Bcl-2 19-kDa Interacting Protein 3 (BNIP3)-Mediated Mitophagy Attenuates Intermittent Hypoxia-Induced Human Renal Tubular Epithelial Cell Injury

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Conflict of interest:
None declared

Background:
As a novel pathophysiological characteristic of obstructive sleep apnea, intermittent hypoxia (IH) contributes to human renal tubular epithelial cells impairment. The underlying pathological mechanisms remain unrevealed. The present study aimed to evaluate the influence of Bcl-2 19-kDa interacting protein 3 (BNIP3)-mediated mitophagy on IH-induced renal tubular epithelial cell impairment.

Material/Methods:
Human kidney proximal tubular (HK-2) cells were exposed to IH condition. IH cycles were as follows: 21% oxygen for 25 min, 21% descended to 1% for 35 min, 1% oxygen sustaining for 35 min, and 1% ascended to 21% for 25 min. The IH exposure lasted 24 h with 12 cycles of hypoxia and re-oxygenation. Both the siBNIP3 and BNIP3 vector were transfected to cells. Cell viability and apoptosis, mitochondrial morphology and function, and mitophagy were detected by cell counting kit-8, flow cytometry and TUNEL staining, transmission electron microscopy, western blotting, and immunofluorescence, respectively.

Results:
In the IH-induced HK-2 cells, inhibition of BNIP3 further aggravated mitochondrial structure damage, and decreased mitophagy level, leading to increased cell apoptosis and decreased cell viability. While overexpression of BNIP3 enhanced mitophagy, which protected mitochondrial structure, it can decrease cell death in HK-2 cells exposed to IH.

Conclusions:
The present study showed that BNIP3-mediated mitophagy plays a protective role against IH-induced renal tubular epithelial cell impairment.

Keywords:
Apoptosis • BNIP3 Protein, Human • Epithelial Cells • Hypoxia • Mitophagy

Abbreviations:
OSA – obstructive sleep apnea; CPAP – continuous positive airway pressure; IH – intermittent hypoxia; RTEC – renal tubular epithelial cells; BNIP3 – Bcl-2 19-kDa interacting protein 3; HIF-1α – hypoxia-induced factor-1α; HK-2 – human kidney proximal tubule; siRNA – small interfering ribonucleic acid; CTL – control; CCK-8 – cell counting kit-8; TUNEL – terminal dUTP nick-end labeling; DAPI – 4', 6-diamidino-2-phenylindole; RIPA – radioimmunoprecipitation assay; Bax – B-cell lymphoma 2-associated protein X; CST – Cell Signaling Technology; TEM – transmission electron microscope; BSA – bovine serum albumin; MMP – mitochondrial membrane potential; ANOVA – analysis of variance

Full-text PDF: https://www.medscimonit.com/abstract/index/idArt/936760
Background

The relationship between obstructive sleep apnea (OSA) and renal dysfunction was found in recent years [1,2]. Especially, OSA can aggravate renal dysfunction in chronic renal diseases and vice versa [3]. Our previous study indicated that compared with non-OSA subjects, cystatin C, the serum biomarker of early renal damage, was increased in severe OSA patients without complications [4]. Further study showed that continuous positive airway pressure (CPAP) treatment decreased serum cystatin C levels in those patients [5]. A meta-analysis by Hwu et al [6] showed that OSA is consistently associated with higher albuminuria/proteinuria and a lower estimated glomerular filtration rate. Evidence has confirmed that OSA accelerates renal injury, but the underlying mechanism is still unknown.

Intermittent hypoxia (IH), one of the novel pathophysiological characteristics of OSA, can result in several physiological changes, including systemic inflammation, endothelial dysfunction, oxidative stress, and apoptosis. Experimental studies proved that IH causes renal damage, inducing high renal oxidative stress [7] and apoptosis levels [8-10], and abnormal mitochondrial dynamics [11]. Consistent with previous studies, a study by our team confirmed that chronic IH contributed to high protein levels associated with autophagy and apoptosis in mice mimicking OSA [12]. Studies have found that renal tubular epithelial cells (RTEC) are the earliest affected site in renal damages caused by several factors [13,14]. These damages in the cells included high oxidative stress and apoptosis levels, and mitochondrial structure destruction, and mitophagy was also evaluated. Emerging evidence elucidated that as one type of autophagy, mitophagy might play a protective role against renal tubular epithelial cell injury [14,15]. Studies have demonstrated that Bcl-2 19-kDa interacting protein 3 (BNIP3) is a Bcl-2 family protein localized in the mitochondrial outer membrane. As a mitophagy receptor, BNIP3 plays a protective role in several pathological conditions. Hypoxia-induced factor-1α (HIF-1α)/BNIP3 signal pathway is involved in hypoxia-relevant renal injury [15,16]. There is little research addressing the effect of BNIP3-mediated mitophagy on IH-induced renal dysfunction. The present study aimed to evaluate the effect of BNIP3-mediated mitophagy on IH-induced RTEC injury.

Material and Methods

Cell Culture and Transfection

Human kidney proximal tubular (HK-2) cells were cultured in DMEM/F-12 and 10% fetal bovine serum at 37°C in a humidified atmosphere containing 5% CO₂. After culturing in 6-well plates for 24 h, cells were transfected with BNIP3 small interfering ribonucleic acid (siRNA) (5′-GCACCTCGAATGGGAATGG-3′) using Lipofectamine 3000 (Invitrogen) and BNIP3 vector (5′-CAGAATCCATGGACGAGAA ACTCATCTCTGAAGAGATCTGTGCAGAAACGGAGGC-3′) using BNIP3 overexpression plasmid (LncBio, shanghai, China) according to the instructions of the manufacturer. Scrambled siRNA and empty vector as negative controls were used in parallel. Western blotting and quantitative RT-PCR were conducted and the relevant results which verified the successful transfection are presented in Figure 1.

Group and Intermittent Hypoxia Exposure

After routine culturing and reaching a certain density (1×10⁶ cells/µl), HK-2 cells were grouped. In step 1, HK-2 cells were divided into the normoxia group (control, CTL) and the IH group. In step 2, when reaching the density of 1×10⁶ cells/µl, HK-2 cells were transfected with siBNIP3 and BNIP3 vector, and the transfected effects were confirmed. Then, cells were divided into CTL, IH, IH+siBNIP3, and IH+BNIP3 (transfected with BNIP3 vector) groups. Cells in the CTL group were cultured as previously described. Cells in the IH, IH+siBNIP3, and IH+BNIP3 groups were kept in a hypoxia-reoxygenation chamber: 21% oxygen for 25 min, 21% decreased to 1% for 35 min, 1% oxygen sustained for 35 min, and 1% increased to 21% for 25 min [17-20]. The IH exposure time was 24 h, with 12 cycles of hypoxia-reoxygenation.

Cell Viability and Apoptosis Detection

Cell viability was assayed by cell counting kit-8 (CCK-8, Beyotime Institutes of Biotechnology, Beijing, China) according to the manufacturer’s instructions. Briefly, cells were seeded in 96-well plates with 1×10⁴ cells/well. Two hours after adding CCK-8 solution, we determined the absorbance at 450 nm with a microplate reader. Cell apoptosis was analyzed using a fluorescence-activated cell sorter. After being transfected with BNIP3 siRNA or vector, cells were exposed to IH or normoxia condition. Then, cells were collected, washed, and stained with Annexin V-FITC and propidium iodide (PI) reagents in binding buffer using the Annexin V-PI double staining kit (BD Biosciences, San Diego, CA). Apoptosis was also studied by terminal dUTP nick-end labeling (TUNEL) assay using a one-step TUNEL kit (Cat# C 1090, Beyotime Institute of Biotechnology, Shanghai, China) according to the manufacturers’ protocols. Briefly, after exposure to IH or normoxia and being fixed with 10% formaldehyde, cells were incubated with TUNEL reaction mixture and counterstained with 4’, 6-diamidino-2-phenylindole (DAPI). The coverslips were photographed using confocal laser scanning microscopy version E-C1 (Nikon, Tokyo, Japan). The percentage of TUNEL-positive cells was calculated and analyzed.
Western Blotting

The total protein of HK-2 cells in each group was lysed by radioimmunoprecipitation assay (RIPA) buffer on ice for 10 min. After centrifuging, protein concentrations in the supernatants were analyzed by bicinchoninic acid protein assay (Beyotime, Beijing, China). The equal proteins (40 µg per lane) were subjected to 12% SDS-PAGE and transferred to polyvinylidene difluoride membranes. After being blocked with TBST for 1 h, the membrane was incubated with primary antibodies overnight at 4°C. The dilutions of each primary antibody were as follows: mouse anti-HIF-1α monoclonal antibody (1: 250; Novus Biologicals, Littleton, CO, USA), mouse anti-BNIP3 monoclonal antibody (1: 1000, Abcam, Cambridge, MA, USA), rabbit anti-Beclin-1 (1: 2000, Cell Signaling Technology, Beverly, MA, USA), rabbit anti-LC3 (1: 1000, Novus Biologicals, Littleton, Colorado, USA), rabbit anti-B-cell lymphoma 2-associated protein X (Bax) (1: 1000; Cell Signaling Technology [CST], Danvers, MA, USA), rabbit anti-Caspase-3 (1: 1000, Cell Signaling Technology, Beverly, MA, USA), and mouse anti-β-actin (1: 2000; Santa Cruz Biotechnology, Dallas, USA). After incubation with goat anti-rabbit or goat anti-mouse IgG horseradish peroxidase-conjugated antibody (1: 10000; Santa Cruz Biotechnology), the membranes were developed with an enhanced chemiluminescence detection system. β-actin was set as the internal control. Image J software version 1.51 (Image J software, National Institutes of Health, Bethesda, MD, USA) was conducted to quantify these protein levels. All experiments were performed in triplicate.

Figure 1. Successful transfection proof of BNIP3 siRNA and vector. RT-PCR verified BNIP3 mRNA expression level after transfection with BNIP3 siRNA and vector (A1-A3). Western blotting confirmed the corresponding change in BNIP3 protein level (B1-B3). The figure was created using Image J software version 1.51 (Image J software, National Institutes of Health, Bethesda, MD, USA) and GraphPad Prism version 5.0 (GraphPad Software, Inc., LaJolla, CA, USA).
Transmission Electron Microscope (TEM) Analysis

After digesting and collecting with the density of 1-5×10⁵/ml in 6-well plates, the HK-2 cells were suspended and fixed with 2.5% glutaraldehyde in 0.1 mM PBS (PH 7.4) at 4°C for 2 h. Subsequently, cells were washed with PBS and treated with dehydration, osmosis, embedding, sectioning, and staining. A Hitachi electron microscope version H7700 (Hitachi, Tokyo, Japan) was used for observing the cellular ultrastructure. The numbers of damaged mitochondria per cell in each group were counted.

Immunofluorescence Colonization Analysis

Cells were fixed in 4% paraformaldehyde, then rinsed 3 times with PBS. All cells were permeabilized with 0.1% Triton X-100 for 10 min and rinsed with PBS. Bovine serum albumin (BSA) was used to block the cells at room temperature. Cells were incubated with the LC3 antibodies overnight at 4°C. COX IV (Abcam, Cambridge, MA) antibody was used to label mitochondria. After being rinsed with PBS, cells were incubated with a fluorescent secondary antibody for 1 h at room temperature. Nuclei were stained with DAPI. Colocalization of LC3 (green) and COX IV (red) was defined as overlapped red and green peaks. These images were observed and analyzed via confocal laser scanning microscopy.

Detection of Intracellular ATP and Mitochondrial Membrane Potential

An ATP assay kit (Beyotime Biotech, Beijing, China) was used to assess the intracellular ATP levels of HK-2 cells according to the manufacturer’s instructions. The ATP content of the cells was measured in μmol/g protein. The mitochondrial membrane potential (MMP) was determined with JC-1 fluorescence dye (Molecular Probe) according to the manufacturer’s instructions. Normal and IH-exposed cells were rinsed twice with PBS, then labeled with JC-1 (300 nM) for 20 min at 37°C. After being washed twice with JC-1 staining buffer (1×), cells were incubated with a complete medium for detection using a confocal microscope. The relative intensity of fluorescence expression was analyzed by using Image J software (Version 1.51). Data are presented as the ratio of red to green fluorescence areas.

Statistical Analysis

GraphPad Prism version 5.0 (GraphPad Software, Inc., LaJolla, CA, USA) and SPSS 22.0 software (SPSS, Inc., Chicago, IL, USA) were used for data analysis. All experiments were performed in triplicate. Data are presented as mean±standard deviation (Mean±SD). The unpaired t test or two-way analysis of variance (ANOVA) followed by Dunnett post hoc tests were performed for 2 or 3 groups, respectively. P<0.05 was considered statistically significant.

Results

Intermittent Hypoxia Decreased Cell Viability and Increased Cell Apoptosis in HK-2 Cells

To evaluate the influence of IH on cell viability and apoptosis, CCK8, flow cytometry, and TUNEL assay were conducted after IH exposure. In comparison with the CTL group, cells in the IH group have lower cell viability (Figure 2A) (P<0.05). Flow cytometry results showed that the rate of apoptotic HK-2 cells in the IH group (12.72±0.95) was higher than that in the CTL group (3.81±0.55) (P<0.001) (Figure 2B) (P<0.001). TUNEL staining results demonstrated that the TUNEL-positive cells rate in the IH group was higher than in the CTL group (26.30±2.76 vs 5.20±0.71, P<0.001) (Figure 2C).

Intermittent Hypoxia Induced Mitochondrial Injury and Promoted Mitophagy in HK-2 Cells

To assess the influence of IH on mitochondrial morphology and mitophagy, TEM and western blot were conducted. TEM results showed that HK-2 cells in the CTL group had normal cytoplasm and no typical autophagosomes, while autophagosomes and mitophagic vesicles were observed; furthermore, the proportion of damaged mitochondria per cell was significantly higher in the IH group than in the CTL group (P<0.01) (Figure 3A). Compared with the CTL group, lower ATP (P<0.01) and MMP (P<0.05) levels were detected in the IH group (Figure 3B). The confocal microscopic analysis showed that compared with the CTL group, cells in the IH group had more LC3 green and COX IV red (P<0.01) (Figure 3C). Western blotting results demonstrated that IH exposure increased the levels of LC3, Beclin-1, Caspase-3, and Bax (Figure 3D) (all P<0.05).

HIF-1α/BNIP3 Signal Pathway is Involved in Intermittent Hypoxia-Induced Mitophagy

It is confirmed that HIF-1α overexpression after IH exposure [20]. The mitophagy regulatory pathway, HIF-1α/BNIP3 signal pathway, is also known to participate in ischemia-reperfusion renal impairment [16]. To determine whether HIF-1α/BNIP3 pathway is involved in IH-induced mitophagy, we detected the protein levels of HIF-1α and BNIP3 via western blot analysis. The results showed that compared with the CTL group, the levels of HIF-1α and BNIP3 in HK-2 cells were upregulated in the IH group (both P<0.001) (Figure 4), which indicates the HIF-1α/BNIP3 signal pathway was involved in the IH-induced mitophagy.

BNIP3 Enhanced Cell Viability and Attenuated Cell Apoptosis in IH-Induced HK-2 Cells

To determine the effect of BNIP3 on cell viability and apoptosis in the IH-induced cells, we inhibited and overexpressed the
levels of BNIP3. CCK8, flow cytometry, and TUNEL assay were performed. The results showed that inhibition of BNIP3 decreased cell viability (48.17±5.78 in the IH group vs 18.17±5.85 in the IH+siBNIP3 group, P<0.05), while overexpression of BNIP3 in the IH condition re-activated cell viability (48.17±5.78 in the IH group vs 80.83±7.25 in the IH+BNIP3 group, P<0.05) (Figure 5A). When compared with the IH group (12.42±1.49), the apoptosis rate was higher in the IH+siBNIP3 group (17.88±1.67) (P<0.05) and lower in the IH+BNIP3 group (3.20±0.75) (P<0.001) (Figure 5B). The proportion of TUNEL-positive cells was also higher after inhibition of BNIP3 (26.30±8.72 in the IH group vs 49.50±10.87 in the IH+siBNIP3 group, P<0.01) and lower after overexpression of BNIP3 (26.30±8.72 in the IH group vs 12.10±2.51 in the IH+BNIP3 group, P<0.05) (Figure 5C).

**BNIP3-Dependent Mitophagy Alleviated Mitochondrial Injury Induced by Intermittent Hypoxia**

IH-induced mitochondrial damage in HK-2 cells was detected by TEM. Compared with the IH group, more markedly swollen
Figure 2. Effect of IH on cell viability and apoptosis in HK-2 cells. Cell viability was detected with the CCK-8 kit (A). Cell apoptosis was determined by flow cytometry (B) and TUNEL staining (C). All detection was performed after 24 h. The figure was created using confocal laser scanning microscopy version E-C1 (Nikon, Tokyo, Japan) and GraphPad Prism version 5.0 (GraphPad Software, Inc., LaJolla, CA, USA).
and damaged mitochondria were found in the IH+siBNIP3 group, while less mitochondrial damage was observed in the IH+BNIP3 group. In comparison with the IH group, the proportion of damaged mitochondria was significantly lower in the IH+siBNIP3 and higher in the IH+BNIP3 group than in the CTL group (P<0.01 and 0.05, respectively) (Figure 6A). The decreased ATP and MMP levels induced by IH exposure were aggravated by inhibition of BNIP3 (IH+siBNIP3 group) and ameliorated by overexpression of BNIP3 (IH+BNIP3 group) (all P<0.05) (Figure 6B). Confocal microscopy showed that in comparison with the IH group, the fluorescence density of LC3 and COX IV was lower in the IH+siBNIP3 group (P<0.001) and higher in the IH+BNIP3 group (P<0.01) (Figure 6C). Western blotting showed that the IH-induced mitophagy-related proteins LC3 and Beclin-1 were decreased by inhibition of BNIP3 (IH+siBNIP3 group vs IH group, both P<0.01) and increased by overexpression of BNIP3 (IH+BNIP3 group vs IH group, both P<0.05). The IH-induced high expression of apoptosis-related proteins, Caspase-3, and Bax were further increased in the IH+siBNIP3 (P<0.01) group and decreased in the IH+BNIP3 group (P<0.001) (Figure 6D). All the evidence indicated that BNIP3 has a protective role against IH-induced HK-2 cell damage.
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Figure 3. Effect of IH on mitochondrial morphology and function in HK-2 cells. (A) Transmission electron microscopy (TEM) demonstrated the morphological changes of mitochondria. Black arrows indicate typical mitophagy visualized as a mitochondria-containing autophagosome. (B) ATP levels were evaluated by an ATP assay kit. Mitochondrial membrane potential (MMP) was determined with JC-1 fluorescence dye. (C) Mitophagy was detected using LC3 green and COX IV red staining and is displayed in the merged image as yellow fluorescence dots; the colocalization coefficient of LC3 and COX IV (%) are presented. (D) Western blotting analysis of LC3, Beclin-1, Caspase-3, and Bax. Densitometry was performed for quantification and the ratio of all the proteins to β-actin was expressed as a fold of control. The figure was created using a Hitachi electron microscope version H7700 (Hitachi, Tokyo, Japan), Confocal laser scanning microscopy version E-C1 (Nikon, Tokyo, Japan), Image J software version 1.51 (Image J software, National Institutes of Health, Bethesda, MD, USA), and GraphPad Prism version 5.0 (GraphPad Software, Inc., LaJolla, CA, USA).
Figure 4. Expression of HIF-1α and BNIP3 between CTL and IH groups in HK-2 cells. The expression of HIF-1α and BNIP3 was determined with western blotting and the ratio of HIF-1α and BNIP3 to β-actin was expressed as a fold of control. The figure was created using Image J software version 1.51 (Image J software, National Institutes of Health, Bethesda, MD, USA) and GraphPad Prism version 5.0 (GraphPad Software, Inc., LaJolla, CA, USA).

Discussion

In the present study, we evaluated the effect of IH on renal injury in HK-2 cells and assessed whether BNIP3 mediated mitophagy was involved in the IH-induced renal injury. The results demonstrated that IH induced low cell viability and high cell apoptosis levels, and IH accelerated mitochondrial damage and stimulated mitophagy. Inhibition of BNIP3 exacerbated IH-induced apoptosis and mitochondrial damage; on the contrary, BNIP3 overexpression relieved IH-induced mitochondrial impairment and reduced apoptosis via BNIP3-mediated mitophagy. The strength of the present study was that we used RNA interference to prove that BNIP3-mediated mitophagy has a protective role against IH-induced cell impairment.

Mitophagy is an essential form of mitochondrial degeneration. Evidence has shown that mitophagy protected tubular cells against ischemia-reperfusion renal injury through attenuation of apoptosis and ROS production [15]. Furthermore, enhancement of mitophagy can prevent iodinated contrast-induced RTEC injury [14]. Several studies confirmed the protective role of mitophagy in IH-induced organ impairment [21-23]. A study revealed that NLRP3 deficiency, which enhanced Parkin-mediated mitophagy, played a protective role in IH-induced neuroinflammation [21]. Pinocembrin has been shown to suppress IH-induced microglial cell damage via BNIP3-dependent mitophagy [23]. Wang et al [24] demonstrated that disturbances of nigrostriatal mitophagy are related to IH-induced mitochondrial morphology and function impairment, and supplementation of adiponectin can relieve the impairment via enhancing mitophagy. A two-way relationship between OSA and renal dysfunction has been proven in recent years [1,2]. As a novel pathophysiologically feature of OSA, IH was demonstrated to be closely associated with renal impairment in experimental studies [7-10,25-28]. Chronic IH reduced renal peri-tubular capillary density and damaged RTEC, and chronic IH increased the expression of HIF-1α, vascular endothelial growth factor, and anti-angiogenesis factor thrombospondin-1 in the renal tissue of rats; furthermore, these increased expressions can be alleviated by losartan [28]. IH can lead to renal tubular injury and increase autophagy, which may participate in IH-induced RTEC injury [27]. IH can induce renal oxidative stress and inflammation which can be attenuated by the antioxidant α-lipoic acid [7]. Wu et al [10] showed that chronic IH accelerated renal histological injury, upregulated inflammation, and induced tubular endothelial apoptosis via P38/JNK signaling.
A

![Graph showing cell viability (%)]

| Condition   | Cell Viability (%) |
|-------------|-------------------|
| CTL         | 100               |
| IH          | 50                |
| IH+siBNIP3  | 75                |
| IH+BNIP3    | 100               |

B

![Flow cytometry plots for CTL and IH conditions]

- **CTL**
  - IH+siBNIP3: 79.84% 15.26%
  - IH+BNIP3: 94.84% 5.16%

- **IH**
  - IH+siBNIP3: 82.98% 12.04%
  - IH+BNIP3: 79.84% 3.06%

![Bar graph showing apoptosis rate (%)]

| Condition   | Apoptosis Rate (%) |
|-------------|--------------------|
| CTL         | 2.40%              |
| IH          | 82.98%             |
| IH+siBNIP3  | 2.58%              |
| IH+BNIP3    | 94.84%             |
Figure 5. Effect of BNIP3 on cell viability and apoptosis in IH-induced HK-2 cells. After being transfected with siBNIP3 and BNIP3 vector, HK-2 cells were exposed to IH for 24 h. Cell morphology was assessed by microscopy and cell viability was quantitatively determined with CCK-8 (A). Flow cytometry (B) and TUNEL staining (C) were conducted to detect cell apoptosis. The figure was created using a Hitachi electron microscope version H7700 (Hitachi, Tokyo, Japan) and Confocal laser scanning microscopy version E-C1 (Nikon, Tokyo, Japan), and GraphPad Prism version 5.0 (GraphPad Software, Inc., LaJolla, CA, USA).

pathways, and a soluble receptor for advanced glycation end-product can relieve the IH-induced renal pathophysiological changes. Bai et al [26] observed thickened and edema of glomerular bottom membrane and mesangium in chronic IH rats and demonstrated that chronic IH induced the pyroptosis of RTECs via activating the nucleotide-binding oligomeric domain-like receptor protein 3 inflammasome. The results of the present study were consistent with the previous studies by showing that IH induced lower cell viability and high apoptosis in RETC. Robust evidence has confirmed that the primary site of renal impairment is RETC [14,15]. However, few studies have assessed the influence of IH on RETC and evaluated the changes in mitochondrial morphology and the function of mitophagy after IH exposure. The present study addressed this issue and demonstrated abnormal mitochondrial structure and increased mitophagy in IH-induced HK-2 cells. Emerging evidence shows that as an important type of autophagy, mitophagy plays a protective role against cell injury [13,14,16]. BNIP3 is an important mediated molecule of mitophagy [15,16,23].

Meanwhile, studies have confirmed that HIF-1α/BNIP3 signal pathway mediates mitophagy in hypoxia-related disease [29]. To detect the function of mitophagy in the IH model in vitro and to assess the mediated function of BNIP3, we inhibited
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**B**  
| Treatment       | ATP concentration (µmol/g) | MMP2 Activity (%) |
|-----------------|---------------------------|-------------------|
| CTL             | 1.50                      | 80                |
| IH              | 1.00 (p<0.05)             | 60                |
| IH+siBNIP3      | 0.50 (p<0.001)            | 40                |
| IH+BNIP3        | 0.00                      | 20                |

**C**  

COX IV  

LC3  

DAPI  

Merge  

Colocalization coefficient of LC3 and COX IV (%):  
- CTL: 20 ± 5  
- IH: 40 ± 5  
- IH+siBNIP3: 60 ± 5  
- IH+BNIP3: 80 ± 5  

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Figure 6. Effect of BNIP3 on mitochondrial morphology and function in IH-induced HK-2 cells. (A) Transmission electron microscope (TEM) demonstrated the morphological changes of mitochondria between groups. Black arrows indicate typical mitophagy visualized as a mitochondria-containing autophagosome. (B) ATP levels was evaluated by an ATP assay kit. Mitochondrial membrane potential (MMP) was determined with JC-1 fluorescence dye. (C) Representative images under the confocal microscope. HK-2 cells were treated with LC3 green and COX IV red to label autophagosomes and mitochondria. A confocal microscope was used to analyze the distribution of different fluorescence. Colocalization of LC3 and COX IV was defined as overlapped green and red peaks. (D) Western blotting was used to detect the expression of LC3, Beclin-1, Caspase-3, and Bax. β-actin was used as the internal control. The figure was created using a Hitachi electron microscope version H7700 (Hitachi, Tokyo, Japan), Confocal laser scanning microscopy version E-C1 (Nikon, Tokyo, Japan), Image J software version 1.51 (Image J software, National Institutes of Health, Bethesda, MD, USA), and GraphPad Prism version 5.0 (GraphPad Software, Inc., Lalolla, CA, USA).
and overexpressed BNIP3 in IH-induced HK-2 cells. Our results showed that inhibition of BNIP3 significantly decreased mitophagy, aggravated the destruction of mitochondrial morphology and function, and enhanced cell apoptosis; meanwhile, IH-induced mitochondrial impairment and cell apoptosis were ameliorated after overexpression of BNIP3. We speculated from the results that BNIP3-mediated mitophagy plays an important role in the protection of RETC against IH-induced pathophysiological changes.

The present study has some limitations. Firstly, we only created the IH model in vitro by using HK-2 cells, and the results need to be confirmed in other cell lines and in vivo animal experiments, even in clinical practice. Secondly, we did not interfere with the expression of BNIP3 under normoxia conditions, which might weaken the reliability of the conclusions of the present study. Thirdly, inflammation biomarkers and oxidative stress levels were not determined in our study. Lastly, OSA, which is considered a systemic disease, can impair both the renal tubular epithelial and glomerular filtration function, but we only assessed IH-induced RETC, and the deterioration of glomerular filtration by IH was not studied. Further study is warranted to assess the influence of IH on glomerular filtration function.

Conclusions
IH induced mitochondrial morphological and functional impairments and facilitated cell apoptosis in HK-2 cells. BNIP3-mediated mitophagy has a protective role against IH-induced RTEC impairment.

Ethics Approval and Consent to Participate
The study was approved by the Ethics Committee of Zhongshan Hospital, School of Medicine, Xiamen University (approval no. 2021-065).

Declaration of Figures’ Authenticity
All figures submitted have been created by the authors, who confirm that the images are original with no duplication and have not been previously published in whole or in part.

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