Rapamycin ameliorates chronic intermittent hypoxia and sleep deprivation-induced renal damage via the mammalian target of rapamycin (mTOR)/NOD-like receptor protein 3 (NLRP3) signaling pathway

Wei Liu, Dong Zhao, Xiaofeng Wu, Fang Yue, Haizhen Yang, and Ke Hu

Department of Respiratory and Critical Care Medicine, Renmin Hospital of Wuhan University, Wuhan, Hubei, China

ABSTRACT

Rapamycin inhibits the activation of NOD-like receptor protein 3 (NLRP3) by regulating the mammalian target of rapamycin (mTOR) to treat obstructive sleep apnea-related renal injury. Sleep deprivation (SD) and chronic intermittent hypoxia (CIH) mouse models were used to assess the effects of autophagy in vivo. Compared with the control, SD, and CIH groups, the SD+CIH group had lower body weight and higher levels of blood urea nitrogen (BUN), creatinine, and urinary albumin (U-Alb) (P < 0.05); renal injury and oxidative damage occurred in the SD+CIH group, the kidney cell nucleus ruptured, and morphological structure of the cells was unclear in the SD+CIH group. The SD+CIH group demonstrated increased apoptosis compared with the control, SD, and CIH groups using Western blot analysis. Compared to the control, SD, and CIH groups, the SD+CIH group showed a higher degree of microtubule-associated protein light chain 3 staining. Compared to the SD+CIH group, BUN, creatinine, and U-Alb levels decreased, and apoptosis increased in the SD+CIH+rapamycin group, and the structure of the kidney after rapamycin treatment was well preserved. The mTOR expression was increased in the kidneys of the SD+CIH group. The NLRP3, Gasdermin D (GMDSD), interleukin (IL)-18, IL-1β, and cleaved-caspase-1 protein levels were higher in the SD+CIH group than the SD+CIH+rapamycin group, and the NLRP3, GMDSD, IL-18, IL-1β, and cleaved-caspase-1 mRNA levels were higher in the SD+CIH group than the SD+CIH+rapamycin group. Following rapamycin treatment, pyroptosis was suppressed. Rapamycin ameliorates renal damage by inhibiting the mTOR/NLRP3 signaling pathway.

Introduction

Obstructive sleep apnea (OSA) is the most common sleep-disordered breathing disease. Epidemiological studies have shown that the incidence rate of adults is 9–38% [1]. OSA is characterized by recurrent upper airway collapse, occlusion leading to OSA or hypopnea, chronic intermittent hypoxia (CIH), and sleep deprivation (SD). During intermittent hypoxia, the body produces a large number of oxygen...
free radicals after repeated hypoxia, resulting in damage to various organs. Respiratory awakening leads to the disintegration of normal sleep structure, SD, and a vicious cycle [2]. SD refers to a lack of sleep or a serious lack of sleep time for various reasons. SD can lead to serious health challenges [3], triggering many short and long-term complications and negative effects [4]. SD leads to physiological clock disorders, shortened sleep time, and reduced sleep depth and continuity. SD affects the systemic inflammatory response, causes an imbalance in antioxidant oxidation products, affects the production and metabolism of a variety of cytokines, and aggravates the process of intermittent hypoxia. The pharyngeal airway of patients with OSA collapses during sleep, resulting in hypoxia and termination after awakening.

Recent studies have shown that OSA may increase the incidence of chronic kidney disease [5] and acute kidney injury [6]. The kidneys are not only regulated by activities, diet, temperature, and the endocrine system but is also regulated by circadian genes. Long-time interference with sleep time in rats leads to peroxidation damage in renal tissue. Wisit et al. found that people with short sleep durations were more prone to proteinuria [7]. Intermittent hypoxia induces oxidative stress and activates inflammatory factors. Studies have found that mice with SD often experience multiple organ inflammatory reactions [8]. The kidney is rich in mitochondria, and the mitochondrial respiratory chain and nicotinamide adenine dinucleotide phosphate oxidase are the main sources of reactive oxygen species (ROS), which are considered to be important factors leading to cardiovascular disease and diabetes. These diseases can also lead to acute kidney injury [9], which indicates that the kidneys are highly susceptible to oxidative stress injury [10]. Continuous positive airway pressure is the main therapy for OSA [11]; however, there is no good way to deal with renal injury caused by OSA. Autophagy refers to the degradation of the cytoplasmic components of the lysosome. Generally, autophagy is an important protective mechanism of the body, and enhancing the autophagy process can appropriately reduce the oxidative damage of the body; however, over-activated autophagy will also lead to cell death and organ damage. Yang et al. found that oxidative stress can mediate the change in mammalian target of rapamycin (mTOR), an important target of autophagy [12]. During the occurrence of renal injury, the level of intracellular autophagy increases to a certain extent with the aggravation of oxidative stress injury. The NOD-like receptor protein 3 (NLRP3) inflammasome is an important target of the classical focal death pathway. The level of NLRP3 is significantly increased in patients with OSA [13], mTOR is an important target mediating autophagy, inhibition of the mTOR signaling pathway can significantly reduce the level of NLRP3 and reduce organ damage caused by OSA [14], and the mTOR/NLRP3 pathway can be used as a potential intervention target to increase autophagy and then reduce cell death to delay renal injury [15].

Rapamycin, an autophagy activator, is involved in various cellular processes by inhibiting mTOR, such as protein translation, ribosome biogenesis, mitochondrial oxygen consumption, proliferation, and differentiation [16]. Rapamycin modulates multiple cellular signaling pathways in OSA. It can improve renal function, decrease proteinuria levels, and alleviate renal tubulointerstitial inflammation, and interstitial fibrosis [17]. In the mesangial proliferative rat model, rapamycin significantly reduced various symptoms of glomerulonephritis [18]. Therefore, we speculate that rapamycin may play an important role in the prevention and treatment of nephropathy through its anti-inflammatory and anti-proliferative effects. However, studies on the influence of rapamycin on OSA-induced renal injury in vivo are limited.

Although the evidence demonstrates that OSA-induced pyroptosis occurs when oxidative stress is overproduced, the mechanism of pyroptosis leading to an inflammatory response in OSA-related renal injury remains unknown. This study aimed to explore whether rapamycin inhibits the activation of NLRP3-dependent pyroptosis by regulating mTOR, to reduce OSA-related renal injury, and to provide a new basis for the diagnosis and treatment of OSA-related renal injury.
**Materials and methods**

**Mouse model experimental protocol**

Female C57BL/6 J mice (6–8 weeks old) were randomly allocated into the following six groups (six mice in each group): control, SD, CIH, SD +CIH, rapamycin, and SD+CIH+rapamycin groups. The SD+intermittent hypoxia device used was a rectangular cabin (24.5 × 21.0 × 19.7 cm) made of 4-mm-thick plexiglass (Biospherix, USA, A84XOV). The cabin was equipped with sleep deprivation-related equipment, such as a metal sweeping rod, electric motor, and frequency modulation knob (1 min/cycle). The metal scanning rod sweeps from one side of the box to the other side to provide the mouse with relatively mild intermittent tactile stimulation. The animals in the SD groups were deprived of sleep for 18 h (4:00 PM to 10:00 AM). Mice can be prevented by gently crossing the rod. Simultaneously, two holes with a diameter of 5 mm were placed in the center of the sidewall of the cabin to input hypoxic gas and air to prepare the hypoxic animal cabin. The mice were exposed to intermittent hypoxia for 8 h/day (08:00 AM to 4:00 PM) [19]. Oxygen concentrations in the chamber were monitored using a heated zirconium sensor connected to solenoid valves that controlled the flow of oxygen and nitrogen. The valves were operated using a microprocessor-controlled timer. The mice were exposed to 30s of 5% inspired oxygen every 2 min. Control mice were housed under normoxic conditions, similar to the chamber. Mice were acclimated for 6 days before starting the experiments, and on the 7th day, the mice in each group were placed in a SD combined with intermittent hypoxia chamber to adapt to the experimental environment for 24 h, and the chamber started operating on days 8–28. In the SD+CIH group, 18 h (4:00 PM to 10:00 AM) of SD and 8 h (8:00–16:00) of intermittent hypoxia were performed every day [20]. In all rapamycin groups, mice received an intraperitoneal injection of rapamycin (MedChemExpress, USA) at a concentration of 4 mg/kg on days 8–28 (once/day). After the study, the mice were sacrificed under general anesthesia; serum and urine [21] were collected; and the left kidney tissue was excised for the following experiment.

**Histological assessments**

The harvested kidneys were fixed in 4% paraformaldehyde for 24 h. The sections were then embedded in paraffin, cut into sections of 5-μm thickness, and stained with hematoxylin and eosin (HE) [22]. Histological assessment was conducted using a light microscope (Olympus, Tokyo, Japan). Paraffin sections were dewaxed, hydrated, and rinsed with distilled water. According to the requirements of the first antibody to tissue antigen repair, the slices were dripped with endogenous peroxidase blocker and rinsed with phosphate-buffered saline (PBS), then the first antibody was dropped on the section, and after incubated at 25°C and rinsed with PBS; primary antibody enhancer (Proteintech, Wuhan, China) was added to the section and incubated at room temperature, then washed with PBS; and secondary antibody-horseradish peroxidase (HRP) polymer (Proteintech, Wuhan, China) reagent was added, incubated at room temperature, and washed. The following antibodies were used: LC3 (Abmart, PA9415); mTOR (Abmart, T55306); and NLRP3 (Cell Signaling Technology, 15101).

**Oxidative stress analysis**

Commercial kits for detection of malondialdehyde (MDA), superoxide dismutase (SOD), catalase (CAT), and glutathione (GSH) were obtained from Nanjing Jiancheng Bioengineering Institute (Nanjing, China). The levels of MDA (A003-2) and CAT (A007-1-1) were monitored by measuring the absorbance at 532 nm and 240 nm, respectively. The enzymatic activities of SOD (A001-3) and GSH (A006-2-1) were measured at 450 and 405 nm, respectively.

**Biochemical assays**

Blood urea nitrogen (BUN), creatinine, and urinary albumin (U-Alb) contents were measured using the respective kits according to the manufacturer’s instructions. BUN (C013-2-1), creatinine (C011-2-1), and urine microalbumin assay kits
(E038-1-1) were purchased from Nanjing Jiancheng Institute of Biotechnology (Nanjing, China). BUN, creatinine, and U-Alb levels were measured at 640 nm, 546 nm, and 340 nm, respectively.

**RNA isolation and quantitative real-time PCR**

Total RNA from mouse kidneys was isolated using TRIzol (Invitrogen), and reverse transcription was performed using 500 ng of total RNA in the first-strand cDNA synthesis reaction with PrimeScript RT reagent Kit (Takara). The expression of each gene was determined using an ABI Prism 7500 PCR system (Applied Biosystems). The reaction conditions were as follows: pre-denaturation stage: 95°C for 30s, 1 cycle; amplification: 95°C for 5s, 60°C for 30s, 40 cycles; and dissolution curve analysis stage: 95°C for 15s, 60°C for 1 min, and 95°C for 15s. The primer sequences (forward and reverse) used in this study were as follows: NLRP3 (forward: CCCTGTCTCTATCAGG; reverse: CATCATGAGATGGAGT), Caspase-1 (forward: TTTCCGCAAGGTTCGATTTC; reverse: GCATCTGTCTCTAGGG), GSDMD (forward: GTGTTGCAACCTGGTCTTACAGG; reverse: CATGGCATGAGAAGTGAAG); IL-1β (forward: CGCTGCTCTCATACATCG; reverse: CTGCCTTGGAGAATCGG); cleaved-caspase-1 (forward: AGCTACGAATCTCCGACCAC; reverse: GGACGACAGTGAGATA); IL-18 (forward: GCATCTGTCTACTGCTC; reverse: CATGACGTGGGATA); Beclin 1 (forward: TCAGGAGAGTATATGTACGC; reverse: GGACGACAGTGAGATA); and mTOR (forward: GGGTGCATTATCTCTAC; reverse: TCTTCATTGACCAAGGAAATCGG). The gene expression was determined using the 2− ΔΔCT method.

**Western blotting**

Approximately 30 mg of mouse kidney tissue was homogenized, the total protein was extracted using RIPA buffer, and the protein concentration was determined using a total protein assay kit. NLRP3, Gasdermin D (GMDSD), cleaved-caspase-1, interleukin IL-1β, IL-1β, microtubule-associated protein light chain 3 (LC3), sequestosome 1 (P62), Beclin 1, mTOR, and phospho-mammalian target of rapamycin (p-mTOR) protein was measured by Western blotting using monoclonal mouse anti-human NLRP3 (Cell Signaling Technology, 15101) and rabbit anti-human GMDSD (Cell Signaling Technology, 97558), cleaved-caspase-1 (Cell Signaling Technology, 4199), IL-18 (Cell Signaling Technology, 57058), IL-1β (Cell Signaling Technology, 12703), LC3 (Cell Signaling Technology, 12741), P62 (Cell Signaling Technology, 16177), Beclin 1 (Cell Signaling Technology, 3495), mTOR (Cell Signaling Technology, 2983), and p-mTOR (Cell Signaling Technology, 5536). The proteins were separated on 10% polyacrylamide gels and transferred onto polyvinylidene difluoride membranes, primary antibody (1:1000) at 4°C overnight with gentle shaking and incubated with an appropriate HRP-labeled secondary antibody for 90 min at room temperature on a shaker. Finally, the membranes were again washed three times with 1× TBST and assayed using a ChemiDoc XRS gel imaging system (Bio-Rad, Hercules, CA, USA). The protein expression levels were quantified using Image J software (NIH Image, Bethesda, MD, USA).

**Transmission electron microscopy**

Kidney tissues were fixed in 2.5% glutaraldehyde and 4% paraformaldehyde dissolved in 100 mM sodium phosphate (pH 7.2) [23]. The tissues were washed using 100 mM sodium cacodylate (pH = 7.4), fixed in 2% osmium tetroxide, and washed again. The fixed tissues were dehydrated across an ethanol gradient and propylene oxide and then embedded in epoxy resin (Tab 812 Resin; Canemco Inc., Montreal, QC, Canada). The resulting ultrathin (60–70 nm) sections were counterstained with uranyl acetate and lead citrate and then viewed using a Hitachi 7600 transmission electron microscope (Hitachi High-Technologies America, Inc., Schaumburg, IL, USA) equipped with a MacroFire monochrome progressive scan CCD camera (Optronics, 10 Inc., Muskogee, OK, USA) with AMTv image capture software (Advanced Microscopy Techniques, Inc., Danvers, MA, USA).

**Statistical analysis**

Data are expressed as mean ± standard deviation. All statistical analyses were performed
using GraphPad Prism 6.0 (GraphPad Software, La Jolla, CA, USA). A two-tailed paired Student’s t-test was used for comparison between two groups and one-way analysis of variance with Tukey’s post hoc test for more than two groups. A p-value < 0.05 was considered as an indicator of a statistically significant difference.

Results

Weight and changes in renal function in chronic intermittent hypoxia and sleep-deprived mice

OSA is characterized by CIH and SD and can lead to kidney injury. The study aimed to explore whether the mechanism of rapamycin inhibits the activation of NLRP3 inflammatory bodies by regulating mTOR, to explore the therapy of OSA-induced kidney injury.

Compared with the control group, the SD, CIH, and SD+CIH groups had lower body weights, and the difference was statistically significant (P < 0.05) (Figure 1(a)). Compared with the control, SD, and CIH groups, the SD+CIH group had higher BUN, creatinine, and U-Alb levels, and the difference was statistically significant (P < 0.05) (Figure 1(b-d)). The obvious renal injury occurred in the CIH + SD group in images of HE-stained kidney tissues, including inflammatory cell infiltration, nucleus abscission, unclear cell boundary, and tissue edema. Renal injury in the SD and CIH groups was between the control group and the CIH + SD group (Figure 1(e)).

Enhanced autophagy in the kidneys of chronic intermittent hypoxia and sleep-deprived mice

Next, we investigated the role of autophagy in SD/CIH-induced kidney injury. To understand the effect of SD/CIH on the microscopic substructure of the kidney, we observed the kidney using transmission electron microscopy. Compared to the control, SD, and CIH groups, the nucleus ruptured, and the morphological structure of the cells was unclear in the SD+CIH group (Figure 3(a)). Western blot analysis revealed that the SD+CIH group showed increased LC3 and Beclin 1 protein expression and reduced P62 protein expression compared with the control, SD, and CIH groups (Figure 3(b,e)). Compared to the control, SD, and CIH groups, the SD+CIH group had a higher degree of LC3 staining (Figure 3(f)).

Rapamycin ameliorates renal damage mediated by autophagy

Treatment with rapamycin, an agonist of autophagy, for 3 successive weeks protected mice against kidney injury. Compared to the SD+CIH group, BUN, creatinine, and U-Alb levels were decreased in the SD+CIH+rapamycin group (Figure 4(a)). The structure and histology of the kidney after rapamycin treatment were well preserved, including decreased nucleus abscission and inflammatory cell infiltration (Figure 4(b,c)). Thus, treatment with rapamycin significantly ameliorated kidney damage. Compared to the SD+CIH group, Western blot analysis revealed that the SD+CIH+rapamycin group showed increased LC3 and Beclin 1 protein expression and reduced P62 protein expression (Figure 4(d)), and it had a higher degree of LC3 staining than the other groups (Figure 4(e)).

Rapamycin ameliorates renal damage mediated by inhibiting autophagy-dependent mTOR/NLRP3 signaling pathway

The autophagy marker mTOR was increased in the kidneys of SD+CIH-exposed mice (Figure 5(a,b)). The NLRP3, GMDSD, IL-1β, IL-18, and cleaved-caspase-1 protein levels were higher in
the SD+CIH group than in the SD+CIH+rapamycin group (Figure 5(b)), and NLRP3, GMDSD, IL-18, IL-1β, and cleaved-caspase-1 mRNA levels were higher in the SD+CIH group than in the SD+CIH+rapamycin group. After rapamycin treatment, pyroptosis was suppressed (Figure 5(c)). Compared to the SD+CIH+rapamycin group, the SD+CIH group showed a higher degree of NLRP3 staining (Figure 5(d)).

**Discussion**

Sleep is an important physiological demand that is regulated by sleep homeostasis and circadian rhythm. Good sleep is conducive to normal body operation [24]. SD refers to a lack of sleep or a serious lack of sleep time for various reasons. Rapid eye movement is an important sleep stage to deal with the metabolic waste of brain tissue and prepare the proteins and neurotransmitters...
required by daytime nerve cells. Sleep in patients with OSA often has obvious rapid eye movement SD [25]. Sleep is the most important external manifestation of maintaining the circadian rhythm. When sleep problems occur, renal function is affected [26]. Intermittent hypoxia has many adverse effects on the body. After repeated hypoxia and reoxygenation, the body reacts similarly to ischemia-reperfusion injury, produces a large number of oxygen free radicals, promotes oxidative stress and inflammatory reactions, and leads to blood glucose and lipid metabolism disorders and cognitive impairment. OSA-related renal diseases can be caused by CIH-induced renal diseases [27], which could be mediated through renal iron overload, apoptosis, and increased oxidative stress [28].

Knutson et al. found that decreased sleep quality and insufficient sleep time can lead to decreased glomerular filtration rate, and decreased sleep quality can also lead to an increased protein/creatinine ratio [29]. Poor sleep status is closely associated with an increased risk of chronic kidney disease [30]. SD is an independent risk factor for early chronic kidney disease [31]. The physiological function of the kidney has a circadian rhythm [32], the excretion of urine volume, glomerular filtration rate, renal blood flow, urinary protein, electrolytes such as sodium [33], potassium, calcium, and phosphorus [34] and some metabolites depend on the circadian rhythm of the kidney. Emans et al. found that the oxygen content of the rat renal cortex and medulla has periodic changes and is high at night and low during

![Figure 2. Effect of chronic intermittent hypoxia and sleep deprivation on the antioxidative status. (a) Calculation of malondialdehyde (MDA) abundances. (B, C, D) Calculation of superoxide dismutase (SOD), catalase (CAT), and glutathione (GSH) activities. *p < 0.05; **p < 0.01; ***p < 0.001; ****p < 0.0001.](image-url)
Figure 3. Enhanced autophagy in the kidneys of chronic intermittent hypoxia and sleep-deprived mice. (a) Representative electron microscope diagram of the kidney after different processing (1 μm); (B, C, D, E) LC3, P62, and Beclin 1 expression levels are determined by immunoblotting. Data are expressed as mean ± standard deviation (n = 6) and are analyzed by one-way analysis of variance. (f) Immunohistochemistry (IHC) images of LC3 degree of staining of mouse kidney (original magnification, ×20). *p < 0.05; **p < 0.01; ***p < 0.001; ****p < 0.0001.
Figure 4. Rapamycin ameliorates renal damage mediated by autophagy. (a) Blood urea nitrogen (BUN), creatinine, urinary albumin (U-Alb); (b) Representative images of hematoxylin and eosin (HE)-stained kidney tissues in four different groups (original magnification, × 40); (c) Representative electron microscope diagram of the kidney after different processing (1 μm); (d) LC3, P62, and Beclin 1 expression levels are determined using immunoblotting. Data are expressed as mean ± standard deviation (n = 6) and are analyzed by one-way analysis of variance. (e) Immunohistochemistry (IHC) images of LC3 degree of staining of mouse kidney (original magnification, × 20). *p < 0.05; **p < 0.01; ***p < 0.001; ****p < 0.0001.
Figure 5. Rapamycin ameliorates renal damage mediated by inhibiting the autophagy-dependent mTOR/NLRP3 signaling pathway. (a) Immunohistochemistry (IHC) images of mTOR degree of staining of mouse kidney (original magnification, × 20); (b) Kidney mTOR, p-mTOR, NLRP3, GMDSD, IL-18, IL-1β, and cleaved-caspase-1 protein levels in mice are determined using immunoblotting; (c) Kidney mTOR, p-mTOR, NLRP3, GMDSD, IL-18, IL-1β, and cleaved-caspase-1 mRNA levels in mice are determined using reverse transcriptase-polymerase chain reaction (RT-PCR). Data are expressed as mean ± standard deviation (n = 6) and are analyzed by one-way analysis of variance. (d) Immunohistochemistry (IHC) images of NLRP3 degree of staining of mouse kidney (original magnification, × 20). *p < 0.05; **p < 0.01; ***p < 0.001; ****p < 0.0001.
the day, indicating that the kidney is more vulnerable to attack during sleep [35]. In the current study, the SD, CIH, and SD+CIH groups had lower body weights. Weight loss due to renal injury is due to the decline of renal function, and many wastes in the body cannot be discharged from the body, resulting in loss of appetite, insufficient energy and calorie intake, and weight loss. Additionally, after kidney injury, the body is often in a high catabolic state, due to accelerated protein catabolism and increased muscle decomposition rate, and weight loss occurs. To explore the effect of OSA on renal injury, we found that the SD +CIH group had higher BUN, creatinine, and U-Alb levels. The obvious renal injury occurred in the SD+CIH group in images of HE-stained kidney tissues, confirming that chronic intermittent hypoxia combined with SD caused renal injury in mice.

ROS are produced during cell metabolism and are important redox signals [36]. Under normal conditions, ROS are safely metabolized by cellular antioxidant mechanisms. When the generation of ROS exceeds the cellular antioxidant capacity, ROS are released during oxidative stress, and non-physiological ROS react abnormally with proteins, nucleic acids, carbohydrates, and lipids [37]. Oxidative stress is a marker of cell dysfunction and cell death. Rodrigues et al. also found that oxidative stress in Drosophila increases significantly after SD [38]. Therefore, we investigated the effect of SD/CIH on the activation of these processes. The results demonstrated that CIH+SD exposure induced oxidative damage, as evidenced by the enhanced MDA levels and decreased SOD, CAT, and GSH activities in mice. A recent study confirmed that OSA predisposed the kidneys to the adverse effects of oxidative stress and phospholipid damage [39]. Research has shown that the levels of SOD and GSH PX in the hippocampus of sleep-deprived mice decrease significantly, and MDA increases significantly, suggesting that SD leads to abnormal metabolism in the hippocampus, produces a large number of free radicals, and enhances the lipid oxidation reaction [40]. SOD is responsible for the antioxidant activity in the body. It is the only known enzyme in the body that can scavenge ROS. The decrease in SOD activity indicated that the antioxidant capacity decreased [41]. Oxidative stress leads to oxidative damage of important cellular components such as proteins, lipids, and DNA; cell death; and multiple organ damage; this will not only lead to a series of morphological changes, such as vascular degeneration of renal parenchymal cells and interstitial cells, but also reduce glomerular filtration function and renal tubular reabsorption function [42]. As important pathogenesis of renal injury, oxidative stress can be detected in animal models such as ischemia-reperfusion injury and sepsis [43]. Accumulating evidence indicates that CIH/SD contributes to oxidative stress and inflammation.

Autophagy refers to the degradation of cytoplasmic components in lysosome [44]. Here, we investigated the role of autophagy in SD/CIH-induced kidney injury. To understand the effect of SD/CIH on the microscopic substructure of the kidney, we observed the kidney using transmission electron microscopy; the nuclei ruptured, and the morphological structure of the cells was unclear in the SD +CIH group. Compared to the control, SD, and CIH groups, the SD+CIH group showed a higher degree of LC3 staining, indicating that autophagy was significantly increased in the SD+CIH group, and Western blot analysis revealed that the SD +CIH group showed increased apoptosis. Generally, autophagy is an important protective mechanism of the body [45], and enhancing the autophagy process to a certain extent can appropriately reduce the oxidative damage of the body [46] and can inhibit pyroptosis; in the process of renal injury, the level of intracellular pyroptosis is an important mechanism of renal injury caused by OSA.

Rapamycin modulates multiple cellular signaling pathways in OSA. It has been shown that rapamycin can improve renal function, decrease proteinuria levels, and alleviate renal tubulointerstitial infiltration and interstitial fibrosis. Compared to the SD +CIH group, BUN, creatinine, and U-Alb levels decreased in the SD+CIH+rapamycin group, and the structure and histology of the kidney after rapamycin treatment were well preserved. Thus, treatment with rapamycin significantly ameliorated kidney damage. Compared to the SD+CIH group, Western blot analysis revealed that the SD+CIH +rapamycin group showed increased LC3 and Beclin 1 protein expression, and reduced P62 protein
expression, while the SD+CIH+rapamycin group had a higher degree of LC3 staining than the other groups, indicating that rapamycin ameliorates renal damage mediated by increased autophagy. Although the level of autophagy increases, autophagy can only maintain cell homeostasis to a certain extent [47]. Simultaneously, ROS with increased concentration will attack organs and tissues in the aircraft, such as the liver and kidney, resulting in corresponding morphological and functional damage. Xu et al. found that upregulation of autophagy can reduce the level of oxidative stress in renal tissue to protect renal function [48]. In our study, treatment with rapamycin, an agonist of autophagy, for 3 successive weeks protected mice against kidney injury. In our study, the autophagy marker mTOR was increased in the kidneys of SD+CIH-exposed mice. The pyroptosis makers such as NLRP3, GMDSD, IL-18, IL-1β, and cleaved-caspase-1 protein levels and the NLRP3, GMDSD, IL-18, IL-1β, and cleaved-caspase-1 mRNA levels were higher in the SD+CIH group than in the SD+CIH+rapamycin group. Following rapamycin treatment, pyroptosis was suppressed. Compared to the SD+CIH+rapamycin group, the SD+CIH group showed a higher degree of NLRP3 staining, indicating that rapamycin ameliorated renal damage mediated by inhibition of the autophagy-dependent mTOR/NLRP3 signaling pathway. OSA can induce the activation of inflammatory factors and promote the occurrence of multi-organ inflammatory responses. The NLRP3 inflammasome is an important target of the classical focal death pathway, mediating the classical pyroptosis pathway [13], and the level of NLRP3 in patients with OSA is significantly increased. mTOR is an important target of autophagy. Inhibiting the mTOR signaling pathway can significantly reduce the levels of NLRP3 and organ damage caused by OSA. When autophagy is enhanced, it can inhibit cell focal death and reduce related organ damage through the mTOR/NLRP3 signaling pathway [49], which can be used as a potential intervention target to increase autophagy and reduce cell death to delay renal injury [14].

**Conclusion**

In conclusion, the present study identified the effects of rapamycin on OSA-associated renal injury. Rapamycin suppression-mediated effects were dependent on its correlation with the mTOR/NLRP3 signaling pathway, consequently resulting in the attenuation of renal injury, and NLRP3 suppression-mediated effects were dependent on its correlation with the activation of mTOR, consequently resulting in the attenuation of renal injury (Figure 6). Our findings suggest that modulation of mTOR/NLRP3 pathways by rapamycin could aid in the identification of novel therapeutic strategies for the treatment of renal injury with OSA.

![Figure 6](image_url)

*Figure 6.* The kidney injured by sleep deprivation (SD) and intermittent hypoxia (IH) is attributed to the activation of the NLRP3 inflammasome. In contrast, inhibition of mTOR/NLRP3 with rapamycin ameliorates chronic intermittent hypoxia and sleep deprivation-induced renal damage.
Author’s contributions

WL, DZ, and KH designed the study; XFW, FY, and HZY assisted in completing the experiment; KH had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis, concept, design, and critical revision of the manuscript for important intellectual content. All authors have revised the manuscript and approved the current version.

Disclosure statement

No potential conflict of interest was reported by the author(s).

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