



# Effects of liver X receptor in O<sub>3</sub>-induced airway inflammation and remodeling in mice

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**Background:** The current clinical treatment of chronic obstructive pulmonary disease (COPD) mainly uses drugs to improve symptoms, but these drugs cannot reverse the progression of the disease and the pathological changes in lung tissue. This study aimed to investigate the effects and mechanisms of Liver X receptors (LXRs) in ozone (O<sub>3</sub>)-induced airway inflammation and remodeling in mice.

**Methods:** Wild mice and LXR deficient mice were exposed to O<sub>3</sub> twice a week for 6 weeks. Some wild mice were intraperitoneally injected with T0901317 (a LXR agonist) before O<sub>3</sub> exposure. Wild mice were exposed to ambient air and intraperitoneally injected with normal saline (NS) as control group. The lung tissues and bronchoalveolar lavage fluid (BALF) were collected to evaluate airway inflammation, airway remodeling and lipid disorder.

**Results:** After O<sub>3</sub> exposure, LXR deficient mice showed severe airway inflammation and airway remodeling compared with the wild mice. There were a lot of foamy macrophages appeared in BALF of LXR deficient mice. The inflammatory proteins such as myeloid differentiation primary response protein 88 (MyD88) and interleukin-1 receptor-associated kinase (IRAK) in the lung tissues of LXR deficient mice were significantly increased compared with the wild mice. In wild mice exposed to O<sub>3</sub>, T0901317 treatment can alleviate airway inflammation, airway remodeling and foamy macrophages in BALF. And MyD88 and IRAK expression in lung tissue were also attenuated by T0901317 treatment.

**Conclusions:** LXRs play protective roles in O<sub>3</sub>-induced lipid accumulation, airway inflammation and airway remodeling.

**Keywords:** Liver X receptor (LXR); O<sub>3</sub>; airway inflammation; airway remodeling

Submitted Nov 29, 2023. Accepted for publication Jun 21, 2024. Published online Aug 14, 2024.

doi: 10.21037/jtd-23-1820

View this article at: <https://dx.doi.org/10.21037/jtd-23-1820>

## Introduction

Ozone (O<sub>3</sub>) is a common air pollutant that can cause various adverse health issues in humans, particularly respiratory diseases. Inhalation of O<sub>3</sub> can result in airway

inflammation and decreased lung function. Clinical studies have confirmed that high-level O<sub>3</sub> exposure can exacerbate symptoms and increase hospitalization rate in patients with asthma (1) and chronic obstructive pulmonary disease

(COPD) (2). Previous studies have demonstrated that chronic O<sub>3</sub> exposure can induce an emphysema-like model in mice, characterized by features such as emphysema, neutrophilic inflammation, and elevated cytokine levels (3-5). O<sub>3</sub> has been employed in various animal species to induce lung inflammation and airway remodeling (6,7). However, the precise mechanisms underlying O<sub>3</sub>-mediated lung injury remain unclear. Inhaled O<sub>3</sub> readily reacts with lung lining fluid, which is distributed throughout the airway, forming a lipid ozonation product (LOP), which may contribute to the pro-inflammatory effect of O<sub>3</sub>. Within the human airway epithelium, LOP can activate pro-inflammatory factors, such as interleukin-6 (IL-6) and interleukin-8 (IL-8) (8,9).

Liver X receptors (LXRs), part of the nuclear receptor superfamily, are transcription factors with two isoforms:  $\alpha$  (NR1H3) and  $\beta$  (NR1H2) (10). When bound to natural ligands, such as oxysterols and desmosterol (11) or synthetic ligands, such as T0901317 (12), LXRs regulate the transcription of genes related to cholesterol efflux and inflammation mediated by macrophage. Cholesterol is a crucial component of cells and macrophages are the primary inflammatory cells in the alveolar microenvironment. Thus, we hypothesized that LXRs play a significant role in O<sub>3</sub>-induced lipid accumulation and airway injury. In this study, we utilized LXR agonist and LXR-deficient mice to investigate the effect and mechanisms of LXRs following O<sub>3</sub> exposure. We present this article in accordance with the ARRIVE reporting checklist (available at <https://jtd.amegroups.com/article/view/10.21037/jtd-23-1820/rc>).

### Highlight box

#### Key findings

- Liver X receptors (LXRs) play protective roles in ozone (O<sub>3</sub>)-induced airway inflammation and airway remodeling in mice.

#### What is known and what is new?

- Previous research revealed the impact of LXR agonists *in vivo* following O<sub>3</sub> exposure, and the susceptibility to O<sub>3</sub>-induced acute inflammation (3-hour exposure) was only observed in LXR $\alpha^{-/-}$ .
- We confirmed the protective role of LXR in O<sub>3</sub>-induced airway inflammation and remodeling through LXR agonists and LXR gene knockout.

#### What is the implication, and what should change now?

- Our research provides an experimental basis for the study of drug targets for ozone-induced airway inflammation and airway remodeling, and we need to further study its mechanism.

## Methods

A protocol including the research question, key design features, and analysis plan was prepared before the study without registration.

### Experimental animal

Animals were provided with unlimited sterile tap water and standard rodent feed at a temperature of 22±24 °C and a relative humidity of 50%±60% within a specific pathogen-free biosafety Level 3 facility. The animals involved in this experiment consisted of wild mice and gene knockout mice (LXR $\alpha^{-/-}$ , LXR $\beta^{-/-}$ , LXR $\alpha^{-/-}$ / $\beta^{-/-}$ ). The LXR-deficient mice, with a C57BL/6J background, were generated by the Nanjing University-Nanjing Biomedical Research Institute. LXR $\alpha^{-/-}$ / $\beta^{-/-}$  mice were bred from LXR $\alpha^{-/-}$  mice and LXR $\beta^{-/-}$  mice using Clustered Regularly Interspaced Short Palindromic Repeats (CRISPR)/Cas9 technique, where specifically designed and transcribed gRNA were used *in vitro*. Cas9 protein and (guide ribonucleic acid) gRNA was concurrently injected into fertilized mouse eggs. Cas9 protein, guided by gRNA, bound to the target site, inducing deoxyribonucleic acid (DNA) double-strand breaks, ultimately resulting in the deletion of base sequences at the target site and the removal of specific genes. The animal experiments were performed under a project license (No. SYXK [Su] 2016-0006) granted by the Animal Experiment Center of Nanjing First Hospital Affiliated to Nanjing Medical University, in compliance with guidelines established by the institution's Animal Care and Use Committee for the care and use of animals. All animal procedures were conducted under anesthesia (pentobarbital sodium at a concentration of 1%), with every effort made to minimize animal discomfort.

### Exposure to O<sub>3</sub> and LXR agonist treatment

Mice were randomly divided into six groups (n=5 per group):

- (I) Control group: wild mice were exposed to ambient air and administrated normal saline (NS);
- (II) O<sub>3</sub> group: wild mice were exposed to O<sub>3</sub> and administrated NS;
- (III) T0901317 group: wild mice were exposed to O<sub>3</sub> and administrated LXR agonist T0901317;
- (IV) LXR $\alpha^{-/-}$  group: LXR $\alpha^{-/-}$  mice were exposed to O<sub>3</sub> and administrated NS;

- (V) LXR $\beta^{-/-}$  group: LXR $\beta^{-/-}$  mice were exposed to O<sub>3</sub> and administrated NS;
- (VI) LXR $\alpha^{-/-}/\beta^{-/-}$  group: LXR $\alpha^{-/-}/\beta^{-/-}$  mice were exposed to O<sub>3</sub> and administrated NS.

With the exception of the control group, mice were exposed to O<sub>3</sub> produced by an Ozonizer (Sander Ozonizer, Germany) mixed with air at a concentration of 2.5 parts per million (ppm) in a glass container for 3 hours, twice a week for 6 weeks. Continuous monitoring of O<sub>3</sub> concentration using O<sub>3</sub> probe (Analytical Technology, UK). T0901317 (Sigma Aldrich, USA) was dissolved in dimethyl sulfoxide (DMSO) to prepare a 100 mg/mL stock solution. One hour before O<sub>3</sub> exposure, T0901317 (30 mL/kg) was administered intraperitoneally. This dose of T0901317 has been shown to activate ATP-binding cassette transporter A1 (ABCA1) and has an inhibitory effect on inflammatory diseases *in vivo* (13,14). Control animals were exposed to ambient air. Mice were euthanized 24 hours after the last O<sub>3</sub> exposure.

#### ***Bronchoalveolar lavage fluid (BALF) collection***

Mice were euthanized, and tracheotomy was performed. Ice-cold PBS (0.5 mL) was instilled into the lungs and the procedure was repeated three times, and BALF was retrieved. Return volume was recorded and was consistently >80% of the instilled volume. Total cell counts and differential cell counts were determined from the slides stained with Wright-Giemsa stain (KeyGEN BioTECH, Nanjing, China) using an optical microscope (Olympus Optical, Tokyo, Japan). A minimum of 200 cells per mouse were counted under  $\times 400$  magnification, and cells were classified as macrophages, lymphocytes, or neutrophils based on their standard morphological characteristics. The BALF supernatant was collected and stored at  $-80^{\circ}\text{C}$  for future biochemical analysis.

#### ***Enzyme linked immunosorbent assay (ELISA)***

Levels of IL-6, tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), and matrix metalloproteinase 9 (MMP9) in BALF were quantified using ELISA kits from CUSABIO, China, following the manufacturer's instructions.

#### ***Histology and Masson trichrome staining***

After collecting the BALF samples, the left lung tissue was fixed in 10% (v/v) neutral buffered formalin. Subsequently, the

tissue was embedded in paraffin, sectioned to a thickness of 5  $\mu\text{m}$ , and stained with hematoxylin and eosin (HE) solution to assess the inflammatory response. Quantitative analysis was conducted following previously described methods (15,16). Peribronchial inflammation severity was assessed on lung images stained with HE, scored from 0 to 5, where 0= no cells, 1= a few cells, 2= a ring of cells one layer deep, 3= a ring of cells two layers deep, 4= a ring of cells three to four layers deep, and 5= a ring more than four cell layers deep. Additionally, Masson Trichrome staining was employed to detect collagen deposition, following the recommended standard protocol provided by the manufacturer. All histological assessments were conducted in a double-blinded manner under  $400\times$  magnification. The results were quantified by calculating the average of each airway.

#### ***Mean linear intercept (Lm)***

The HE-stained tissues were examined under an optical microscope at a magnification of  $200\times$ , and ten random peripheral visual fields were selected from each section. The section images were imported into Photoshop, where a reference line was drawn passing through the central positive region of the image (while avoiding the areas containing airways and blood vessels as much as possible). The number of alveolar space (n) intersected by these two-reference line was counted, and the total length of the reference line (L) was measured. The average alveolar diameter (Lm) was then calculated using the formula  $Lm = L/N$ . This value serves as an indicator of the average alveolar size and reflects the alveolar damage (5).

#### ***Immunostaining for $\alpha$ -smooth muscle actin ( $\alpha$ -SMA) and MMP9***

The paraffin-embedded tissue sections were initially dewaxed in water. Subsequently, they were immersed in a 3% H<sub>2</sub>O<sub>2</sub> solution at room temperature for 10 minutes to quench the activity of the endogenous peroxidase. Afterward, they were thoroughly rinsed with distilled water and soaked in phosphate buffered saline (PBS) for 5 minutes, repeating this step twice. Following this, the sections were blocked with 5% normal goat serum (diluted with PBS) and left to incubate at room temperature for 10 minutes, after which the excess serum was drained without further washing.

Next, primary antibodies against MMP9 (Servicebio GB11132) and  $\alpha$ -SMA (Servicebio GB13044) were applied to the sections. The section was incubated either for two

hours at 37 °C or overnight at 4 °C. Subsequently, the sections were rinsed with PBS for 5 minutes, repeating this step three times.

Afterward, an appropriate horseradish peroxidase (HRP)-labeled secondary antibody working solution (Servicebio GB23303) was applied to the sections, and they were incubated at 37 °C for 30 minutes. The sections were once again rinsed with PBS for 5 minutes, repeating this step three times. Following this, an adequate amount of horseradish enzyme-labeled streptomycin working solution was applied to the sections, and they were incubated at 37 °C for 30 minutes. The sections were then rinsed with PBS for 5 minutes, repeating this step three times.

Finally, the chromogenic agent (DAB) was applied to the section for 8 minutes. Subsequent steps included thorough flushing, re-staining, dehydration, clearing, and sealing with tap water. The results were quantified based on the averaged optical density within the airway.

#### *Oil red O assay*

BALF was collected from mice and plated on slides with Dulbecco's modified eagle medium (DMEM). The cells were incubated for four hours to allow for adhesion, washed three times with sterile PBS, and fixed with 4% paraformaldehyde fixative solution for 20 minutes. Subsequently, the cells were rinsed three times with distilled water, and the slides were briefly immersed in 60% isopropanol for 15 seconds.

Next, Oil Red O stain solution (KeyGEN BioTECH, Nanjing, China) was applied to each slide, and the slides were incubated for 30 minutes at 37 °C. After 30 minutes, the slides were briefly washed with 60% isopropanol for 15 seconds (with precise timing control). Following this, the cells, were washed with distilled water three times, treated with the hematoxylin dye solution for 5 minutes, covered with a water-based sealing film, and imaged using a Nikon Eclipse Ti microscope (Nikon, Tokyo, Japan).

The results were analyzed by calculating the percentage of stained cells relative to the total number of cells counted.

#### *RNA extraction and real-time polymerase chain reaction (PCR) analysis*

Total RNA from lung tissues were extracted using TRIzol Reagent (Invitrogen). Subsequently, equal amounts of RNA were used to synthesize complementary deoxyribonucleic acid (cDNA) with a reverse transcriptase reagent kit

(TakaRa Biotechnology, Inc., Shiga, Japan). To determine messenger ribonucleic acid (mRNA) levels, real-time PCR analysis was conducted using SYBR Premix Ex Taq (TakaRa Biotechnology, Inc., Shiga, Japan).

This method was used to assess the mRNA levels of the target gene in lung tissues.

#### *Sample preparation and protein analysis*

Lung tissues were used radio-immunoprecipitation assay (RIPA) lysis buffer. Subsequently, the tissue homogenate was subjected to centrifugation at 12,000 rpm for 15 minutes, and the resulting supernatant was carefully collected. The total protein content within the lung tissue was quantified using a bicinchoninic acid (BCA) protein analysis kit (KGP902, KeyGEN BioTECH). The supernatant was mixed with sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) loading buffer in a 4:1 ratio and heated in boiling water for 15 minutes to denature the proteins. Those proteins then transferred onto a polyvinylidene fluoride (PVDF) membrane. The PVDF membrane was subsequently blocked with 5% skim milk at room temperature for 2 hours to prevent nonspecific binding of antibodies. The primary antibodies [ $\beta$ -actin (CST, 4970S), MyD88 (CST, 4283S), interleukin-1 receptor-associated kinase (IRAK) (CST, 4504S)] were diluted in tris buffered saline tween (TBST) buffer and incubated with the membrane overnight at 4 °C. Afterwards, the membrane was washed three times with TBST buffer to remove excess primary antibodies. The membrane was then incubated with the second antibody for 2 hours. Finally, the proteins were visualized using the ECL Key-GEN system.

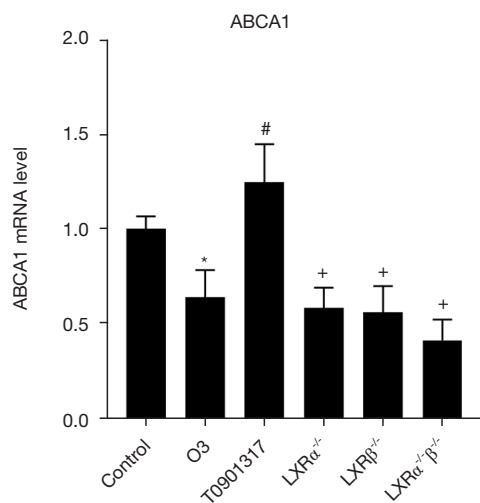
#### *Statistical analysis*

The data were presented as mean  $\pm$  standard error of the mean. Statistical differences among the six groups were analyzed using one-way analysis of variance (ANOVA) (with nonparametric or mixed test) with multiple comparison testing for all datasets (GraphPad Prism 8.0). A significance level of  $P < 0.05$  was considered statistically significant.

## **Results**

### *The effect of LXRs on ABCA1 mRNA in lung after O3 exposure*

Quantitative PCR was used to assess mRNA expression in

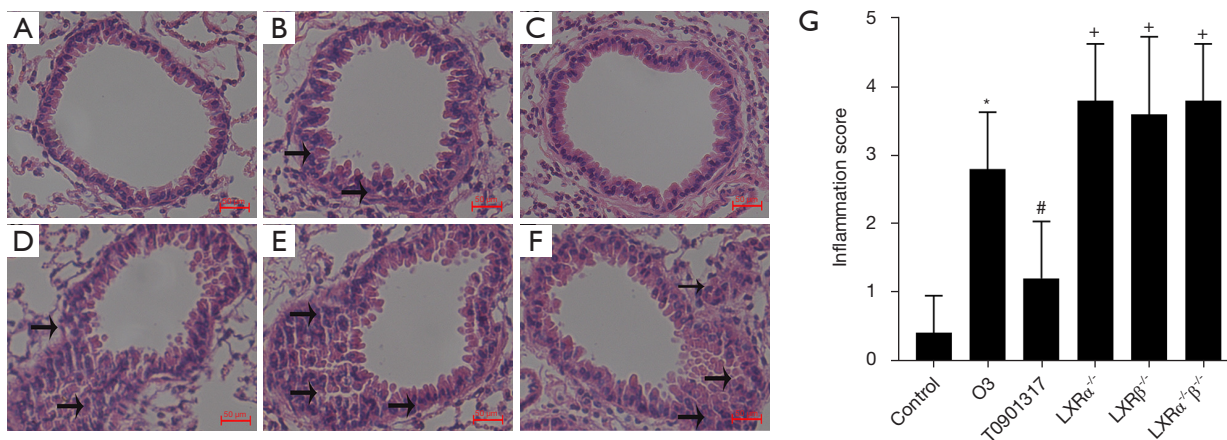


**Figure 1** The effect of LXRs on ABCA1 mRNA in lung after O<sub>3</sub> exposure. Deletion of O<sub>3</sub> and LXR resulted in lower ABCA1 expression compared to the control group. However, mice treated with T0901317 displayed significantly higher ABCA1 levels compared to the O<sub>3</sub> group. \*, P<0.05 O<sub>3</sub> group vs. Control group; #, P<0.05 T0901317 group vs. O<sub>3</sub> group; +, P<0.05 LXR-deficient group vs. O<sub>3</sub> group. Three independent experiments were conducted, with 5 mice in each group for each experiment. ABCA1, ATP-binding cassette transporter A1; LXRs, liver X receptors; mRNA, messenger ribonucleic acid; O<sub>3</sub>, ozone.

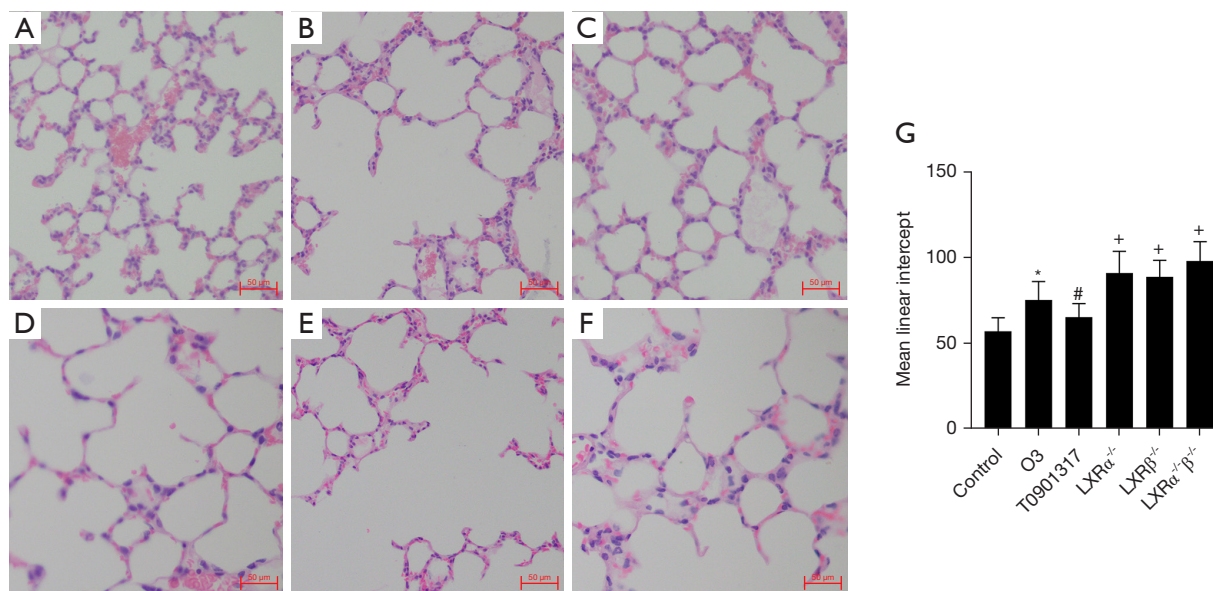
the whole lung. Compared to the control group, ABCA1 mRNA expression decreased significantly after O<sub>3</sub> exposure and increased significantly following T0901317 treatment (P<0.05). LXR $\alpha^{-}/\beta^{-}$  mice exhibited the lowest levels of ABCA1 compared to wild mice (Figure 1). As an essential target gene of LXRs, ABCA1 expression was suppressed by O<sub>3</sub> exposure and LXR deletion, while LXR agonist treatment reversed the inhibition of ABCA1 in wild mice after O<sub>3</sub> exposure.

#### *The effect of LXRs on airway inflammatory cells and alveolar enlargement after O<sub>3</sub> exposure*

Compared with the lungs of mice in the control group, those of mice in the O<sub>3</sub>-exposed group exhibited severe emphysema and neutrophil and macrophage infiltration after O<sub>3</sub> exposure. Mice treated with T0901317 showed a marked reduction in inflammatory cells and alveolar enlargement compared to the O<sub>3</sub>-exposed group (P<0.05). Furthermore, the LXR-deficient mice exhibited more severe airway inflammation and alveolar dilation than the wild mice after O<sub>3</sub> exposure (P<0.05). There were no significant differences in O<sub>3</sub>-induced airway inflammation and alveolar injury among LXR $\alpha^{-}$ , LXR $\beta^{-}$  and LXR $\alpha^{-}/\beta^{-}$  mice (Figures 2,3).



**Figure 2** The effect of LXRs on airway inflammatory cells after O<sub>3</sub> exposure. Representative HE-stained images for the (A) Control group, (B) O<sub>3</sub> group, (C) T0901317 group, (D) LXR $\alpha^{-}/\beta^{-}$  group, (E) LXR $\beta^{-}$  group and (F) LXR $\alpha^{-}/\beta^{-}$  group (magnification,  $\times 200$ ). (G) Inflammation score. The black arrows point to inflammatory cells. \*, P<0.05 O<sub>3</sub> group vs. Control group; #, P<0.05 T0901317 group vs. O<sub>3</sub> group; +, P<0.05 LXR-deficient group vs. O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). LXRs, liver X receptors; O<sub>3</sub>, ozone; HE, hematoxylin and eosin.



**Figure 3** The effect of LXRs on alveolar enlargement after O<sub>3</sub> exposure. (A-F) The representative HE-stained sections from the lungs of the different groups of mice (original magnification, ×200). (A) Control group, (B) O<sub>3</sub> group, (C) T0901317 group, (D) LXRα<sup>-/-</sup> group, (E) LXRβ<sup>-/-</sup> group and (F) LXRα<sup>-/-</sup>/β<sup>-/-</sup> group. (G) Quantitative analysis of the lung destruction, as represented by mean linear intercept. \*, P<0.05 O<sub>3</sub> group vs. control group; #, P<0.05 T0901317 group vs. O<sub>3</sub> group; +, P<0.05 LXR-deficient group vs. O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). LXRs, liver X receptors; O<sub>3</sub>, ozone; HE, hematoxylin and eosin.

#### *The effect of LXRs on inflammatory cells in BALF after O<sub>3</sub> exposure*

After O<sub>3</sub> exposure, the total number of leukocytes, predominantly consisting of macrophages, neutrophils, and lymphocytes, in BALF significantly increased compared to the control group (P<0.05). Pre-treatment with T0901317 before O<sub>3</sub> exposure significantly reduced the number of macrophages, neutrophils, and lymphocytes in BALF (P<0.05). LXR knockout mice, on the other hand, had higher levels of these inflammatory cells compared to wild mice after O<sub>3</sub> exposure (P<0.05). However, no significant differences in inflammatory cell counts were observed among LXRα<sup>-/-</sup>, LXRβ<sup>-/-</sup> and LXRα<sup>-/-</sup>/β<sup>-/-</sup> mice (Figure 4).

#### *The effect of LXRs on the formation of foamy macrophages*

Following O<sub>3</sub> exposure, lipid accumulation was observed in macrophages in both wild mice and LXR knockout mice. This expression of foamy macrophage was significantly higher in absence of LXR mice (P<0.05). However, there was no difference in the percentage of foamy macrophages among the three groups: LXRα<sup>-/-</sup>, LXRβ<sup>-/-</sup> and LXRα<sup>-/-</sup>/β<sup>-/-</sup> mice. Treatment with T0901317 sharply reduced the

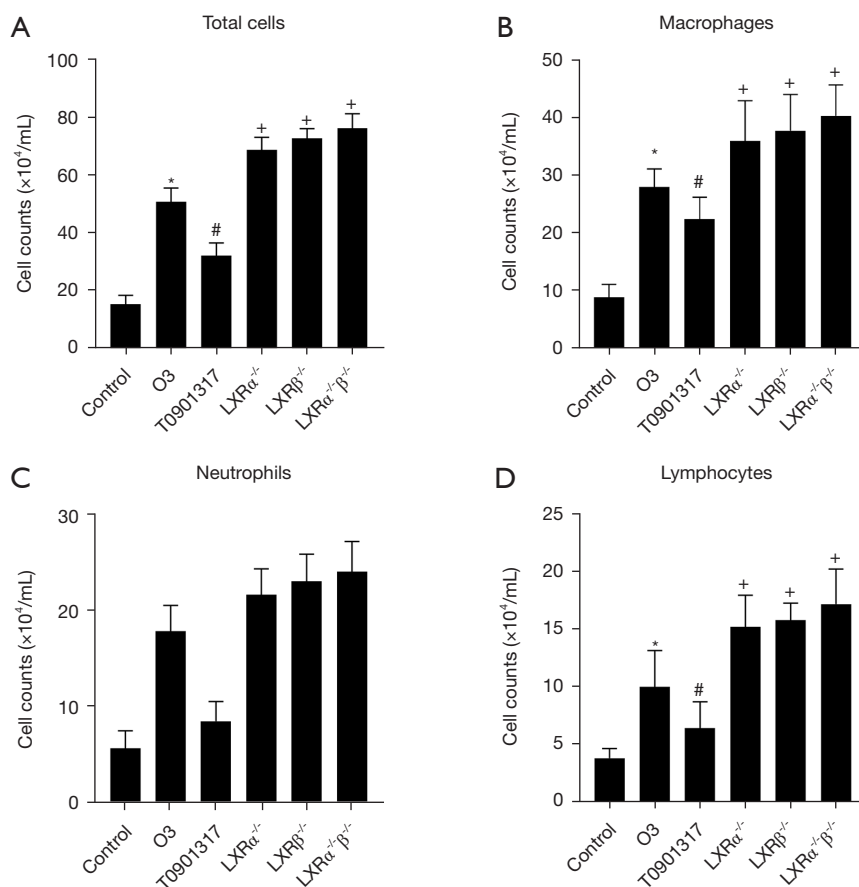
proportion of foamy cells (P<0.05) (Figure 5).

#### *The effect of LXRs on inflammatory cytokines in the BALF after O<sub>3</sub> exposure*

Levels of IL-6, TNF-α, and MMP9 in BALF were significantly higher in the O<sub>3</sub>-exposed group compared to the control group (P<0.05). Levels of these inflammatory cytokines were lower in the T0901317-treated group compared to the O<sub>3</sub>-exposed group (P<0.05). LXR-deficient exhibited higher levels of these cytokines in BALF than wild mice (P<0.05). No changes in inflammatory cytokine expression were observed in LXRα<sup>-/-</sup>, LXRβ<sup>-/-</sup> and LXRα<sup>-/-</sup>/β<sup>-/-</sup> mice (Figure 6).

#### *The effect of LXRs on MyD88 and IRAK after O<sub>3</sub> exposure*

Expression of MyD88 and IRAK increased after O<sub>3</sub> exposure in wild mice. Pre-treatment with T0901317 before O<sub>3</sub> exposure attenuated the levels of MyD88 and IRAK in wild mice (P<0.05). LXR-deficient mice showed increased expression of MyD88 and IRAK compared to



**Figure 4** The effect of LXRs on inflammatory cells in BALF after O<sub>3</sub> exposure. Total cells in BALF were counted using a cell counting plate and stained with Giemsa dye (A). Then, different cell numbers were counted at 400 $\times$  magnification using a microscope ( $\geq 200$  cells were counted) (B-D). \*,  $P < 0.05$  O<sub>3</sub> group vs. control group; #,  $P < 0.05$  T0901317 group vs. O<sub>3</sub> group; +,  $P < 0.05$  LXR-deficient group vs. O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). LXRs, liver X receptors; BALF, bronchoalveolar lavage fluid; O<sub>3</sub>, ozone.

wild mice after O<sub>3</sub> exposure ( $P < 0.05$ ) (Figure 7). Expression of MyD88 and IRAK did not differ among LXR $\alpha$ <sup>-/-</sup>, LXR $\beta$ <sup>-/-</sup> and LXR $\alpha$ <sup>-/-</sup> $\beta$ <sup>-/-</sup> groups.

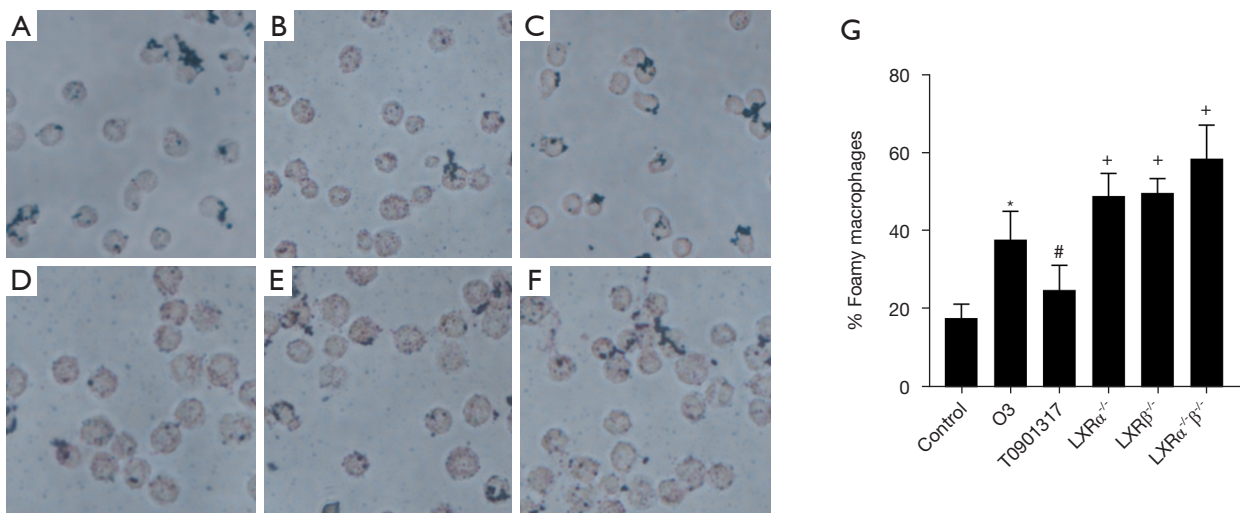
#### The effect of LXRs on O<sub>3</sub>-induced submucosal fibrosis and smooth muscle hyperplasia

O<sub>3</sub> exposure induced submucosal fibrosis (Figure 8) and smooth muscle hyperplasia (Figure 9) in wild mice. Pre-treatment with T0901317 before O<sub>3</sub> exposure inhibited submucosal fibrosis and smooth muscle hyperplasia. LXR deficient mice exhibited more severe submucosal fibrosis and smooth muscle hyperplasia than wild mice ( $P < 0.05$ ). Compared to the control group, O<sub>3</sub> stimulation increased the deposition of fibers around the bronchi, which was

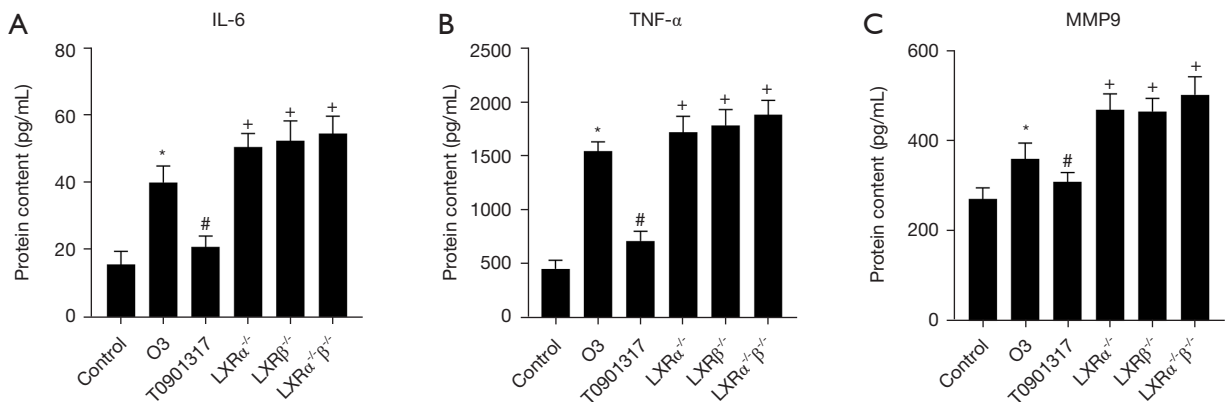
pronounced in LXR-deficient mice. However, T0901317 treatment significantly reduced the lung collagen levels ( $P < 0.05$ ) (Figure 8A-8G). Immunohistochemistry showed that  $\alpha$ -SMA and MMP9 protein were highly expressed in wild mice and LXR knockout mice after O<sub>3</sub>, but this expression was attenuated by treating with the T0901317 ( $P < 0.05$ ). Compared to wild mice, LXR-deficient mice exhibited even higher expression of  $\alpha$ -SMA and MMP9, although no differences were observed among LXR $\alpha$ <sup>-/-</sup>, LXR $\beta$ <sup>-/-</sup> and LXR $\alpha$ <sup>-/-</sup> $\beta$ <sup>-/-</sup> groups (Figure 9A-9G).

#### Discussion

O<sub>3</sub> is considered a critical air pollutant that causes lung injuries and reduction in lung function (17). When inhaled,



**Figure 5** The effect of LXRs on the formation of foamy macrophages. The representative Oil red O staining for (A) Control group, (B) O<sub>3</sub> group, (C) T0901317 group, (D) LXR $\alpha^{-/-}$  group, (E) LXR $\beta^{-/-}$  group and (F) LXR $\alpha^{-/-}\beta^{-/-}$  group (magnification,  $\times 200$ ). (G) Differential BALF macrophage cell count describing regular macrophages and foamy macrophages. \*,  $P < 0.05$  O<sub>3</sub> group *vs.* control group; #,  $P < 0.05$  T0901317 group *vs.* O<sub>3</sub> group; \*,  $P < 0.05$  LXR-deficient group *vs.* O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). LXRs, liver X receptor; O<sub>3</sub>, ozone; BALF, bronchoalveolar lavage fluid.

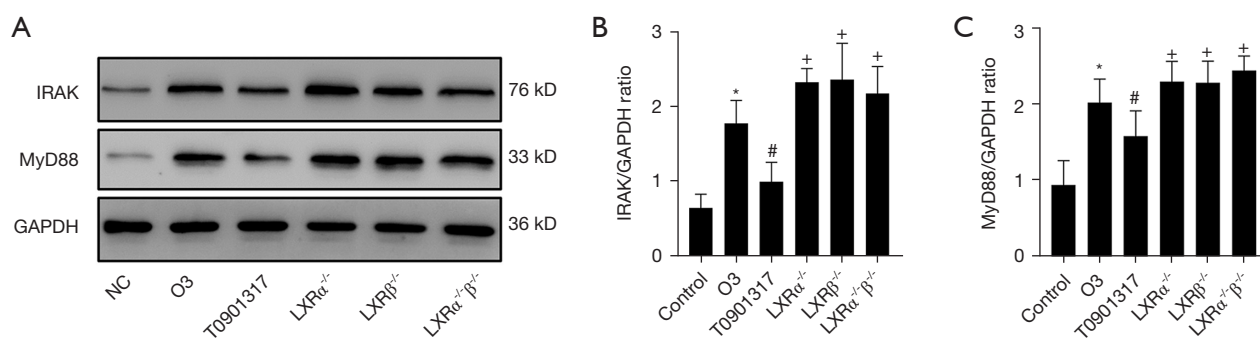


**Figure 6** The effect of LXRs on inflammatory cytokines in the BALF after O<sub>3</sub> exposure. The levels of IL-6 (A), TNF- $\alpha$  (B) and MMP9 (C) were determined by ELISA. \*,  $P < 0.05$  O<sub>3</sub> group *vs.* control group; #,  $P < 0.05$  T0901317 group *vs.* O<sub>3</sub> group; \*,  $P < 0.05$  LXR-deficient group *vs.* O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). O<sub>3</sub>, ozone; LXRs, liver X receptors; IL-6, interleukin-6; TNF- $\alpha$ , tumor necrosis factor- $\alpha$ ; MMP9, matrix metalloproteinase 9; BALF, bronchoalveolar lavage fluid; ELISA, enzyme linked immunosorbent assay.

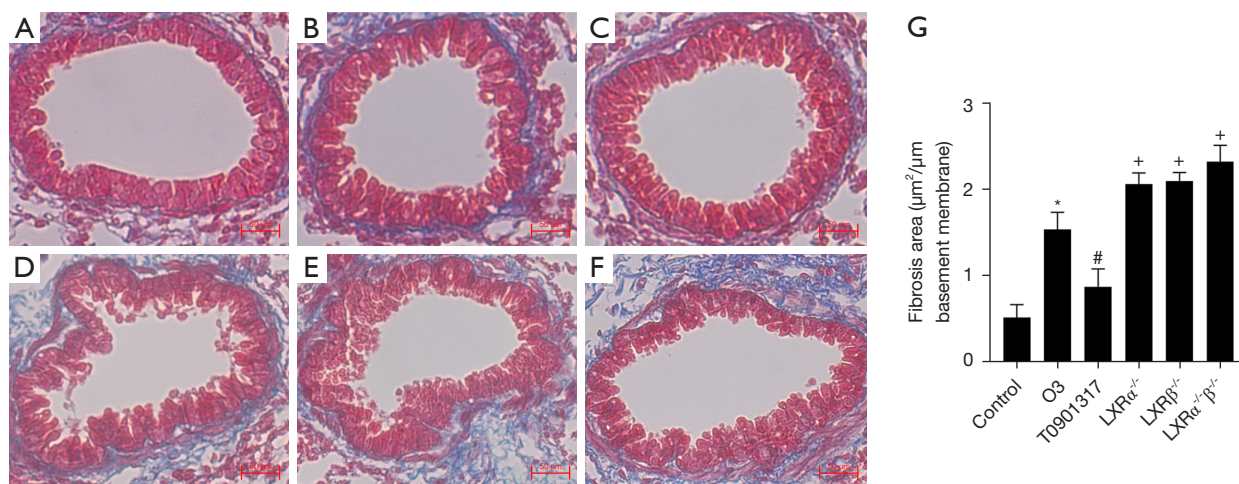
O<sub>3</sub> reacts with the lung lining fluid to form a LOP that can activate pro-inflammatory factors, such as IL-6 and IL-8. However, the precise mechanisms through which LOP mediates airway inflammation remain unclear. Previous studies have shown that O<sub>3</sub> oxidizes cholesterol in lung-lining fluid, leading to the generation of reactive oxysterols

that exhibit pro-inflammatory effects *in vitro* (8,18). Interestingly, instead of activating LXRs, these oxysterols form adducts with LXRs and inhibit their activation by traditional agonists *in vitro* (19). Nevertheless, previous research did not investigate the impact of LXR agonists *in vivo* following O<sub>3</sub> exposure, and the susceptibility to O<sub>3</sub>-





**Figure 7** The effect of LXRs on O<sub>3</sub>-induced IRAK and MyD88 expression in lung. (A) IRAK and MyD88 protein expression in lung tissues of each group of mice. (B,C) Quantitative analysis of IRAK and MyD88 protein expression (n=5). \*, P<0.05 O<sub>3</sub> group vs. control group; #, P<0.05 T0901317 group vs. O<sub>3</sub> group; +, P<0.05 LXR-deficient group vs. O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). NC, negative control; O<sub>3</sub>, ozone; LXRs, liver X receptors; IRAK, interleukin-1 receptor-associated kinase; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; MyD88, myeloid differentiation primary response protein 88.

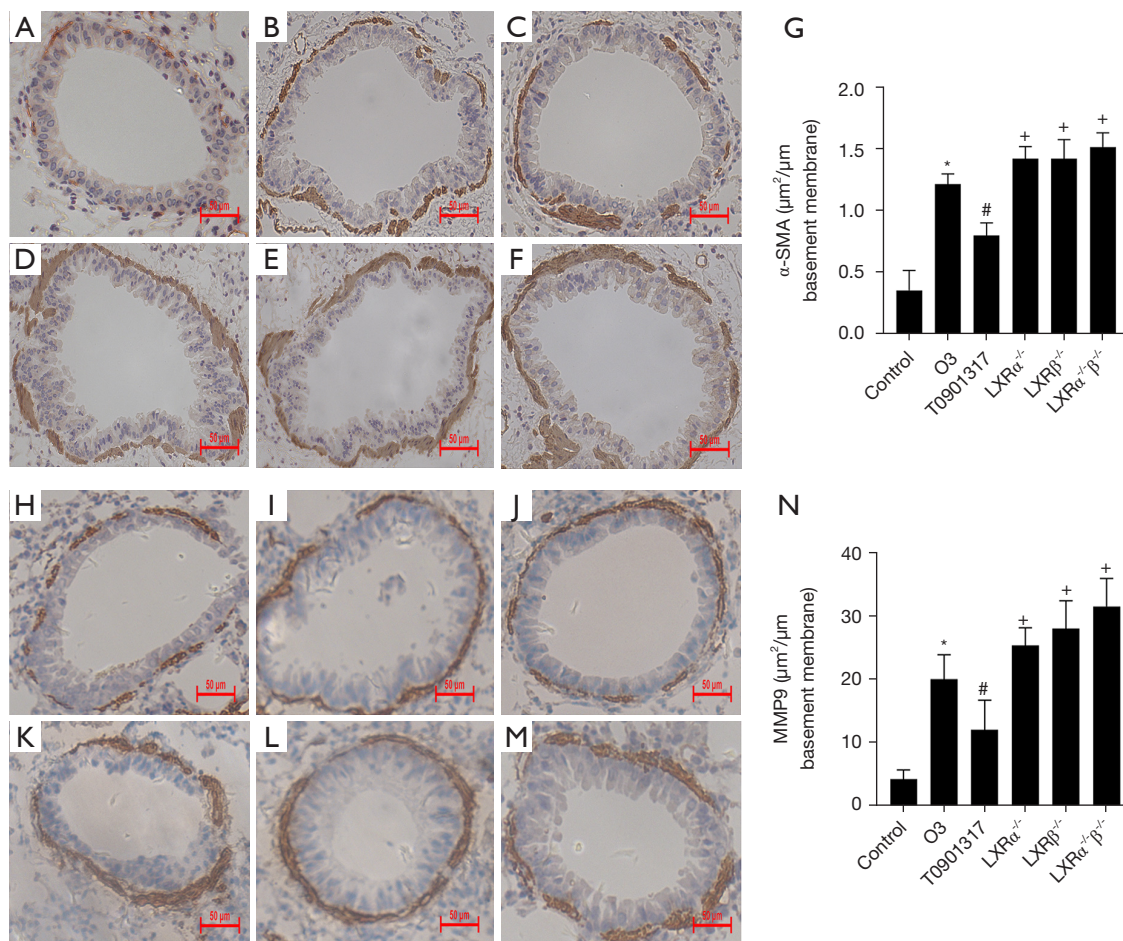


**Figure 8** The effect of LXRs on O<sub>3</sub>-induced submucosal fibrosis. The representative photomicrographs demonstrate the peribranchial collagen deposition detected by Masson's trichrome staining (blue color; original magnification, ×200). (A) Control group, (B) O<sub>3</sub> group, (C) T0901317 group, (D) LXR $\alpha^{-/-}$  group, (E) LXR $\beta^{-/-}$  group and (F) LXR $\alpha^{-/}\beta^{-/}$  group. (G) Image analysis assessing extend of Masson's trichrome staining expressed as means. \*, P<0.05 O<sub>3</sub> group vs. control group; #, P<0.05 T0901317 group vs. O<sub>3</sub> group; +, P<0.05 LXR-deficient group vs. O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). LXRs, liver X receptors; O<sub>3</sub>, ozone.

induced acute inflammation (3-hour exposure) was only observed in LXR $\alpha^{-/-}$ . In this study, we aimed to describe the role of LXR in airway injuries and remodeling induced by intermittent O<sub>3</sub> exposure in emphysema-like model in mice.

Myeloid differentiation primary response protein 88 (MyD88) serves as the key adaptor in inflammatory signaling pathways triggered by members of the Toll-like receptor (TLR) and interleukin-1 (IL-1) receptor

families. MyD88 facilitates the connection between IL-1 receptor (IL-1R) or TLR family members and IRAK family kinases through homotypic protein-protein interactions. The activation of IRAK family kinases results in various functional outcomes, including the activation of nuclear factor-kappa B (NF $\kappa$ B), mitogen-activated protein kinases, and activator protein 1, positioning MyD88 as a central component in inflammatory pathways (20-22). Many studies



**Figure 9** The effect of LXRs on O<sub>3</sub>-induced smooth muscle hyperplasia. The representative immunohistochemical staining for (A-F)  $\alpha$ -SMA expression and (H-M) MMP9 expression. (A,H) Control group, (B,I) O<sub>3</sub> group, (C,J) T0901317 group, (D,K) LXR $\alpha^{-/-}$  group, (E,L) LXR $\beta^{-/-}$  group and (F,M) LXR $\alpha^{-/-}\beta^{-/-}$  group (magnification,  $\times 200$ ). The area of  $\alpha$ -SMA and MMP9 per micrometer length of basement membrane of bronchiole (G,N) were calculated. \*,  $P < 0.05$  O<sub>3</sub> group *vs.* control group; #,  $P < 0.05$  T0901317 group *vs.* O<sub>3</sub> group; +,  $P < 0.05$  LXR-deficient group *vs.* O<sub>3</sub> group. Three independent experiments were examined (5 mice in each group of one experiment). LXRs, liver X receptors; O<sub>3</sub>, ozone;  $\alpha$ -SMA,  $\alpha$ -smooth muscle actin; MMP9, matrix metalloproteinase 9.

have shown that MyD88 is involved in the pathogenesis of COPD (23,24). Indoor and outdoor air pollution induces inflammatory response activating epithelial cells and alveolar macrophages via MyD88. In this study, we found that O<sub>3</sub> exposure increased the protein expression of MyD88 and IRAK, especially in LXR deficient mice. We also found that LXR agonist can inhibit O<sub>3</sub> induced MyD88 signaling. In addition, the experiment showed that expression of cytokines in BALF were correlated with LXR deletion or activation, So we believe that LXR may mediate O<sub>3</sub>-induced pulmonary inflammation via MyD88-IRAK signaling pathway.

LXR regulates cholesterol metabolism through ABCA1.

ABCA1 is positioned on the cell membrane and facilitates the efflux of lipids, including intracellular cholesterol. This process initiates reverse cholesterol transport, which helps lower intracellular cholesterol levels. We observed that foamy macrophages increased in BALF after O<sub>3</sub> exposure, concomitant with a significant decrease in ABCA1 expression in lung, which suggested that the cholesterol efflux capacity of macrophages was impaired by O<sub>3</sub>. The presence of foamy macrophages has been noticed in arteries of patients with atherosclerosis. The accumulation of cholesterol and ABCA1 inhibition in macrophages are associated with releasing inflammatory

mediators which are crucial for local inflammation (25). The endoplasmic reticulum (ER) is involved in protein and lipid metabolism (26), and disruption of lipid homeostasis can trigger endoplasmic reticulum stress (ERS) (27,28). Sun *et al.* discovered that ERS influences lipid catabolism in macrophages by enhancing cholesterol uptake, suppressing cholesterol efflux, and modulating the expression of associated transport proteins (29). Consequently, we propose that the emergence of foamy cells is due to the accumulation of cholesterol in macrophages, which in turn triggers ERS. Our study suggests that LXRs plays an important role in maintaining lipid homeostasis and may contribute to lung inflammation in O<sub>3</sub> environment. Further research is needed to explore the potential involvement of ERS in this process.

COPD is characterized by irreversible fibrosis and small airway remodeling. Our findings indicate that LXRs have the capacity to mitigate O<sub>3</sub>-induced alveolar enlargement and submucosal fibrosis. Airway remodeling is closely linked to the overexpression of MMP9, an enzyme involved in extracellular matrix degradation and tissue repair (30,31). LXRs can effectively suppress the O<sub>3</sub>-induced upregulation of MMP9 levels, which may contribute to the attenuation of airway remodeling (32,33). Previous research has demonstrated that LXRs inhibit MMP9 production in macrophages through ubiquitination (34). Macrophage activation is widely recognized as a significant contributor to lung structural damage and a prominent source of MMP9 (35). Hence, LXRs may partially ameliorate airway remodeling by restraining MMP9 production in macrophages. Furthermore, epithelial-mesenchymal transition (EMT) plays a crucial role in the remodeling of small airways and the development of subepithelial fibrosis (36,37). Cigarette smoke extract and cigarette smoking can stimulate the expression of MMP9 (2,38,39), multiple studies have demonstrated a strong correlation between MMP9 and the EMT process (40-42). In our study, LXRs may have affected the process of EMT and thereby alleviated airway remodeling. In future experiments, we will strengthen relevant research to confirm this idea.

## Conclusions

Our results suggest that activated LXRs confer protective effects against O<sub>3</sub>-induced lipid accumulation, airway inflammation, and remodeling in mice. The protective mechanisms of LXRs may be attributed to the activation of

ABCA1 and the suppression of MyD88-IRAK pathway.

## Acknowledgments

The authors acknowledge all people who contributed to this article.

*Funding:* This study was supported by Nanjing Medical Science and Technique Development Foundation of China (No. JQX16028 to Y.S.).

## Footnote

*Reporting Checklist:* The authors have completed the ARRIVE reporting checklist. Available at <https://jtd.amegroups.com/article/view/10.21037/jtd-23-1820/rc>

*Data Sharing Statement:* Available at <https://jtd.amegroups.com/article/view/10.21037/jtd-23-1820/dss>

*Peer Review File:* Available at <https://jtd.amegroups.com/article/view/10.21037/jtd-23-1820/prf>

*Conflicts of Interest:* All authors have completed the ICMJE uniform disclosure form (available at <https://jtd.amegroups.com/article/view/10.21037/jtd-23-1820/coif>). The authors have no conflicts of interest to declare.

*Ethical Statement:* The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The animal experiments were performed under a project license (No. SYXK [Su] 2016-0006) granted by the Animal Experiment Center of Nanjing First Hospital Affiliated to Nanjing Medical University, in compliance with guidelines established by the institution's Animal Care and Use Committee for the care and use of animals.

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**Cite this article as:** Yu F, Ma J, Xu K, Tang Y, Wu B, Gu W, Shi Y. Effects of liver X receptor in O<sub>3</sub>-induced airway inflammation and remodeling in mice. *J Thorac Dis* 2024;16(8):5005-5017. doi: 10.21037/jtd-23-1820