



# In vitro study on effect of bardoxolone methyl on cisplatin-induced cellular senescence in human proximal tubular cells

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## Abstract

Bardoxolone methyl [methyl-2-cyano-3, 12-dioxooleana-1, 9(11)dien-28-oate (CDDO-Me)], an activator of the nuclear factor erythroid-derived 2-related factor2 pathway, is a potential therapeutic candidate for the treatment of kidney diseases. However, its effect against cellular senescence remains unclear. This study aimed to investigate whether CDDO-Me protects cells against cisplatin-induced cellular senescence using an in vitro model. The human renal proximal tubular epithelial cell line HK-2 was treated with cisplatin for 6 h, followed by treatment with or without CDDO-Me (0.1 or 0.2  $\mu\text{mol/L}$ ). Senescence markers were analyzed using western blotting and real-time PCR. Apoptosis was evaluated through TUNEL staining. Cisplatin induced changes in the levels of markers specific for proliferation, cell cycle, and senescence in a time- and dose-dependent manner. Furthermore, IL-6 and IL-8 levels in the culture medium increased markedly. These data suggested that cellular senescence-like alterations occurred in HK-2 cells exposed to cisplatin. CDDO-Me treatment reversed the cisplatin-mediated alterations in the levels of cellular senescence markers. The antioxidant enzymes, *HO1*, *NQO1*, *GPX1*, and *CAT* were upregulated by CDDO-Me treatment. Furthermore, CDDO-Me treatment induced apoptosis in cisplatin-exposed HK-2 cells. Pretreatment with Ac-DEVD-CHO, the caspase inhibitor, suppressed the reversal effect of CDDO-Me against cisplatin-induced cellular senescence-like alterations. This study showed that CDDO-Me attenuated cisplatin-induced premature senescence of HK-2 cells. This beneficial effect may be related to Nrf2 activation. Our findings also showed that CDDO-Me induced apoptosis in cisplatin-treated HK-2 cells, potentially protecting the kidneys from cellular senescence. CDDO-Me appears to be a candidate treatment for acute kidney injury.

**Keywords** CDDO-Me · Cellular senescence · Acute Kidney injury · Nrf2 · Apoptosis

## Introduction

Acute kidney injury (AKI) is a complex syndrome characterized by renal dysfunction and is associated with high morbidity and mortality worldwide. Patients with post-AKI are at an increased risk for chronic kidney disease, CKD [1–3]. Proximal tubule injury caused by AKI triggers several features of CKD by inducing inflammation and interstitial fibrosis [4]. Therefore, it is important to protect the proximal tubules for preventing the progression of AKI to CKD; however, the definitive evidence for this effect is lacking.

There is an association between cellular senescence in proximal tubular epithelial cells (PTCs) and CKD in the animal models of hypertension [5] or diabetes [6]. Senescence is a tumor suppressor mechanism by which cells adapt to DNA damage, oxidative stress, and telomere shortening. Senescence induces cell cycle arrest in cells exposed to various stresses

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[7, 8]. Senescent cells secrete pro-inflammatory molecules in order to recruit immune cells. This behavior is termed as senescence-associated secretory phenotype, SASP [9]. These pro-inflammatory molecules, including cytokines and chemokines, can lead to the clearance of senescent cells in the damaged tissue [10, 11]. In contrast, the chronic secretion of these SASP factors promotes inflammation and epithelial-to-mesenchymal transition, thereby reducing the functional reserve of tissues. Thus, senescence is protective in the acute phase; however, chronic senescence can be harmful to tissue homeostasis. Senescent cells undergo an anti-apoptotic state [12–14], leading to the constitutive secretion of SASP factors. Chronic secretion of SASP factors can spread senescence to nearby cells [15]; therefore, the removal of senescent cells by activating apoptosis seems to be a critical mechanism for protecting the proximal tubule and preventing CKD progression. In fact, much effort has been made to eliminate senescent cells using novel drugs called senolytics. These drugs remove senescent cells by inducing apoptosis [16] or by blocking their resistance to apoptosis [17, 18]. Therefore, protecting PTCs from cellular senescence could prevent CKD development; however, effective treatments that target senescence are lacking.

Methyl-2-cyano-3, 12-dioxooleana-1, 9(11)dien-28-oate (CDDO-Me), also known as bardoxolone methyl, is a semi-synthetic triterpenoid and the most potent activator of the nuclear factor erythroid-derived 2-related factor2 (Nrf2) pathway [19–21]. Besides activating Nrf2, CDDO-Me upregulates the antioxidant response and suppresses pro-inflammatory signaling reducing oxidative stress and inflammation, and promoting mitochondrial function [22, 23]. CDDO-Me and its analogs have beneficial effects on CKD associated with type 2 diabetes [24, 25], obesity [26], and angiotensin-induced kidney injury [27]. Furthermore, CDDO-Me induces apoptosis [28]; therefore, CDDO-Me could be beneficial for eliminating cellular senescence. CDDO-Me is an attractive therapeutic candidate for managing cellular senescence in kidney diseases, but the precise effects remain unknown.

Cisplatin is one of the most widely used anti-cancer drugs; however, its use is limited because of its nephrotoxicity, which causes AKI [29]. Cisplatin treatment is widely used as a model for AKI; it induces cellular senescence, both in vitro [30, 31] and in vivo [32, 33]. The aim of this study was, therefore, to investigate whether CDDO-Me protects PTCs against cisplatin-induced cellular senescence to explain the beneficial effect of CDDO-Me.

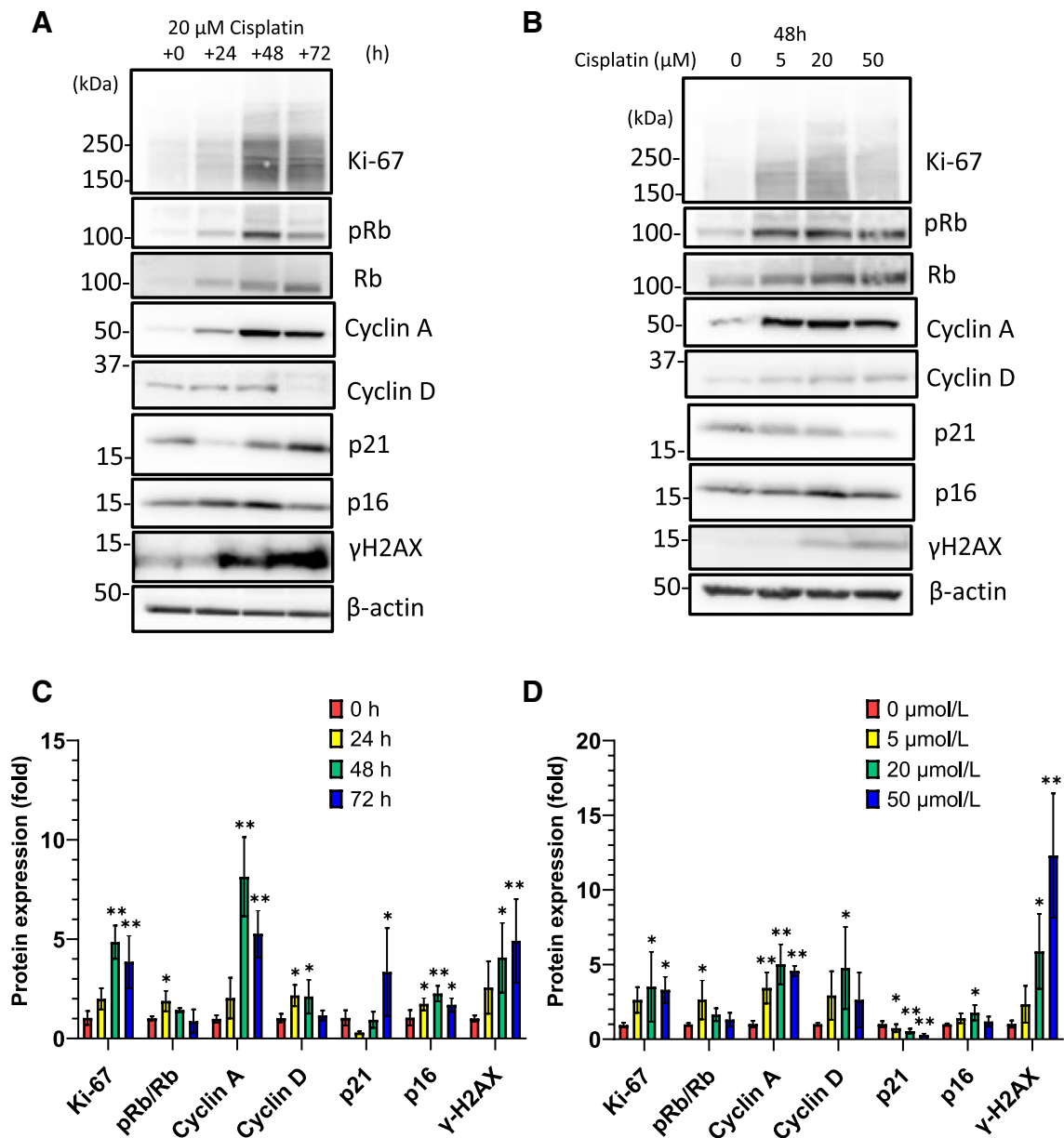
## Materials and methods

### Cell culture

The human renal proximal tubular epithelial cell line HK-2 was cultured in Dulbecco's Modified Eagle's Medium: Nutrient Mixture F-12 (DMEM/F12, Fujifilm Wako Chemical Co., Osaka, Japan) supplemented with 10% fetal bovine serum (FBS, Biosera, Inc., Nuaille, France) and 1% penicillin–streptomycin, as previously described [34]. Cells were treated with 0–50  $\mu\text{mol/L}$  cisplatin in low-glucose DMEM containing 0.1 mg/mL human serum albumin for 6 h. The medium was then replaced with fresh medium with or without 0.1–0.2  $\mu\text{mol/L}$  CDDO-Me (Sigma-Aldrich Co., St. Louis, MO, USA). The cells were treated with 5 or 50  $\mu\text{mol/L}$  of the caspase inhibitor, Ac-DEVD-CHO (Selleckchem, Houston, TX, USA), for 60 min before adding CDDO-Me to the medium. Cells were cultured for 24–72 h, and proteins or mRNA were extracted at the indicated points.

### Western blot (WB) analysis

The cultured cells were solubilized in lysis buffer (150 mmol/L NaCl, 50 mmol/L Tris–HCl, 5 mmol/L EDTA–2Na, 1% Triton X-100, and 1 tablet/10 mL complete mini EDTA-free) and centrifuged at  $15,000 \times g$  at 4 °C for 30 min. Samples were separated by SDS-PAGE and then transferred to PVDF membranes. The membranes were first blocked in a buffer containing 25 mmol/L Tris–HCl (pH 7.4), 150 mmol/L NaCl, 0.1% Tween 20, and 4% skim milk for 1 h and then incubated with primary antibodies at 4 °C overnight. This was followed by incubation with horseradish peroxidase-conjugated secondary antibodies for 1 h. Primary antibodies against human p21<sup>Waf1/Cip1</sup>, p16<sup>INK4a</sup>, phosphorylated H2AX (Ser139,  $\gamma$ -H2AX), retinoblastoma (Rb), phosphorylated Rb (Ser780, pRb), cyclin D, and caspase-3 were all purchased from Cell Signaling Technology, Inc. (Beverly, MA, USA). Antibodies against human cyclin A (Novocastra Laboratories Ltd., Newcastle, UK), p16<sup>INK4a</sup> (BD Biosciences, Inc., Farmingdale, NY, USA), Ki-67 (Dako from Agilent, Santa Clara, CA, USA), p62/SQSTM1, and  $\beta$ -actin (Santa Cruz Biotechnology, Santa Cruz, CA, USA) were also used. The immunoreactive proteins were then detected by enhanced chemiluminescence (GE Healthcare, Fairfield, CT, USA). Immunoblots were quantified using the CS Analyzer 3.0 software (ATTO, Tokyo);  $\beta$ -actin expression was used as the internal control.



**Fig. 1** Cisplatin induces cellular senescence-like alterations in HK-2 cells. HK-2 cells were treated with 0–50  $\mu\text{mol/L}$  cisplatin, and western blot analysis revealed changes in the expression levels of prolifer-

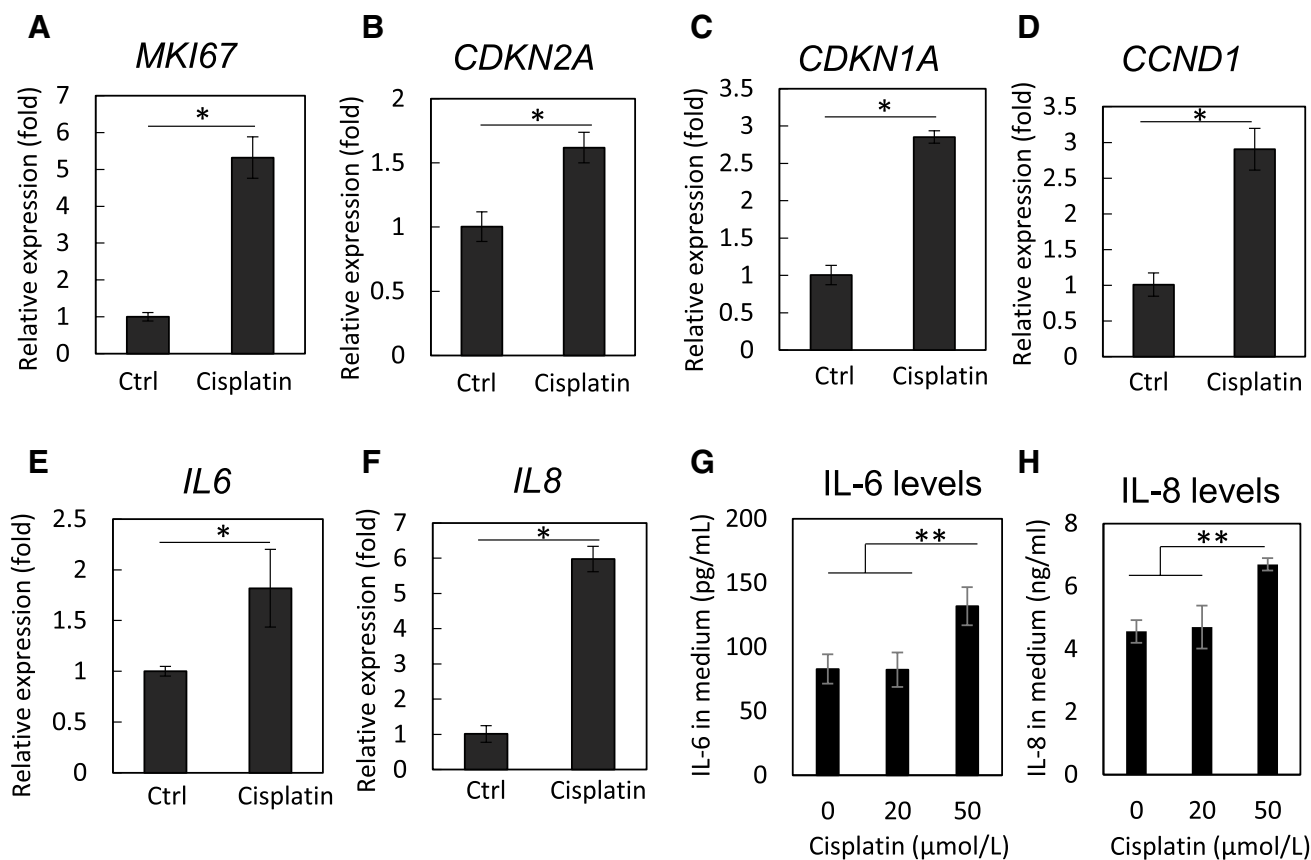
ation, cell cycle, and senescence markers in a time- (A) and dose- (B) dependent manner. Densitometric analysis is shown in C–D.  $n=4-6$ ,  $*P < 0.05$ ,  $**P < 0.01$

### Real-time reverse transcription-PCR (RT-qPCR)

Total RNA and cDNA from HK-2 cells were prepared using the ISOGEN-II (Nippon Gene, Tokyo) and Prime Script RT-PCR kits (Takara Bio, Shiga), according to the manufacturer's instructions. The primers used for qPCR are shown in Supplementary Table 1. qPCR was performed on a 7500 Real-Time PCR System (Applied Biosystems) in a 96-well reaction plate using Power SYBR Green. The mRNA expression of the target genes was normalized to that of *GAPDH*, using the delta–delta  $C_t$  method.

### Cell cycle analysis using flow cytometry

HK-2 cells were harvested, washed, and resuspended in phosphate buffered saline. Cells were then fixed with 70% ethanol and stored at 4 °C overnight. Subsequently, cells were incubated with 100  $\mu\text{g/mL}$  RNase for 30 min at 37 °C and stained with 5  $\mu\text{g/mL}$  propidium iodide (PI) for 10 min. Flow cytometry analyses were performed using a BD FACSCalibur (BD Biosciences). Cell cycle phase distributions were determined using Modfit LT software version 3.0 (Verity Software House, Topsham, ME, USA).



**Fig. 2** Cisplatin treatment induces the increase in senescence-related gene transcriptions and cytokine secretion. The mRNA expression of senescence-related genes, *MKI67* (A), *CDKN2A* (B), *CDKN1A* (C), *CCND1* (D), *IL-6* (E), and *IL-8* (F), after cisplatin (20 μmol/L) treat-

ment was determined by qPCR. Cytokine levels, IL-6 (G) and IL-8 (H), in cultured medium of cisplatin (20 or 50 μmol/L)-exposed cells were quantified by ELISA.  $n=3$ , \* $P<0.05$ , \*\* $P<0.01$

### Determination of cytokine levels by ELISA

Following exposure to cisplatin, supernatants were collected from the culture medium. Cytokine levels were determined using ELISA kits for IL-6 and IL-8 (R&D Systems, Minneapolis, MN, USA) as per the manufacturer's instructions.

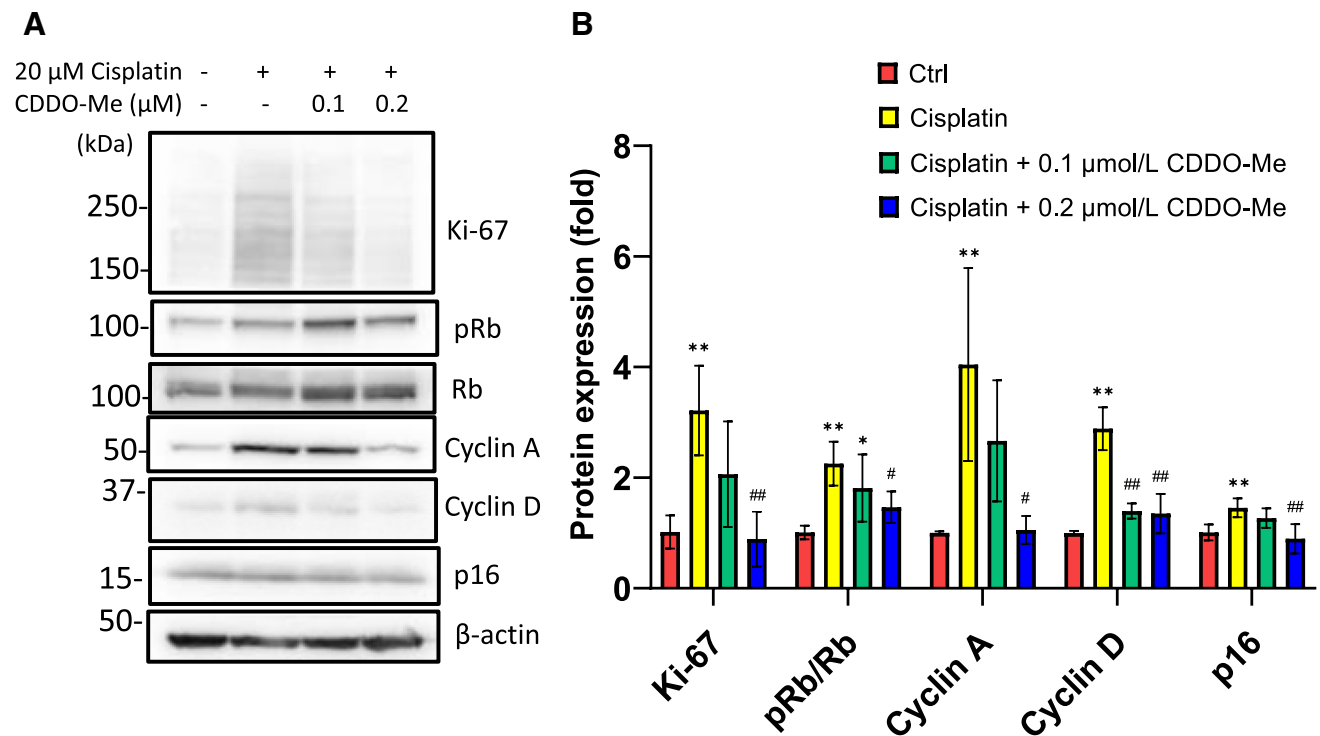
### TdT-mediated dUTP-biotin nick end labeling (TUNEL) staining

TUNEL staining was performed using a commercial kit (Takara Bio), following each treatment described above. The percentage of apoptotic cells was determined by counting the TUNEL-positive cells and the total number of cells (nucleus) in four to six photomicrographs ( $\times 200$  magnification, approximately 1500 cells) by a blinded observer.

This experiment was performed thrice, on different days. The apoptosis rate was expressed as the means  $\pm$  standard deviation (SD) from three independent experiments.

### Statistical analysis

All data are reported as the mean  $\pm$  SD. The RT-qPCR and WB analysis data are presented as the fold change relative to the controls (untreated cells). The RT-qPCR data were evaluated using Students *t*-test to assess differences between the control and cisplatin-treated cells. Other data were analyzed using analysis of variance (ANOVA) with post hoc comparisons using the Student–Newman–Keuls method.  $P$  values  $<0.05$  were considered statistically significant.



**Fig. 3** CDDO-Me treatment reversed the cisplatin-induced cellular senescence phenotype in HK-2 cells. HK-2 cells were exposed to 20  $\mu$ mol/L cisplatin for 6 h, followed by treatment with or without CDDO-Me (0.1 or 0.2  $\mu$ mol/L). Proteins were collected 48 h after

treatment and analyzed by western blotting (A), and protein bands were quantified by densitometry (B).  $n=4-6$ , \* $P<0.05$ , \*\* $P<0.01$  vs. control (Ctrl); + $P<0.05$ , ++ $P<0.01$  vs. cisplatin

## Results

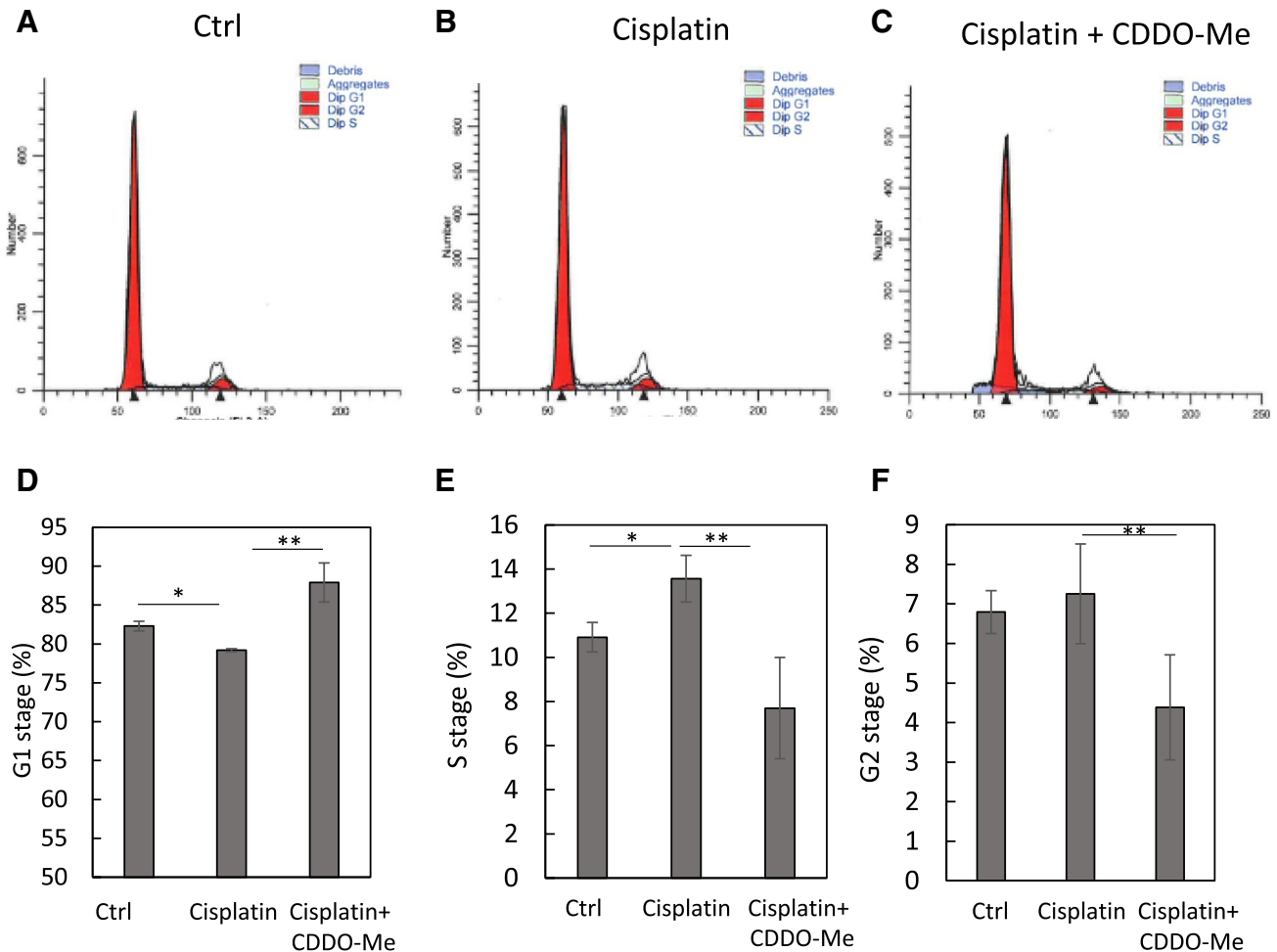
### Cisplatin treatment induces cellular senescence-like alterations in HK-2 cells

First, we examined whether cisplatin exposure induces cellular senescence in HK-2 cells. Cisplatin treatment induced alterations in the expression of proliferation, cell cycle, and senescence markers in a time- and dose-dependent manner (Fig. 1). The expression of Ki-67, a proliferation marker, significantly increased after exposure to 20  $\mu$ mol/L cisplatin; it decreased after exposure to 50  $\mu$ mol/L cisplatin compared to that after exposure to 20  $\mu$ mol/L cisplatin. The expression of the cell cycle markers pRb, Rb, cyclin D, and cyclin A increased in cisplatin-exposed HK-2 cells. Cisplatin treatment increased the levels of p16<sup>INK4a</sup> in a time-dependent manner. The expression level of p21<sup>Waf1/Cip1</sup> was decreased at 48 h; however, it was increased at 72 h. Cisplatin treatment increased the level of  $\gamma$ -H2AX, a DNA damage marker. The mRNA expression of *MKI67* (encoding Ki-67), *CDKN1A* (encoding p21<sup>Waf1/Cip1</sup>), *CDKN2A* (encoding p16<sup>INK4a</sup>), and *CCND1* (encoding cyclin D1) increased 48 h after exposure to 20  $\mu$ mol/L cisplatin (Fig. 2A–D). Furthermore, the mRNA levels of *IL-6* and *IL-8*

increased markedly (Fig. 2E, F). An increase in the cytokine levels was observed in the culture medium of cells exposed to 50  $\mu$ mol/L cisplatin (Fig. 2G, H). These data suggest that cisplatin treatment induced proliferation, DNA damage, and subsequent cellular senescence-like alterations in HK-2 cells.

### CDDO-Me reverses cisplatin-induced cellular senescence-like alterations in HK-2 cells

CDDO-Me treatment normalized the cisplatin-induced alterations in the expression of the cellular senescence markers Ki-67, cyclin A, cyclin D, pRb/Rb, and p16<sup>INK4a</sup> in HK-2 cells (Fig. 3). Cell cycle analysis using cytometry revealed that CDDO-Me treatment ameliorated cisplatin-induced cell cycle abnormalities (Fig. 4). Cisplatin markedly decreased the percentage of cells in the G1 phase and increased that in the S phase. The cells in the G2 phase also tended to increase after cisplatin treatment. However, CDDO-Me treatment increased the number of G1-phase cells and decreased that of S- and G2-phase cells.



**Fig. 4** CDDO-Me protects cells against cisplatin-induced abnormalities in cell cycle. Cisplatin (20  $\mu\text{mol/L}$ )-exposed cells were treated with 0.2  $\mu\text{mol/L}$  CDDO-Me for 48 h. Cells were harvested and

stained with propidium iodide, and then analyzed by flow cytometry (A–C). The statistics of each cell cycle phase (D–F).  $n=3$ , \* $P<0.05$ , \*\* $P<0.01$

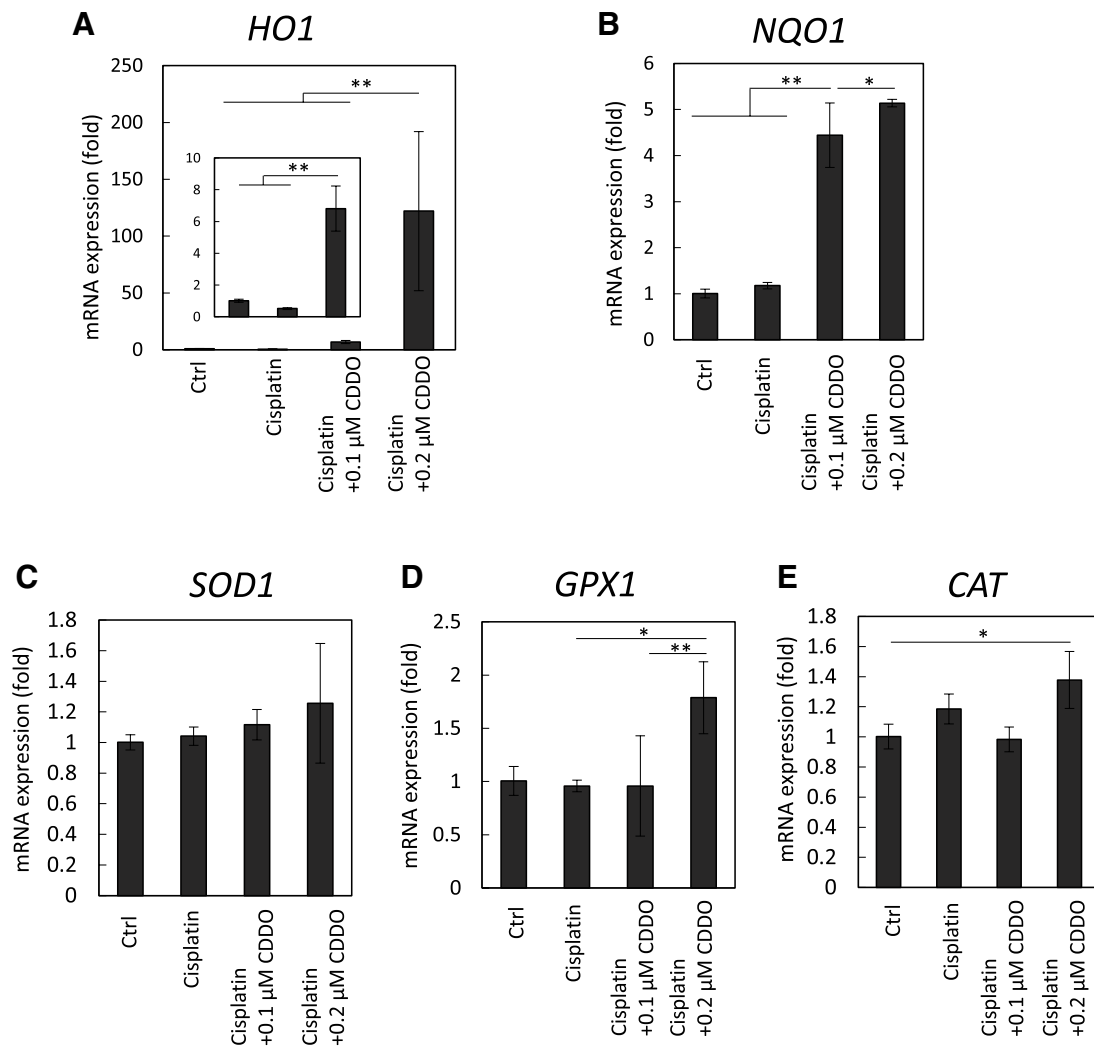
### CDDO-Me upregulates the expression of antioxidant enzymes

To evaluate the mechanisms by which CDDO-Me protected the cisplatin-exposed HK-2 cells against cellular senescence-like alterations, we evaluated whether the expression of antioxidant enzyme gene were upregulated under the present conditions (Fig. 5). The mRNA of antioxidant enzymes, heme oxygenase-1 (*HO1*), NAD(P)H:quinone oxidoreductase (*NQO1*), glutathione peroxidase 1 (*GPX1*), and catalase (*CAT*), were upregulated by CDDO-Me treatment. However, CDDO-Me treatment did not significantly increase the mRNA expression of superoxide dismutase 1 (*SOD1*).

### CDDO-Me induces apoptosis in cisplatin-treated HK-2 cells

The effect of CDDO-Me on apoptosis was evaluated because CDDO-Me treatment significantly decreased the cell number in cisplatin-exposed HK-2 cells (Fig. 6A). CDDO-Me induced apoptosis in cisplatin-exposed HK-2 cells; however, only cisplatin treatment also induced apoptosis, as observed using TUNEL staining (Fig. 6B, C). WB analysis confirmed that CDDO-Me treatment induced apoptosis, decreased the level of pro-caspase-3, and increased the level of cleaved caspase-3 and p62/SQSTM1 in HK-2 cells (Fig. 7A, B). CDDO-Me accelerated the phosphorylation of H2AX in cisplatin-treated HK-2 cells. To elucidate whether CDDO-Me





**Fig. 5** CDDO-Me induces ARE-related gene transcription by activating the Keap1-Nrf2 pathway. HK-2 cells were exposed to 20  $\mu\text{mol/L}$  cisplatin for 6 h, followed by treatment with or without CDDO-Me (0.1 or 0.2  $\mu\text{mol/L}$ ). The mRNA expression levels of Nrf2-related

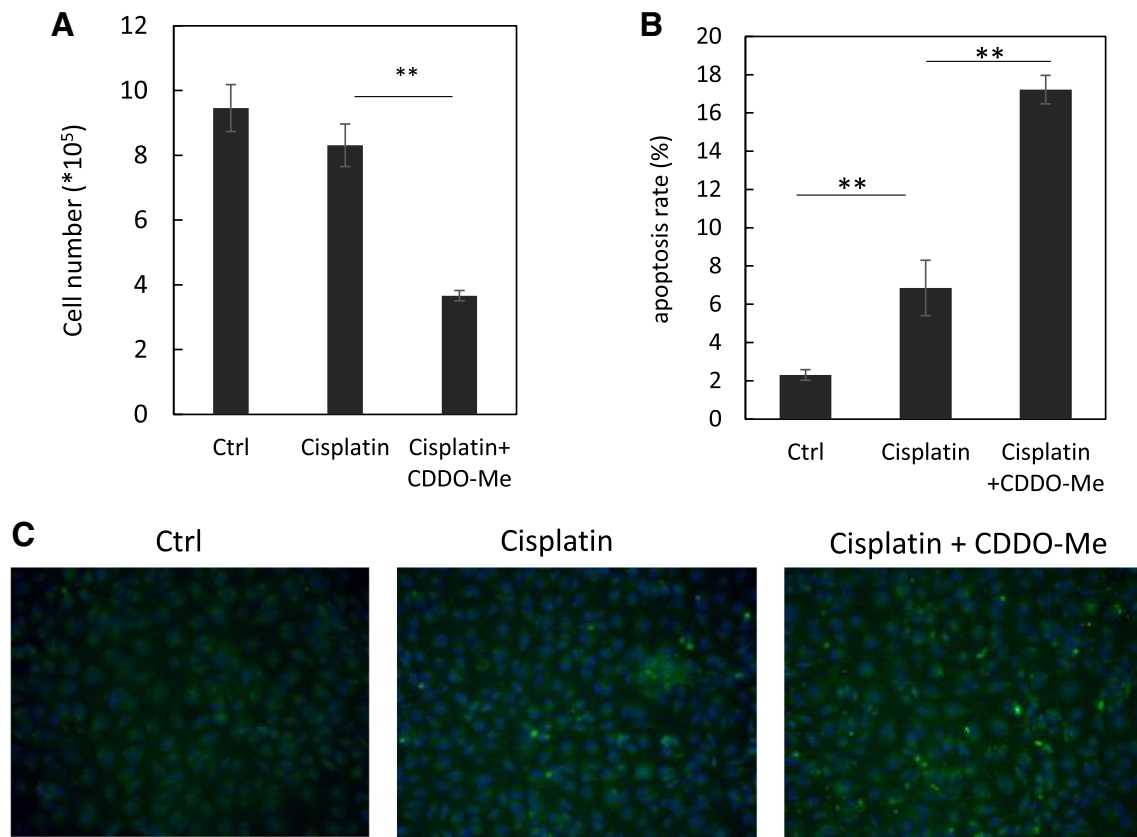
genes were analyzed by qPCR 48 h after treatment: heme oxygenase-1 (A), NAD(P)H:quinone oxidoreductase (B), superoxide dismutase 1 (C), glutathione peroxidase 1 (D), and catalase (E).  $n=3$ , \* $P<0.05$ , \*\* $P<0.01$

reverses cisplatin-induced cellular senescence-like alterations via apoptosis, cisplatin-exposed cells were treated with Ac-DEVD-CHO, a caspase inhibitor, before the CDDO-Me treatment. Ac-DEVD-CHO treatment inhibited the decrease in Ki-67, cyclin A, cyclin D, and p16<sup>INK4a</sup> expression levels induced by CDDO-Me (Fig. 7C, D).

## Discussion

CDDO-Me therapy improves renal function in CKD patients, in clinical trials [24, 25, 35]. However, the renal protective mechanism of CDDO-Me on the AKI-CKD

transition remains unknown. Protecting the proximal tubules from AKI, which causes cellular senescence, is important for preventing CKD development. This study elucidated the protective effect of CDDO-Me on cisplatin-induced cellular senescence in cultured human PTCs, HK-2, which is one of the most used cells for the study of senescent PTCs [36–38]. In this study, WB analysis revealed an increased expression of Ki-67 in HK-2 cells following cisplatin treatment; chronic increase in proliferation induces cellular senescence in PTCs [39]. Accelerated proliferation may be one mechanism by which cellular senescence is induced in cisplatin-treated cells. In addition to a pool of proliferating cells, elevated cell cycle arrest markers were observed, indicating the presence



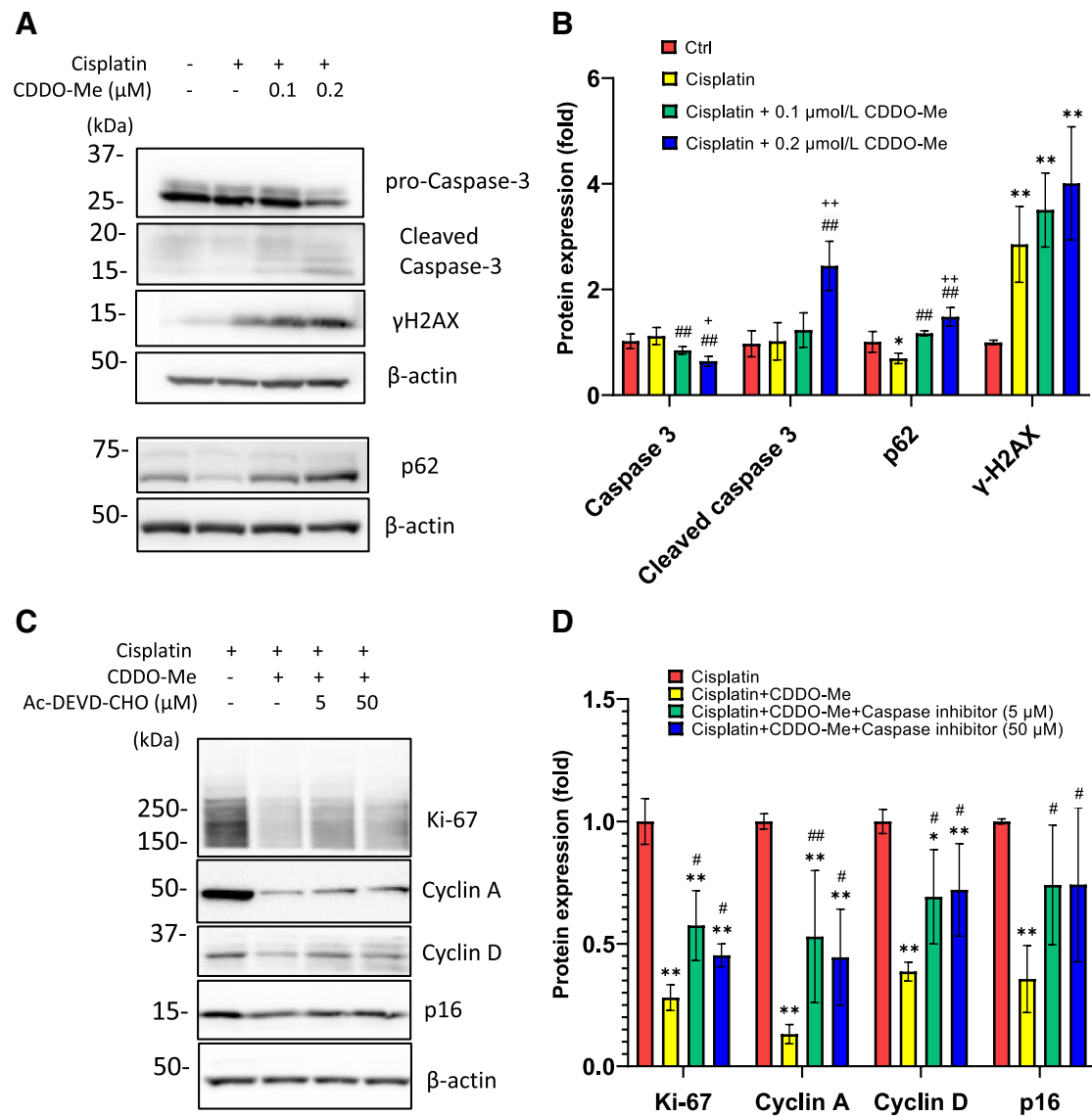
**Fig. 6** CDDO-Me treatment decreases the cell number in cisplatin-exposed HK-2 cells by inducing apoptosis. Cisplatin (20  $\mu\text{mol/L}$ )-exposed HK-2 cells were cultured in medium with or without 0.2  $\mu\text{mol/L}$  CDDO-Me. Cell numbers were counted at 48 h (A).  $n=3$ ,  $**P<0.01$ . Apoptotic cells were detected in cisplatin (20  $\mu\text{mol/L}$ )-

exposed cultures after treatment with or without 0.2  $\mu\text{mol/L}$  CDDO-Me by TUNEL staining (original magnification  $\times 200$ ) (B), and the number of apoptotic cells was expressed as the apoptosis rate (C). Data are presented as mean  $\pm$  SD from three independent experiments.  $**P<0.01$

of a pool of senescent cells following cisplatin treatment. This was demonstrated by the upregulation of cyclin-dependent kinase inhibitors, such as p16<sup>INK4a</sup>, p21<sup>Waf1/Cip1</sup>, and phosphorylated Rb in cells exposed to cisplatin for 72 h. An increase in p21<sup>Waf1/Cip1</sup> and p16<sup>INK4a</sup> induces G2/M arrest through the phosphorylation of Rb [40]. Flow cytometric analysis showed an increase in the number of S-phase cells in cisplatin-treated cultures. Thus, cisplatin treatment seemed to be associated with S- or G2/M-phase arrest by regulating p16<sup>INK4a</sup> and/or p21<sup>Waf1/Cip1</sup>. Cisplatin stalls cell proliferation by damaging DNA and inhibiting DNA synthesis, which is cytotoxic during the S phase [41]. In fact,  $\gamma$ -H2AX, a DNA damage marker, was elevated in cisplatin-treated cells. Furthermore, cisplatin treatment induced the upregulation and secretion of IL-6 and IL-8 in the medium. These results indicate that cisplatin-exposed cells undergo premature cellular senescence, consistent with the results from previous studies [30, 31].

This study showed that CDDO-Me treatment decreased senescence markers in PTCs exposed to cisplatin. Upon activation by CDDO-Me, Nrf2 translocates to the nucleus and induces the transcription of antioxidant enzyme genes via the antioxidant response element, ARE [42]. This was demonstrated by the upregulation of the antioxidant enzymes following the CDDO-Me treatment of the cisplatin-treated HK-2 cells. Aleksunes et al. elucidated using *in vivo* studies that cisplatin accelerates cell proliferation and secretion of IL-1 $\beta$ , IL-6, and TNF- $\alpha$  in the kidney; importantly, this phenotype was inhibited by CDDO-Im, an analog of CDDO-Me, through the increase of Nrf2 signaling [43]. Treatment with the antioxidant N-acetylcysteine ameliorates cellular senescence in PTCs induced by multiple cisplatin treatments [32]. Furthermore, an *in vitro* study using human PTCs showed that the upregulation of the antioxidant genes by Nrf2 enhanced cell viability [44]. These studies suggest that removing oxidative stress via the Keap1-Nrf2 pathway protects PTCs from cellular senescence.





**Fig. 7** CDDO-Me induces apoptosis in cisplatin-treated HK-2 cells. HK-2 cells were exposed to 20  $\mu\text{mol/L}$  cisplatin for 6 h, followed by treatment with or without CDDO-Me (0.1 or 0.2  $\mu\text{mol/L}$ ). Apoptotic marker proteins were analyzed by western blotting (A), and protein bands were quantified by densitometry (B).  $n=4-6$ ,  $*P<0.05$ ,  $**P<0.01$  vs. control (Ctrl),  $\#P<0.05$ ,  $\#\#P<0.01$  vs. cisplatin,  $+P<0.05$ ,  $++P<0.01$  vs. cisplatin+CDDO-Me (0.1  $\mu\text{mol/L}$ ). Cisplatin-

atin-exposed cells were treated with 5 or 50  $\mu\text{mol/L}$  Ac-DEVD-CHO, a caspase inhibitor, for 60 min prior to adding 0.2  $\mu\text{mol/L}$  CDDO-Me to the medium. Senescence markers were analyzed using western blotting (C) and the expression was quantified using densitometry (D).  $n=4-5$ ,  $*P<0.05$ ,  $**P<0.01$  vs. cisplatin,  $\#P<0.05$ ,  $\#\#P<0.01$  vs. cisplatin + CDDO-Me

Overdosing with CDDO-Me induces apoptosis [28]. Eliminating the senescent cells by activating apoptosis has a protective effect on tissue aging [16–18]. Therefore, we evaluated whether CDDO-Me induced apoptosis and reversed senescence via an apoptotic mechanism. We showed that CDDO-Me decreased the cell number, which occurs through the apoptotic mechanism. CDDO-Me treatment increased the level of cleaved caspase-3 and the number of TUNEL-positive cells. Furthermore, CDDO-Me accelerated the phosphorylation of H2AX, which occurs in

response to DNA damage and induces apoptosis by mitochondrial cytochrome C release [45]. Additionally, CDDO-Me increases p62/SQSTM1 expression, which plays a critical role in both autophagy and apoptosis [46, 47]. Inhibition of apoptosis suppressed the reversal effect of CDDO-Me against cisplatin-induced cellular senescence-like alterations. Therefore, CDDO-Me induces apoptosis and suppressed cellular senescence in cisplatin-treated cells. The activation of apoptosis could contribute to the removal of senescent cells and improved kidney functions.

The use of only one cell line was a limitation of this study. HK-2 cells were used in this study, because senescence could be easily induced by treating the HK-2 cells with D-serine, indoxyl sulfate, and hydrogen peroxide [36–38]. Additionally, HK-2 cells maintain biochemical properties similar to that in the *in vivo* proximal tubule cells [48, 49]. Therefore, this model was well-suited for this study. However, HK-2 cells lack expression of organic anion transporter 1, 3, and organic cation transporter 2, which are important for drug metabolism [50]. Therefore, drug metabolism in HK-2 might not be completely similar to that in the *in vivo* state. It is unclear whether CDDO-Me removes senescent cells selectively *in vivo*; therefore, we cannot conclude that activation of apoptosis by CDDO-Me protects against kidney injury in humans. Further *in vivo* studies are needed to evaluate the protective effect of CDDO-Me against cellular senescence in PTCs.

In conclusion, our results demonstrated a beneficial effect of CDDO-Me on cellular senescence in HK-2 cells. It is hypothesized that this beneficial effect is related to Nrf2 activation. CDDO-Me appears to be a candidate therapeutic for AKI. Our findings also showed that CDDO-Me induced apoptosis in cisplatin-treated HK-2 cells and potentially protects the kidneys from cellular senescence. Future studies are needed to assess the effects of CDDO-Me-induced apoptosis on cellular senescence and kidney function in AKI animal models.

**Supplementary Information** The online version contains supplementary material available at <https://doi.org/10.1007/s11010-021-04295-y>.

**Author contributions** YK, MO, RN, and NI provided conception and design of research; YK, AI, FK, and AM performed experiments; YK, AM, and MY analyzed data; YK, FK, MO, TT, TI, MK, and NI interpreted results of experiments; YK, AM, and MY prepared figures; YK drafted manuscript; YK, MO, TT, TI, MK, RN, and NI edited and revised manuscript; YK, AI, FK, MO, AM, MY, TT, TI, MK, RN, and NI approved final version of manuscript.

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**Data availability** The data used to support the findings of this study are included within the article. All data generated or analyzed during this study are included in this published article and its Supplementary Information.

**Code availability** Not applicable.

## Declarations

**Conflict of interest** Not applicable.

**Ethical approval** Not applicable.

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