Hyperoxia induces alveolar epithelial cell apoptosis by regulating mitochondrial function through small mothers against decapentaplegic 3 (SMAD3) and extracellular signal-regulated kinase 1/2 (ERK1/2)

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ABSTRACT

Oxygen therapy and mechanical ventilation are widely used to treat and manage neonatal emergencies in critically ill newborns. However, they are often associated with adverse effects and result in conditions such as chronic lung disease and bronchopulmonary dysplasia. Hence, a clear understanding of the mechanisms underlying hyperoxia-induced lung damage is crucial in order to mitigate the side effects of oxygen-based therapy. Here, we have established an in vitro model of hyperoxia-induced lung damage in type II alveolar epithelial cells (AECIIs) and delineated the molecular basis of oxygen therapy-induced impaired alveolar development. Thus, AECIIs were exposed to a hyperoxic environment and their cell viability, cell cycle progression, apoptosis, mitochondrial integrity and dynamics, and energy metabolism were assessed. The results showed that hyperoxia has no significant effect as an inhibitor of SMAD3 and ERK1/2 in AECIIs, but leads to significant inhibition of cell viability. Further, hyperoxia was found to promote AECII apoptosis and mitochondrial, whereas chemical inhibition of SMAD3 or ERK1/2 further exacerbated the detrimental effects of hyperoxia in AECIIs. Overall, these findings presented herein demonstrate the critical role of SMAD/ERK signaling in the regulation of AECII behavior in varying oxygen environments. Thus, this study offers novel insights for the prevention of neonatal lung dysfunction in premature infants.

1. Introduction

The survival rate of critically ill newborns, especially premature babies, has shown continuous improvement with the advancement of intensive care technology. Among the methods currently in use, oxygen therapy and mechanical ventilation are the prevalent procedures in the neonatal intensive care unit. However, long-term application of these methods has led to increased incidences of chronic lung disease and bronchopulmonary dysplasia in the newborns [1]. These conditions are the main cause of chronic respiratory diseases in infancy, which severely affect the quality of life in premature infants. Clinical studies have revealed that the etiology of these adverse conditions is closely related to immature lung development and oxidative stress-related injury in the lung [2,3]. The main pathological features of damaged lung tissues is stagnant or disordered alveolar development, which is manifested as a decrease in the number of alveoli, an increase in size, simplification of alveolar structure, and pulmonary interstitial fibrosis [4,5]. It is speculated that the attributable cause underlying these phenomena may be the obstruction of lung tissue maturity due to alveolar growth arrest [6]. To further understand the in-depth etiology and combat this issue, it is essential to investigate the molecular basis underlying the relevant stages of alveolar development. Specific molecular mechanisms regulating processes such as the proliferation, differentiation, and rearrangement of type II alveolar epithelial cells and lung fibroblasts are thus currently under research to promote the development of strategies against chronic lung disease and bronchopulmonary dysplasia in children.

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There are two types of alveolar epithelial cells: type I and type II. Type I cells cover most of the alveolar surface and can be replaced by type II cells upon damage. Additionally, type II alveolar epithelial cells (AECIIs) play an important role in maintaining alveolar integrity and assisting immune defense [7–10]. AECII apoptosis is a pathological event associated with hyperoxia-induced lung injury [11], involving a variety of molecular pathways. Mitochondrial dysfunction has been identified as an important contributor to impaired alveolar development in mice subjected to hyperoxia treatment [12]. In addition, mitochondrial dysfunction is one of the factors associated with poor alveolar development and failure of lung maturation [13]. In particular, loss of small mothers against decapentaplegic 3 (SMAD3) is associated with impaired alveogenesis in the lungs of neonatal mice [14]. Moreover, SMAD3 has also been shown to be linked to vascular remodeling in pulmonary arterial hypertension [15].

Previous studies on hyperoxia-induced lung injury have shown that SMAD3 inhibitor is involved in regulating leukocyte infiltration, fibrin deposition, activation of nicotinamide adenine dinucleotide phosphate (NADPH) oxidase (NOX), and the expression of MMP-2 and MMP-9 [16]. In addition, recent studies of sepsis induced lung injury in rats have demonstrated that drug-mediated inhibition of transforming growth factor -β1 (TGF-β1) and SMAD3 signaling pathways significantly improved the epithelial and endothelial microvascular permeability [17]. Extracellular signal-regulated kinase 1/2 (ERK1/2) signaling has also been implicated in lung development. It has been documented that activation of ERK1/2 signaling via phosphorylation is crucial for lung fibroblast proliferation, transdifferentiation, and migration in neonatal rats [18]. The ERK cascade is generally considered to be a pathway mediator associated with the protective effect of growth factors on cell death. While many studies have reported pro-survival effects of the ERK cascade in hyperoxia, there are a few studies that have also documented pro-apoptotic effects of the same under certain conditions. Hyperoxic exposure has been reported to increase ATP release in human lung endothelial cells, which is also apparently involved in ERK activation.

Further, the ERK activation results in a cell survival response that prevents hyperoxia-induced cell cycle arrest in these cells [19]. However, whether a direct relationship between hyperoxia-induced lung damage and SMAD/ERK signaling in addition to the possible involvement of mitochondrial dynamics exists in AECIIs currently remain unknown.

In this study, we have investigated whether hyperoxia impairs alveolar development and induces lung damage via SMAD3 and/or ERK1/2 signaling. Accordingly, we subjected rat AECIIIs to in vitro hyperoxia treatment and/or inhibition of SMAD3 or ERK1/2, and examined the changes in cell viability, cell cycle progression, apoptosis, mitochondrial dynamics, and energy metabolism in these cells. The findings of this study aim to establish the molecular basis underlying hyperoxia-induced lung impairment and provide novel insights into the prevention of neonatal lung dysfunction in premature infants.

2. Materials and methods

2.1. Cell culture and treatment

The rat AECII cells RLE-6TN were cultured in RPMI-1640 medium (SH30809.01B, Hyclone Laboratories Inc., UT, USA) containing 10% fetal bovine serum (10,270–106, Gibco, Thermo Fisher Scientific Corporation., MA, USA) and 1% penicillin-streptomycin at 37°C in an atmosphere containing 5% CO₂. For culture in normoxia, the cells were incubated in an atmosphere containing 21% oxygen, 74% nitrogen, and 5% CO₂ for 24 h. For culture in hyperoxia, the cells were incubated in an atmosphere containing 95% oxygen and 5% CO₂ for 24 h. For the inhibition of SMAD3 or ERK1/2, RLE-6TN cells were respectively pre-treated for 4 h with 3 μM SIS3HCL [20], an inhibitor of SMAD3 (S7959, Selleck Chemicals., TX, USA) or 1 h with 10 μM SCH772984 [21], an inhibitor of ERK1/2 (S7101, Selleck Chemicals.) before normoxic or hyperoxic treatment.

2.2. Western blotting

After normoxic or hyperoxic culture, cell samples were lysed to extract total proteins and the protein
concentration was measured using a bicinchoninic acid assay (PC0020, Beijing Solarbio Science & Technology Co., Ltd., Beijing, P. R. China). For Western blotting, 20 μg of protein samples to be loaded per lane were first denatured by boiling in sample loading buffer for 10 min. The samples were then loaded and resolved onto a 12% polyacrylamide gel for sodium dodecyl sulfate-polyacrylamide gel electrophoresis at 120 V for 50 min. At the end of the reaction, the proteins were transferred onto polyvinylidene fluoride membranes (pre-soaked in acetone for 5 min) at 90 V for 50 min. The membranes were blocked with 5% skimmed milk overnight at 4°C and incubated overnight at 4°C with rabbit primary antibodies against phosphorylated (P-) SMAD3 (PAB43521-P, 1:1000, Bioswamp, Wuhan, P. R. China), P-ERK1/2 (PA36335-P, 1:1000, Bioswamp), B-cell lymphoma 2 (BCL-2, PAB33482, 1:1000, Bioswamp), Bad (PAB32756, 1:1000, Bioswamp), mitofusin 1 (MFN1, PAB34056, 1:1000, Bioswamp), Mfn2 (PAB31663, 1:1000, Bioswamp), or GAPDH (PAB36369, 1:1000, Bioswamp). After three washes with phosphate-buffered saline (PBS) with 0.1% Tween20 (PBS-T) for 5 min each, the membranes were incubated at room temperature for 1 h with goat anti-rabbit IgG secondary antibodies (SAB43714, 1:10,000, Bioswamp) and washed again three times with PBS-T for 5 min each. The membranes were then immersed in enhanced chemiluminescence reaction solution in the dark and the protein bands were scanned using a Tanon-5200 automatic analyzer (Tanon Science & Technology, Shanghai, China). The band gray values were analyzed using Tanon GIS software.

Thereafter, 10 μL of CCK-8 solution (CA1210, Beijing Solarbio Science & Technology Co., Ltd., P. R. China) was added to each well and the cells were further cultured for 4 h. Subsequently, the absorbance of the wells was measured using a microplate reader (AMR-100, Hangzhou Alsheng Instrument Co., Ltd., Zhejiang, P. R. China) at 450 nm.

2.4. Flow cytometry

2.4.1. Detection of cell cycle progression

After normoxic or hyperoxic culture, 1 × 10⁷ RLE-6TN cells were resuspended in 1 mL of culture medium and centrifuged for 5 min at 400 × g. The supernatant was discarded and the cells were resuspended in 300 μL of PBS. Subsequently, 700 μL of anhydrous ethanol was added and the cells were fixed at −20°C for 24 h. The cells were then centrifuged at 4°C for 5 min at 700 × g and the supernatant was removed. The cell pellet was washed twice with cold PBS and resuspended in 100 μL of 1 mg/mL RNase A solution (550,825, BD Bioscience). After 30 min, 400 μL of 50 μg/mL propidium iodide (PI, 550,825, BD Bioscience) was added and the cells were incubated in the dark at 4°C for 10 min. The cells were thereafter subjected to flow cytometry (NovoCyte, ACEA Biosciences, Agilent Technologies, Inc. CA, USA) to measure the proportion of cells in each phase of the cell cycle. The data were analyzed using NovoExpress software.

2.4.2. Detection of cell apoptosis

After exposure to normoxic or hyperoxic conditions, 1 × 10⁶ RLE-6TN cells were resuspended in culture medium and centrifuged at 4°C for 5 min at 400 × g. The supernatant was discarded and the cells were resuspended in 1 mL of cold PBS. The cells were centrifuged again at 4°C for 5 min at 400 × g and the supernatant was removed. The cells were resuspended in 200 μL of PBS with 10 μL of Annexin V conjugated to fluorescein isothiocyanate (Annexin V-FITC) and 10 μL of PI (556,547, BD Biosciences). After gentle mixing, the cells were incubated for 30 min at 4°C in the dark, followed by the addition of 300 μL of PBS to

2.3. Cell Counting Kit-8 (CCK-8) assay

RLE-6TN cells (3 × 10³ per well) were resuspended in 100 μL of medium and seeded into the wells of a 96-well plate. After overnight culture at 37°C in an atmosphere containing 5% CO₂, the cells were subjected to SMAD3 or ERK1/2 inhibition or/and 24 h of normoxic/hyperoxic culture.
the cells [22]. Flow cytometry (NovoCyte, ACEA Biosciences) was performed to evaluate cell apoptosis and the data were analyzed using NovoExpress software.

2.4.3. Detection of mitochondrial membrane potential (MMP)

The 5,5′;6,6′;3,3′-tetachloro-1,1′;3,3′-tetrachryl-imidacarbocyanine iodide (JC-1) assay was performed to examine MMP using a commercial kit (C2006, Beyotime Biotechnology, Shanghai, P. R. China). After normoxic or hyperoxic culture, 1 × 10^6 RLE-6TN cells were resuspended in 0.5 mL of culture medium, to which 500 μL of JC-1 working solution was added. The cells were gently agitated and incubated at 37°C for 20 min, followed by centrifugation at 4°C for 3 min at 400 × g. The supernatant was then removed and the cell pellet was washed twice with JC-1 buffer. The cells were resuspended in 1 mL of JC-1 buffer and centrifuged at 4°C for 3 min at 400 × g, and the supernatant was discarded twice. The cells were finally resuspended in 400 μL of JC-1 buffer and flow cytometry (NovoCyte, ACEA) was performed to evaluate the MMP in the indicated cells. The data were analyzed with NovoExpress software.

2.5. Transmission electron microscopy (TEM)

After normoxic or hyperoxic culture, 1 × 10^7 RLE-6TN cells were centrifuged for 5 min at 400 × g. The supernatant was discarded and the cells were fixed with 2 mL of 2.5% glutaraldehyde at 4°C for 30 min. After fixation, the cells were washed three times with 0.1 mol/L PBS for 10 min each and fixed with 1% osmium acid for 1 h. The cells were washed three times with 0.1 mol/L PBS for 10 min each and dehydrated in a graded series of ethanol. Subsequently, the cells were embedded and cut into ultrathin sections (~60 nm in thickness). The sections were stained with uranyl acetate in the dark for 20 min and washed three times with double-distilled water [23]. After the sections were dried with filter paper, they were observed by using a Hitachi HT7700 transmission electron microscope.

2.6. Biochemical assays

After normoxic or hyperoxic culture, ATP production (A095, Nanjing Jiancheng Bioengineering Inc.), lactic acid production (A019-2, Nanjing Jiancheng Bioengineering Inc.), and glucose consumption (KA4086, Abnova) were measured in RLE-6TN cells subjected to various indicated treatments using corresponding commercial assay kits, in accordance with the instructions included in the manufacturer’s manual.

2.7. Statistical analysis

All experiments were performed in three replicates (n = 3) and the data are presented as the mean ± standard deviation. One-way analysis of variance followed by Tukey’s post hoc test was performed to evaluate the significance of the difference between the mean values. A P value of less than 0.05 was considered statistically significant.

3. Results

In this study, RLE-6TN cells were cultured in normoxic or hyperoxic conditions to evaluate the phosphorylation of SMAD3 and ERK1/2, and to study the relationship between hyperoxia and SMAD/ERK signaling pathway. Secondly, the cell cycle progression and apoptosis of RLE-6TN cells were detected in the presence/absence of hyperoxia and/or inhibition of SMAD3 or ERK1/2, and their effects on mitochondrial function and cell energy metabolism were evaluated. Overall, this study aimed to offer novel insights that may serve to be helpful for the prevention of neonatal lung dysfunction in premature infants.

3.1. Hyperoxia and SMAD3/ERK inhibition synergistically suppress the viability of RLE-6TN cells

To delineate the relationship between hyperoxia and SMAD/ERK signaling, the phosphorylation of SMAD3 and ERK1/2 was evaluated in RLE-6TN cells that were cultured in normoxic or hyperoxic conditions (Figure 1a). Our results revealed that hyperoxia exerts no significant effect on the expression of SMAD3 and ERK1/2. In
addition, the effect of SMAD3/ERK inhibition was investigated by pretreating the RLE-6TN cells for 4 h with an inhibitor of SMAD3 (3 μM SIS3HCL or SMAD3-i) or 1 h with an inhibitor of ERK1/2 (10 μM SCH772984 or ERK1/2-i) and followed by assessment of cell viability post exposure to normoxic or hyperoxic conditions for 24 h (Figure 1b). Our results showed that under normoxia, inhibition of neither SMAD3 nor ERK1/2 affected the viability of RLE-6TN cells. Moreover, the significant decrease in the cell viability observed upon exposure to hyperoxia was further exacerbated upon inhibition of either SMAD3 or ERK1/2. These findings suggest that inhibition of SMAD3 or ERK1/2 acts in a synergistic with hyperoxia to reduce the viability of RLE-6TN cells.

3.2. Hyperoxia and SMAD3/ERK inhibition regulate cell cycle progression and apoptosis of RLE-6TN cells

We next investigated the role of hyperoxia and/or inhibition of SMAD3 or ERK1/2 on the regulation of cell cycle progression and apoptosis in RLE-6TN cells. Our flow cytometry studies revealed that under normoxia, SMAD3 or ERK1/2 inhibition does not affect the proportion of cells in each phase of the cell cycle. However, exposure to hyperoxia was found to promote cell cycle arrest at the G1 phase, and this effect was further accentuated by SMAD3 or ERK1/2 inhibition (Figure 2a). Similarly, SMAD3 or ERK1/2 inhibition showed no effect on the apoptosis of RLE-6TN cells under normoxic conditions. In contrast, hyperoxia was found to significantly enhance the apoptosis of RLE-6TN cells, and this effect was further accentuated upon pre-treatment of the cells with SMAD3-i or ERK1/2-i under hypoxia as compared to non-pretreated cells (Figure 2b). These results were further complemented by Western blotting results, which showed that hyperoxia results in downregulation of the anti-apoptotic protein BCL-2 and upregulation of the pro-apoptotic proteins BAD and cleaved Caspase-3 (Figure 2c). In addition, SMAD3 or ERK1/2 inhibition resulted in exacerbation of these apoptotic effects only under hyperoxia. Moreover, our results also showed that SMAD3-i and ERK1/2-i have no effect on the expression of apoptosis associated proteins under normoxia.

3.3. Hyperoxia and SMAD3/ERK inhibition alter mitochondrial dynamics in RLE-6TN cells

After determining the effect of hyperoxia and SMAD3/ERK inhibition on the cell viability, cell cycle progression, and apoptosis, its effect on mitochondrial function was evaluated in the RLE-6TN cells. Our TEM observations showed that under normoxia, the RLE-6TN cells exhibited
the typical morphology of healthy mitochondria both, in the absence or presence of SMAD3-i or ERK1/2-i (Figure 3a). However, hyperoxia apparently led to mitochondrial damage, and this damage was more pronounced when the RLE-6TN cells were cultured in the presence of SMAD3-i or ERK1/2-i as compared to their absence under hyperoxia. Further, Western blotting was performed to detect the expression of proteins associated with mitochondrial fission (DRP1) and fusion (MFN1 and MFN2) in the RLE-6TN cells pretreated with SMAD3-i or ERK1/2-I under normoxic or hyperoxic conditions (Figure 3b). Our results revealed that hyperoxia significantly downregulates the expression of MFN1 and MFN2 and upregulates the expression of DRP1 in RLE-6TN cells as compared to normoxia. This effect of hypoxia on the expression of mitochondrial proteins associated with fusion and fission was further enhanced by both SMAD3 and ERK1/2 inhibition. Furthermore, the MMP in the RLE-6TN cells was also found to be significantly decreased under hyperoxia as compared to normoxia, and this effect became more apparent upon SMAD3 or ERK1/2 inhibition (Figure 3c). Thus, in both the cases, inhibition of SMAD3 and ERK1/2 exerted less significant effect on the levels of mitochondrial proteins and MMP in the RLE-6TN cells under normoxia than hyperoxia.

3.4. Hyperoxia and SMAD3/ERK inhibition regulated energy metabolism in RLE-6TN cells

Finally, the effect of hyperoxia and SMAD/ERK inhibition on energy metabolism was examined...
in RLE-6TN cells pretreated with SMAD3-i or ERK1/2-I under normoxic or hyperoxic conditions. Our results demonstrated that under hyperoxia, ATP production (Figure 4a) was significantly reduced, whereas lactic acid production (Figure 4b) and glucose consumption (Figure 4c) were significantly promoted in the RLE-6TN cells as compared to those cells assayed under normoxia. Moreover, inhibition of SMAD3 or ERK1/2 further suppressed the ATP production but did not affect lactic acid production and glucose consumption in RLE-6TN cells subjected to hyperoxia.
hyperoxic as compared to the normoxic conditions. Additionally, inhibition of either SMAD3 or ERK1/2 did not exert any effect on energy metabolism under normoxia.

4. Discussion

In this study, we have successfully established an *in vitro* model of hyperoxia-induced AECII
damage by culturing RLE-6TN cells in the presence of excess oxygen. In addition, we have shown that chemical inhibition of SMAD3 or ERK1/2 exerts no effect on RLE-6TN cell behavior under normoxic conditions, but aggravates the damaging effects of hyperoxia. In particular, hyperoxia-induced suppression of RLE-6TN cell viability, promotion of RLE-6TN cell apoptosis, induction of mitochondrial damage (increased fission and decreased fusion), and induction of MMP (indicative of increased mitochondrial depolarization) were all exacerbated by the inhibition of SMAD3 or ERK1/2 as compared to normoxia.

As a key oxygen-sensing organelle in the cell, mitochondria adapt to varying oxygen levels in order to maintain homeostasis [24]. Even subtle changes in cellular oxygen levels can affect cellular function and fate. Thus, the balance between hyperoxia and hypoxia is of critical importance in cellular physiology, and an in-depth understanding of the role of mitochondria and the mechanisms involved therein is particularly imperative. Mitochondrial dysfunction has been shown to accompany the development and progression of lung diseases such as idiopathic pulmonary fibrosis and bronchopulmonary dysplasia [25,26]. A number of studies have uncovered the relationship between hyperoxia and mitochondrial dysfunction. For instance, Ma et al. have demonstrated that hyperoxia induces mitochondrial damage by prompting fragmentation (fission), which is evident by the upregulation of pro-fission proteins under hyperoxia as compared to normoxia [27]. Mu et al. have reported that hyperoxia leads to AECII apoptosis and reduced MMP in a model of acute lung injury [28]. Further, Yu et al. have shown that hyperoxia-exposed AECIIs exhibit abnormal mitochondrial morphology and decreased MMP, which were possibly associated with impaired alveolarization in neonatal rats [29]. In the present study, our data are consistent with previously published results showing that hyperoxia induces mitochondrial damage by promoting fission and mitochondrial depolarization in AECIIs. This was confirmed by the upregulation of the fission-related protein DRP1, downregulation of the fusion-related proteins MFN1 and MFN2, and suppression of MMP. Concurrently, the changes in mitochondrial dynamics also led to corresponding alterations in cell cycle progression, and mitochondrial fission was possibly the reason for the enhanced apoptosis observed under hyperoxic conditions.

In our study, we have also revealed that the detrimental effect of hyperoxia-induced mitochondrial dysfunction in AECIIs was associated with the regulation of SMAD3 and ERK1/2 signaling. Changes in oxygen level affect the activation of signaling cascades involved in a variety of cellular processes. Exposure of cells to hypoxic conditions (low levels of oxygen) has been shown to induce the phosphorylation of SMADs [30] and ERK1/2 [31–33]. Accordingly, we hypothesized that hyperoxia, a condition wherein oxygen exists in over-abundance, exerts the opposite effects. More specifically, we hypothesized that, hyperoxia inhibits the activation of SMAD/ERK signaling by suppressing the phosphorylation of associated proteins in the pathway, which was also observed in our results. SMAD3 deficiency reportedly leads to impaired mitochondrial biogenesis in skeletal muscle regeneration [34], and the regulation of mitochondrial ERK1/2 activation has been shown to be an important aspect of rat brain development [35]. However, the role and effect of SMAD3 and ERK1/2 signaling in alveolar development and pulmonary remodeling remain unclear. In this study, we have verified that SMAD3 and ERK1/2 signaling are crucial in the maintenance of alveolar integrity, as their inhibition results in AECII apoptosis due to alterations in mitochondrial stability and dynamics in conjunction with hyperoxia.

An interesting observation in our study was that inhibition of SMAD3 or ERK1/2 only affected AECII viability, cell cycle progression, apoptosis, and mitochondrial dynamics under hyperoxic conditions. In normoxia, neither SMAD-i nor ERK-i showed any prominent effect, suggesting that chemical inhibition of SMAD3/ERK is detrimental to AECIIs only when SMAD3 or ERK1/2 signaling was already partially suppressed by hyperoxia. In other words, chemical inhibition of SMAD3 and ERK1/2 using SIS3HCL and SCH772984, respectively, accentuated the effect of hyperoxia in a synergistic manner. In addition, the balance between mitochondrial fission and fusion is only affected by ERK1/2 inhibition but not by SMAD3 inhibition, suggesting that the actions of
SMAD3 and ERK1/2 are mutually exclusive. We thus propose that mitochondrial function in AECII is influenced by hyperoxia in an ERK1/2-dependent but SMAD3-independent manner. Finally, assessment of energy metabolism revealed that inhibition of either SMAD3 or ERK1/2 lowers ATP production in the presence of hyperoxia, but does not affect the lactic acid content and glucose consumption even under hyperoxia.

Our study faces some limitations that will be addressed in the prospective work. First, it is unclear whether compensation of SMAD3 or ERK1/2 via, for instance, overexpression can counteract or circumvent the negative effects of hyperoxia on AECII. Moreover, the crosstalk between SMAD3 and ERK1/2, if any, was not explored. Thus, there may be additive or synergistic interplay between these pathways that has yet to be elucidated. These questions will therefore form the basis of future study designs that aim to expand on the current topic. The specific involvement of SMAD3/ERK signaling in neonatal lung development also needs to be verified in vivo using animal models in follow-up studies.

5. Conclusion

The findings reported in this study demonstrate that hyperoxia impairs the growth of AECIIIs (RLE-6TN cells) by promoting cell apoptosis, inducing mitochondrial damage, and affecting energy metabolism. In addition, we have also reported that hyperoxia acts as an inhibitor of SMAD3 and ERK1/2, suggesting that SMAD3/ERK signaling is critically involved in the regulation of AECII behavior in environments with varying oxygen content. This study may thus offer innovative insights in developing therapeutic strategies for the prevention of neonatal lung dysfunction in premature infants.

Disclosure statement

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

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Data availability

The data that support the findings of this study are available from the corresponding author upon reasonable request.

Author contributions

DH designed the study and supervised the project. JJ, JW, CL, LM, and DH performed the experiments. JJ and DH analyzed the experimental data. DH wrote and revised the manuscript.

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