Roles of the mammalian target of rapamycin (mTOR) signaling pathway in the repair of hyperoxia-induced acute lung injury

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A – research concept and design; B – collection and/or assembly of data; C – data analysis and interpretation; D – writing the article; E – critical revision of the article; F – final approval of the article

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Conflict of interest
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Abstract

Background. Rapamycin inhibits the mammalian target of rapamycin (mTOR) activity and has been proven effective for the treatment of lung injury.

Objectives. The objective of this study was to investigate the roles of the mTOR pathway and its inhibitor rapamycin in the repair of hyperoxia-induced acute lung injury (ALI).

Material and methods. Firstly, premature rat lung fibroblast L929 cells were cultured under different oxygen concentrations (40%, 60%, and 90%). At day 3, 7 and 14 after exposure, MTT assay and flow cytometry were used to evaluate the effect of oxygen stress on cell viability and apoptosis of L929 cells, respectively. Secondly, microscopy, MTT assay and flow cytometry was used to investigate the effect of 10 nM rapamycin on 90% O₂ exposed L929 cells. We also used small interfering RNAs (siRNAs) to abrogate the expression of mTOR in 90% O₂ exposed L929 cells, and then evaluated the apoptosis and cell viability using flow cytometry and the MTT assay, respectively. In addition, western blot was used to detect the protein expression of Bcl-2, p53, TGF-β and connective tissue growth factor (CTGF). A hyperoxia-induced lung injury model was established in Sprague Dawley (SD) rats in order to evaluate the histopathological changes in lung tissues and expression of the mTOR pathway and fibrosis related factors.

Results. Exposure to 40%, 60% or 90% oxygen all significantly inhibited the growth of L929 cells. Application of 10 nM rapamycin was found to effectively promote apoptosis of 90% O₂ exposed L929 cells. We also used small interfering RNAs (siRNAs) to abrogate the expression of mTOR in 90% O₂ exposed L929 cells, and then evaluated the apoptosis and cell viability using flow cytometry and the MTT assay, respectively. In addition, western blot was used to detect the protein expression of Bcl-2, p53, TGF-β and connective tissue growth factor (CTGF). A hyperoxia-induced lung injury model was established in Sprague Dawley (SD) rats in order to evaluate the histopathological changes in lung tissues and expression of the mTOR pathway and fibrosis related factors.

Conclusions. Rapamycin may be useful for the treatment of hyperoxia-induced acute lung injury (ALI) by inhibiting the activation of mTOR signaling pathway.

Key words: mTOR, rapamycin, siRNA interference, hyperoxia-induced lung injury

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Introduction

Bronchopulmonary dysplasia (BPD), a form of chronic lung disease (CLD), is characterized by the abnormal formation of alveoli and chronic pulmonary vascular changes, especially in preterm infants. BPD has a reported incidence of >60% in preterm infants, and is one of the leading causes of death and disability in preterm infants in China. Oxygen therapy is a common method for the treatment of preterm infants, and appropriate oxygen therapy can effectively save the lives of preterm infants suffering from hypoxia. However, sustained oxygen inhalation at high concentrations can lead to extensive and non-specific inflammatory responses in lung tissue, followed by pulmonary stromal hyperplasia, and pulmonary fibrosis, or even acute lung injury (ALI). Lung fibroblasts (LFS) are the main cells involved in the repair of lung injury. The accumulation of the abnormal extracellular matrix (ECM) that is produced by LFs plays an important role in the pathogenesis of BPD fibrosis. Currently, one of the major problems in the prevention and early treatment of lung injury is the excessive proliferation of LFs in lung repair and reconstruction, which may end up replacing the normal terminal bronchioles and alveoli, resulting in irreversible damage of both the structure and function of the lung.

As a serine/threonine protein kinase, mammalian target of rapamycin (mTOR) is a member of the phosphoinositide-3-kinase-related kinase family (PI3K). The mTOR signaling pathway can phosphorylate multiple target proteins for the regulation of transcription, translation initiation, protein synthesis, and degradation, mainly through the activation of p70S6K/S6 and the inhibition of 4EBP1/eIF4E. Mammalian target of rapamycin signaling pathway has been shown to be able to regulate embryonic development, and lung development, and is involved in a variety of lung diseases, such as chronic obstructive pulmonary disease (COPD), cystic fibrosis, and pulmonary fibrosis. Pulmonary fibrosis is characterized by the activation of the mTOR and its down-stream target, the ribosomal S6 kinase (p70S6K), pulmonary fibrosis generated fibrotic foci with abundant activated hepatic stellate cells and excessive collagen deposition, it has been reported that rapamycin, an inhibitor of p70S6K phosphorylation, could inhibit hepatic stellate cell proliferation and limits fibrogenesis. Blocking the mTOR signaling pathway has been shown to suppress the proliferation of fibroblasts and the over-production of extracellular matrix (ECM) in liver tissue, which may reverse pulmonary fibrosis. Rapamycin is a macrolide immunosuppressive agent, an inhibitor of mTOR, and it has been proven effective for the treatment of pulmonary fibrosis. Nevertheless, the detailed molecular mechanisms underlying the role of the mTOR signaling pathway in the repair of hyperoxia-induced lung injury have not yet been fully elucidated and require further investigation.

In this study, we aim to investigate the roles of mTOR signaling pathway in the repair of hyperoxia-induced acute lung injury in vivo and in vitro.

Material and methods

Ethics statement

All animal handling procedures were carried out in accordance with the protocols approved by the Institutional Animal Care and Use Committee of University of South China (No. 2014).

Animals and cell lines

The mouse lung fibroblast cell line L929 was purchased from the American Type Culture Collection (ATCC) (Manassas, USA). The L929 cells were thawed at 37°C, centrifuged at 1000 g for 5 min, and rinsed twice with RPMI-1640 medium containing 10% fetal bovine serum (Gibco, USA). Cells were harvested by centrifugation at 1000 g for 5 min, and cultured in RPMI-1640 medium containing 10% fetal bovine serum (Gibco, USA), which was replaced every day. Fifty-four specific-pathogen-free (SPF) SD rats (weighing 225–240 g and aged 8 weeks), including 36 females and 18 males, were obtained from the Experimental Animal Center of Nanhua University (Hunan, China). Female and male rats were co-housed at a ratio of 2:1 for mating. Females were checked for the identification of vaginal plugs each morning. The day a vaginal plug was identified was the 1st day of pregnancy. On the 21st day of pregnancy, 66 neonatal rats were obtained as the premature rats in this study. These 66 premature rats were used for 2 parts of the in vivo studies. In the first part of animal study, 42 premature rats were included and divided into 7 groups (n = 6 per group): (1) control, 2) 90% O2, 3) 90% O2 + rapamycin, 4) 90% O2 + rapamycin 7d, 5) 90% O2 + rapamycin 3d, 6) 90% O2 + rapamycin 7d, 7) 90% O2 + rapamycin 14d. In the second part of animal study, 24 premature rats were included and divided into 4 groups (n = 6 per group): (1) 90% O2, 2) 90% O2 + rapamycin 3d, 3) 90% O2 + rapamycin 7d, 4) 90% O2 + rapamycin 14d.

Oxygen exposure on L929 cells

L929 cells were cultured in 37°C with 5% of carbon dioxide and in saturated humidity until moisture began adhering to the wall. After aspiration of the culture medium, the cells were rinsed twice with 37°C pre-warmed PBS, and cultured in fresh complete medium. The L929 cells were cultured in distinct media and were randomly divided into 4 groups (control, 40% O2, 60% O2, and 90% O2). The control group cells were cultured in fresh complete medium with regular air at 37°C. The other 3 groups were cultured in fresh complete medium under different oxygen concentrations (40%, 60%, and 90%) in independent
culture chambers. Air, including the specific concentrations of oxygen and nitrogen, as well as 5% CO₂, was injected into the culture chamber at a speed of 1 L/min for 30 min. The culture chamber was sealed once the oxygen monitor showed expected oxygen concentrations, and placed in an incubator for culture. The 40% oxygen group was cultured in 40% O₂, 55% N₂, and 5% CO₂. The 60% oxygen group was cultured in 60% O₂, 35% N₂, and 5% CO₂. The 90% oxygen group was cultured in 90% O₂, 5% N₂, and 5% CO₂. The corresponding concentration air was 21% O₂, 78% N₂, and 5% CO₂.

To prepare the 60%, 90% oxygen groups, 55% N₂, and 5% CO₂, primers sequences (5'-3') for 24 h. To prepare 25%, 5% N₂, and 5% CO₂, were cultured under different conditions were in air, 35% N₂, and 5% CO₂. The qPCR was conducted according to the following program: 94°C for 5 min, 25 cycles of denaturation at 94°C for 1 min, annealing at 55°C for 1 min, and extension at 72°C for 4–6 h. The supernatant was removed and 3 mL RPMI1640 was added to each well. The siRNA-transfected L929 cells were divided into 4 groups: control, 90% O₂, 90% O₂ + rapamycin (10 nM), and the mTOR siRNA groups.

Quantitative real-time PCR

The mTOR siRNA transfected cells were harvested for RNA isolation using the Trizol reagent (Takara, Japan). The Takara kits were used for cDNA synthesis and qPCR, which was conducted in a 20 μL reaction system with 10 μL 2 × Mix buffer (Aidlab Biotechnologies, Beijing, China), forward and reverse primers (Table 1) (0.4 μL each), 1 μL DNA, and 15.4 μL double distilled water. The primers used in this study were shown in Table 1. The qPCR was conducted according to the following program: 94°C for 5 min, 25 cycles of denaturation at 94°C for 1 min, annealing at 55°C for 1 min, and extension at 72°C for 40 s. Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) was used as the internal reference. The relative mRNA level was calculated using the 2⁻ΔΔCt method.

Table 1. The primers for real-time quantitative PCR

<table>
<thead>
<tr>
<th>Gene names</th>
<th>Primers sequences (5'-3')</th>
</tr>
</thead>
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<tr>
<td>GAPDH</td>
<td>F: CCGGAGAATGGGAAAGCTGTGC</td>
</tr>
<tr>
<td></td>
<td>R: AAGCACAACAGAGGAGAA</td>
</tr>
<tr>
<td>mTORC1</td>
<td>F: TCAACTGGGAAAAGTGACC</td>
</tr>
<tr>
<td></td>
<td>R: CTGATGGTCTGCTTCAACT</td>
</tr>
<tr>
<td>p70S6K</td>
<td>F: AAATCTCCATGGCTTTGGGG</td>
</tr>
<tr>
<td></td>
<td>R: AGGGGCCATGTATTCTATTG</td>
</tr>
<tr>
<td>4EBP1</td>
<td>F: CCAGGATTCTATGACCGCG</td>
</tr>
<tr>
<td></td>
<td>R: AATTGTGACTTTCACCGCC</td>
</tr>
</tbody>
</table>

Inverted phase contrast microscopy

L929 cells exposed to 90% O₂ concentrations of oxygen, with rapamycin intervention, were harvested after 3 d. The morphology and number of L929 cells were evaluated under an inverted phase contrast microscope to investigate the effects of rapamycin on the morphology of oxygen exposed L929 cells.

Inhibition of mTOR expression using siRNA

The cells were digested to a concentration of 1 × 10⁶ cells/mL and inoculated in a 6-well plate (3 mL/well). They were then cultured at 37°C and 5% CO₂ for 24 h. To prepare siRNA-Lipofectamine for (Invitrogen, Carlsbad, USA) L929 cell transfection, 75 pmol synthesized mTOR siRNA diluted in 100 μL serum-free RPMI1640 medium (Invitrogen Life Technologies, New York, USA), and 5 μL lipofectamine 2000 diluted in 100 μL serum-free RPMI1640 medium were mixed and incubated at room temperature for 20 min. After cells were washed with PBS, 200 μL siRNA-Lipofectamine 2000 was added to each well and cells were incubated at 37°C and 5% CO₂ for 4–6 h. The supernatant was removed and 3 mL RPMI1640 was added to each well. The siRNA-transfected L929 cells were divided into 4 groups: control, 90% O₂, 90% O₂ + rapamycin (10 nM), and the mTOR siRNA groups.

Flow cytometry for apoptosis

To evaluate the apoptosis of L929 cells, cells were double stained with annexin V-FITC and PI, and analyzed by flow cytometry (BD Biosciences, New York, USA). The L929 cells (10⁶/mL) cultured under different conditions were incubated at 37°C and 5% CO₂ for 24 h before being digested by trypsin without EDTA. The cells were then rinsed with pre-cooled PBS, centrifuged at 1500 g for 5 min, and resuspended in 300 μL of ×1 binding buffer. After incubation with Annexin V-FITC (5 μL) for 15 min and PI (10 μL) for 5 min, the L929 cells were analyzed by FACSCalibur flow cytometry (BD Biosciences, USA).

MTT assay for cell viability

To evaluate the cell viability of L929 cells, the cell concentration of each group was adjusted to 3 × 10⁴ cells/mL. The L929 cells were inoculated into a 96-well culture plate and 15 μL MTT was added to each well. Cells were incubated at 37°C in a humidified chamber and 5% CO₂ for 3–4 h. After aspiration, 200 μL DMSO was added to each well and incubated at room temperature in a shaker for 10 min; the absorbance of each well was measured at 492 nm. All samples were assayed in duplicate.

Table 1. The primers for real-time quantitative PCR
calculated according to the $2^{-\Delta\Delta Ct}$ method with ABI Prism 7500 SDS software (Applied Biosystems, Foster City, USA).

**Western blot**

Harvested cells were mixed with 500 μL RIPA lysate with PMSF (Sigma-Aldrich, St. Louis, USA) and incubated on ice for 2 h, before being centrifuged at 12,000 g at 4°C for 10 min to obtain the supernatant. Western blotting was performed using the BCA method. Protein quantification was performed using the BCA method. Western blotting was used to evaluate the protein expressions of p70S6K and 4EBP1 proteins in the control and mTOR siRNA groups. The expressions of Bcl-2, p53, TGF-β, and connective tissue growth factor (CTGF) in the control, 90% O$_2$, 90% O$_2$ + rapamycin, and mTOR siRNA groups were also evaluated using western blot. Relative density of western blot band was analyzed by using Image J software (National Institutes of Health, Bethesda, USA; http://imagej.nih.gov/ij/) and protein expression level of each group was normalized with β-actin.

**Establishment of a model of hyperoxia-induced lung injury in SD rats and rapamycin intervention**

On the 21$^{st}$ day of pregnancy, the rats were sacrificed in order to obtain premature neonatal rats. After 3, 7, and 14 days, the premature rats were sacrificed for follow-up experiments (n = 6 rats per group). The hyperoxia-induced lung injury rat model was established according to the study conducted by Zhang, et al.[22] Briefly, the premature rats were raised in 90% oxygen (5–6 L/min), at 22–25°C, and 65–75% humidity with unlimited food and water. The CO$_2$ level in the rat house was set to 5%, using the appropriate amount of barium hydroxide. Sun et al.[23] reported that mTOR signaling was blocked by intraperitoneal injection of rapamycin (1.5 mg/kg) in mice, thus 1.5 mg/kg dose of rapamycin was chosen in the animal study. Next, the 90% O$_2$ + rapamycin group received intraperitoneal injection of rapamycin (1.5 mg/kg) once a day. To avoid hyperoxicosis, the maternal rats of the control and hyperoxia groups were interchanged every 24 h.

**Histopathological examination**

Six rats were randomly selected from each group and sacrificed for lung dissection. The left lung was fixed in 4% paraformaldehyde to prepare paraffin sections for histopathological examination. The paraffin sections were stained with hematoxylin and eosin and examined under light microscopy (XDS-1A; Supore, Shanghai, China). Other lung tissue was washed with PBS at 4°C for 3 min and frozen in liquid nitrogen for follow-up experiments. In order to evaluate the effects of 90% oxygen and intraperitoneal injection of 1.5 mg/kg rapamycin on lung tissues of rats, histopathological scores were evaluated according to the protocol developed by Murakami et al.[24]

**Enzyme-linked immunosorbent assay**

Five lung tissue samples (10 mg per sample) randomly selected from each group were homogenized in cold PBS (pH = 7.4). The homogenate was centrifuged at 10,000 g for 20 min at 4°C. Total protein concentration of each sample was assayed using a BCA protein quantification kit (Sangon, Shanghai, China). The contents of collagen I, collagen III, and FN in ECM were measured using an ELISA kit (Cosmo Bio, Tokyo, Japan) according to the manufacturer’s instructions. The absorbance was measured at 450 nm on a microtiter reader (Thermo Scientific, USA). In addition, the levels (ng/mg) of TGF-β, CTGF, and collagen I in the lung tissue of premature rats were also determined with an ELISA kit (Sangon, Shanghai, China) according to the manufacturer’s instructions. Absorbance values were normalized to the standard curve.

**Western blot analysis for the expression of mTORC1, p70S6K, and 4EBP1 in lung tissues**

The lung tissue was washed with pre-cooled saline to remove residual blood and was then dried. One gram of lung tissue was used to prepare a 10% lung homogenate. After centrifugation at a low temperature, the supernatant from the lung homogenate was used for western blot to quantify the protein levels of mTORC1, p70S6K, and 4EBP1.

**Statistical analyses**

The data was expressed as mean ± standard deviation (x ±SD) and analyzed using SPSS V. 19.0 software (IMB Corp., Armonk, USA). One way analysis of variance (ANOVA) for repeated measures and post-hoc Tukey’s test for pairwise comparison were performed. A p-value less than 0.05 was considered to be statistically significant.

**Results**

Oxygen exposure induced apoptosis and inhibited proliferation in L929 cells in a time and concentration-dependent manner. The L929 cells exposed to 40%, 60%, and 90% oxygen exhibited significantly higher apoptosis rates than those cultured in regular air for the same time period (3, 7, or 14 days) in a time and concentration-dependent manner (p < 0.05) (Fig. 1A–B). The L929 cells exposed to 40%, 60%, and 90% oxygen also exhibited significantly lower cell viability than those cultured in regular air for the same time period (3, 7, or 14 days) in a time and concentration-dependent manner (p < 0.05) (Fig. 1C). These results suggest that exposure to 40%, 60% and 90% oxygen all could induced apoptosis and inhibited cell viability of L929 cells in a time and concentration-dependent manner.
Effects of 10 nM rapamycin on morphology of L929 cells exposed to 90% O₂

According to the results of above, 90% oxygen caused more severe damage to the L929 cells than 40% and 60% oxygen. Therefore, 90% oxygen was used in the follow-up experiments. We added 10 nM rapamycin to L929 cells exposed to 90% O₂ and cultured for 3 days, to investigate the effects of rapamycin on the morphology of L929 cells. As shown in Fig. 2, the majority of L929 cells cultured with regular air grew well and had bright, translucent, and compact cell bodies. With exposed to 90% O₂, cells cultured with rapamycin exhibited a reduced refractive index and had increased numbers of granules in the cytoplasm, while most cells were still viable and they exhibited a slightly different morphology (Fig. 2).

Rapamycin inhibited the growth and promoted apoptosis of L929 cells exposed to 90% O₂.

The L929 cells cultured with 10 nM rapamycin and exposed to 90% O₂ exhibited significantly higher apoptosis...
Fig. 3. A, B – Flow cytometry analysis for apoptosis of L929 cells exposed to 90% O₂ with rapamycin intervention; C – MTT assay for cell viability of L929 cells 90% O₂ with rapamycin intervention vs control group; #p < 0.05 vs 90% O₂ group.

Fig. 4. A – qPCR analysis for the mRNA expression of mTORC1, 4EBP1, and p70S6K in L929 cells transfected with mTOR siRNA; B – western blot analysis for the protein expressions of mTORC1, 4EBP1, and p70S6K in L929 cells transfected with mTOR siRNA; C – flow cytometry analysis for the apoptosis of L929 cells transfected with mTOR siRNA; D – MTT assay for cell viability of L929 cells transfected with mTOR siRNA.

* p < 0.05, ** p < 0.01 vs control group.
rates than the control group (*p < 0.05, Fig. 3A,B), and 90% O₂ exposure significantly inhibited the cell viability of L929 cells (*p < 0.05, Fig. 3C). In addition, the apoptosis ratio was increased and the cell viability was inhibited in the 90% O₂ + rapamycin group compared with 90% O₂ group (#p < 0.05) (Fig. 3A–C).

Validation of mTOR siRNA

After transfected with mTOR siRNA in L929 cells, we found that the mRNA and protein expression levels of mTORC1, 4EBP1 and p70S6K in L929 cells transfected with mTOR siRNA were significantly lower than in control cells (*p < 0.05, Fig. 4A,B). In addition, the rates of apoptotic cells transfected with mTOR siRNA (1.2%) were significantly increased than in the control (20.1%) (***p < 0.01, Fig. 4C), and the cell viability of cells transfected with mTOR siRNA was significantly decreased compared with control group (*p < 0.05, Fig. 4D). These results suggest that mTOR siRNA could down-regulate the expression of mTORC1, 4EBP1 and p70S6K, inhibit the cell viability and promote apoptosis of L929 cells.

Effects of rapamycin and mTOR siRNA on the protein expressions of Bcl-2, p53, TGF-β, and CTGF in L929 cells

We evaluated the protein expressions of apoptosis-related genes, including p53 and Bcl-2, as well as fibrosis-related genes, including TGF-β and CTGF, in L929 cells which were transfected with mTOR siRNA, treated with 10 nM rapamycin, and exposed to 90% O₂. As shown in Fig. 5, the expression of Bcl-2 in cells exposed to 90% oxygen and cultured with 10 nM rapamycin or transfected with mTOR siRNA was significantly lower than in cells exposed to regular air (*p < 0.05), and mTOR siRNA significantly increased the expression of Bcl-2 compared with 90% O₂ group (#p < 0.05). The expression level of p53 in L929 cells exposed to 90% oxygen and cultured with rapamycin or those transfected with mTOR siRNA was significantly higher than in cells exposed to regular air (*p < 0.05); however, the application of rapamycin or mTOR siRNA groups all exhibited lower expression level of p53 compared with 90% O₂ group (#p < 0.05). The expression levels of TGF-β and CTGF in L929 cells exposed to 90% oxygen was significantly higher than in the control group (*p < 0.05); however, the application of 10 nM rapamycin or mTOR siRNA groups...
all exhibited lower expression levels of TGF-β and CTGF compared with 90% \(O_2\) groups. These results suggest that 90% \(O_2\) exposure could promote L929 cell apoptosis and successfully inhibit the expression of CTGF and TGF-β.

**The contents of collagen I, III and fibronectin in extracellular matrix**

The contents of collagen I, collagen III (col I, col III), and fibronectin in extracellular matrix (ECM) of L929 cells exposed to 90% \(O_2\) was significantly higher than those in cells exposed to regular air \((p < 0.05)\), suggesting that 90% \(O_2\) could induce the production of FN and collagen. However, the contents of col I, col III, and FN in L929 cells treated with rapamycin or transfected with mTOR siRNA were significantly lower than that in 90% \(O_2\) \((#p < 0.05)\), suggesting that blocking the mTOR signaling pathway suppressed collagen deposition and decreased the production of FN (Fig. 6).

**Histopathological changes in the lung tissues**

The lung tissues of control group rats exhibited normal histopathology, without exudation of inflammatory cells (Fig. 7). Exudation of a few inflammatory cells and red cells was observed in the lungs exposed to 90% \(O_2\) for 3, 7 and 14 days, and in a time-dependent manner. After 7 days of exposure to 90% oxygen, we observed alveolar rupture and fusion, and disordered distribution of cells along the tracheal arteries. After 14 days of exposure to 90% oxygen, extensive exudation of inflammatory cells and thickening blood vessels and airway walls were observed. The lungs of rats that had received intraperitoneal rapamycin injections and were exposed to 90% oxygen for 3 days exhibited normal histopathology. Alveolar rupture and fusion, as well as thickening blood vessels and airway walls, were observed in the lungs of rats that had received intraperitoneal injections of rapamycin and were exposed to 90% oxygen for 7 days. After 14 days of exposure to 90% oxygen, we observed more severe alveolar rupture and fusion in the lungs of rats that had received intraperitoneal injection of rapamycin.

The pathological scores of lung injury are shown in Table 2. The pathological scores of lung injury in the rats exposed to 90% oxygen for 3, 7, and 14 days were significantly higher than those of the control \((p < 0.05)\). After 3 and 7 days, the pathological lung injury scores of the 90% \(O_2\) + rapamycin group were \((3.50 \pm 0.84)\) and \((9.67 \pm 1.97)\), respectively, which were significantly lower than those of the 90% \(O_2\) group for 3 \((6.33 \pm 2.34)\) and 7 days \((14.0 \pm 2.45)\), respectively \((p < 0.05)\).

**Dynamic changes of col I, TGF-β, and CTGF in lung tissues**

The col I concentration in the lungs of rats exposed to 90% oxygen for 3, 7, and 14 days were \((471.87 \pm 5.72 \text{ng/mg})\), \((529.72 \pm 6.97 \text{ng/mg})\), and \((556.44 \pm 8.52 \text{ng/mg})\), which were significantly higher than those of the air control group \((414.43 \pm 8.97 \text{ng/mg})\) \((p < 0.05)\). The TGF-β concentration in the lung tissues of rats exposed to 90% oxygen for 3, 7, and 14 days were \((33.74 \pm 2.84 \text{ng/mg})\), \((58.65 \pm 3.10 \text{ng/mg})\), and \((90.87 \pm 4.23 \text{ng/mg})\), which were significantly higher than those of the air control group \((28.34 \pm 2.56 \text{ng/mg})\) \((p < 0.05)\). The CTGF concentration in the lung tissues of rats exposed to 90% oxygen for 3, 7, and 14 days were \((23.83 \pm 3.71 \text{ng/mg})\), \((36.17 \pm 4.15 \text{ng/mg})\), and \((50.01 \pm 4.73 \text{ng/mg})\), which were significantly higher than those of the air control group \((18.74 \pm 2.45 \text{ng/mg})\) \((p < 0.05)\).

<table>
<thead>
<tr>
<th>Group</th>
<th>Pathological scores</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>1.95 ±0.34</td>
</tr>
<tr>
<td>90% (O_2) 3d</td>
<td>6.33 ±2.34*</td>
</tr>
<tr>
<td>90% (O_2) + rapamycin 3d</td>
<td>3.50 ±0.84*</td>
</tr>
<tr>
<td>90% (O_2) 7d</td>
<td>14.0 ±2.45*</td>
</tr>
<tr>
<td>90% (O_2) + rapamycin 7d</td>
<td>9.67 ±1.97*</td>
</tr>
<tr>
<td>90% (O_2) 14d</td>
<td>23.83 ±3.71*</td>
</tr>
<tr>
<td>90% (O_2) + rapamycin 14d</td>
<td>24.0 ±3.74*</td>
</tr>
</tbody>
</table>

* compared with control group, \(p < 0.05\); * compared with 90% \(O_2\) groups at the same time, \(p < 0.05\).

Fig. 7. Histopathological changes of the lung tissues from hyperoxia-induced lung injury rats. Hematoxylin and eosin stain; bar 50 μm; magnification ×200.
ng/mg), and (98.81 ±1.55 ng/mg), which were significantly higher than those of the control group (25.50 ±1.86 ng/mg) (\( p < 0.05 \)). The CTGF concentration in the lung tissues of rats exposed to 90% oxygen for 3, 7, and 14 days were (50.72 ±1.80 ng/mg), (68.65 ±2.24 ng/mg), and (94.39 ±2.48 ng/mg), which were significantly higher than those of the control group (41.23 ±1.08 ng/mg) (\( p < 0.05 \)). These results suggest that the production of pulmonary fibrosis factors was induced by 90% oxygen. Compared with the 90% oxygen group, rapamycin significantly reduced the concentrations of col I, TGF-β1, and CTGF in the lung tissues of rats exposed to 90% oxygen for 3, 7, and 14 days (\( p < 0.05 \)) (Table 3).

**Rapamycin inhibits the mTOR pathway in the lung tissue of rats**

The above results indicated that rapamycin attenuates acute lung injury induced by 90% oxygen. The objective of western blot was to determine whether the mTOR pathway was involved in the repair of lung injury. As shown in Fig. 8, the expression levels of mTORC1, p70S6K, and 4EBP1 in the lung tissue of rats in the 90% \( O_2 \)+ rapamycin group were significantly decreased compared with control animal at day 7 and 14 (*\( p < 0.05 \)), which suggested that the application of rapamycin could inhibit the activation of mTOR signaling pathway and inhibit the expression of downstream proteins p70S6K and 4EBP1.

**Discussion**

In the present study, we found that exposure to oxygen concentrations of 40%, 60%, and 90% all promoted the apoptosis of L929 cells in a time and concentration-dependent manner. Application of 10 nM rapamycin or transfected with mTOR siRNA could inhibit the activation of mTOR signaling pathway and promote the apoptosis of L929 cells exposed to 90% \( O_2 \), and down-regulate the expression of downstream proteins 4EBP1, and p70S6K. Exposure to 90% oxygen increased the production of col I, col III, and FN in ECM of L929 cells; however, inhibition of the mTOR signaling pathway decreased the production of col I, III and FN, which could reduce hypoxia-induced pulmonary fibrosis. In lung tissues of hyperoxia-induced lung injury rats, we found that rapamycin inhibited

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**Table 3.** The concentrations of TGF-β, CTGF and col I in the lung tissue determined by ELISA (n = 6)

<table>
<thead>
<tr>
<th>Group</th>
<th>TGF-β [ng/mg]</th>
<th>CTGF [ng/mg]</th>
<th>Col I [ng/mg]</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>25.50 ±1.86</td>
<td>41.23 ±1.08</td>
<td>414.43 ±8.97</td>
</tr>
<tr>
<td>90% ( O_2 ) 3d</td>
<td>33.74 ±2.84</td>
<td>50.72 ±1.80</td>
<td>471.87 ±5.72</td>
</tr>
<tr>
<td>90% ( O_2 )+rapamycin 3d</td>
<td>25.72 ±1.44</td>
<td>43.45 ±1.71</td>
<td>425.11 ±3.50</td>
</tr>
<tr>
<td>90% ( O_2 ) 7d</td>
<td>58.65 ±3.10</td>
<td>68.65 ±2.24</td>
<td>529.72 ±6.97</td>
</tr>
<tr>
<td>90% ( O_2 )+rapamycin 7d</td>
<td>34.39 ±3.32</td>
<td>50.92 ±2.42</td>
<td>466.17 ±0.60</td>
</tr>
<tr>
<td>90% ( O_2 ) 14d</td>
<td>98.81 ±1.55</td>
<td>94.39 ±2.48</td>
<td>556.44 ±8.52</td>
</tr>
<tr>
<td>90% ( O_2 )+rapamycin 14d</td>
<td>54.31 ±4.25</td>
<td>63.64 ±2.78</td>
<td>511.72 ±13.72</td>
</tr>
</tbody>
</table>

* compared with control group, \( p < 0.05 \); \( ^b \) compared with 90% \( O_2 \) groups at the same time, \( p < 0.05 \); col I – collagen I.
the activation of the mTOR pathway, which protected the lung from the oxidative injury and fibrosis induced by high concentration of oxygen.

Lung fibroblasts play an important role in the repair of lung injury. Yang et al. found that high oxygen concentrations inhibited the proliferation of fibroblasts isolated from embryonic lungs. The mTOR signaling pathway is essential for cell proliferation, survival, and migration by regulating the transcription and translation of proteins. Rapamycin is a specific inhibitor of the mTOR pathway. Sun et al. found that 0.01–10 nM rapamycin could inhibit the proliferation of lung cancer cells; however, only >1 nM rapamycin quickly inhibited the phosphorylation of downstream proteins and the growth of lung cancer cells. In the present study, mTOR siRNA down-regulated the expression of mTORC1, 4EBP1, and p70S6K, suggesting that mTOR can specifically regulate the expression of 4EBP1 and p70S6K. The extracellular matrix is synthesized and secreted by animal cells to form a network structure between cells, which consists of protein and polysaccharide. In the present study, we found that blocking the mTOR signaling pathway inhibited collagen deposition in 1L29 cells, which prevented the lung fibrosis caused by oxidative injury. Collagen deposition in the lung is one of the most important causes and major characteristics of pulmonary fibrosis. In the pathology of pulmonary fibrosis, apoptosis-related proteins, such as Bcl-2, P53, and caspase-3, as well as other cytokines, interplay to promote the proliferation of lung fibroblasts, which replace alveolar type II epithelial cells (AT II), subsequently causing pulmonary fibrosis. TGF-β and CTGF could activate the PI3K/Akt/mTOR signaling pathway, promoting the proliferation, collagen synthesis, and anti-apoptotic ability of lung fibroblasts. TGF-β, which is a key component in a number of cytokine networks, is currently recognized to be one of the most powerful factors causing fibrosis. Lee and Kim reported that CTGF was closely associated with TGF-β in pulmonary fibrosis. Connective tissue growth factor is a newly identified cytokine closely involved in pulmonary fibrosis and is widely distributed in human tissues and organs; it plays important roles in cell adhesion, fibroblast proliferation, and ECM synthesis. In the present study, we observed increased expression of CTGF and TGF-β, which was induced by 90% oxygen. Additionally, we found that blocking the mTOR pathway could down-regulate the expression of CTGF and TGF-β, which suggests that rapamycin could decrease the expression CTGF and TGF-β through inhibiting the activation of mTOR signaling pathway. In other words, the mTOR pathway is involved in lung fibrosis by regulating the expression of CTGF and TGF-β.

The mammalian target of rapamycin is widely found in mammals in 2 complex forms: mTORC1 and mTORC2. As the target protein of rapamycin, mTORC1 is sensitive to rapamycin; however, mTORC2 is not. The mammalian target of rapamycin complex 1 regulates cell growth, proliferation, and metabolism by phosphorylating 2 major downstream proteins, S6K1 and eukaryotic initiation factor 4E-BP1. In the present study, we found that 90% O2 induced lung injury in premature rats in a time-dependent manner. We also observed increased expression of fibrosis-related proteins, including TGF-β, CTGF, col II and col III, in the lung tissue of rats, also in a time-dependent manner. These results suggest that lung injury induced by a high concentration of oxygen is consistent with lung fibrosis. High concentrations of oxygen induced changes in the expression of mTORC1, p70S6K, and 4EBP1, which are key components of the mTOR pathway, suggesting that the mTOR pathway is activated by high concentrations of inhaled oxygen. It has been reported that the mTOR pathway is activated in lipopolysaccharide (LPS)-induced acute lung injury (ALI) in mice and rapamycin reduced the expression of inflammatory factors in bronchoalveolar lavage fluid (BLF). Lorne et al. reported that the phosphorylation of rpS6 and 4EBP1, 2 effectors of the mTOR signaling pathway, induced the over-production of inflammatory cytokines by neutrophils in ALI, which could be inhibited by rapamycin. It has been reported that rapamycin administration causes a significant reduction of p70S6K phosphorylation, increased autophagy, decreases neuronal cells apoptosis and significantly reduces brain damage in neonatal rats, which testifies to the neuroprotective effect of rapamycin in neonatal hypoxia-ischemia. Furthermore, it was found that the inhibition of the mTOR signaling pathway by rapamycin prevented the development and progression of lung fibrosis in a rat pulmonary fibrosis model that was induced by TGF-α. Tulek et al. also found that rapamycin could inhibit the progress of lung fibrosis in the early stage of bleomycin-induced pulmonary fibrosis in rats. Therefore, we speculate that the mTOR signaling pathway plays an important role in the repair of hyperoxia-induced lung injury by regulating the expression of TGF-β and CTGF. Rapamycin can effectively inhibit the activation of the mTOR signaling pathway and downregulate downstream target proteins P70S6K and 4EBP1, and reduce the production of TGF-β and CTGF, inhibiting the development of pulmonary fibrosis induced by high concentration oxygen exposure.

Conclusions

In conclusion, our results suggest that rapamycin prevents the progression of lung fibrosis induced by high concentrations of oxygen by inhibiting the mTOR signaling pathway, including suppressing the expression of p70S6K, 4EBP1, TGF-β, and CTGF. Therefore, rapamycin may be useful for the treatment of hyperoxia-induced ALI. Our results help elucidate the molecular mechanism underlying hyperoxia-induced lung injury and may contribute to the identification of novel targets for its treatment.
References