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Curcumin nanocrystals ameliorate ferroptosis of diabetic nephropathy through glutathione peroxidase 4

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Objective: The aim of this study was to investigate the effect of curcumin nanocrystals (Cur-NCs) on ferroptosis in high-glucose (HG)-induced HK-2 cells and streptozotocin (STZ)-induced diabetic nephropathy model (DN) rats. The purpose is to determine whether Cur NCs can become a promising treatment option for diabetes nephropathy by reducing ferroptosis.

Methods: Cur-NCs were prepared using microfluidic technology and studied using dynamic light scattering and transmission electron microscopy. HK-2 cells were treated with 30 mM HG to create a renal tubule damage cell model. Then, cell viability was evaluated in HK-2 cells treated with varying concentrations of Cur-NCs (0.23, 0.47, 0.94, 1.87, 3.75, 7.5, 15, and 30 $\mu\text{g}/\text{mL}$) using Cell Counting Kit-8 (CCK-8). Furthermore, *in vivo* experiments were carried out to investigate the roles of Cur-NCs in STZ-induced DN rats.

Results: The results showed that HG treatment greatly enhanced the levels of LDH, MDA, Iron, lipid ROS, apoptosis, NCOA4, TFR-1, while decreasing the expression of GSH, GPX4, SLC7A11, and FTH-1. These effects induced by HG could be attenuated by Cur-NCs. Cur-NCs also reduced the HG-induced decrease in cell viability, as well as the increase in lipid ROS and cell apoptosis, however erastin could inhibit their effects. Furthermore, the *in vivo* results showed that Cur-NCs reduced ferroptosis and inhibited renal damage in DN rats.

Conclusion: This study demonstrates that Cur-NCs can significantly attenuate ferroptosis in a STZ-induced renal damage model by recovering GPX4, implying that Cur-NCs may be a promising therapy option for DN.

KEYWORDS

curcumin, nanocrystals, ferroptosis, diabetic nephropathy, GPX4

Introduction

The primary cause of end-stage renal disease (ESRD) worldwide is diabetes nephropathy (DN), which not only seriously compromises patient quality of life but also places a significant financial and social strain on the world's healthcare system (Samsu, 2021; Hu et al., 2023). Its prevalence rate is rising annually, particularly in the population with a high incidence of diabetes, making it an urgent public health concern. Worldwide, an estimated 537 million people have been diagnosed with diabetes (Sun et al., 2022). Up to 30%–40% of people in these sizable diabetes patient groups are predicted to develop DN (Selby and Taal, 2020). The pathogenesis of DN is complex and multifaceted, including multiple factors such as metabolic abnormalities, hemodynamic disorders, and inflammatory responses (Lin et al., 2018; Chen et al., 2022a). DN is primarily caused by chronic hyperglycemia, which sets off a cascade of events in the kidneys that include aberrant activation of protein kinase C (PKC) subtypes, build-up of advanced glycation end products (AGEs), and excessive production of reactive oxygen species (ROS) (Tanase et al., 2022; Feng et al., 2023). These molecular level changes gradually erode kidney health, leading to dysfunction of renal endothelial cells, apoptosis of podocytes, mesangial dilation, ultimately resulting in glomerulosclerosis and tubulointerstitial fibrosis (Tanase et al., 2022; Yao et al., 2022). At the same time, renal hemodynamic dysregulation characterized by glomerular filtration and hypertension further exacerbates kidney damage (Tuttle et al., 2022). Furthermore, the kidneys' inflammatory response and fibrosis process are made worse by the involvement of several pro-inflammatory cytokines, including interleukin (IL), tumor necrosis factor- α (TNF- α), and transforming growth factor- β (TGF- β) (Wang et al., 2021b; Ahmed et al., 2022). While significant progress has been made in exploring the pathophysiological mechanisms of DN, there is still a dearth of efficient treatments to slow the disease's progression. This necessitates further research to fully understand its complex mechanisms and create innovative therapies that can effectively intervene and delay the devastating effects of DN.

In recent years, the growing idea of ferroptosis, a new type of controlled cell death marked by iron-dependent lipid peroxidation, has received a lot of attention in the field of DN (Li et al., 2023b; Mengstie et al., 2023). The disorders of iron homeostasis and lipid metabolism are the key driving factors of DN pathogenesis, which would further trigger and exacerbate ferroptosis. In various studies, quite a number of molecular pathways have been confirmed to be implicated in it, including the iron-regulatory proteins, the phospholipid metabolism pathway involving the enzyme acyl-CoA synthetase long-chain family member 4 (ACSL4), and the glutathione peroxidase 4 (GPX4)-regulated antioxidant defense system (Li et al., 2022a; Wu et al., 2023). Furthermore, diabetic conditions promote oxidative stress and inflammation, which may amplify ferroptotic cell death in renal cells (Shen et al., 2022). Thus, investigating the function of ferroptosis in DN is expected to provide new therapeutic targets for minimizing renal damage and maintaining renal function in diabetics. However, further research into the specific biochemical processes involved and the development of appropriate therapeutics are still needed to fully realize the potential of addressing ferroptosis in the treatment of diabetic nephropathy.

Traditional Chinese medicines (TCM) provide a holistic approach to attenuate renal damage and delay disease progression with antioxidant, anti-inflammatory and antifibrotic properties (Chen et al., 2023; Liang et al., 2024). For example, salvinorin B and tanshinone IIA synergistically attenuate early DN by upregulating the PI3K/Akt pathway and downregulating the NF- κ B signaling pathway (Xu et al., 2024b). In addition, mullein isoflavone extracts have anti-ferroptosis sagging effects by downregulating lipid ROS in DN (Huang et al., 2022). Ginkgolide B ameliorates DN oxidative stress and ferroptosis by inhibiting GPX4 ubiquitination (Chen et al., 2022b). Therefore, researchers are increasingly focusing on ferroptosis-suppressive drugs as they show good potential to treat human diseases.

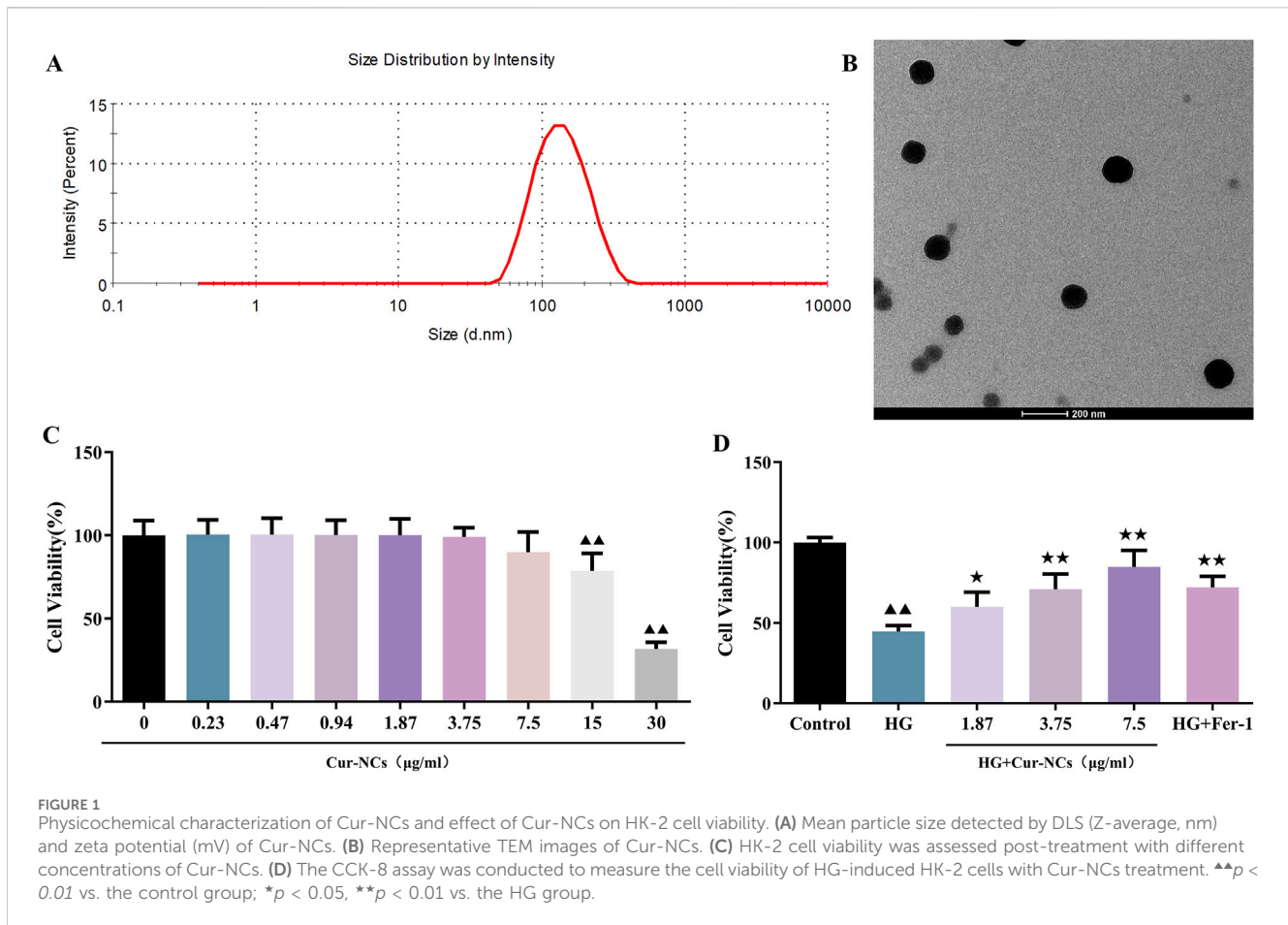
In contemporary scientific research, the exploration of novel therapeutic approaches for DN has led to a growing interest in the application of curcumin as a potential treatment modality (Zhu et al., 2022). A naturally occurring polyphenol chemical, curcumin, is extracted from the rhizome of *Curcuma longa* L. With its wide range of pharmacological characteristics, such as its anti-inflammatory, anti-fibrosis, and antioxidant activities, curcumin has a lot of promise for DN intervention (Wang et al., 2023; Yang et al., 2024). Accumulating evidence has shown that curcumin attenuated apoptosis of podocyte cells and accelerated cell autophagy in DN via regulating Beclin1/UVRAG/Bcl2 pathway (Zhang et al., 2020). Curcumin also decreases oxidative stress by keeping the Nrf2 pathway in a steady state, dramatically improving retinal damage in diabetes (Xie et al., 2021). Furthermore, curcumin can suppress ERK/JNK phosphorylation in insulin resistance cells generated by high glucose (HG) and promote the PI3K-AKT-GSK3B signaling pathway, hence increasing insulin sensitivity, providing a novel avenue for targeted diabetes treatment (Hussain et al., 2022). These results suggest that curcumin could be a new therapy option for diabetic nephropathy.

Curcumin's fast metabolism, low bioavailability, and poor water solubility, however, make it difficult to translate clinically (Chen et al., 2022b). Fortunately, the rapid advancement of nanotechnology has created more opportunities for the clinical application of curcumin. Curcumin nanocrystals (Cur-NCs) meticulously created by nanotechnology not only considerably improve the solubility and bioavailability of the medicine (Wang et al., 2021a), but also lengthen its retention time in the body, significantly increasing the therapeutic effect (Huang et al., 2023; Wang et al., 2023). This novel achievement has broadened the application possibilities for Cur-NCs in the treatment of DN. However, our current exploring of Cur-NCs is insufficient, necessitating more in-depth studies *in vitro* and *in vivo* to further elucidate their potential mechanism of action, optimize the formulation parameters of their clinical translation, so that they can function as effectively as possible in therapeutic practice.

Materials and methods

Preparation of Cur-NCs

A total of 40 mg of curcumin (HXJHS20200710, purity: 98%, Xi'an Haoxuan Biotech Co., Ltd., China) was dissolved in 4 mL of acetone to form a 10 mg/mL solution and then passed through



0.22 μm microporous filter to obtain the organic phase solution. Polyvinylpyrrolidone K30 (PVP-K30, 50 mg, Shanghai yuanye Bio-Technology Co., Ltd., China) was accurately weighed and dissolved in 10 mL of water to obtain the aqueous phase solution. The organic and aqueous phase solutions were fed into a stainless steel microfluidic chip with a split-and-recombine (SAR) structure by MPE-L2 microfluidic instrument (Suzhou Aitesen Pharmaceutical Equipment CO., Ltd., China) through a syringe pump, and the flow rate of the two phases was controlled by a syringe pump connected to the system at 9 and 1 mL/min, respectively. After the system was stabilized, the curcumin suspension was collected with a beaker at the outlet and freeze-dried to obtain Cur-NCs powder.

Particle size, Zeta potential, and morphology

The particle size and zeta potential of Cur-NCs (Zheng et al., 2024) were detected in purified water by Zetasizer Nano ZS (Malvern Instruments, United States). For morphological examination, Cur-NCs were mounted on 300-mesh Cu grids and negatively stained with 1% uranyl acetate for 2 min. Transmission electron microscope (TEM) images were taken using a Talo L120C G2 TEM (United States) operating at 200 keV. The average length and width of 200 randomly selected Cur-NCs were identified from the TEM images using ImageJ software (v1.53, NIH, United States).

Cell culture and treatment

Human renal proximal tubular epithelial cells (HK-2) were obtained from the iCell Bioscience Inc (Shanghai, China). The cells were cultured in Dulbecco's Modified Eagle Medium/Nutrient Mixture F-12 (11330032, Gibco, United States) supplemented with 10% fetal bovine serum (A5669701, Gibco, United States) in a humidified incubator at 37°C with 5% CO₂. To induce HG conditions, HK-2 cells were treated with 30 mM D-glucose, while the normal glucose (control) group was exposed to 5.5 mM D-glucose and 24.5 mM mannitol for 48 h. HK-2 cells were exposed to control or HG, with or without Cur-NCs (1.87, 3.75, 7.5 $\mu\text{g}/\text{mL}$), Ferrostatin-1 (1 μM , S81461, Shanghai yuanye Bio-Technology Co., Ltd., China), Erastin (35 μM , S80804, Shanghai yuanye Bio-Technology Co., Ltd., China) treatment alone or in combination for the specified time (0 h, 12 h, 24 h, or 48 h).

siRNA transfection

When HK-2 cells were seeded in 6-well plates at 80% confluence. Then, Lipofectamine™ 3,000 (L3000015, Invitrogen, United States) was diluted with Opti-MEM medium and mixed thoroughly. The short interfering RNAs (siRNAs) used were siRNA_GPX4_NC (5'-UUCUCC GAACGUGUCACGUTT-3'), siRNA_GPX4_#1 (5'-ACCAAGTTT GGACACCGTCTCTCCA-3'), siRNA_GPX4_#2 (5'-CAAGTTTGG ACACCGTCTCTCCACA-3'), and siRNA_GPX4_#3 (5'-ACACCG

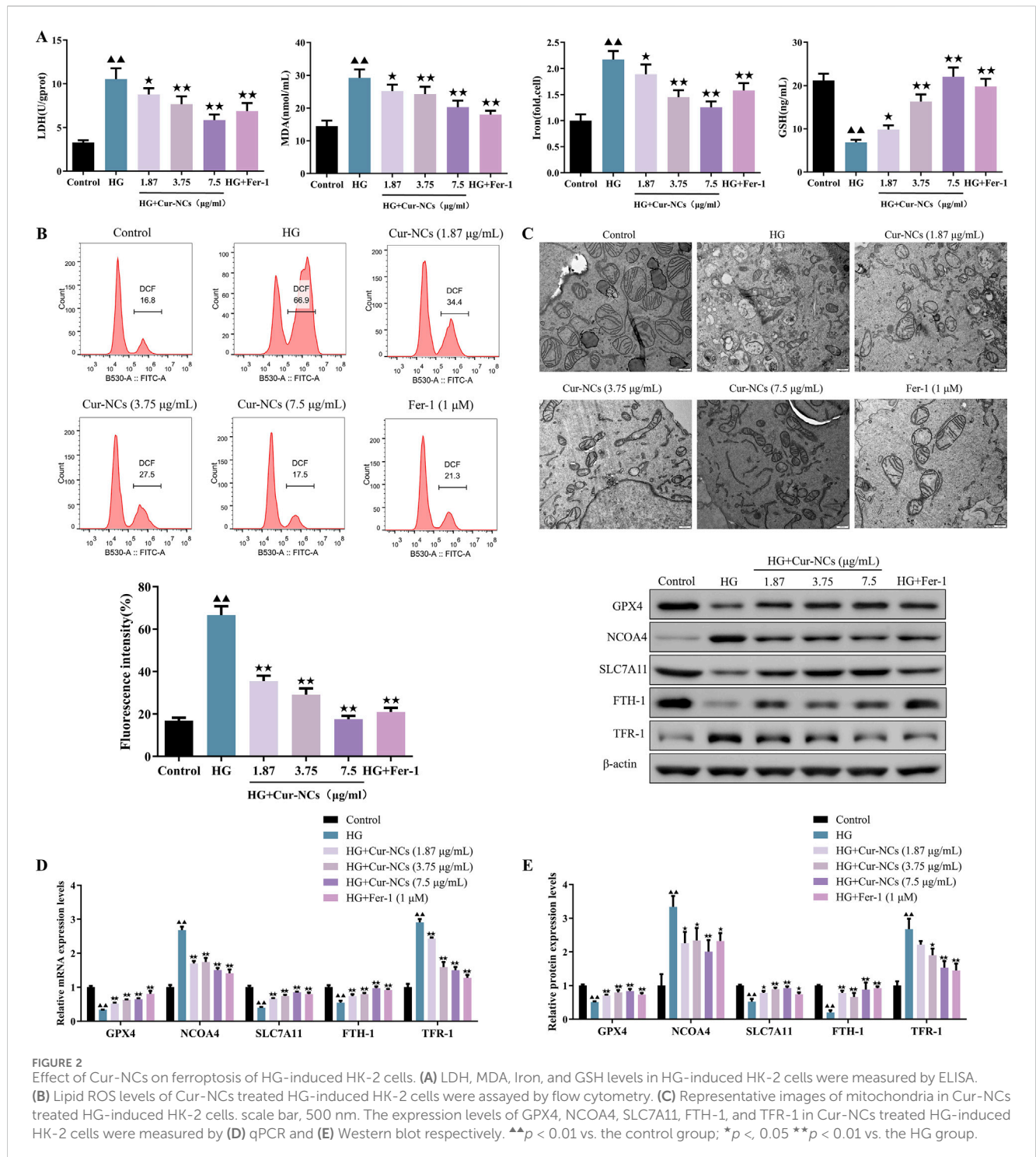


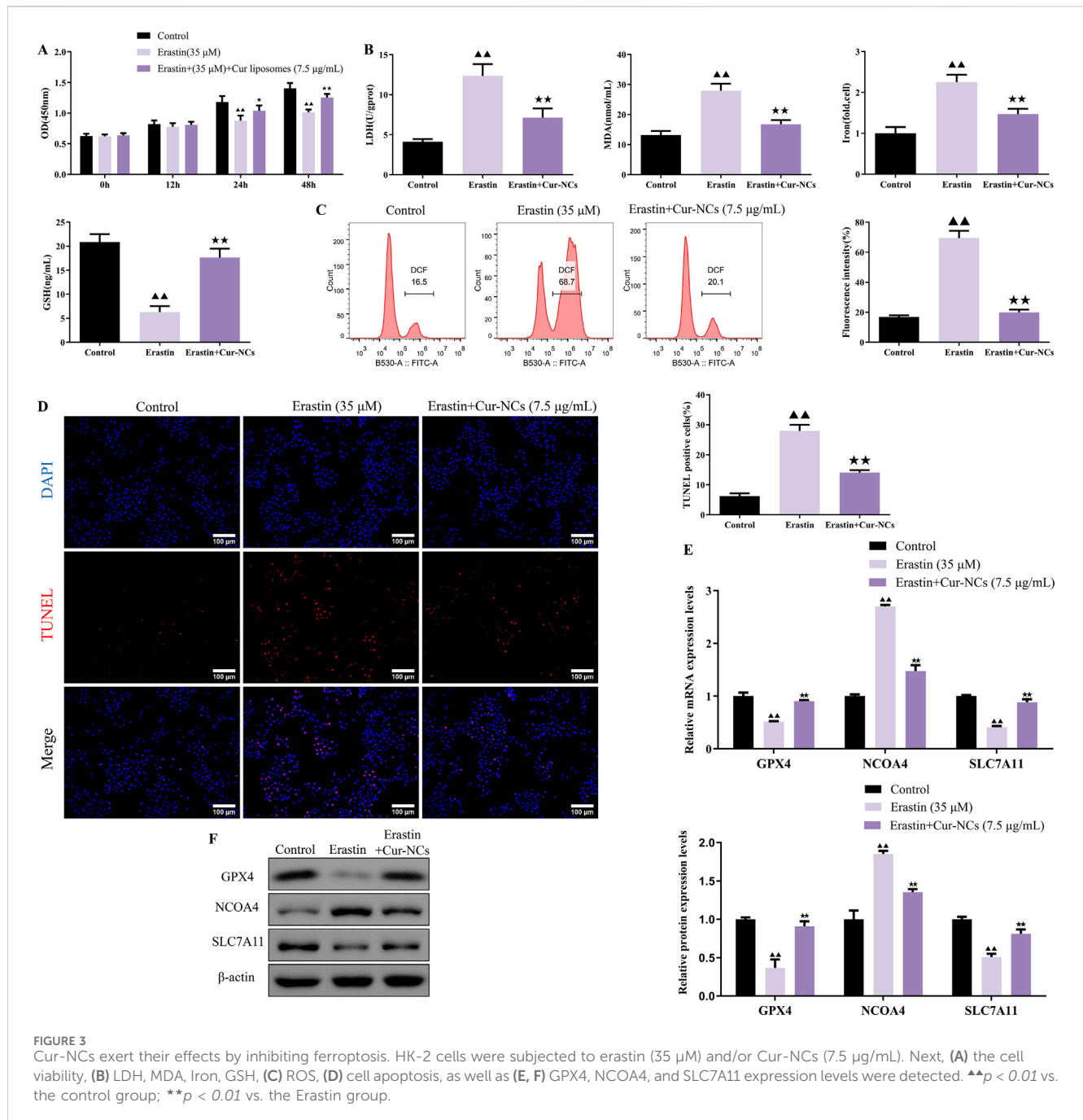
FIGURE 2 Effect of Cur-NCs on ferroptosis of HG-induced HK-2 cells. **(A)** LDH, MDA, Iron, and GSH levels in HG-induced HK-2 cells were measured by ELISA. **(B)** Lipid ROS levels of Cur-NCs treated HG-induced HK-2 cells were assayed by flow cytometry. **(C)** Representative images of mitochondria in Cur-NCs treated HG-induced HK-2 cells. scale bar, 500 nm. The expression levels of GPX4, NCOA4, SLC7A11, FTH-1, and TFR-1 in Cur-NCs treated HG-induced HK-2 cells were measured by **(D)** qPCR and **(E)** Western blot respectively. $\Delta\Delta p < 0.01$ vs. the control group; $*p < 0.05$ $**p < 0.01$ vs. the HG group.

TCTCTCCACAGTTCCTCAT-3'). These siRNAs were also diluted with Opti-MEM medium in the same way and mixed thoroughly. Then, the diluted siRNAs and the diluted Lipofectamine™ 3,000 were mixed at a ratio of 1:1 at room temperature and incubated for 10 min. The original culture medium was removed, and the cells were washed twice with serum-free medium. Subsequently, fresh serum-free medium was added, and the complexes were added into the wells and evenly distributed. The cells were incubated for 4 h, and then the medium containing serum was replaced. After transfection for 48 h,

the transfection efficiency was measured by using quantitative real-time PCR (qRT-PCR) and Western blotting assays.

Cell counting kit-8 (CCK-8) assay

Cell viability was assessed using the CCK-8 assay according to the manufacturer's instructions (C0039, Beyotime Biotechnology, China). Briefly, HK-2 cells were seeded in 96-well plates at a density of 5,000 cells



per well and allowed to adhere overnight. Following treatment with various experimental conditions, 10 μ L of CCK-8 solution was added to each well and incubated for 2 h at 37°C. The absorbance was measured at 450 nm using a microplate reader (Molecular Devices).

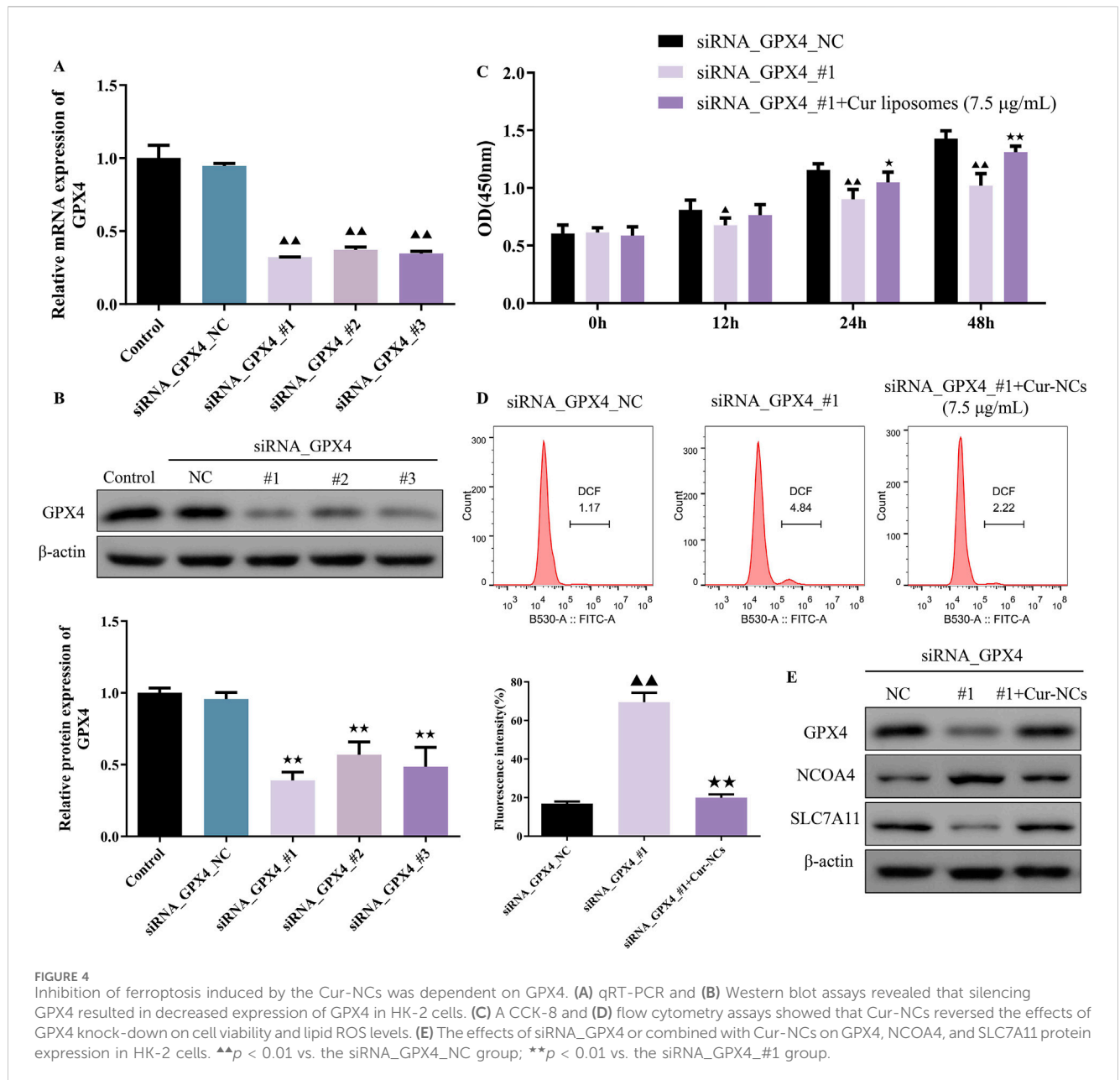
Flow cytometry assay

ROS levels were quantified using flow cytometry analysis. After treatment with experimental conditions, the treated HK-2 cells were harvested and washed with phosphate-buffered saline (FG701-01, TransGen, China). Subsequently, cells were incubated with 10 μ M 2',7'-dichlorofluorescein diacetate (DCFH-DA) probe (S0033M,

Beyotime Biotechnology, China) for 30 min at 37°C in the dark to detect intracellular ROS. Following incubation, cells were washed and resuspended in PBS. Flow cytometry analysis was performed using a BD Accuri™ C6 flow cytometer (BD Biosciences, United States), and ROS levels were quantified based on the fluorescence intensity of oxidized DCFH. Data were analyzed using Tree Star FlowJo software.

TEM assay

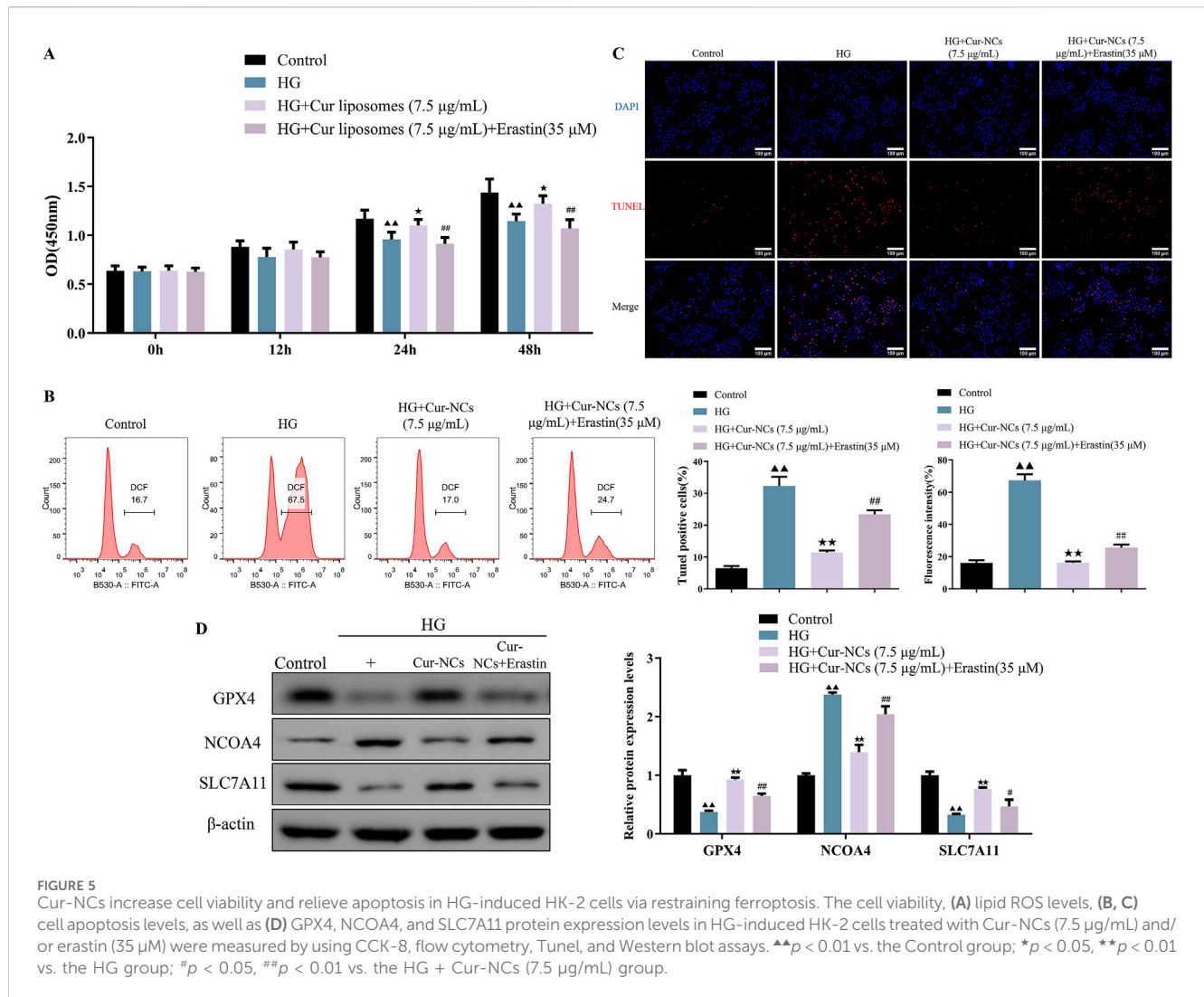
TEM was employed to investigate the ultrastructural characteristics of HG-induced HK-2 cells treated with Cur-



NCs or Fer-1. Before imaging, HK-2 cells were cultured in appropriate growth media and seeded onto carbon-coated electron microscopy grids. Following fixation with 2.5% glutaraldehyde in 0.1 M phosphate buffer (pH 7.4) for 2 h at room temperature, cells were post-fixed with 1% osmium tetroxide in the same buffer for 1 h. Dehydration was achieved through a graded series of ethanol solutions, and cells were then embedded in Epon resin. Ultrathin sections (70 nm) were cut using a diamond knife on an ultramicrotome and collected on copper grids. Sections were contrasted with uranyl acetate and lead citrate before examination under a TEM (H-7650, Hitachi, Japan) operating at an accelerating voltage of 120 kV. Images were captured using a CCD camera (Gatan Orius SC1000, United States) at various magnifications to assess cellular morphology and subcellular structures.

Terminal deoxynucleotidyl transferase dUTP nick end labeling (Tunel) staining

Tunel staining (C1090, Beyotime Biotechnology, China) was performed to assess the apoptosis levels in HK-2 cells. Cultured HK-2 cells were subjected to various experimental conditions as appropriate and subsequently fixed with 4% paraformaldehyde in PBS for 15 min at room temperature. Following fixation, cells were permeabilized with 0.1% Triton X-100 in PBS for 5 min and then incubated with the 50 μ L of Tunel reaction mixture containing terminal deoxynucleotidyl transferase enzyme and fluorescein-labeled dUTP for 1 h at 37°C in a humidified chamber. After Tunel staining, cells were counterstained with DAPI (4',6-diamidino-2-phenylindole) for nuclear visualization. The stained cells were imaged using a fluorescence microscope (Nikon DS-Fi2, Japan) equipped with appropriate filter sets. Quantification of

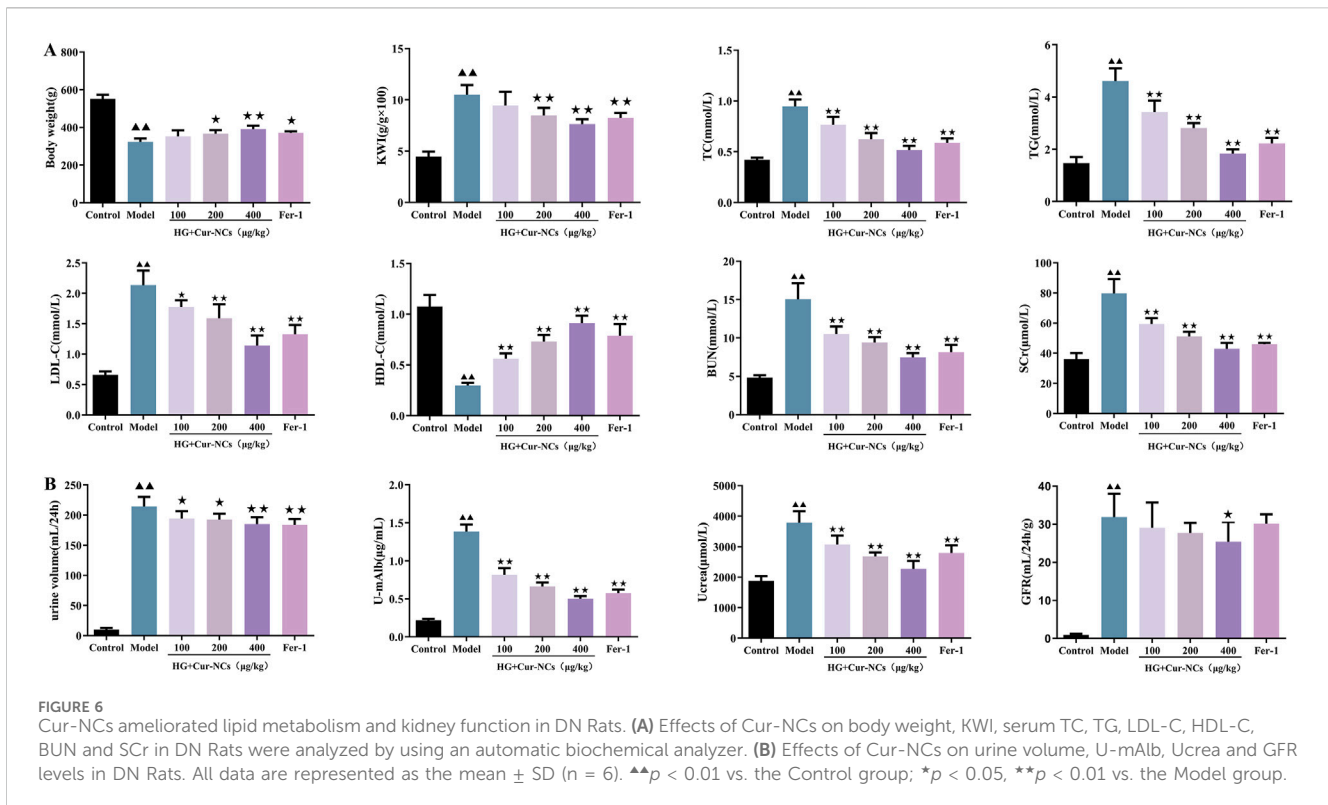


TUNEL-positive cells was performed by counting labeled nuclei in randomly selected fields of view.

Animals and treatment

All animal experiments were approved by the Institutional Animal Care and Use Committee of Zhejiang Center of Laboratory Animals (ZJCLA) (Approval number: ZJCLA-IACUC-20010530). A total of 42 6-week-old male Sprague-Dawley rats (Tu et al., 2019; Valdivielso et al., 2019; Noshahr et al., 2020; Ganugula et al., 2023) were obtained from Zhejiang Center of Laboratory Animals. Rats were housed at $22^{\circ}\text{C} \pm 2^{\circ}\text{C}$, a 12 h light-dark cycle, $50\% \pm 5\%$ humidity, with free access to food and water. After acclimatization for 7 days, 6 rats were randomly assigned to the control group. Following overnight fasting, the rats were received a single intraperitoneal injection of streptozotocin (S0130, Sigma, Germany), which was dissolved in sodium citrate buffer solution (pH 4.0, R21466, Shanghai yuanye Bio-Technology Co., Ltd., China) at a dosage of 65 mg/kg (ALTamimi et al., 2021; Machado et al., 2022). Three days after the injection, the blood glucose levels were measured by using a blood glucose meter (Accu-Chek). Rats with fasting blood

glucose levels exceeding 16.7 mmol/L for three consecutive days were diagnosed with diabetes (Supplementary Figure S1) and then included in the subsequent research. To investigate the therapeutic efficacy of Cur-NCs, DN rats were randomly assigned to five groups: Model, Cur-NCs low-dose, Cur-NCs middle-dose, Cur-NCs high-dose, and Fer-1 groups, with 6 rats in each group. Rats in the control and model groups were intraperitoneally injected with an equal volume of buffered saline solution daily; Rats in Fer-1 group received daily intraperitoneal injection of Fer-1 (2.5 $\mu\text{mol}/\text{kg}$, S81461, Shanghai yuanye Bio-Technology Co., Ltd., China); and rats in the low-dose group, medium-dose group, and high-dose group of Cur-NCs were intraperitoneally injected with 100, 200, and 400 $\mu\text{g}/\text{kg}$ of Cur-NCs daily (Zheng et al., 2024), respectively. The rats were injected once daily for 8 weeks. Animal body weight were determined weekly until the end of the experiment. At the end of the experiment, all were housed in metabolic chambers for 24 h to collect urine samples and record urine volume. U-mAlb (Urinary microalbumin) and Ucrea (Urinary creatinine) levels in urine samples were detected using a biochemical analyzer (Hitachi 7020, Japan). After that, the rats were fasted overnight and anesthetized with isoflurane. Then blood was collected from the heart, and centrifuged at 3,000 rpm for 15 min to obtain the serum. The



levels of TC, TG, LDL-C, HDL-C, BUN, and SCr in each group of rats were determined using a biochemical analyzer (Hitachi 7020, Japan). Based on the obtained data of urinary creatinine, plasma creatinine, urine volume and body weight, glomerular filtration rate (GFR) was calculated according to the formula “GFR = (Ucrea/SCr) * urine volume/body weight” (Gao et al., 2020; Zhao et al., 2022). Finally, the kidneys were removed immediately, weighed, and used to calculate the kidney/weight index (KWI). The freshly dissected kidneys were preserved at -80°C for subsequent experiments.

Enzyme-linked immunosorbent assay (ELISA)

Levels of LDH (A020-2-2, Nanjing Jiancheng, China), MDA (MM-0436H, Jiangsu Meimian Industrial Co., Ltd., China), Iron (Fe^{2+} , BC5310, Solarbio, China), and GSH (MM-0458H, Jiangsu Meimian Industrial Co., Ltd., China) in the cultured HK-2 cells, as well as the levels of MDA, SOD, Cat, and GSH (MM-0385R2, MM-0386R1, MM-20447R1, MM-0602R1; Jiangsu Meimian Industrial Co., Ltd., China) in kidney tissue samples were measured using the commercial kits according to the manufacturer’s instructions.

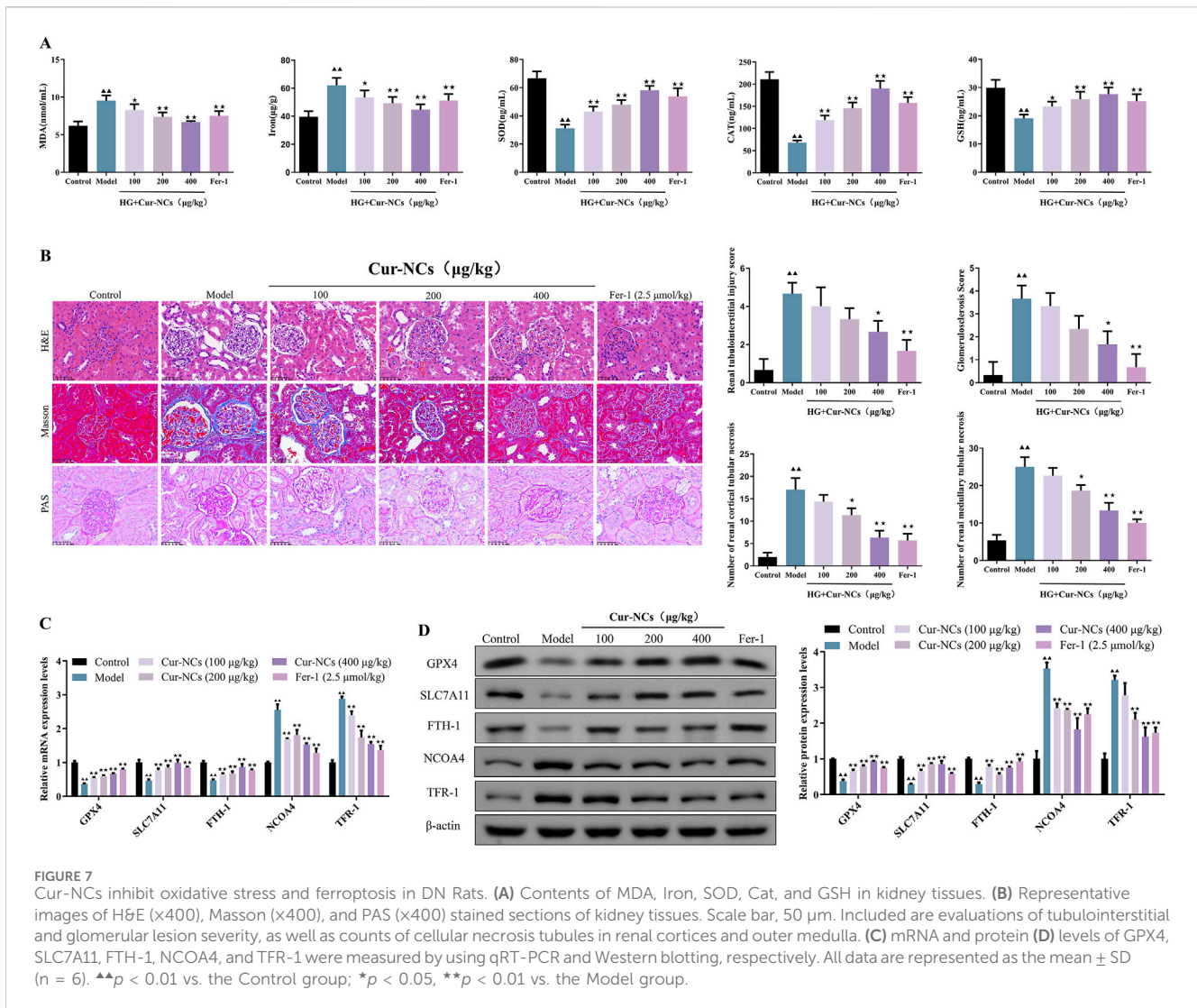
Renal histological analysis

Histological analyses were performed on kidney tissue samples using a combination of hematoxylin and eosin (H&E), Masson’s trichrome, and Periodic Acid-Schiff (PAS) staining techniques. Briefly, kidney tissues were fixed in 10% paraformaldehyde for 24 h, embedded in paraffin, and sectioned into 5- μm -thick slices.

For H&E staining, tissue sections were deparaffinized, rehydrated, and stained with Hematoxylin (H3136, Sigma, Germany) for 5 min followed by Eosin (E4009, Sigma, Germany) for 2 min. Masson’s trichrome staining was conducted to assess collagen deposition and fibrosis, with tissue sections subjected to Weigert’s iron hematoxylin (R20387, Shanghai yuanye Bio-Technology Co., Ltd., China), Ponceau S acid fuchsin stain (71019360, 71033761, Sinopharm Chemical Reagent Co., Ltd., China), and Aniline blue (71003644, Sinopharm Chemical Reagent Co., Ltd., China) staining solutions. For PAS staining to evaluate glycogen and basement membranes, tissue sections were oxidized with periodic acid (C0142S, Beyotime Biotechnology, China), treated with Schiff’s reagent (C0142S, Beyotime Biotechnology, China), and counterstained with Hematoxylin (Bry-0001-01, Runner bio, China). Stained sections were examined under a light microscope ($\times 400$; Nikon, Japan).

qRT-PCR assay

Total RNA was extracted from the treated HK-2 cells and kidney samples using an EZ-10 Total RNA Mini-Preps Kit (B618583-0100, Sangon, China) according to the manufacturer’s instructions. The quantity and quality of RNA were determined using a spectrophotometer. Complementary DNA (cDNA) was synthesized from the purified RNA using a high-capacity cDNA reverse transcription kit (CW2569, CoWin Biotech Co., Ltd., China). qRT-PCR was performed on a real-time PCR system (LightCycler[®] 96, Roche, Switzerland) with specific primers designed for the target genes and a SYBR Green PCR master mix (11201ES08, YeasenBiotechnology Co., Ltd., China). The detailed information of the related primer sequences is presented in



Supplementary Table S1. The amplification conditions included an initial denaturation step at 95°C for 10 min, followed by 40 cycles of denaturation at 95°C for 15 s and annealing/extension at 60°C for 1 min. The relative expression levels of the target genes were calculated using the $2^{-\Delta\Delta Ct}$ method.

Western blot analysis

Western blot analysis was conducted to assess the protein expression levels of GPX4, NCOA4, FTH-1, TFR-1, and SLC7A11 in both treated HK-2 cells and kidney tissues. Total protein was extracted from the treated HK-2 cells and kidney samples using a RIPA buffer (P0013B, Beyotime Biotechnology, China) supplemented with protease and phosphatase inhibitors (CW2200S, CoWin Biotech Co., Ltd., China). Protein concentrations were determined using a Bradford protein assay kit (P0012, Beyotime Biotechnology, China). Equal amounts of protein were resolved on sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred onto polyvinylidene fluoride (PVDF) membranes. Membranes were blocked with 5% non-fat milk in

Tris-buffered saline with 0.1% Tween-20 (TBST) for 1 h at room temperature and then incubated overnight at 4°C with primary antibodies specific for GPX4 (1:1,000, ab125066, Abcam, Britain), NCOA4 (1:1,000, ab86707, Abcam, Britain), FTH-1 (1:1,000, ab75973, Abcam, Britain), TFR-1 (1:1,000, ab214039, Abcam, Britain), SLC7A11 (1:1,000, ab175186, Abcam, Britain), and β -actin (1:10,000, AF7018, Affinity, United States). Following primary antibody incubation, membranes were washed with TBST and incubated with appropriate secondary antibodies conjugated to horseradish peroxidase (HRP) for 1 h at room temperature. Protein bands were visualized using an enhanced chemiluminescence (ECL) detection system, and densitometric analysis was performed using ImageJ software. β -actin was used as a loading control to normalize protein expression levels.

Statistical analysis

All results are shown as the mean \pm SD. A significant difference was analyzed using SPSS software (Version 22.0, IBM Corp., United States). Statistical analysis was carried out using two-paired-tailed *t*-tests or one-way analysis of variance (ANOVA)

with Tukey's *post hoc* test. $p < 0.05$ was considered statistically significant.

Results

Cur-NCs ameliorated the cell viability of HG-induced HK-2 cells

Cur-NCs had an average size of 122.0 ± 2.464 nm, a PDI of 0.167 ± 0.024 , and a zeta potential of -17.36 ± 2.38 mV (Figure 1A). Concurrently, TEM showed that the Cur-NCs were circular-shaped with a uniform size (Figure 1B). The CCK-8 test was used to measure cell viability in order to ascertain the effects of various doses of Cur-NCs on HK-2 cells. The viability of HK-2 cells did not significantly decrease when the concentration of Cur-NCs was increased to $15 \mu\text{g}/\text{mL}$, as Figure 1C illustrates. Furthermore, we discovered that the viability of HG-induced HK-2 cells was considerably increased by dosages of 1.87, 3.75 and $7.5 \mu\text{g}/\text{mL}$ of Cur-NCs and Fer-1 (Figure 1D). Therefore, 1.87, 3.75 and $7.5 \mu\text{g}/\text{mL}$ of Cur-NCs were selected for subsequent treatment.

Cur-NCs ameliorated oxidative damage in HG-induced HK-2 cells

To investigate the role of Cur-NCs in HG-induced oxidative damage, HK-2 cells were treated with 30 mM glucose to establish a cell model. Then, the cell death was detected using LDH release. We found that the LDH levels of the HG group increased when compared with the group control, suggesting that the HG condition induced the damage of HK-2. Furthermore, Cur-NCs exhibited a dose-dependent reduction in LDH levels (Figure 2A). As we know, ferroptosis is involved in the death of renal tubular cells in DN (Li et al., 2023a; Li et al., 2023b). Then, the levels of MDA, Iron, GSH and ROS were determined in HG-induced HK-2 cells treated with Cur-NCs. The results showed that the levels of MDA, Iron, and ROS in the different concentrations of Cur-NCs were significantly decreased from that of the HG group, while there was an increase of GSH levels in the Cur-NCs and Fer-1 groups (Figures 2A, B). Importantly, electron microscopic observation revealed that HG-induced mitochondrial damage of ferroptosis in HK-2 cells was characterized by the disappearance of mitochondrial cristae and rupture of the mitochondrial membrane, which was significantly attenuated by Cur-NCs and Fer-1 treatment (Figure 2C). In addition, the expression of ferroptosis-related markers GPX4, NCOA4, SLC7A11, FTH-1 and TFR-1 was detected by Western blotting. The results showed that HG treatment induced ferroptosis, as evidenced by increased expression of NCOA4 and TFR-1, and decreased expression of GPX4, SLC7A11, and FTH-1, which were reversed by Cur-NCs and Fer-1 treatment (Figures 2D, E).

Cur-NCs evoked its functions through the inhibition of ferroptosis by targeting GPX4

To investigate the role of ferroptosis in Cur-NCs' function, HK-2 cells were treated with erastin ($35 \mu\text{M}$), a ferroptosis inducer, and/

or Cur-NCs ($7.5 \mu\text{g}/\text{mL}$). Compared to the control, erastin dramatically increased LDH, MDA, Iron, ROS, and apoptosis levels while decreasing cell viability and GSH content in HK-2 cells, but Cur-NCs reversed the alterations (Figures 3A–D). In addition, erastin therapy drastically reduced the mRNA and protein expression levels of GPX4 and SLC7A11, although Cur-NCs treatment restored them. Interestingly, Cur-NCs treatment considerably reduced NCOA4 levels (Figures 3E, F). To further identify the potential mechanism of Cur-NCs, the expression levels of GPX4 were markedly decreased in HK-2 cells after transfection with si-GPX4 (Figures 4A, B). In addition, GPX4 knockdown significantly reduced the viability of the HK-2 cells, increased the formation of lipid ROS and NCOA4 expression, and reduced the expression of SLC7A11. However, treatment with Cur-NCs nearly eliminated this effect in the HK-2 cells (Figures 4C–E). Moreover, erastin process reversed the Cur-NCs treatment's improvement in cell viability as well as its suppression of cell apoptosis and ROS generation in HG-induced HK-2 cells (Figures 5A–C). To verify whether curcumin could inhibit HG-induced ferroptosis in HK-2 cells, we detected the expression of ferroptosis regulatory proteins GPX4, NCOA4 and SLC7A11 by Western blotting. The experimental results showed that the protein expression levels of GPX4 and SLC7A11 were significantly higher and that of NCOA4 was significantly lower in the Cur-NCs-treated group compared with the high-glucose group. However, after adding Erastin for intervention, we observed that the above protective effects of Cur-NCs were significantly suppressed (Figure 5D). Therefore, the results showed that Cur-NCs had the potential to suppress ferroptosis in HG-induced HK-2 cells through the activation of GPX4.

Cur-NCs ameliorated the hyperlipidemia and renal function in a rat DN model

To explore the protective roles of Cur-NCs in DN *in vivo*, the body weight, KWI, serum TC, TG, LDL-c, HDL-C, BUN, SCr, as well as the urine volume, U-mAlb, Ucrea and GFR were recorded and detected. The results showed that administration with Cur-NCs (100, 200, $400 \mu\text{g}/\text{kg}$) or Fer-1 significantly increased the body weight and resulted in a significant decrease in KWI, serum TC, TG, LDL-C, HDL-C, BUN, SCr, urine volume, U-mAlb, Ucrea and GFR levels when compared with model group (Figures 6A, B). Notably, the Ucrea and GFR levels in the model group were higher than those in the control group. It suggests that the kidneys were still in the state of ultrafiltration at this stage. In conclusion, these results suggested that Cur-NCs could ameliorate hyperlipidemia and renal function in STZ-induced DN rats.

Cur-NCs improved oxidative stress and glomerular structure in a rat DN model

It is generally acknowledged that iron buildup and oxidative stress are important factors in the etiology of ferroptosis in DN. Therefore, we looked at how Cur-NCs affected iron accumulation in the renal tissues of DN rats as well as oxidative stress markers (MDA, SOD, Cat, and GSH). Our findings demonstrated that,

following treatment with Cur-NCs or Fer-1, MDA was considerably decreased in the renal tissues of DN rats, whereas SOD, Cat, and GSH production increased (Figure 7A), indicating a decrease in oxidative stress. Furthermore, we saw a decrease in iron accumulation, which adds more credence to Cur-NCs' antioxidant and iron-regulating abilities. Since renal structural abnormalities are the main cause of renal insufficiency, we evaluated the renal microstructure using H&E, PAS and Masson staining. In control rats, no signs of glomerulosclerosis, interstitial hyperplasia, or basement membrane thickening were observed. In contrast, DN rats exhibited marked dilatation of the thylakoid matrix and tubulointerstitial fibrosis. However, the kidneys of DN rats treated with Cur-NCs or Fer-1 showed a trend toward normalization. To quantify the extent of renal injury, we assessed the severity of tubulointerstitial and glomerular lesions and counted the number of tubules with cellular necrosis in the renal cortex and outer medulla. Our findings revealed that Cur-NC treatment dramatically decreased the severity of tubulointerstitial and glomerular lesions while decreasing the number of necrotic tubules in the renal cortex and outer medulla when compared to untreated DN rats (Figure 7B). In addition, Western blotting results confirmed that Cur-NCs upregulated the protein levels of GPX4, SLC7A11, and FTH-1, while suppressing the expression of NCOA4 and TFR-1 compared with the model group (Figures 7C, D). These findings suggest that Cur-NCs regulate ferritin-related proteins and iron homeostasis, thereby attenuating DN-induced kidney injury.

Discussion

DN is a severe complication of diabetes mellitus and a leading cause of end-stage renal disease worldwide (Jin et al., 2023; Zhao et al., 2023). Despite extensive research, effective therapeutic strategies for DN remain limited. In this study, we investigated the potential of Cur-NCs to improve DN by inhibiting ferroptosis, a newly recognized form of regulated cell death associated with oxidative stress and inflammation. The results of our study revealed that Cur-NCs significantly ameliorated the pathological manifestations of DN in animal models. This included improvements in renal function and structural preservation of renal tissue. Importantly, we elucidated the molecular mechanism underlying these effects, showing that Cur-NCs targeted GPX4, a key regulator of ferroptosis (Xu et al., 2024a), thereby attenuating oxidative stress in HG-induced HK-2 cells and renal tissues and consequently mitigating DN progression.

Research has shown that the overproduction of lipid peroxides can lead to cell death via ferroptosis (Jiang et al., 2021). Ferroptosis is closely associated with the development of DN (Li et al., 2023b; Mengstie et al., 2023). Ferroptosis is primarily related to iron metabolism disorders, GPX4 antioxidation, and lipid peroxidation (Liu et al., 2022). Under physiological conditions, Fe³⁺ enters cells via transferrin (TF) and is reduced to Fe²⁺ by STEAP3. Fe²⁺ can bind to ferritin heavy chain 1 (FTH1) for storage in cells or be oxidized to Fe³⁺ and transported out of cells via membrane transferrin receptor 1 (TFR1), participating in the body's iron circulation to maintain iron homeostasis (Dong et al., 2020; Plays et al., 2021). When iron is overloaded, high concentrations of

Fe²⁺ undergo the Fenton reaction, generating ROS that cause oxidative damage to cell membranes, ultimately resulting in cell death via ferroptosis (Zhao et al., 2021). Our study found that the levels of Fe in HG-induced HK-2 cells and kidney tissues of DN rats were significantly higher than those in the control group, which may be associated with a significant decrease in FTH1 levels. However, intervention with Cur-NCs alleviated iron-mediated cell death in HG-induced HK-2 cells and kidney tissues of DN rats in this study. GPX4 plays a crucial role in cellular defense against oxidative stress by catalyzing the reduction of lipid hydroperoxides to their corresponding alcohols. Its importance in kidney physiology and pathology has been increasingly recognized. Studies have demonstrated that GPX4 deficiency exacerbates renal injury in various experimental models, highlighting its protective role against oxidative damage and inflammation in the kidney (Hosohata et al., 2022; Li et al., 2022b). Our findings add to this body of knowledge by showing that modulation of GPX4 activity with Cur-NCs can mitigate DN progression by inhibiting ferroptosis.

Furthermore, our study underscores the potential of nanotechnology in drug therapy. Nanocrystals possess advantages such as a large specific surface area, rapid dissolution rate, and high drug-loading capacity, which can enhance the bioavailability of drugs and improve therapeutic efficacy. The practical value of nanocrystals is demonstrated by the fact that multiple nanocrystal-based drugs are already in clinical use. Therefore, the development of Cur-NCs holds promise as an effective treatment for DN by modulating GPX4 activity. Despite the encouraging results of our study, several limitations warrant consideration. At present, our research has primarily focused on exploring the potential of Cur-NCs using cell and animal models. However, we acknowledge that a rigorous preclinical evaluation of Cur-NCs, including safety assessments such as acute and potential long-term toxicity studies, as well as pharmacodynamic investigations across various animal species, has yet to be conducted. This comprehensive preclinical assessment represents a critical next step in our research, and we plan to undertake it in the future with the aim of laying a solid foundation for eventual clinical trials in human subjects, which are essential for validating the safety and efficacy of Cur-NCs in the treatment of DN. Additionally, while we identified GPX4 as a key mediator of Cur-NCs effects in DN, the precise molecular mechanisms underlying GPX4 modulation and ferroptosis inhibition require further investigation. Finally, the complexity of DN pathogenesis suggests that a multifaceted approach combining various therapeutic strategies may be necessary to fully address the disease's progression.

Conclusion

In conclusion, the current study provided compelling evidence that Cur-NCs are effective in slowing the progression of DN by inhibiting ferroptosis. This therapeutic effect was achieved through the upregulation of GPX4, a key regulator of ferroptosis, which attenuated oxidative stress and protected renal tissues from further damage in a cellular model of renal tubular injury and an animal model of DN. These findings highlight the potential application of Cur-NCs in the treatment of DN, highlighting the great potential of nanotechnology in drug therapy.

Data availability statement

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found below: <https://figshare.com/>, <https://doi.org/10.6084/m9.figshare.27193149.v2>.

Ethics statement

The animal study was approved by the Institutional Animal Care and Use Committee of Zhejiang Center of Laboratory Animals (ZJCLA) (Approval number: ZJCLA-IACUC-20010530). The study was conducted in accordance with the local legislation and institutional requirements.

Author contributions

MX: Formal Analysis, Investigation, Methodology, Writing—original draft. YT: Formal Analysis, Investigation, Resources, Writing—review and editing. HZ: Methodology, Visualization, Writing—review and editing. SD: Resources, Writing—review and editing. YW: Resources, Writing—review and editing. JJ: Conceptualization, Funding acquisition, Project administration, Supervision, Writing—review and editing. JC: Conceptualization, Resources, Supervision, Writing—review and editing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fphar.2024.1508312/full#supplementary-material>

SUPPLEMENTARY FIGURE S1

Establishment of STZ induced diabetes model in SD rats.

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