves an antioxidant role by activating the Nrf2 pathway (14,15). ROS is a key upstream mechanism of NLRP3 inflammasome activation and pyroptosis (16). Therefore, the present study hypothesized that melatonin could inhibit ROS production by activating Nrf2 pathway and further inhibit NLRP3 inflammasome activation and pyroptosis in diabetes-induced vascular endothelial cell injury.

MT1 and MT2 melatonin membrane receptors are G protein-coupled receptors widely expressed in the cardiovascular system. Previous studies have shown that melatonin can bind to the melatonin receptor (MT1/MT2) on the cell membrane surface, activate the phosphorylated kinase pathway in the cytoplasm, and then participate in the regulation of normal physiological functions (17). The present study determined whether melatonin attenuates high glucose (HG)-induced pyroptosis of vascular endothelial cells through the Nrf2-ROS-NLRP3 pathway.

Meanwhile, whether melatonin protects vascular endothelial cells from HG-induced damage through MT1/MT2-mediated Nrf2 signaling was also investigated.

Materials and methods

Cell culture and experimental design. Primary human umbilical vein endothelial cells (HUVECs; cat. no. 8000) were purchased from ScienCell Research Laboratories (Carlsbad, CA, USA) and cultured in endothelial cell medium (ScienCell Research Laboratories, Inc.) supplemented with 5% fetal bovine serum under 5% CO₂ at 37°C. All experiments were performed between cell passages three and five. Melatonin (100 μ M; MilliporeSigma), a specific NLRP3 inhibitor MCC950 (50 μ M; Selleck Chemicals), and a melatonin receptor antagonist luzindole (5 μ M; MilliporeSigma) were added to the cell cultures 3 h in advance before being co-incubated with HG (30 mM) for 72 h. All experiments were performed at least three times and representative results are presented in the present study.

Small interfering RNA (siRNA) treatment. The predesigned siRNA duplexes for Nrf2 and the negative control (NC) were purchased from HippoBio Co., Ltd. Table I lists the sequences of all Nrf2 siRNAs and the non-targeting NC-siRNA used. Cells were incubated in 6-cm dishes until they were 80% confluent and the original medium was replaced with fresh basal medium 2 h before siRNA transfection. Two sterile 1.5-ml centrifuge tubes were prepared and 100 μ l Opti-MEM (Invitrogen; Thermo Fisher Scientific, Inc.) was added to each. siRNA (2 μ l, 20 μ M) was added to one of the centrifuge tubes and was thoroughly mixed to dilute the RNA. LipoRNAi MAX (2 μ l; Invitrogen; Thermo Fisher Scientific, Inc.) was added to the other centrifuge tube and was mixed to dilute the LipoRNAi MAX. Both centrifuge tubes were maintained at room temperature for 5 min. Subsequently, siRNA and LipoRNAi MAX were transferred to the same centrifuge tube and were thoroughly mixed; the transfection complex was then maintained for 25 min at room temperature. The HUVECs to be transfected were removed from the incubator, the transfection complex was added to the culture dish in drops, and the cells were gently shaken. The transfection complex was incubated with cells at 37°C for 48 h. Finally, the transfected cells were harvested for subsequent verification and experiments.

Enzyme-linked immunosorbent assay (ELISA). The HUVEC culture supernatants were collected following an intervention for 72 h. The concentrations of IL-1 β (cat. no. SEA563Hu) and IL-18 (cat. no. SEA064Hu) in the cell culture supernatants were determined using ELISA kits (both from Cloud-Clone Corp.) according to the manufacturer's instructions.

Detection of ROS production. ROS levels in the HUVECs were measured using an ROS assay kit (Beijing Solarbio Science & Technology Co., Ltd.) in accordance with the manufacturer's instructions.

Cell viability assay. A Cell Counting Kit (CCK-8) assay (Nanjing Jiancheng Bioengineering Institute) was used to measure the level of cell viability.

Table I. The sequences of all Nrf2 siRNAs and the siRNA-NC used.

Name	Sequence
hNRF2 siRNA-1 sense	GGUUGAGACUACCAUGGUUTT
hNRF2 siRNA-1 antisense	AACCAUGGUAGUCUCAACCAG
hNRF2 siRNA-2 sense	GCCCAUUGAUGUUUCUGAUTT
hNRF2 siRNA-2 antisense	AUCAGAAACAUCA AUGGGCCC
hNRF2 siRNA-3 sense	GCAGUCAAUGAAGCUCAATT
hNRF2 siRNA-3 antisense	UUGAGCUUCAUUGAACUGCTC
si-NC sense	GUCUACUGCUAUGUCUGUATT
si-NC antisense	AAUACAGACAUAGCAGUAGAC

Nrf2, nuclear factor erythroid 2-related factor 2; siRNA, small interfering RNA; NC, negative control.

Cell death assay. Pyroptotic cell death was assessed using a lactate dehydrogenase (LDH) release assay and Hoechst 33342/propidium iodide (PI) staining. For LDH release, the LDH activity of the cell culture supernatants was measured using an LDH assay kit (Nanjing Jiancheng Bioengineering Institute). For Hoechst 33342/PI double staining, the collected cells were washed with PBS and subsequently incubated with PI (5 μ l) for 10 min at 37°C in the dark. After three washes with PBS, the nuclei were stained with Hoechst 33342 (10 μ l) for 30 min at 37°C in the dark. Images were captured under a fluorescence microscope. The proportion of PI-positive cells was calculated using Image J software (version 1.46; National Institutes of Health).

Reverse transcription-quantitative (RT-q) PCR. The experimental method, which had been described in our previous study (4), was as follows: Total RNA was extracted from cells (1 \times 10⁶) using TRIzol® (Invitrogen; Thermo Fisher Scientific, Inc.) and quantified with a NanoDrop 2000 Spectrophotometer (Thermo Fisher Scientific, Inc.). Complementary DNA (cDNA) was synthesized with 1 μ g total RNA using a High-Capacity cDNA Reverse Transcription kit (Takara Bio, Inc.); the temperature protocol was: 42°C for 60 min, 70°C for 15 min, and then 4°C for 5 min. qPCR was then performed in a CFX96 Touch Real-Time PCR Detection System (Bio-Rad Laboratories, Inc.) using SYBR Premix (Shanghai Yeasen Biotechnology Co., Ltd.). The thermocycling conditions were as follows: Predenaturation step at 95°C for 3 min, followed by 40 cycles of 95°C for 10 sec and 60°C for 40 sec. Each sample was analyzed in triplicate. Table II lists the primer sequences used. The relative mRNA expression levels were normalized to β -actin expression using the 2^{- $\Delta\Delta$ C_q} method (18).

Western blotting. Total protein or nuclear protein (nucleopro-

tein extraction kit, Beijing Solarbio Science & Technology Co., Ltd.) was extracted from the collected HUVECs, and protein concentration was determined using a BCA Protein Assay Kit (cat. no. P0012A; Beyotime Institute of Biotechnology). Subsequently, equal amounts of protein lysates (30 μ g/lane) were separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis and transferred to polyvinylidene fluoride (PVDF) membranes (MilliporeSigma). After transfer, the PVDF membranes was incubated in 25 ml blocking buffer (1X Tris-buffered saline and 0.1% Tween-20 with 5% non-fat dry milk) for 1 h at room temperature and incubated overnight at 4°C with the following primary antibodies: β -actin (1:1,000; MDL; cat. no. MD6553), Histone H3 (1:1,000; Abcam; cat. no. ab1791), Nrf2 (1:1,000; CST; cat. no. 12721T), heme oxygenase-1 (HO-1; 1:1,000; Affinity; cat. no. AF5393), NQO1 (1:1,000; Novus; cat. no. NB200-209SS), superoxide dismutase 2 (SOD2; 1:1,000; Novus; cat. no. NB100-1992SS), NLRP3 (1:1,000; Abcam; cat. no. ab214185), ASC (1:1,000; ABclonal Biotech Co., Ltd.; cat. no. A1170), cleaved-caspase-1 (1:1,000; CST; cat. no. 4199T), GSDMD (1:1,000; Abcam; cat. no. ab210070), and cleaved N-terminal GSDMD (1:1,000; Abcam; cat. no. ab215203). The membranes were then incubated with HRP-linked anti-mouse IgG (cat. no. 7076; 1:3,000; Cell Signaling Technology, Inc.) or HRP-linked anti-rabbit IgG (cat. no. 7074; 1:3,000; Cell Signaling Technology, Inc.) for 1 h at room temperature. The antibody-antigen complexes were detected using enhanced electrochemiluminescence reagents (Santa Cruz Biotechnology, Inc.) and densitometric analysis was conducted using Image J software (version 1.46; National Institutes of Health).

Statistical analysis. All data were analyzed using GraphPad Prism version 9.0 (Dotmatics). All experimental data are represented as the means \pm standard deviation. The Shapiro-Wilk test was used to evaluate Gaussian distributions, while the Brown-Forsythe test was used to evaluate homogeneity of variance. All data fitted the Gaussian distribution and the variances were equal. One-way ANOVA followed by Šídák's multiple comparisons test was used to compare differences between three or more groups. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Melatonin activates the Nrf2 pathway in HUVECs under HG conditions. In this part of the experiment, an optimal melatonin protective concentration was first determined through CCK-8 by setting control group, HG (30 mM) + different melatonin concentrations (0 μ M-1 M) treatment group. It was found that a melatonin concentration of 100 μ M was the most effective in alleviating the loss of cell viability caused by HG and 100 μ M was selected as the treatment concentration for subsequent experiments (Fig. 1A). Next, in order to select an appropriate time for HG intervention, HUVECs were incubated with 30 mM HG medium and the expression of Nrf2 protein in the nucleus of HUVECs was observed by western blotting with prolonged HG intervention time. It was found that the expression of n-Nrf2 increased to the highest level after 6 h of HG incubation and then gradually decreased as HG incubation time was prolonged (Fig. 1B and C). In order to observe more marked therapeutic effect of melatonin in the follow-up experiments,

Table II. Primer sequences.

Gene	Forward (5'-3')	Reverse (5'-3')
β -actin	TCCTCCTGAGCGCAAGTACTCC	CATACTCCTGCTTGCTGATCCAC
Nrf2	TCAGCGACGGAAAGAGTATGA	CCACTGGTTTCTGACTGGATGT
NLRP3	AAGGAAGTGGACTGCGAGAA	AACGTTCGTCCTTCCTTCCT
Caspase-1	GGCATGACAATGCTGCTACA	TCTGGGACTTGCTCAGAGTG

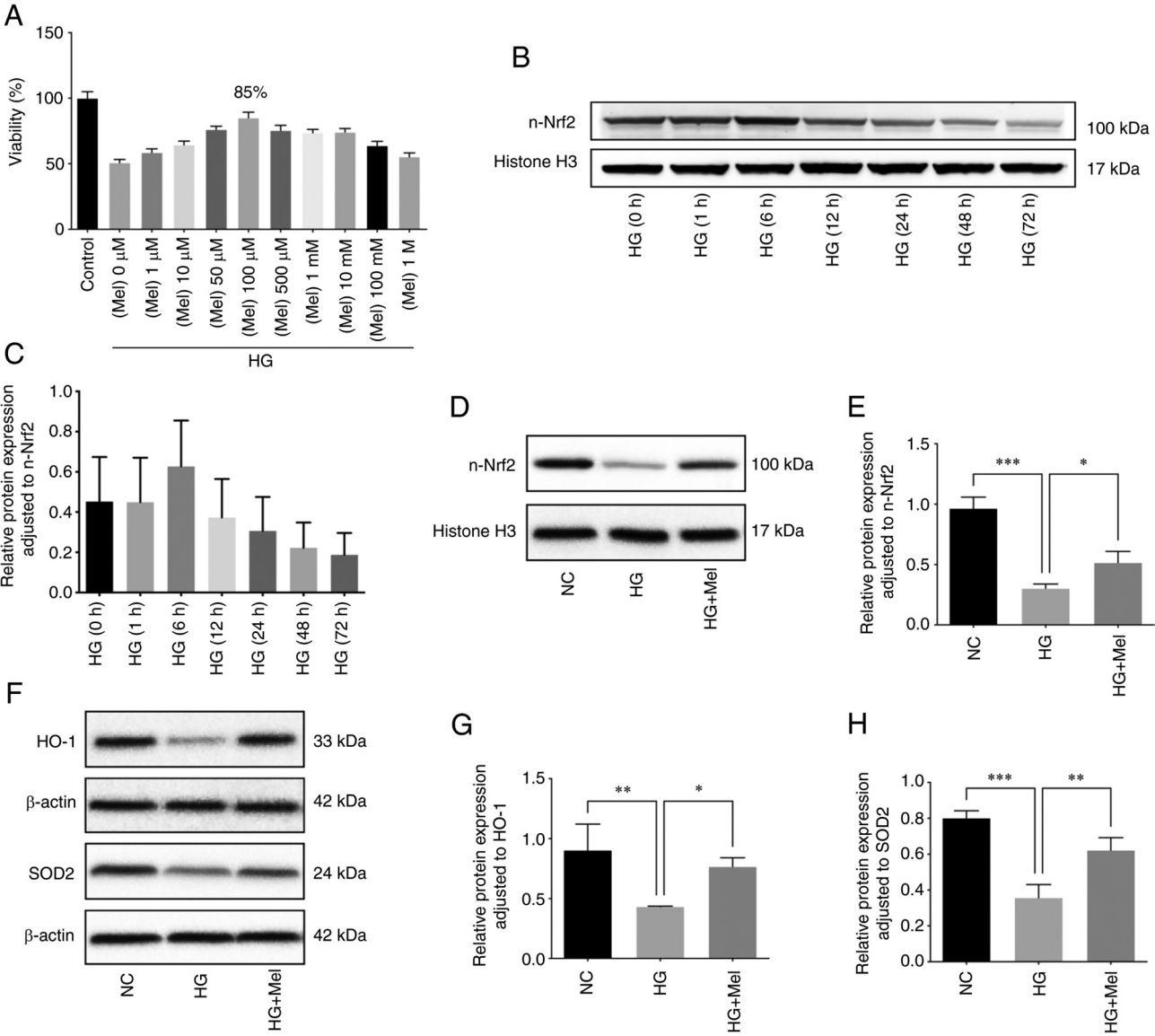


Figure 1. Mel activates the Nrf2 pathway in HUVECs under HG conditions. (A) CCK-8 cell viability test confirmed that the optimal protective concentration of melatonin was 100 μ M. (B) Western blot analysis of n-Nrf2 expression in HUVECs at different HG incubation time. (C) Semi-quantitative analysis of n-Nrf2 relative expression. (D) Western blot analysis of n-Nrf2 expression in HUVECs of each group. (E) Semi-quantitative analysis of n-Nrf2 relative expression. (F) Western blot analysis of HO-1 and SOD2 expression in the HUVECs of each group. Semi-quantitative analysis of (G) HO-1 and (H) SOD2 relative expression. Results are presented as the mean \pm standard deviation. * P <0.05, ** P <0.01 and *** P <0.001. Mel, melatonin; Nrf2, nuclear factor erythroid 2-related factor 2; HUVECs, human umbilical vein endothelial cells; HG, high glucose; NC, negative control; HO-1, heme oxygenase-1; and SOD2, superoxide dismutase 2.

72 h was chosen as the intervention time for HG. Finally, it was confirmed by western blotting that melatonin treatment could activate Nrf2 signaling pathway in HUVECs under HG condition. The result showed that melatonin promoted the expression of n-Nrf2, HO-1 and SOD2 (Fig. 1D-H).

Melatonin activates the Nrf2 pathway in HUVECs mainly through the MT1/MT2 pathway under HG conditions. Luzindole is an effective competitive MT1/MT2 membrane receptor antagonist and is widely used in melatonin research. In this part of the experiment, it was found that the protective effect

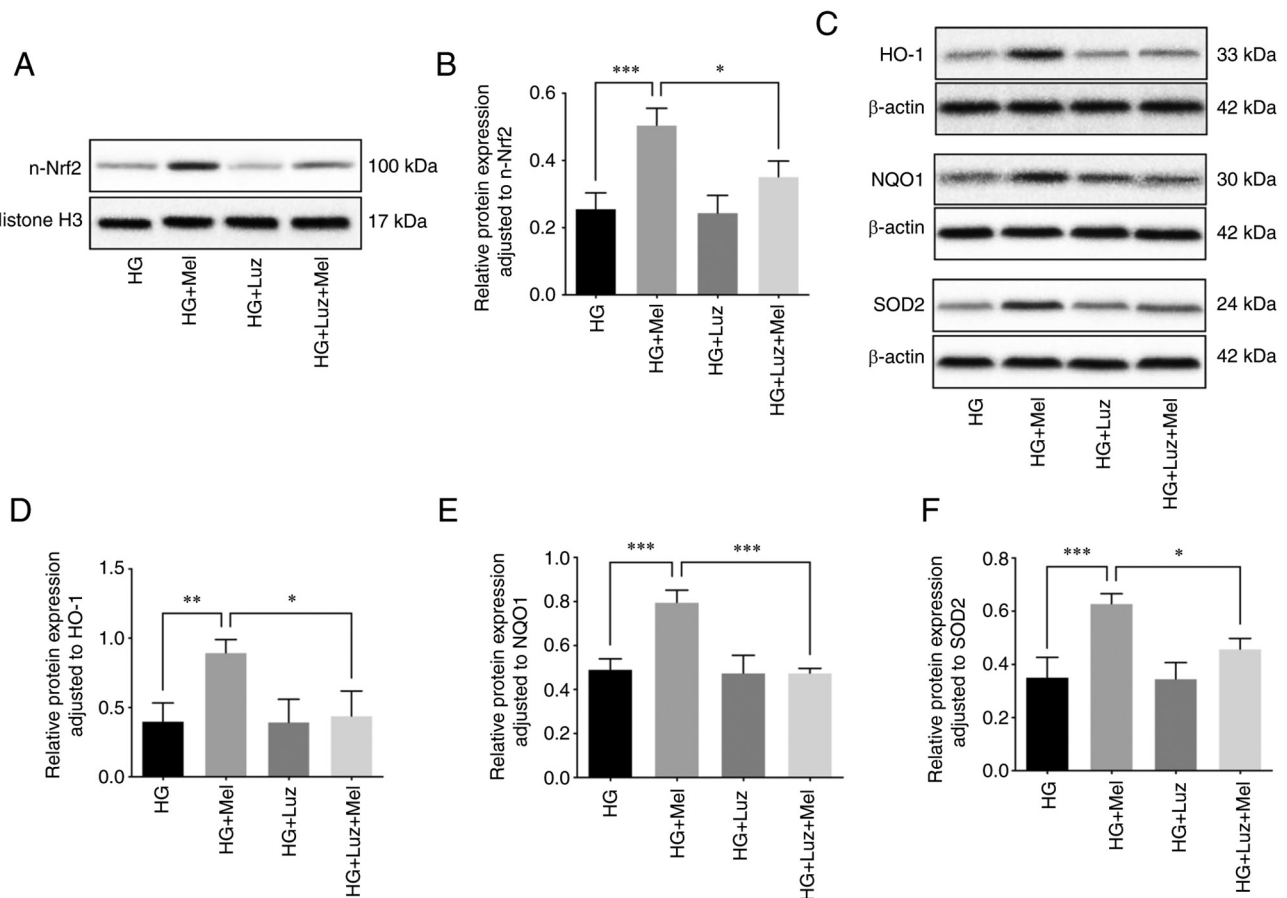


Figure 2. Mel activates the Nrf2 pathway in HUVECs mainly through the MT1/MT2 pathway under HG conditions. (A) Western blot analysis of n-Nrf2 expression in HUVECs of each group. (B) Semi-quantitative analysis of n-Nrf2 relative expression. (C) Western blot analysis of HO-1, NQO1 and SOD2 expression in HUVECs of each group. (D-F). Semi-quantitative analysis of HO-1, NQO1 and SOD2 relative expression. Results are presented as the mean \pm standard deviation. * P <0.05, ** P <0.01 and *** P <0.001. Mel, melatonin; Nrf2, nuclear factor erythroid 2-related factor 2; HUVECs, human umbilical vein endothelial cells; HG, high glucose; HO-1, heme oxygenase-1; NQO1, oxygenase-1 (HO-1), NADPH: Quinone Oxidoreductase 1; SOD2, superoxide dismutase 2; Luz, Luzindole.

of melatonin in promoting n-Nrf2 expression was significantly reduced when MT1/MT2 was blocked by Luzindole under HG conditions (Fig. 2A and B). Similarly, the effect of melatonin on promoting the expression of HO-1, NQO1 and SOD2 proteins under HG conditions was also significantly reduced when MT1/MT2 was blocked by Luzindole (Fig. 2C-F).

Melatonin or MCC950 inhibits HG-induced endothelial cell pyroptosis. Hoechst 33342/PI staining and LDH release assay are often used to observe pyroptotic cell death (19,20). In this part of the experiment, it was found that the proportion of PI-positive cells and LDH release of HUVEC in the HG group were significantly increased compared with the control group. However, following treatment with the NLRP3 specific inhibitor MCC950 and melatonin, HG-mediated pyroptosis was significantly reduced (Fig. 3A-C). Then the protein expression levels of Gasdermin D (GSDMD), a key executioner of the pyroptotic pathway and the N-terminal GSDMD cleavage product (GSDMD-N), which can induce pyroptosis was examined. Western blotting results showed that GSDMD and GSDMD-N were significantly increased in HG-incubated HUVEC compared with the control group. However, following treatment with MCC950 and melatonin, HG-mediated increases in GSDMD and GSDMD-N were significantly reduced (Fig. 3D-F).

Melatonin inhibits ROS/NLRP3 inflammasome pathway in HUVECs under HG conditions by activating the Nrf2 pathway. In this part of the experiment, in order to confirm that melatonin inhibits the activation of ROS/NLRP3 inflammasome pathway in vascular endothelial cells induced by HG mainly through or at least partly through Nrf2 signaling pathway, Nrf2 expression was knocked down by siRNA interference, which was verified by RT-qPCR. hNrf2 siRNA-2 was selected as the interference sequence (Fig. 4A). Next, western blotting results showed that melatonin treatment could significantly reduce the overexpression of NLRP3, ASC and cleaved-caspase-1 protein induced by HG, and the protective effect of melatonin was also significantly decreased after knockdown of Nrf2 expression (Fig. 4B-E). In addition, melatonin treatment could significantly attenuate the increase of NLRP3 mRNA and caspase-1 mRNA expression induced by HG, while the protective effect of melatonin was significantly reduced after Nrf2 knockdown (Fig. 4F and G). It was also found by ELISA that melatonin treatment significantly attenuated the increased secretion of IL-1 β and IL-18 in culture supernatants induced by HG, and this protective effect of melatonin was also significantly decreased after knockdown of Nrf2 expression (Fig. 4H and I). Finally, ROS, a key activator upstream of the NLRP3 inflammasome was measured. Results showed that melatonin treatment significantly attenuated the increase of

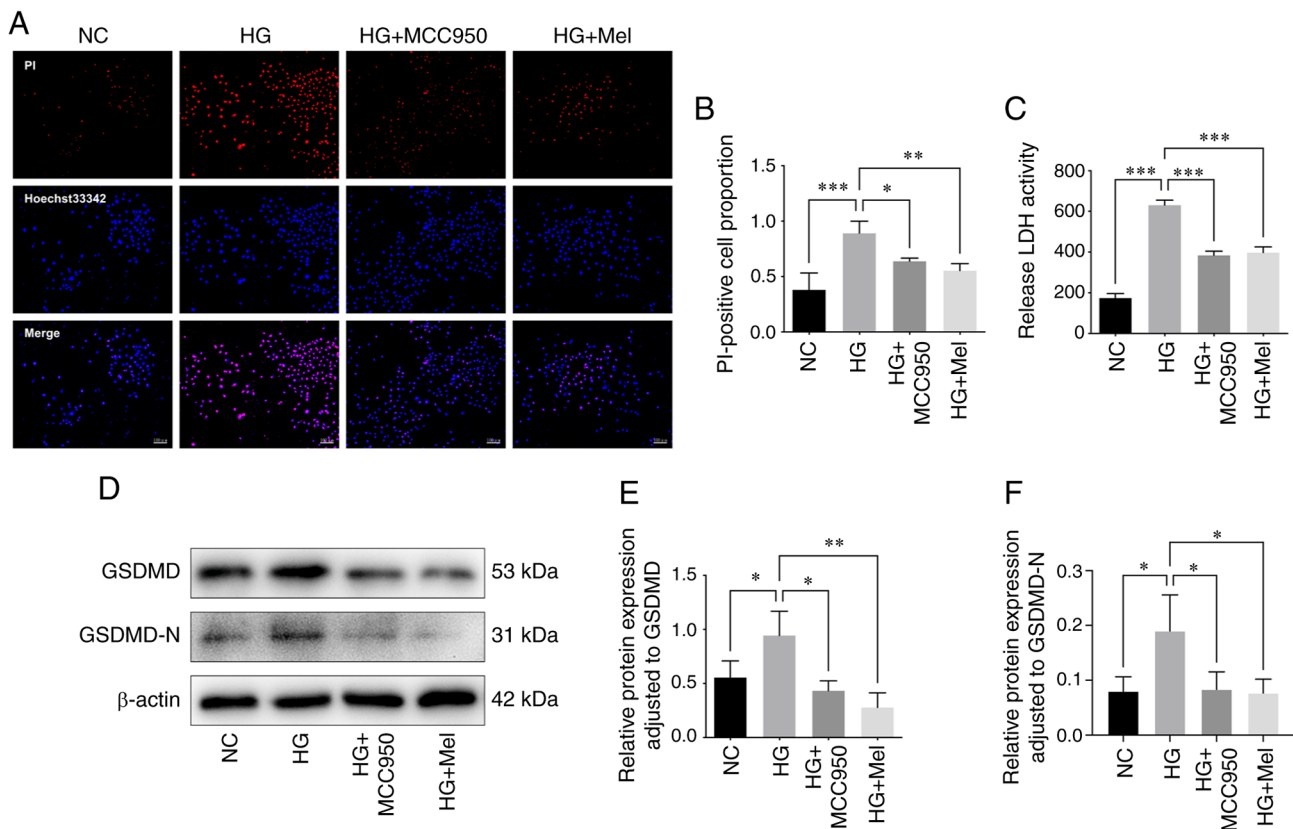


Figure 3. Mel or MCC950 inhibits HG-induced endothelial cell pyroptosis. (A) Representative pyroptotic cell death images of each group obtained using Hoechst33342/PI staining (original magnification, $\times 100$; scale bar, $100\ \mu\text{m}$). (B) The proportion of PI-positive cells in each group. (C) LDH release activity in each group. (D) Western blot analysis of GSDMD and GSDMD-N expression in HUVECs of each group. Semi-quantitative analysis of (E) GSDMD and (F) GSDMD-N relative expression. Results are presented as the mean \pm standard deviation. * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$. Mel, melatonin; HG, high glucose; PI, propidium iodide; LDH, lactate dehydrogenase; GSDMD, gasdermin D; GSDMD-N, gasdermin D-N; HUVECs, human umbilical vein endothelial cells; NC, negative control.

ROS production induced by HG, and this protective effect of melatonin was also significantly decreased after knockdown of Nrf2 expression (Fig. 4J).

Discussion

The present study found that melatonin inhibited HG-induced pyroptosis of vascular endothelial cells by activating the Nrf2 pathway to inhibit NLRP3 inflammasome activation. It was further found that melatonin protects vascular endothelial cells from HG-induced injury through the activation of Nrf2 pathway, which is mediated by MT1/MT2.

Previous studies have found that melatonin is beneficial for a variety of cardiovascular diseases, such as ischemia-reperfusion injury (21), hypertension (22), and atherosclerosis (23). However, some clinical studies have come to different conclusions. For example, several randomized controlled trials have found benefits of melatonin in heart failure (24), ST segment elevation myocardial infarction (STEMI) (25), coronary artery bypass grafting (26), and percutaneous coronary intervention (27). However, some randomized controlled trials have found no significant positive effect of melatonin in STEMI (28), acute coronary syndromes (29) and macrovascular surgery (30) despite its safety. Therefore, more experiments are needed to further evaluate the underlying molecular mechanisms of melatonin activity.

The current study, found through CCK-8 experiment under the condition of high glucose of $30\ \text{mM}$ that the therapeutic concentration of melatonin of $100\ \mu\text{M}$ could minimize the adverse effects of high glucose on the vitality of HUVECs. However, higher concentrations of melatonin showed similar or even reduced protective effects. In fact, one of our previous studies found that a therapeutic concentration of $100\ \mu\text{M}$ of melatonin was optimal for alleviating the adverse effects of cigarette smoke extractants on the vitality of HUVECs (11). In addition, in a number of studies on the protection of vascular endothelial cells with melatonin, $100\ \mu\text{M}$ is used as the therapeutic concentration of melatonin and it showed an ideal cellular protective effect (31,32). Therefore, the present study set the concentration gradient of melatonin treatment at around $100\ \mu\text{M}$ and found the aforementioned results. As for why melatonin showed similar or even reduced protective effect with higher concentration under the experimental conditions of the present study, it was hypothesized that it might be related to the pharmacological properties of melatonin itself, which still needs to be verified in further studies.

Endothelial cell pyroptosis is involved in the pathogenesis of atherosclerosis (19,33). Studies have found that melatonin reduces endothelial cell pyroptosis. For example, a recent study revealed that melatonin prevents endothelial cell pyroptosis by upregulating TET2 to inhibit the methylation of UQCRC1 and improving mitochondrial function (34). Zhang *et al* (20)

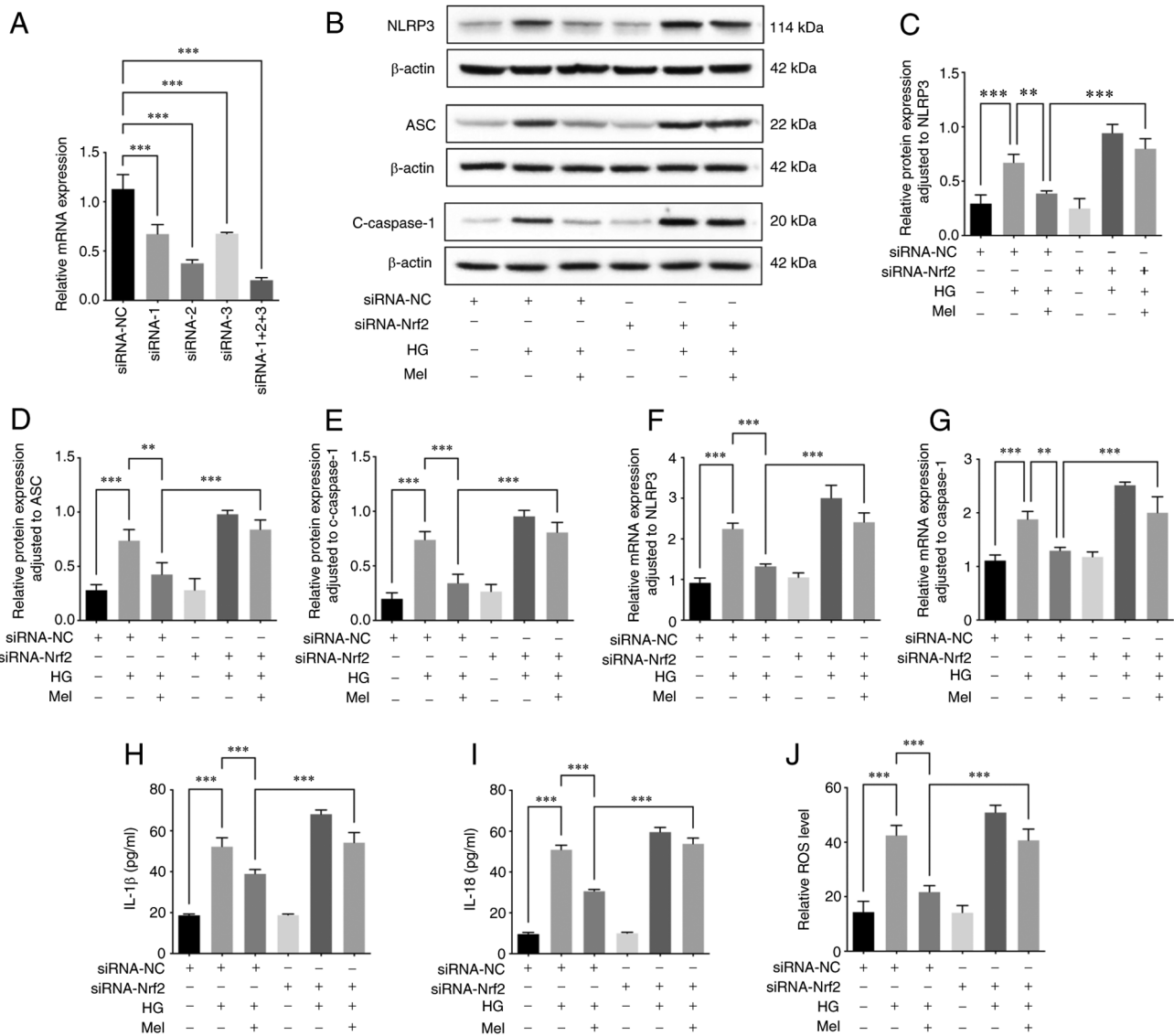


Figure 4. Mel inhibits ROS/NLRP3 inflammasome pathway in HUVECs under HG conditions by activating the Nrf2 pathway. (A) Nrf2 knockdown was verified by reverse transcription-quantitative PCR. (B) Western blot analysis of NLRP3, ASC and cleaved-caspase-1 expression in HUVECs of each group. Semi-quantitative analysis of (C) NLRP3, (D) ASC and (E) cleaved-caspase-1 relative expression. Expression levels of (F) NLRP3 and (G), caspase-1 mRNA in HUVECs of each group; H-I. The levels of (H) IL-1β and (I) IL-18 in the supernatant of HUVEC cultures in each group. (J) The levels of ROS in HUVECs of each group. Results are presented as the mean ± standard deviation. **P<0.01 and ***P<0.001. Mel, melatonin; ROS, reactive oxygen species; NLRP3, pyrin domain-containing 3; HUVECs, human umbilical vein endothelial cells; HG, high glucose; Nrf2, nuclear factor erythroid 2-related factor 2; NC, negative control.

reported that melatonin prevents endothelial cell pyroptosis via MEG3/miR-223/NLRP3 axis in atherosclerosis. Our previous study has also demonstrated that melatonin alleviates cigarette smoke-induced endothelial cell pyroptosis through inhibiting ROS/NLRP3 axis (11). However, the role of melatonin in diabetes-induced pyroptosis of vascular endothelial cells is rarely reported. The current study has clarified the protective role of melatonin in HG-induced pyroptosis of vascular endothelial cells.

Nrf2 is a key nuclear transcription factor that regulates antioxidant gene expression. Studies (35,36) have reported that during the early stages of diabetes, the expression of endothelial nitric oxide synthase (eNOS) and antioxidant genes may be upregulated as a compensatory mechanism in response to ROS/reactive nitrogen species, however, as diabetes progresses, NADPH oxidase continues to be activated and the continued

overexpression of ROS will lead to eNOS uncoupling, mitochondrial dysfunction and decreased expression of antioxidant genes mediated by Nrf2. Finally, the cells enter a vicious circle of antioxidant imbalance. Consistent with the results of previous studies, the present study found that HUVEC initially showed an increased compensatory expression of Nrf2 under the intervention of high glucose, but with the extension of high glucose injury time, Nrf2 expression also gradually decreased.

ROS is a key regulator of inflammasome activation (37,38). Our previous study also confirmed that cigarette smoke causes endothelial pyroptosis through the ROS/NLRP3 axis (11). Therefore, it is particularly important to find suitable ROS inhibitors to reduce diabetes-induced vascular injury associated with NLRP3 inflammasome. Increasing evidence has shown that the Nrf2 signaling pathway interacts with inflammasomes at multiple points. A recent study revealed

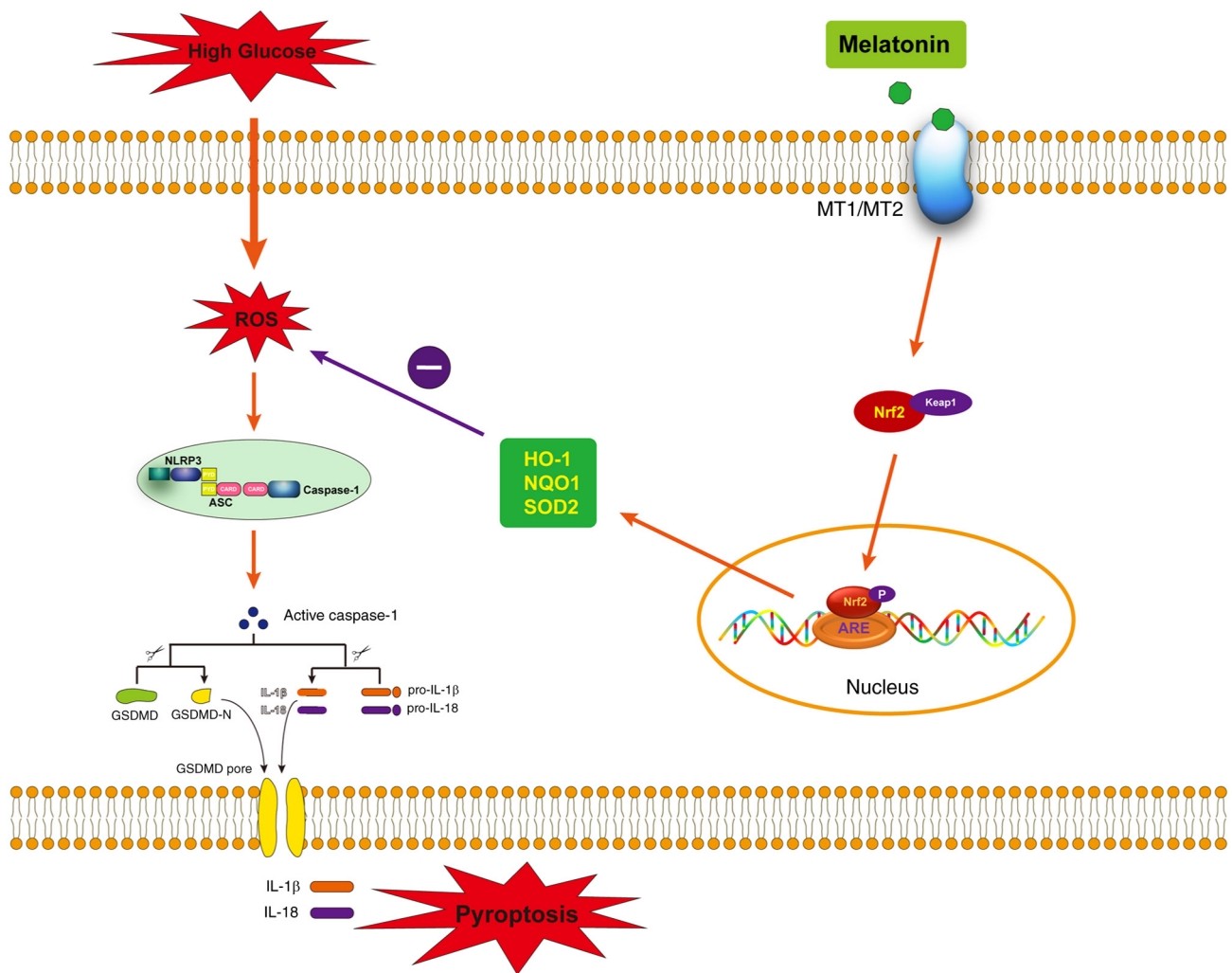


Figure 5. A schematic hypothetical model showing the specific signaling mechanisms by which melatonin attenuates HG-induced endothelial cell pyroptosis through MT1/MT2-Nrf2-ROS-NLRP3 signaling pathway. HG, high glucose; Nrf2, nuclear factor erythroid 2-related factor 2; ROS, reactive oxygen species; NLRP3, NOD-like receptor family, pyrin domain-containing 3; HO-1, heme oxygenase-1; NQO1, oxygenase-1 (HO-1), NADPH: Quinone Oxidoreductase 1; SOD2, superoxide dismutase 2; Keap1, kelch-like ECH-associated protein 1.

that in renal ischemia/reperfusion injury, Nrf2 regulates the inhibition of NLRP3 inflammasome activity (39). In addition, Hou *et al* (40) reported that Nrf2 inhibits NLRP3 inflammasome activity through regulating Trx1/TXNIP complex in cerebral ischemia reperfusion injury. However, only a few studies have evaluated the interaction between Nrf2 and NLRP3 inflammasomes in diabetes-induced vascular injury. Previous studies have shown that melatonin may serve a therapeutic role in different diseases, such as ischemia-reperfusion injury (41), oxidative stress injury (42) and neurotoxicity (43), through activation of Nrf2 signaling pathway. Previous studies have found that melatonin can exert anti-inflammatory effects through Nrf2-NLRP3 pathway. One study found that melatonin attenuates LPS-induced acute depression-like behavior and NLRP3 inflammasome activity in microglia via the SIRT1/Nrf2 pathway (44). Another study found that melatonin alleviates sepsis-induced cardiac injury by activating Nrf2 pathway and inhibiting NLRP3 inflammasome (45). In addition, our previous study also found that melatonin may effectively protect smoking-caused vascular injury and atherosclerosis through Nrf2/ROS/NLRP3 signaling pathway (46).

However, it remains to be clarified whether melatonin serves a protective role through Nrf2/NLRP3 pathway in vascular endothelial injury induced by HG. Considering different disease states, the effect and mechanism of drugs may change. The present study confirmed that melatonin alleviated HG-induced endothelial cell pyroptosis by activating the Nrf2 pathway to inhibit NLRP3 inflammasome activation.

Studies have shown that melatonin activates the Nrf2 pathway through various other pathways. Chen *et al* (47) found that melatonin protects murine cortical astrocytes from hemin-induced toxicity through activation of protein kinase C α /Nrf2 pathway. It has also been reported that SIRT1-dependent melatonin activation of Nrf2 pathway serves a neuroprotective role in BV2 microglia (42). Melatonin exerts its cytoprotective effects through receptor and non-receptor pathways. The present study found that melatonin may activate Nrf2 pathway through MT1/MT2 pathway. Similar to this result, a previous study found that melatonin serves a protective role in methamphetamine-induced rat brain microvascular endothelial cell inflammation by interacting with its receptors (MT1/MT2)

to promote Nrf2 signaling and inhibit NF- κ B signaling (48). However, the mechanistic basis of melatonin-mediated Nrf2 pathway activation in diabetic vascular injury needs to be further investigated.

MT1/MT2 is a G-protein-coupled receptor existing in cell membrane and widely exists in the nervous and cardiovascular systems (49,50). Cui *et al* confirmed the normal expression of MT1/MT2 in HUVEC by western blotting (51). Therefore, the current study did not conduct experiments to verify it again. However, the lack of verification of MT1/MT2 expression in HUVECs remains a potential limitation of the present study. As for the model of vascular endothelial cells with high glucose injury, the glucose concentrations in most previous studies were controlled within the range of 20-40 mM, among which 25 and 30 mM are the most used at present (52-55) because they are closer to the blood glucose values of severe hyperglycemia in humans. In addition, in order to observe pyroptosis and the protective effect of melatonin more markedly, the present study selected a glucose concentration of 30 mM. However, the lack of an addition concentration to examine any dose-dependent effect is also a potential limitation of the present study.

In conclusion, the current study showed that melatonin may serve a protective role in HG-induced vascular endothelial cell pyroptosis by activating the Nrf2 pathway to inhibit NLRP3 inflammasome activation. In addition, it was further found that melatonin attenuates HG-induced vascular endothelial cell injury by interacting with its receptors (MT1/MT2) to promote activation of Nrf2 pathway (Fig. 5). These findings provide a new theoretical basis and a promising therapeutic strategy for the early prevention and treatment of diabetic vasculopathy.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

XW, HF and CL conceived and designed the present study. XW, RZ and BM performed the experiments. XW, WW and RZ interpreted the results, analyzed the data and wrote the paper. XW, WW and LN analyzed the data and designed the figures and table. HF and CL reviewed and edited the manuscript. XW, HF and CL confirmed the authenticity of all the raw data. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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