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# LncRNA OIP5-AS1 reduces renal epithelial cell apoptosis in cisplatin-induced AKI by regulating the miR-144-5p/PKM2 axis



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# ABSTRACT

Background: The abnormal expression of long non-coding RNA (lncRNA) Opa-interacting protein 5 antisense RNA 1 (OIP5-AS1) has been observed in many human cancers and the underlying mechanisms have been well studied. However, the function of OIP5-AS1 in acute kidney injury (AKI) remains unclear.

*Methods*: To explore the role of OIP5-AS1 in the progression of AKI, the cisplatin-induced AKI mouse and cell model were established. To confirm the potential protective effect of OIP5-AS1 during cisplatin-induced AKI, rescue experiments were performed. Targetscan was used to predict the potential targets of miR-144-5p. To further determine whether the effect of miR-144-5p during cisplatin-induced AKI was mediated by PMK2, the recuse experiments using PMK2 overexpressing vector was applied.

Results: OIP5-AS1 was significantly downregulated both in cisplatin-induced AKI mice and human renal tubular cell line HK-2 cells. Moreover, overexpression of OIP5-AS1 efficiently promoted cell growth and reduced cisplatin-induced apoptosis of HK-2 cells. Furthermore, OIP5-AS1 was identified as a sponge of miR-144-5p, and upregulation of miR-144-5p could significantly reverse overexpression of OIP5-AS1-induced protective effect on the damage of cisplatin to HK-2 cells. In addition, pyruvate kinase M2 (PKM2) was found to be a direct target of miR-144-5p, and overexpression of PKM2 efficiently reversed the effect of miR-144-5p mimics on the damage in cisplatin-stimulated HK-2 cells.

Conclusions: OIP5-AS1 reduced the apoptosis of cisplatin-stimulated renal epithelial cells by targeting the miR-144-5p/PKM2 axis, which extended the regulatory network of lncRNAs in cisplatin-induced AKI and also provided a novel therapeutic target for AKI treatment.

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#### At a glance commentary

#### Scientific background on the subject

Various lncRNAs have been reported to be involved in the development and progression of acute kidney injury (AKI). However, the biological role and underlying mechanisms of lncRNA OIP5-AS1 in AKI remain unknown.

#### What this study adds to the field

This study demonstrated that OIP5-AS1 reduced the apoptosis of cisplatin-induced renal epithelial cells by targeting the miR-144-5p/PKM2 axis. These findings imporved our understanding of the pathogenesis of cisplatin-induced AKI and provided a potential therapeutic target for AKI treatment.

Acute kidney injury (AKI) is not a single disease but a loose collection of multiple syndromes such as sepsis, cardiorenal syndrome, and urinary tract obstruction [1]. AKI occurs in approximately 20% of hospitalized patients and more common in critically ill patients [2]. AKI causes about 2 million deaths every year, and increasing evidence indicated that the incidence of AKI in intensive care is more than 50% of patients [3]. The pathophysiological process of AKI includes acute tubular epithelial cell damage, excessive inflammation and acute vascular dysfunction [4]. During AKI, the loss of kidney function can be evaluated via monitoring the changes of renal functionrelated markers, such as serum creatinine (SCr) and blood urea nitrogen (BUN) levels [5,6]. Although clinical treatments are available, the development of more efficient therapies for AKI is still urgent. Better understanding of the underlying molecular mechanisms in the progression of AKI would contribute to the identification of potential therapeutic targets.

Long chain non-coding RNAs (lncRNAs) are a class of noncoding RNAs with approximately 200 nucleotides in length and lack protein coding function, but can regulate a series of biological processes in human eukaryotic cells [7]. LncRNA OIP5-AS1 (Opa-interacting protein 5 antisense RNA 1), has been found to be abnormally expressed and closely associated with a variety of progressions in various human cancers. For instance, OIP5-AS1 promotes the resistance of doxorubicin through targeting the miR-137-3p/PTN axis in osteosarcoma [8]. Overexpression of OIP5-AS1 can inhibit cell viability, colony formation, invasion and migration, while induce cell cycle arrest at G1 phase and apoptosis of multiple myeloma cells in vitro, and can also suppress tumorigenesis in vivo [9]. OIP5-AS1 promotes cell proliferation, migration and induce angiogenesis through regulating the miR-3163/VEGFA axis in hepatocellular carcinoma [10]. Overexpression of OIP5-AS1 can aggravate the progression through targeting the miR-153-3p/ZBTB2 axis in gastric cancer cells [11]. In pancreatic cancer, OIP5-AS1 promotes cell growth by sponging miR-342-3p via the AKT/ERK pathway [12]. In addition, the expression of OIP5-AS1 is also closely related to the prognosis of cancer patients, in which upregulation of OIP5-AS1 predicts a poor prognosis of patients with thyroid cancer due to its regulation of cell proliferation and migration [13].

These evidences confirmed the important functions of OIP5-AS1 in human cancers. Interestingly, a recent study reported that OIP5-AS1 could induce epithelial-to-mesenchymal transition (EMT) and renal fibrosis by binding to miR-30c-5p in diabetic nephropathy [14]. Meanwhile, EMT played essential roles in the pathogenesis of renal fibrosis [15], and the mechanism of chronic kidney disease (CKD) post-AKI is characterized by progressive renal fibrosis [16]. Hence, we thought that OIP5-AS1 might play a potential role in AKI, and paid our attentions on it.

# Materials and methods

#### The mouse model of cisplatin-induced AKI

A total of 20 BALB/c mice (SPF, male, 1–8 weeks old and approximately 20 g) were purchased from the Institute of Medical Laboratory Animals, Chinese Academy of Medical Sciences. Mice were kept under the following conditions: normal grade, free access to food and drink. The cisplatin-induced AKI mice model was established as previously described [17]. Briefly, mice were intraperitoneally injected with 20 mg/kg cisplatin and kept for different time periods, while mice in the vehicle group were injected with equal amount of saline simultaneously. All experiment procedures were performed following the Animal Experimental Guide of the First Affiliated Hospital of Zhengzhou University and approved by the First Affiliated Hospital of Zhengzhou University.

#### The measurement of renal function and tissue morphology

The renal function of mice was evaluated by the serum creatinine and BUN level of mice using the respective detection assay kit (built, Nanjing, China). For the observation of renal tissue morphology, the renal tissues were removed from mice and fixed with 10% formaldehyde aqueous solution, then paraffin-embedded, sliced and strained by hematoxylin-eosin (HE). Subsequently, the pathological changes were observed using a light microscope. In addition, the score of renal tissue damage was evaluated according to the percentage of damaged tubules as previously reported [18]. 0 indicates no obvious damage.1 indicates the damage was less than 25%. 2 indicates the damage was 50–75%. 4 indicates the damage was more than 75%.

#### Cell culture and treatment

Human renal tubular cell line HK-2 was obtained from the American Type Culture Collection (ATCC, USA). Cells were cultured in DMEM medium containing 10% fetal bovine serum at 37 °C with 5% CO<sub>2</sub>. To construct the AKI cell model, different concentration of cisplatin ( $10^{-9}$ ,  $10^{-8}$ ,  $10^{-7}$  and  $10^{-6}$  M) was added into cultured HK-2 cells for 24 h as previously reported [19].

#### Cell transfection

Cell transfection was performed using the RNAi Max and Lipofectamine 3000 with Plus Reagent. The cDNA sequence of OIP5-AS1 was amplified and cloned into the expression vector



Fig. 1 Altered expression of OIP5-AS1 and miR-144-5p in cisplatin-induced AKI model. The mice were induced with or without cisplatin for 1 d, 3 d and 7 d. (A–C) The serum level of creatinine (A) and BUN (B). (C) The morphology of renal tissues was evaluated by HE staining assay. Scale bar = 50  $\mu$ m. (D) The tubular injury score. (E and F) The mRNA level of OIP5-AS1 (E) and miR-144-5p (F) were evaluated by qRT-PCR. (G and H) The HK-2 cells were treated with or without different concentrations of cisplatin for 24 h. The mRNA expression level of OIP5-AS1 (G) and miR-144-5p (H) was evaluated by qRT-PCR. \*p < 0.05, \*\*p < 0.01.

pcDNA3.1 (ABM, Canada) to construct the OIP5-AS1 overexpressing plasmid. MiR-144-5p mimics and miR-NC were obtained from RiboBio (Guangzhou, China). For overexpression of PMK2, the cDNA sequence of PMK2 was also amplified and cloned into pcDNA3.1 vector. The empty pcDNA3.1 vector was used as the negative control.

# qPCRs

Total RNAs were extracted from cultured cells using TRIzol reagent (Invitrogen, Carlsbad, CA). PrimeScript Reverse Transcriptase (TaKaRa) was used for reverse transcription. Then qRT-PCR assay was performed with the platform LightCycler 480 (Roche, Basel, Switzerland). The relative expression of targets was calculated using the  $2^{-\Delta\Delta CT}$  method with GAPDH and U6 as the internal reference. The primer sequences were

as follows: miR-144-5p forward: 5'-TGCCGTTTGGCCATTG-TAAAAC-3', reverse: 5'-CAAGTCCAGGGTCCGAGGT-3'; OIP5-AS1 forward: 5' CAAGAATCCCAGGGCTGATA 3', reverse: 5' GATGTTGGGAAGCATCTGGT 3''; GAPDH forward: 5'-CCACC-CAGAAGACTGTGGAT-3', reverse: 5'-TTCAGCTCAGGGAT-GACCTT-3'; U6 forward: 5'-CTCGCTTCGGCAGCACA-3', reverse: 5'-AACGCTTCACGAAATTGCGT-3'.

# Western blot

Total proteins were isolated from cultured cells using RIPA lysis buffer (Beyotime, Nantong, China). Protein concentrations were determined using the BCA Protein Assay Kit. Then approximately equal amount of protein samples were separated by 12% SDS-PAGE and transferred onto PVDF membranes. After blocking in 3% skim milk, the membranes were incubated with



Fig. 2 The effect of overexpression of OIP5-AS1 on the cell viability and apoptosis of cisplatin-stimulated HK-2 cells. (A) The transfection efficiency of pc-DNA-OIP5-AS1 and pcDNA-vector was evaluated by qRT-PCR. (B) HK-2 cells were transfected with pc-DNA-OIP5-AS1 and pcDNA-vector, then induced with cisplatin for 24, 48, 72 and 96 h. Cell viability was evaluated by CCK-8 kit. (C–E) The HK-2 cells were transfected with pc-DNA-OIP5-AS1 and pcDNA-vector, then induced with cisplatin for 24 h. (C) The apoptosis rate was evaluated by flow cytometry. (D) The protein expression levels of apoptosis-related makers C-caspase 3 and C-caspase 9 were detected by Western blot. (E) The protein expression levels of Bax and Bcl-2 were evaluated by Western blot. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001 vs. pcDNA-vector (–) CP group.

primary antibodies at 1:500 against PMK2, C-caspase 3, C-caspase 9, Bax and Bcl-2 at 4 °C overnight. After blocking, the membranes were submitted to horseradish peroxidase-conjugated secondary antibody for 1 h.  $\beta$ -actin (1:2000) was used as the internal reference. All antibodies were obtained from Abcam, Cambridge, MA. Signals were detected using the EasyBlot ECL Kit (Sangon Biotech, Shanghai, China), and the grey-scale value of target proteins was evaluated using the ChemiDoc XRS System.

#### Luciferase reporter assay

StarBase (http://starbase.sysu.edu.cn/, v2.0) [20] and Targetscan database (http://www.targetscan.org/vert\_72/, v7.2) [21] were used to predict the potential targets of OIP5-AS1 and miR-144-5p, respectively. The fragment of OIP5-AS1(GACATTC) and PMK2 (CTATAGTA) wild type (WT) for miR-144-5p binding site were cloned into the firefly luciferase gene in the pMIR Basic vector (OBiO Biology, Shanghai).



Fig. 3 **OIP5-AS1** acts as a sponge of miR-144-5p. (A) The putative binding site between OIP5-AS1 and miR-144-5p was predicted by Starbase database. (B) HK-2 cells were transfected with miR-144-5p mimics or miR-NC, and the mRNA level of miR-144-5p was evaluated by qRT-PCR. (C) The relative luciferase activity of WT or MUT OIP5-AS1 was detected by the dual luciferase assay system. (D) The enrichment fold of OIP5-AS1 was evaluated by RNA pull down assay. (E) HK-2 cells were transfected with pc-DNA-OIP5-AS1 or pcDNA-vector, the mRNA expression of miR-144-5p was evaluated by qRT-PCR. \*p < 0.05, \*p < 0.01, \*\*p < 0.001.

In addition, the OIP5-AS1 (GATCGTC) and PMK2 mutant plasmids (CTACTATA) were designed in the putative miR-144-5p binding sites. Then the luciferase reporter plasmids were co-transfected with miR-144-5p mimics or miR-NC into HK-2 cells using the Lipofectamine 3000 kit. After transfection for 48 h, cells were lysed, and the relative luciferase activity was detected using a dual luciferase assay system.

# RNA pull-down assay

RNA pull down assay was performed as previously described [22]. The probe of WT and MUT miR-144-5p was biotinylated (Sangon, Shanghai, China) and transfected into HK-2 cells (WT miR-144-5p: 5'-GGAUAUCAUCAUAUACUGUAAG-3'; MUT miR-144-5p: 5'-CCTUAUCAUCAUAUAGACATTC-3'). After 48 h, cells were collected and lysed, then incubated with Dynabeads M–280 streptavidin (Invitrogen, Carlsbad, CA, USA) for 10 min. Finally, the enriched RNA was analyzed by qRT-PCR.

#### Apoptosis analysis

Cell apoptosis was evaluated by flow cytometry after treatment using the Annexin V: FITC Apoptosis Detection Kit II (BD Biosciences, San Jose, CA) following the manufacturer's instructions. Briefly, after transfection for 48 h, cells were centrifuged, and re-suspended with binding buffer. Then 5  $\mu$ L of Annexin V-FITC and 10  $\mu$ L of propidium iodide (PI) were added. Subsequently, apoptosis rate was evaluated by Flow cytometry (BD Biosciences).

#### CCK-8 assay

To measure cell viability, approximately 5  $\times$  10<sup>3</sup> HK-2 cells were seeded into 96-well plates. Then 100  $\mu L$  of CCK8 solution (Dojindo Molecular Technologies, Japan) was added to each well at 24, 48, 72, and 96 h. Finally, the optical density (OD) at 450 nm was detected using a microplate reader (Bio Tek Instruments, USA).

#### Statistical analysis

All experiments were repeated at least 3 times. Data were presented as the mean  $\pm$  standard deviation (SD). Difference between two groups was determined by t test, and difference among multiple groups was determined by one-way ANOVA using the SPSS 19.0 software. p < 0.05 was considered as the significant threshold.



Fig. 4 MiR-144-5p mimics significantly reversed the effects of overexpression of OIP5-AS1 on cell viability and apoptosis in cisplatin-stimulated HK-2 cells. HK-2 cells were transfected with pcDNA-OIP5-AS1, or ac-transfected with pcDNA-OIP5-AS1 and miR-144-5p mimics, then cells were treated with or without cisplatin. (A) Cell viability for different times was evaluated by CCK-8 kit. (B) The apoptosis rate for 24 h was evaluated by flow cytometry. (C) The protein expression of C-caspase 3 and C-caspase 9 for 24 h was detected by Western blot. (D) The protein expression levels of Bax and Bcl-2 for 24 h were detected by Western blot. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001 vs. control group without cisplatin. ##p < 0.01 vs. pcDNA-OIP5-AS1 group with cisplatin.

# Results

# OIP5-AS1 was significantly downregulated in cisplatininduced AKI model both in vitro and in vivo

To explore the role of OIP5-AS1 in the progression of AKI, the cisplatin-induced AKI mice model was firstly established. The results indicated that the expression levels of renal function-related makers such as serum creatinine (p < 0.001, p < 0.01) [Fig. 1A] and serum BUN (p < 0.001, p < 0.01) [Fig. 1B] were significantly increased in cisplatin-induced AKI mice compared with that in vehicle group, and their expression levels were highest on the third day. Meanwhile, HE staining results in the renal tissues showed that the morphology of renal tissues in the vehicle group was almost normal, while tubular epithelial

cells were edematous and necrotic, and exhibited obvious inflammatory cell infiltration in the cisplatin-induced AKI mice [Fig. 1C]. The injury scores of renal tissues were also evaluated, and the results showed that renal tissue injury scores were significantly increased after treatment with cisplatin at day 1 (p < 0.05), day 3 (p < 0.01) and day 7 (p < 0.001) compared with the vehicle group [Fig. 1D]. These data suggested that the cisplatin-induced AKI mice model was successfully established and could be used for the subsequent experiments. Moreover, the expression of OIP5-AS1 in the renal tissues of cisplatininduced mice was significantly downregulated at day 1 (p < 0.05), day 3 (p < 0.01) and day 7 (p < 0.05) compared with that in the vehicle group [Fig. 1E]. Interestingly, the expression of miR-144-5p was oppositely upregulated at day 3 (p < 0.001) and day 7 (p < 0.01) compared with that in the vehicle group [Fig. 1F]. To confirm this, HK-2 cells were treated with different



Fig. 5 **PMK2 was a target of miR-144-5p.** (A) The putative binding site between miR-144-5p and PMK2 mRNA was predicted by Targetscan. (B) The relative luciferase activity of WT or MUT 3'-UTR of PMK2 was detected by the dual luciferase assay system. (C and D) HK-2 cells were transfected with miR-144-5p mimics or miR-NC, then the mRNA level (C) and protein level (D) of PMK2 were evaluated. (F) HK-2 cells were transfected with pcDNA-OIP5-AS1 or pcDNA-vector, then the mRNA level (E) and protein level (F) of PMK2 were evaluated. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001.

concentrations of cisplatin for 24 h, and a dose-dependent decrease of the expression levels of OIP5-AS1 (p < 0.01, p < 0.001) [Fig. 1G] and a dose-dependent increase of the expression levels of miR-144-5p (p < 0.05, p < 0.01, p < 0.001) [Fig. 1H] were observed in the cisplatin treatment group compared with that in the vehicle group. These results suggested that OIP5-AS1 and miR-144-5p play important roles during the progression of cisplatin-induced AKI.

# Overexpression of OIP5-AS1 promoted cell viability and suppressed apoptosis of cisplatin-induced HK-2 cells

To investigate the specific function of OIP5-AS1, the overexpressing vector pc-DNA-OIP5-AS1 was constructed and transfected into HK-2 cells. The transfection efficiency was evaluated by qRT-PCR assay and the results showed that pc-DNA-OIP5-AS1 significantly increased the expression levels of OIP5-AS1 compared with negative control (pc-DNA-vector) (p < 0.001) [Fig. 2A]. Cisplatin obviously reduced cell viability of pc-DNA-vector transfected HK-2 cells (p < 0.05), while overexpression of OIP5-AS1 markedly increased cell viability of cisplatin-stimulated HK-2 cells (p < 0.05) [Fig. 2B]. In addition, cisplatin significantly induced the apoptosis of pc-DNAvector transfected HK-2 cells (p < 0.05), while overexpression of OIP5-AS1 markedly suppressed the apoptosis of cisplatinstimulated HK-2 cells (p < 0.05) [Fig. 2C]. Furthermore, the expression levels of apoptosis-related proteins were evaluated and the results showed that the expression levels of Ccaspase 3 and C-caspase 9 were all significantly increased in cisplatin-stimulated HK-2 cells transfected with pc-DNAvector (p < 0.001), while overexpression of OIP5-AS1 markedly decreased the expression levels of C-caspase 3 and Ccaspase 9 in the cisplatin-stimulated HK-2 cells (p < 0.01) [Fig. 2D]. Meanwhile, cisplatin significantly increased the expression levels of Bax and decreased the expression levels of Bcl-2 in HK-2 cells transfected with pc-DNA-vector (p < 0.001), while overexpression of OIP5-AS1 obviously reversed the effects of cisplatin on the expression of Bax and Bcl-2 in HK-2 cells (p < 0.01) [Fig. 2E]. These results confirmed a potential inhibitory effect of OIP5-AS1 on cisplatin-induced acute kidney injury.



Fig. 6 Overexpression of PMK2 significantly reversed the effects of miR-144-5p mimics on cell viability and apoptosis in cisplatin-stimulated HK-2 cells. HK-2 cells were transfected with miR-144-5p mimics, or co-transfected with miR-144-5p mimics and pcDNA-PMK2, then induced with or without cisplatin. (A) The cell viability for different times was evaluated by CCK-8 kit. (B) The apoptosis rate for 24 h was evaluated by flow cytometry. (C) The protein expression of C-caspase 3 and C-caspase 9 for 24 h was detected by Western blot. (D) The protein expression levels of Bax and Bcl-2 for 24 h were detected by Western blot. \*\*p < 0.001 vs. control group without cisplatin. \*\*p < 0.01 vs. miR-144-5p mimics group with cisplatin.

#### LncRNA OIP5-AS1 acts as a sponge of miR-144-5p

To explore the underlying molecular mechanisms of OIP5-AS1 in cisplatin-induced AKI, the potential targets of OIP5-AS1 were predicted using Starbase online database, and the results showed that miR-144-5p might be a target of OIP5-AS1 [Fig. 3A]. Then HK-2 cells were transfected with miR-144-5p mimics or miR-NC, and qRT-PCR assay showed that overexpression of miR-144-5p compared with miR-NC (p < 0.001) [Fig. 3B]. To determine the relationship between OIP5-AS1 and miR-144-5p, the fragments of OIP5-AS1 containing wild type(WT) or mutant (MUT) putative binding sites against miR-144-5p were cloned into luciferase reporter plasmid and luciferase reporter assay was performed. The results showed that overexpression of miR-144-5p markedly decreased the relative luciferase activity of WT OIP5-AS1 compared with miR-NC (p < 0.01), while

showed no obvious change of MUT OIP5-AS1 [Fig. 3C]. Furthermore, RNA pull-down assay was performed using biotinylated miR-144-5p and the results showed that OIP5-AS1 was significantly enriched by biotinylated miR-144-5p (p < 0.01), but exhibited no change by biotinylated MUT miR-144-5p [Fig. 3D]. In addition, the expression levels of miR-144-5p were markedly decreased in pcDNA-OIP5-AS1 transfected HK-2 cells compared with that in pcDNA-vector transfected cells (p < 0.01) [Fig. 3E]. These results indicated that OIP5-AS1 might function though targeting miR-144-5p during cisplatin-induced AKI.

# Upregulation of miR-144-5p obviously reversed the effects of overexpression of OIP5-AS1 on cell viability and apoptosis in cisplatin-stimulated HK-2 cells

To confirm the potential protective effect of OIP5-AS1 during cisplatin-induced acute kidney injury, the rescue assays were

performed. Cisplatin treatment significantly inhibited cell viability of HK-2 cells (p < 0.001), and overexpression of OIP5-AS1 markedly attenuated the inhibitory effect of cisplatin on cell viability, while co-transfection with pcDNA-OIP5-AS1 and miR144-5p mimics partially reversed the effect of overexpression of OIP5-AS1 on cell viability in cisplatin-stimulated HK-2 cells (p < 0.01) [Fig. 4A]. For apoptosis, cisplatin treatment significantly induced cell apoptosis of HK-2 cells (p < 0.001), and overexpression of OIP5-AS1 inhibited cisplatin-induced apoptosis (p < 0.01), while co-transfection with pcDNA-OIP5-AS1 and miR144-5p mimics partially reversed the effect of overexpression of OIP5-AS1 on the cell apoptosis in cisplatinstimulated HK-2 cells (p < 0.01) [Fig. 4B]. Meanwhile, the expression of apoptosis-related proteins was detected and the results showed that overexpression of OIP5-AS1 significantly decreased the protein expression levels of C-caspase 3, Ccaspase 9, Bax and increased the expression levels of Bcl-2 in cisplatin-stimulated HK-2 cells (p < 0.05), while cotransfection with pcDNA-OIP5-AS1 and miR144-5p mimics obviously and partially reversed the effects of overexpression of OIP5-AS1 on the expression of these apoptosis-related makers (p < 0.05) [Fig. 4C and D]. These results suggested that upregulation of miR-144-5p could obviously reverse the effects of overexpression of OIP5-AS1 on cell viability and apoptosis in cisplatin-stimulated HK-2 cells.

#### PMK2 was a target of miR-144-5p

Next, Targetscan was used to predict the potential targets of miR-144-5p, and the results showed that there was a potential binding site between miR-144-5p and 3'-UTR of PMK2 mRNA [Fig. 5A]. To validate this, we performed the luciferase reporter assay using either WT or MUT 3'-UTR of PMK2 mRNA lacking the miR-144-5p binding site in HK-2 cells. When miR-144-5p was overexpressed, the relative luciferase activity was significantly reduced in HK-2 cells transfected with the luciferase reporter vector containing WT 3'-UTR of PMK2 (p < 0.01), but exhibited no obvious change with that containing the MUT 3'-UTR of PMK2 [Fig. 5B]. Meanwhile, miR-144-5p mimics decreased the expression levels of PMK2 at both mRNA (p < 0.001) [Fig. 5C] and protein levels (p < 0.001) [Fig. 5D] compared with miR-NC group in HK-2 cells. In addition, overexpression of OIP5-AS1 increased the expression levels of PMK2 at both mRNA (p < 0.001) [Fig. 5E] and protein levels (p < 0.001) [Fig. 5F] compared with negative group (pcDNA-vector) in HK-2 cells. These results indicated that PMK2 was a direct target of miR-144-5p, and regulated by OIP5-AS1 and miR-144-5p.

# Overexpression of PMK2 significantly reversed the effects of miR-144-5p mimics on cell viability and apoptosis in cisplatin-stimulated HK-2 cells

To further explore whether the effects of miR-144-5p during cisplatin-induced AKI were mediated by PMK2, the recuse assay using PMK2 overexpressing vector was performed. For cell viability, miR-144-5p mimics significantly decreased the cell viability of cisplatin-stimulated HK-2 cells (p < 0.05), while co-transfection with miR-144-5p mimics and pcDNA-PMK2 obviously reversed the effect of overexpression of miR-144-5p

on cell viability in cisplatin-stimulated HK-2 cells (p < 0.01) [Fig. 6A]. For apoptosis, miR-144-5p mimics significantly induced apoptosis of cisplatin-stimulated HK-2 cells (p < 0.01), while co-transfection with miR-144-5p mimics and pcDNA-PMK2 obviously reversed the effect of overexpression of miR-144-5p on cell apoptosis in cisplatin-stimulated HK-2 cells (p < 0.01) [Fig. 6B]. Meanwhile, the expression levels of apoptosis-related proteins were evaluated by Western blot, and the results showed that miR-144-5p mimics significantly increased the expression levels of C-caspase 3, C-caspase 9, Bax and decreased the expression levels of Bcl-2 (p < 0.01), while cotransfection with miR-144-5p mimics and pcDNA-PMK2 obviously reversed the effects of miR-144-5p mimics on the expression of these apoptosis-related makers (p < 0.01) [Fig. 6C and D]. These data suggested that the effects of miR-144-5p during cisplatin-induced AKI were partially mediated by PMK2.

# Discussion

The kidneys are the primary targets for toxic effects of various chemical agents, so drug-induced AKI is common in clinical medicine [23]. Cisplatin, a commonly used chemotherapeutic agent, has been demonstrated to have a major limitation due to its severe nephrotoxicity [24]. After a single dose of cisplatin (50–100 mg/m<sup>2</sup>), approximately one-third of the patients develop nephrotoxicity [25,26]. Therefore, cisplatin has been frequently used to induce AKI model to investigate the molecular mechanisms in AKI progression [27]. Here, the cisplatin-induced AKI mice model was established. We found that the expression levels of renal function-related makers including serum creatinine and serum BUN were significantly increased in cisplatin-induced AKI mice, and injury scores of renal tissues were significantly increased after treatment with cisplatin.

In the last decades, various lncRNAs have been identified to be closely associated with the progression of AKI. For instance, HOTAIR is significantly downregulated in kidney tissues in sepsis rats and overexpression of HOTAIR efficiently attenuates renal injury through inhibiting the apoptosis of kidney tissues by targeting the miR-34a/Bcl-2 signaling pathway [28]. CRNDE can reduce sepsis-induced renal injury by inhibiting the activation of the TLR3/NF- $\kappa$ B signaling pathway [29]. Overexpression of NEAT1 can aggravate the LPSinduced renal injury through targeting miR-204 and activating the NF- $\kappa$ B signaling pathway [30]. The expression of LINC00520 was markedly upregulated and overexpression of LINC00520 can aggravate ischemia/reperfusion induced acute kidney injury by targeting the miR-27b-3p/OSMR axis [31]. Downregulation of TUG1 contributes to the progression of sepsis-associated acute kidney injury through targeting the miR-142-3p/sirtuin 1 axis and regulating the NF-kB signaling pathway [32]. PVT1 has been identified to promote LPSinduced septic acute kidney injury through modulating the TNFα and JNK/NF-κB pathways in human renal epithelial HK-2 cells [33]. In addition, several lncRNAs including PRINS, PlncRNA-1, XIST and CCAT1 have been demonstrated to play essential roles during the progression of AKI [34-37]. These reports confirmed the crucial roles of lncRNAs during AKI development. Although previous studies indicated that OIP5AS1 might play potential role in AKI [14,15], further details remain unclear. In this study, we found that OIP5-AS1 was significantly downregulated in both the renal tissues of cisplatin-induced AKI mice model and cisplatin-induced HK-2 cell model. Moreover, overexpression of OIP5-AS1 could efficiently attenuate cisplatin-induced renal injury through promoting cell viability and inhibiting apoptosis of HK-2 cells. These data demonstrated that OIP5-AS1 participated in AKI through regulating renal epithelial cell apoptosis, suggesting that OIP5-AS1 might be a potential therapeutic target for AKI.

It has been reported that lncRNAs always function as competing endogenous RNAs (ceRNAs) of microRNAs (miR-NAs) in human diseases [38]. In this study, luciferase reporter assay and RNA pull-down assays were performed to confirm the binding relationship between OIP5-AS1 and miR-144-5p. Previous studies have demonstrated that miR-144-5p plays important roles in various human diseases including lung squamous cell carcinoma, bladder cancer and sickle cell disease [39-41]. MiR-144-5p limits the formation of experimental abdominal aortic aneurysm by inhibiting M1 macrophage-mediated inflammation through downregulating TLR2 and OLR1 [42]. MiR-144-5p inhibits cell proliferation, invasion and migration of colorectal cancer cells through targeting CDKL1 [43]. Inhibition of miR-144-5p can enhance the radio-sensitivity of non-small cell lung cancer cells through directly targeting ATF2 [44]. In addition, abnormal expression of miR-144-5p was also observed in a series of other human cancers including bladder cancer, oral cavity squamous cell carcinoma, esophageal cancer, and lung adenocarcinoma [45-48]. Here, we also found that the expression of miR-144-5p was markedly upregulated in both the renal tissues of cisplatin-induced AKI mice model and cisplatin-induced HK-2 cell model. Meanwhile, overexpression of miR-144-5p obviously reversed the effects of overexpression of OIP5-AS1 on cell viability and apoptosis in cisplatin-stimulated HK-2 cells. Specifically, overexpression of OIP5-AS1 significantly decreased the expression levels of C-caspase 3, C-caspase 9, Bax and increased the expression levels of Bcl-2, while additional miR-144-5p partially reversed the effects of overexpression of OIP5-AS1. Our findings demonstrated that the function of OIP5-AS1 during AKI was partially mediated by miR-144-5p.

PKM2 (pyruvate kinase M2 isoform) has been identified to play an anti-apoptotic role in the progression of many human cancers. PKM2 is a rate-limiting terminal glycolytic enzyme and can catalyze the last step of glycolysis [49]. Increasing evidence has demonstrated that PKM2 acts as an antiapoptotic factor against the AKT signaling pathway, which controls the cell survival and apoptosis [50]. It was reported that PKM2 could efficiently attenuate heat stress-induced damage in mouse kidneys through regulating the AKT and HIF-1a signaling pathway [51]. Knockdown of PKM2 can induce autophagic cell death in prostate cancer cells through affecting the AKT/mTOR signaling pathway [52]. PKM2 serves as the target of miRNAs to participate in the progression of human cancers. For example, miR-338-3p can suppress the growth, metastasis and glycolysis of glioma cells via downregulating PKM2 [53]. MiR-374b re-sensitizes hepatocellular carcinoma cells to sorafenib therapy through antagonizing PKM2mediated glycolysis pathway [54]. PKM2, targeted by lncRNA FEZF1-AS1, can promote tumor proliferation and metastasis in colorectal cancer [55]. In addition, it was demonstrated that mitochondrial PKM2 could regulate oxidative stress-induced apoptosis through stabilizing Bcl2 [56]. In this study, PKM2 was predicted to be a direct target of miR-144-5p. Furthermore, the relative luciferase activity was significantly reduced in HK-2 cells transfected with the luciferase gene containing WT 3'-UTR of PMK2, but exhibited no obvious change with that containing the MUT 3'-UTR of PMK2. And overexpression of miR-144-5p obviously decreased the expression levels of PKM2, while overexpression of OIP5-AS1 increased the expression levels of PKM2. Moreover, overexpression of PMK2 obviously reversed the effects of miR-144-5p mimics on cell viability and apoptosis in cisplatin-stimulated HK-2 cells. All these data suggested that OIP5-AS1 affected the progression of AKI through regulating the miR-144-5p/PKM2 axis. Although our in vitro experiments indicated that OIP5-AS1 significantly reduced cisplatin-induced renal epithelial cell apoptosis through targeting the miR-144-5p/PKM2 axis, in vivo experiments should also be performed to determine the roles of the OIP5-AS1/miR-144-5p axis in AKI.

#### Conclusion

In summary, we first demonstrated that OIP5-AS1 was significantly downregulated in cisplatin-induced AKI both in vitro and in vivo. Specifically, OIP5-AS1 efficiently reduced the apoptosis of renal epithelial cells in cisplatin-induced AKI through targeting the miR-144-5p/PKM2 axis, suggesting that OIP5-AS1 might be a potential therapeutic target for AKI.

#### **Compliance with ethical standards**

#### Research involving animals

All experiment protocols were performed according to the First Affiliated Hospital of Zhengzhou University Animal Experimental Guide and approved by the First Affiliated Hospital of Zhengzhou University.

#### Informed consent

Not applicable.

# **Conflicts of interest**

The authors declare that they have no conflict of interest.

# Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.bj.2021.07.005.

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