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#### RESEARCH ARTICLE



# Extracellular HSP90 $\alpha$ promotes cellular senescence by modulating TGF- $\beta$ signaling in pulmonary fibrosis

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#### Abstract

Recent findings suggest that extracellular heat shock protein  $90\alpha$  (eHSP90 $\alpha$ ) promotes pulmonary fibrosis, but the underlying mechanisms are not well understood. Aging, especially cellular senescence, is a critical risk factor for idiopathic pulmonary fibrosis (IPF). Here, we aim to investigate the role of eHSP90 $\alpha$  on cellular senescence in IPF. Our results found that eHSP90 $\alpha$  was upregulated in bleomycin (BLM)-induced mice, which correlated with the expression of senescence markers. This increase in eHSP90 $\alpha$  mediated fibroblast senescence and facilitated mitochondrial dysfunction. eHSP90 $\alpha$  activated TGF- $\beta$  signaling through the phosphorylation of the SMAD complex. The SMAD complex binding to p53 and p21 promoters triggered their transcription. In vivo, the blockade of eHSP90 $\alpha$  with 1G6-D7, a specific eHSP90 $\alpha$  antibody, in old mice attenuated the BLM-induced lung fibrosis. Our findings elucidate a crucial mechanism underlying eHSP90 $\alpha$ -induced cellular senescence, providing a framework for aging-related fibrosis interventions.

#### K E Y W O R D S

cellular senescence, HSP90 heat-shock proteins, mitochondria, pulmonary fibrosis, reactive oxygen species, transforming growth factor beta

**Abbreviations:** ATP, adenosine triphosphate; BALF, bronchoalveolar lavage fluid; BLM, bleomycin; eHSP90 $\alpha$ , extracellular heat shock protein 90 $\alpha$ ; IL6, interleukin 6; IPF, idiopathic pulmonary fibrosis; MMP, mitochondrial membrane potential; mt-ROS, mitochondrial-derived reactive oxygen species; PAI1, plasminogen activator inhibitor-1; rHSP90 $\alpha$ , recombinant human HSP90 $\alpha$ ; ROS, reactive oxygen species; SA- $\beta$ gal, senescence-associated beta-galactosidase; SASP, senescence-associated secretory phenotype; TGF- $\beta$ , transforming growth factor  $\beta$ .

Wenshan Zhong, Weimou Chen, and Yuanyuan Liu contributed equally to this work.

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cence is unknown.

## 1 | INTRODUCTION

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Idiopathic pulmonary fibrosis (IPF) is a chronic, progressive and age-related lung disease with a median age at diagnosis of 66 years and estimated survival of 3–4 years.<sup>1</sup> IPF is characterized by repetitive injury to the lung epithelium, activation, and proliferation of (myo)fibroblasts, and extracellular matrix deposition, which destroy lung architecture and function.<sup>2</sup> However, current IPF treatment regimens have limited efficacy. A better understanding of the pathological mechanisms of fibrosis is essential to investigate effective therapies.

Heat shock protein 90 (HSP90) is a ubiquitously expressed chaperone involved in the posttranslational folding and stability of proteins. In addition, HSP90, which has an extracellular function, is found in two isotypes: inducible alpha and beta, which is constitutive.<sup>3</sup> Recently, extracellular HSP90 $\alpha$  (eHSP90 $\alpha$ ) was shown to be upregulated in pulmonary fibrosis and interact with adjacent cells' surfaces, leading to increased myofibroblast activation.<sup>4</sup> Our previous experiments also showed that eHSP90 $\alpha$  blockade was beneficial for pulmonary fibrosis.<sup>5</sup> Overall, eHSP90 $\alpha$ is an essential factor for the development of pulmonary fibrosis, but the associated mechanisms are unclear.

Emerging studies have implicated aging as a significant risk factor for IPF, and elderly individuals are also more sensitive to lung injuries.<sup>6</sup> Notably, HSP90 inhibitors have been reported as senolytic agents, indicating a new role in aging.<sup>7</sup> However, whether eHSP90 $\alpha$  involve in the process of aging remains unknown. Cellular senescence, a hallmark of aging, has been thought to be involved in the pathogenesis of lung fibrosis.<sup>8–10</sup> A type of cell fate, cellular senescence, is characterized by permanent growth arrest and other phenotypic alterations, including the development of a pro-inflammatory secretome.<sup>11</sup> Moreover, elevated expression of p16, p21, and p53 are observed in senescent cells and used as miscellaneous senescence biomarkers.<sup>12</sup> Upregulated expression of p16 and p21, Senescence-associated beta-galactosidase (SA-βgal) activity, and a senescence-associated secretory phenotype (SASP) were observed in human samples and mice models of lung fibrosis.<sup>10,13</sup> Recent studies have demonstrated that the accumulation of senescent fibroblasts drives pulmonary fibrosis.<sup>10</sup> It is well established that mitochondria that accumulate in senescence often show a decreased membrane potential, increased mitochondrial mass, and simultaneously produce increased reactive oxygen species (ROS).<sup>14,15</sup> Mitochondrial dysfunction refers to the damage of mitochondrial structure, respiratory chain defects, biogenic dysfunction, gene damage, and changes in oxidative protein activity in cells and tissues.<sup>16</sup> However, the mechanisms of fibroblast senescence are poorly understood.

Transforming growth factor  $\beta$  (TGF- $\beta$ ) is one of the most potent profibrotic factors in the development of pulmonary fibrosis.<sup>17-19</sup> Active TGF- $\beta$  binds to its receptors leading to activation of canonical SMAD signaling.<sup>20</sup> The SMAD complex translocates to the nucleus to regulate the transcription of target genes.<sup>21</sup> Although SMADs contribute to p21 transcription,<sup>22-24</sup> their role in fibroblast senes-

In this study, we assessed the pathophysiological role of eHSP90 $\alpha$  in cellular senescence and the impact of eHSP90 $\alpha$  blockade in the bleomycin (BLM)-induced injury model for understanding how TGF- $\beta$  signaling activation in senescence controls the expression of p21 and p53. We further demonstrate that the function of eHSP90 $\alpha$  may offer new opportunities for targeting eHSP90 $\alpha$  as a therapeutic option for age-related fibrotic disease.

#### 2 | MATERIALS AND METHODS

#### 2.1 | Cell culture and treatment

IMR90 human fibroblasts (ATCC, CCL-186) were used between 25 and 40 population doublings. The cells were cultured in EMEM (ATCC, 30-2003) with 10% FBS (GIBCO, 10100147) at 37°C in a humidified atmosphere containing 5% CO<sub>2</sub>. The laboratory of Dr. Wei Li donated the F-5 fragment and anti-HSP90a monoclonal antibody (1G6-D7). For experiments employing extracellular HSP90αinduced senescence, IMR90 human lung fibroblasts were treated 2 h with 1G6-D7(20µg/ml) before F-5 (10 µg/ml) for 48 h. 1G6-D7 was developed against the dual lysine region of human Hsp90 $\alpha$  and raised in mice. The isotype is IgG1ĸ. Species Reactivity was tested in humans and mice.<sup>5,25</sup> For experiments utilizing etoposide-induced senescence, IMR90 human lung fibroblasts were treated for 48h with 20mM etoposide (Sigma-Aldrich, E1383). For experiments abrogating mtROS, cells were treated for 48 h with F-5 after Mito-Tempo (10 µM, MedChemExpress, HY-112879) treatment. For experiments inhibiting, cells were treated for 48h with F-5 after RepSox (7.5 µM, Selleck, S7223) treatment.

#### 2.2 | Animals

All the animal experiments were approved by the Committee on the Ethics of Animal Experiments of Southern Medical University in Guangzhou, China and performed under standard guidelines for the Care and Use of Laboratory Animals. Male mice (C57BL/6) were kept on a 12h light– dark cycle with free access to food and water. Mice were randomly distributed to indicated groups.

# 2.2.1 | Mouse model for bleomycin-induced lung fibrosis

For bleomycin administration, C57BL/6 mice, 6 to 8 weeks of age were anesthetized with 2, 2, 2-tribromoethanol (Sigma-Aldrich) followed by intratracheal instillation of bleomycin (5 U/kg, i. t.) in 50 $\mu$ l PBS, as previously described.<sup>26</sup> Mice received the first dose of PBS (control group) or 1G6-D7 8 days after bleomycin, as described in the main text. PBS or 1G6-D7 was then administered every day for an additional 13 days.

# 2.2.2 | Mouse model for age-related lung fibrosis

For bleomycin administration, 2-month-old and 18-month-old C57BL/6 mice were anesthetized with 2, 2, 2-tribromoethanol (Sigma-Aldrich) followed by intratracheal instillation of bleomycin (5 U/kg, i. t.) in  $50 \,\mu$ l PBS. 18-month-old mice received the first dose of PBS (control group) or 1G6-D7 8 days after bleomycin treatment. PBS or 1G6-D7 was then administered every day for an additional 13 days.

# 2.3 | Hydroxyproline assay

Hydroxyproline content of the whole lung was measured by a hydroxyproline (HYP) kit (Nanjing Jian Cheng Institute, Nanjing, China) in accordance with the manufacturer's recommendations.

## 2.4 | Lung histology and Immunohistochemistry

Harvested lungs were formaldehyde fixed, paraffinembedded, and stained with H&E or Masson's trichrome stain as previously described.<sup>27</sup>

Sections (5-µm-thick) were deparaffinized and rehydrated, endogenous peroxidases were inactivated with methanol containing 0.3% hydrogen peroxide for 30 min. Antigen retrieval was performed by steaming in citrate buffer (pH 8.0) for 20 min. After incubation in blocking solution for 10 min at room temperature, the slides were incubated with an anti-fibronectin antibody (Abcam, ab2413), anti-p16 antibody (Abcam, ab51243), or anti- $\alpha$ -SMA antibody (Abcam, ab5694) overnight at 4°C. After three washes, the sections were incubated with biotinylated anti-IgG at 37°C for 60 min, followed by Streptavidin-peroxidase conjugate (Zhongshan Golden Bridge Biotechnology). Immunoreactivity was detected using 3, 3' diaminobenzidine, and the sections were counterstained with hematoxylin for observation by microscopy.

# 2.5 | Pulmonary function test

Pulmonary function was characterized at the endpoint using a Buxco FinePointe RC system as previously described.<sup>28,29</sup> Mice were euthanized by an intraperitoneal injection of 2, 2, 2-tribromoethanol and connected to the Buxco system after tracheotomy. Dynamic lung compliance and resistance values were obtained from the respiratory system.

# 2.6 | ELISA

Bronchoalveolar lavage fluid (BALF) and serum samples from mice and cell culture medium from cells were collected as described previously.<sup>30</sup> The HSP90aA1 (Cloud-Clone, Buckingham, UK) and HSP90 $\alpha$  (Enzo) ELISA kits were used following the manufacturer's recommendations.

# 2.7 | Western blot

Western blot analyses were performed as described previously.<sup>5</sup> Briefly, cell lysates or lung homogenates were prepared using a protein extraction kit (KeyGEN BioTECH). After SDS-PAGE, PVDF membranes were probed with specific antibodies described in the figure legends followed by a corresponding secondary antibody (LI-COR). Bands were quantified with ImageJ software. Antibodies are listed as below: anti-fibronectin antibody (Abcam, ab2413); anti-p16 antibody (Abcam, ab51243); anti- $\alpha$ -SMA antibody (Abcam, ab5694); anti-p21 antibody (Santa Cruz, sc-6246); anti-β-actin antibody (Proteintech, 66009-1-Ig); anti-phospho-SMAD3 antibody (Ser423/425, Cell Signaling Technology, 9520T); anti-phospho-SMAD2 antibody (Ser465/Ser467, Cell Signaling Technology, 18338T); anti-SMAD3 antibody (Proteintech, 25494-1-AP); anti-SMAD2 antibody (Proteintech, 12570-1-AP); anti-CollagenI antibody (Affinity, AF7001); anti-SMAD4 antibody (Cell Signaling Technology, 46535). anti-GAPDH antibody (Proteintech, 10494-1-AP).

# 2.8 | SA-β-gal

The SA- $\beta$ -gal activity was detected as described<sup>31</sup> using a commercial Senescence Cells Histochemical Staining Kit (Sigma–Aldrich).

# 2.9 | RNA extraction and RT-qPCR

Mouse lung tissue RNAs were extracted using TRIzol reagent (Takara). cDNA was synthesized using the iScript cDNA Synthesis Kit (Bio-Rad). The iTaq Universal SYBR Green Supermix qPCR Kit (Bio-Rad) was used to perform RT-qPCR. Primers used are listed as follows:

IL6 (Forward: TAGTCCTTCCTACCCCAATTTCC, Reverse:TTGGTCCTTAGCCACTCCTTC); TGF- $\beta$ (Forward: CCACCTGCAAGACCATCGAC, Reverse: CTG GCGAGCCTTAGTTTGGAC); PAI1 (Forward: TCTGGGA AAGGGTTCACTTTACC, Reverse: GACACGCCATAGGG AGAGAAG).

#### 2.10 | Transmission electron microscopy

Cells were fixed in 2.5% glutaraldehyde in 0.05M cacodylate buffer with 0.1 M NaCl, pH 7.5 for 20 min at room temperature, as previously described.<sup>32</sup>

### 2.11 | Mitochondrial mass

The mitochondrial mass was assessed using MitoTracker (Invitrogen, M22425) in accordance with the manufacturer's recommendations. Briefly, cells were incubated with 20 nM Mito Tracker for 20 min at 37°C. Images were obtained using Olympus FV1000 Confocal Laser Scanning Microscopy (Tokyo, Japan).

#### 2.12 Mitochondrial ROS assessment

The levels of mitochondrial ROS were detected using the fluorescent probes MitoSOX<sup>™</sup> Red mitochondrial superoxide indicator (Invitrogen, M36008), and fluorescent intensity was obtained by Olympus FV1000 Confocal Laser Scanning Microscopy (Tokyo, Japan).

### 2.13 | Chromatin immunoprecipitation

Chromatin immunoprecipitation (ChIP) experiments were conducted using the SimpleChIP Enzymatic Chromatin IP Kit (CST, 9003) in accordance with the manufacturer's recommendations. Chromatin was immunoprecipitated with the immunoglobulin G (CST, #2729; as a negative control) or SMAD2/3 (Abcam, ab202445). In all, 10% of total DNA was used for input evaluation. DNA enrichment in the ChIP samples was determined by reverse transcription and semi-quantitative PCR (RT-PCR) with PrimeScript<sup>™</sup> RT reagent Kit with gDNA Eraser (Takara, China) following the manufacturer's protocol. PCR products of immunoprecipitated and input samples were analyzed on a 2% agarose gel. Primers used are listed as follow: p16 promoter P1 (Forward: GCTACGTAAGAGTGATCGCTAAA, Reverse: CTCGGTACAAACCCAAGACAA); p16 promoter P2 (Forward: CCTTTCCTTGCCCTGCTTT, Reverse: CTTCCA CATCACCGATCCTTTC); p21 promoter P1 (Forward: CAT GGGAGGAGCTACAACTATATG, Reverse: CTTCCTGTTC CTGGCTCTAAC); p21 promoter P2 (Forward: CACCTGAA TACCTGGGACTACA, Reverse: GCCTGGCCAATATGGT GAAA); TP53 promoter P1 (Forward: GACCAGGAACCAC TGAGAAATC, Reverse: CTTGGCGACCCAGGTTTATT); TP53 promoter P2 (Forward: GAGTCCCGCGGTAATTCT TAAA, Reverse: CTGAAGCCTGGAGAATGAGATG).

#### 2.14 Statistical analysis

Prism 8.0 software (GraphPad) was used to analyze data. The experiment was set up to use 3 to 7 samples/repeats for each experiment/group/condition. Representative images for IF staining, IHC staining, and immunoblot are shown. Student's *t* test was used to analyze the differences between 2 groups, one-way ANOVA analysis was used to compare the differences between more than 2 groups. *p*-values of less than .05 were regarded as statistically significant.

#### 3 | RESULTS

### 3.1 | Extracellular HSP90α is upregulated in pulmonary fibrosis and associated with cellular senescence

A recent study demonstrated that HSP90a was increased in serum from patients with IPF compared with healthy controls.<sup>4</sup> To further characterize HSP90 $\alpha$  and cellular senescence in IPF, we generated a BLM-induced mouse model (Supporting Information Figure S1). Bodyweight monitoring, an essential indicator of pathological severity following the BLM challenge, revealed that mice that received bleomycin lost substantial body mass relative to mice that received PBS (Figure 1A). HSP90 $\alpha$  levels were increased in the bronchoalveolar lavage fluid (BALF) of the BLM-treated mice from day14 to day21 (Figure 1B). Western blot analysis revealed that senescence marker (p16 and p21) levels were upregulated in lung tissue from the BLM-treated mice (Figure 1C,D). Interestingly, the level of senescence markers p21 gradually increased from day 7 (the inflammatory phase) up to day 21 (established fibrosis), accompanied by the upregulation of



**FIGURE 1** The level of extracellular HSP90 $\alpha$  was increased and in the same trend as senescence markers in the BLM-induced model. C57/BL6 mice were treated with BLM or PBS as control. (A) Body weight (BW) is depicted as a change in grams relative to baseline. Data presented as mean ± SD; PBS: n = 8; BLM day 7: n = 8; BLM day 14: n = 7; BLM day 21: n = 8. (B) Bronchoalveolar lavage fluid (BALF) levels of HSP90 $\alpha$  were measured by ELISA. Data presented as mean ± SD; PBS: n = 5; BLM day 7: n = 5; BLM day 14: n = 5; BLM day 21: n = 5. (C, D) Western blot analysis of p16, p21, COL1A1, Fibronectin, and  $\alpha$ -SMA expression release by mice lung slices. Data presented as mean ± SD. IL6 (E), TGF- $\beta$  (F), and PAI1 (G) expression by qPCR, normalized to  $\beta$ -actin expression. Data presented as mean ± SD. \*\*\*\*p < .0001, \*\*\*p < .001, \*\*p < .01, and \*p < .05 between indicated groups.

fibronectin, type collagen I, and  $\alpha$ SMA (Figure 1C,D), suggesting that cellular senescence is associated with disease severity.

To further confirm the relationship between  $HSP90\alpha$  and cellular senescence in BLM-treated mice, we next detected several SASP factors via qPCR. Results showed that

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expression of Interleukin 6 (IL6), transforming growth factor  $\beta$ (TGF- $\beta$ ), and plasminogen activator inhibitor-1 (PAI1) were upregulated (Figure 1E–G), which has the same trend as the level of HSP90 $\alpha$ . Taken together, these results demonstrate that eHSP90 $\alpha$  is upregulated in pulmonary fibrosis in correlation with cellular senescence.

# 3.2 | Extracellular HSP90α promotes fibroblast senescence

In order to explore the role of eHSP90 $\alpha$  in cellular senescence, we established an etoposide-induced senescent model.<sup>10</sup> Exposure of human IMR90 fibroblasts to etoposide treatment-induced senescence observable after 48 h, which was confirmed by staining for SA- $\beta$ -gal (Figure 2A,B) and expression of senescence effectors, p21 (Figure 2C,D). The secretion of HSP90 $\alpha$  was next assessed in vitro in etoposideinduced fibroblasts. Etoposide-induced senescent cells exhibited a 2-fold increase in HSP90 $\alpha$  compared to control (Figure 2E). This result was also confirmed by western blot analysis (Supporting Information Figure S2A). However, no change in the total level of the HSP90 $\alpha$  protein was observed (Figure 2C,D), suggesting intracellular HSP90 $\alpha$  might not contribute to cellular senescence. Therefore, we hypothesise that eHSP90 $\alpha$  might mediate cellular senescence.

To identify whether eHSP90 $\alpha$  regulates cellular senescence, we treated IMR90 fibroblasts with recombinant human HSP90 $\alpha$  (rHSP90 $\alpha$ ). rHSP90 $\alpha$ -treated fibroblasts exhibited increased SA- $\beta$ -gal activity and p16 and p21 expression relative to those of control fibroblasts (Supporting Information Figure S2B–D). These results suggest that eHSP90 $\alpha$  regulates cellular senescence. To further confirm this phenotype, we used a monoclonal antibody, 1G6-D7, which binds to the F-5 region of HSP90 $\alpha$  and inhibits the extracellular HSP90 $\alpha$  function.<sup>25,33</sup> The previous study has demonstrated that F-5, a 115-amino acid fragment within the linker region and middle domain of Hsp90 $\alpha$ , retains the extracellular function of HSP90 $\alpha$ .<sup>34</sup>

Similarly, in our study F-5 treatment increased SA- $\beta$ -gal activity, while 1G6-D7 reduced F-5-induced SA- $\beta$ -gal activity (Figure 2F,G). F-5 treatment increased p16 and p21 protein expression, but this effect was blocked by 1G6-D7 (Figure 2H,I). Taken together, our results demonstrate that eHSP90 $\alpha$  stimulates a senescence phenotype in healthy human fibroblasts.

#### 3.3 | Extracellular HSP90α-induced fibroblast senescence is associated with mitochondrial dysfunction

Previous studies have demonstrated that mitochondrial function is responsible for cellular senescence in IPF.<sup>35–37</sup>

To clarify whether eHSP90 $\alpha$  affects mitochondrial function, we used MitoTracker Red (MitoRed) to assess mitochondrial mass. Confocal microscopy revealed an increase in mitochondrial mass and the formation of highly elongated mitochondria after F-5 treatment (Figure 3A,B), while 1G6-D7 reduced the effect of eHSP90 $\alpha$ . Fibroblasts treated with F-5 exhibited an increased number of abnormal mitochondria compared to cells treated with F-5 combined 1G6-D7 (Figure 3C,D). Significantly, the F-5 treatment enhanced mitochondrial ROS production shown by MitoSOX Red staining (Figure 3E,F). These results demonstrate that eHSP90 $\alpha$  leads to mitochondrial dysfunction and damaged mitochondria in senescent cells.

### 3.4 | Extracellular HSP90α activates TGF-β signaling to promote fibroblast senescence

TGF- $\beta$  signaling is the master regulator of pulmonary fibrosis<sup>18,19</sup> and has reportedly been linked to the phenotypes of cellular senescence.<sup>38</sup> We next investigated whether eHSP90 $\alpha$  activates TGF- $\beta$  signaling to promote fibroblast senescence. We found that F-5 treatment induced more activation of TGF- $\beta$  signaling, including phosphorylation of SMAD2 and SMAD3 and expression of SMAD4 (Figure 4A–D). Notably, the induction of TGF- $\beta$  by F-5 was entirely abolished by the blockade of extracellular HSP90 $\alpha$ with 1G6-D7 (Figure 4A-D), suggesting eHSP90a improves TGF-β signaling activation. To further validate that eHSP90α promotes cellular senescence by TGF-β signaling, we stimulated IMR90 with RepSox, an inhibitor of TGF- $\beta$ signaling, in the presence or absence of F-5. We found that expression of p16 and p21 (Figure 4E,F) and activity of SAβ-gal (Figure 4G,H) were abrogated by RepSox. Similarly, SMAD4 knockdown also attenuated senescence induced by eHSP90 $\alpha$  (data not shown). Taken together, eHSP90 $\alpha$ promotes cellular senescence through TGF-β signaling.

# 3.5 | mtROS is involved in TGF-β activation

As we observed mtROS upregulation was accompanied by eHSP90 $\alpha$ -induced senescence and ROS has been reported to be involved in TGF- $\beta$  activation.<sup>39</sup> To elucidate whether mtROS is involved in eHSP90 $\alpha$ -TGF- $\beta$  signaling activation, we cultured IMR90 in media containing Mito-Tempo, an inhibitor of mtROS, together with F-5. Results showed that Mito-Tempo suppressed the activation of TGF- $\beta$  signaling induced by F-5 (Figure 5A–D), which suggests mtROS partly help with TGF- $\beta$  signaling activation.



**FIGURE 2** Extracellular HSP90 $\alpha$  induces senescence in fibroblasts. (A, B) Cytochemical evaluation of the senescence marker SA- $\beta$ -gal activity of IMR90 lung fibroblasts exposed to etoposide. (C, D) Representative immunoblots of p21, HSP90 $\alpha$ , and  $\beta$ -actin protein levels in IMR90 lung fibroblasts exposed to etoposide. Data presented as mean  $\pm$  SD. (E) supernatant levels of HSP90 $\alpha$  were measured by ELISA. Data presented as mean  $\pm$  SD. SA- $\beta$ -gal activity (F, G) and (H, I) representative immunoblots of p16, p21, and  $\beta$ -actin protein levels in IMR90 lung fibroblasts exposed to F-5 for 48 h after pre-treat with 1G6-D7. Data presented as mean  $\pm$  SD. \*\*\*\*p <.0001, \*\*\*p <.001, \*\*p <.01, and \*p <.05 between indicated groups.

# 3.6 | The SMAD2/3 complex regulates p53 and p21 transcription

The SMAD complex involved in TGF- $\beta$  signaling has been reported as a vital transcript factor to regulate gene expression.<sup>18</sup> Some researchers found that p21 transcription can be mediated by SMADs.<sup>22,24</sup> These results inspired

us to investigate whether the expression of senescence markers in fibroblasts exposed to  $eHSP90\alpha$  is also under regulation of the SMAD complex. To validate the interaction between SMADs and senescence markers, chromatin immunoprecipitation analysis was performed utilizing IMR90 cells treated with F-5. First, using the NCBI website and JASPAR program, we analyzed the p16, p21,



FIGURE 3 Extracellular HSP90a contributes to abnormal mitochondrial structure and mitochondrial ROS accumulation. (A) Immunofluorescence and (B) quantitative analysis for Mito-Tracker in IMR90 after 1G6-D7 or PBS exposure and treatment with or without F-5. Ten fields were randomly selected and acquired from each group. Data presented as mean  $\pm$  SD (n = 3). (C, D) Representative transmission electron microscopy (TEM) images. Red arrows indicate damaged and swollen mitochondria. Ten fields were randomly selected and acquired from each group. Data presented as mean  $\pm$  SD (n = 3). (E) MitoSOX Red staining and (F) quantitative analysis for mitochondrial ROS production. Fluorescence microscopy detection was performed. Ten fields were randomly selected and acquired from each group. Data presented as mean  $\pm$  SD (n = 3). \*\*\*\*p < .0001, \*\*\*p < .001, \*\*p < .01, and \*p < .05 between indicated groups.



**FIGURE 4** Extracellular HSP90 $\alpha$  promotes cellular senescence through mtROS-dependent TGF- $\beta$  signaling. Representative western blots (A) and quantitative analysis (B–D) show the amounts of p-SMAD2, p-SMAD3, SMAD2, SMAD3, SMAD4, and  $\beta$ -actin in IMR90 fibroblasts stimulated with F-5 in the presence or absence of 1G6-D7. Data presented as mean ± SD. Representative western blots (E) and quantitative analysis (F) show the amounts of p16, p21, and  $\beta$ -actin in IMR90 fibroblasts stimulated with F-5 in the presence or absence of RepSox. Data presented as mean ± SD. (G, H) SA- $\beta$ -gal activity in IMR90 lung fibroblasts exposed to F-5 for 48 h after pre-treat with RepSox. \*\*\*\*p < .0001, \*\*\*p < .001, \*\*\*p

and p53 promoter sequences to identify predicted transcription factor-binding sites (Figure 6A). As shown in Figure 6B–D, P1 and P2 regions from senescence markers could be amplified. Our data showed that F-5 increased the enrichment of SMADs at the binding sites in p21 (Figure 6F) and p53 (Figure 6G), but no increase at the p16 promoter (Figure 6E). They suggested that the SMAD complex regulates p21 and p53 transcription to improve cellular senescence.

### 3.7 | Targeting eHSP90α with 1G6-D7 prevented pulmonary senescence and fibrosis

Our results strongly support the hypothesis that eHSP90 $\alpha$  promotes cellular senescence in vitro. We further quantify whether eHSP90 $\alpha$  regulates senescence in a BLM-induced lung fibrosis model. Indicated dose of 1G6-D7 was administered from day 8 after BLM injury (Figure 7A). H&E



FIGURE 5 mtROS is involved in TGF- $\beta$  signaling activation. Representative western blots (A) and quantitative analysis(B–D) show the amounts of p-SMAD2, p-SMAD3, SMAD2, SMAD3, SMAD4, and  $\beta$ -actin in IMR90 fibroblasts stimulated with F-5 in the presence or absence of Mito-Tempo. Data presented as mean ± SD.

staining showed lungs from BLM-treated mice exhibited more severe pulmonary fibrosis compared with those from BLM plus 1G6-D7-treated mice (Figure 7B).

1G6-D7 treatment attenuated BLM-induced body weight loss and lung function (Figure 7C–E). Furthermore, mice that received 1G6-D7 demonstrated significant reductions in several pro-fibrotic markers, including total lung hydroxyproline (Figure 7F)and the expression of  $\alpha$ -SMA and fibronectin (Figure 7G–K). Similar results were shown in immunohistochemical analysis (Figure 7O).

In order to illustrate the role of eHSP90 $\alpha$  in TGF $\beta$  activation in vivo, we performed western blot to test the phosphorylation of SMAD2 and SMAD3. We observed that 1G6-D7 inhibits the effect of bleomycin in the phosphorylation of SMAD2 and SMAD3 (Figure 7L–N). These data suggest that blockade of extracellular HSP90 $\alpha$  by 1G6-D7 attenuates BLMinduced lung fibrosis even though fibrosis was established.

Importantly, immunohistochemical and western blot analyses revealed that 1G6-D7 treatment reduced the amounts of p16 and p21 (Figure 7C–H). Moreover, as shown in Figure 7P, the expression of  $\gamma$ H2AX was decreased after 1G6-D7 administration. Taken together, these findings suggest that 1G6-D7 attenuated lung fibrosis and cellular senescence in vivo.

# 3.8 | 1G6-D7 reduced lung fibrosis in older mice

Aging is a well-known risk factor for IPF and cellular senescence has been reported as a crucial hallmark of aging.<sup>1,40</sup> Thus, we next investigate whether blocking

eHSP90α by 1G6-D7 attenuates age-related lung fibrosis in vivo. The levels of circulating HSP90α were measured in the sera of old (18 months) and young (2 months) mice. HSP90α levels were upregulated in aged mice compared to young mice (Figure 8A), suggesting that eHSP90α might contribute to aging-induced fibrosis. However, histological analysis showed that the lungs from both old and young mice appeared without inflammation or fibrosis (data not shown).

In order to examine the effect of 1G6-D7 in older mice, 18-month-old and 2-month-old mice were administered low-dose BLM and received 1G6-D7 from day8. H&E and Masson trichrome staining (Figure 8B,C) further confirmed increased fibrosis in the lungs of the 18-month-old mice, while blockade by 1G6-D7 reduced lung fibrosis. Similarly, we observed increased expression of  $\alpha$ -SMA in 18-month-old mice through immunohistochemistry (Figure 8D) and western blot analysis (Figure 8E-G) compared to 2-month-old mice. These results suggested that the older mice were more sensitive to BLM, and aging is involved in fibrosis development. In contrast, 1G6-D7 abrogated the fibrotic effects of BLM in the older mice (Figure 8B,C). Immunohistochemistry (Figure 8D) and western blot (Figure 8E-G) show diminished amounts of  $\alpha$ -SMA and fibronectin. These data suggest that 1G6-D7 intervention prevents severe lung fibrosis in older mice.

#### 4 | DISCUSSION

In this study, we demonstrate that  $eHSP90\alpha$  contributes to fibroblast senescence, accompanied by abnormal



**FIGURE 6** The SMAD2/3 complex binds to p21 and TP53 promoters. (A) The schematic representation of p16, p21, and TP53 promoter with the indicated regions. ChIP analysis of p16 promoter (B), p21 promoter (C) and p53 promoter (D) of control cells. Immunoprecipitation was performed using anti-SMAD2/3 and IgG antibodies followed by PCR analysis. ChIP analysis of p16 promoter (E), p21 promoter (F), and p53 promoter (G) of cells exposed to F-5 or PBS. Immunoprecipitation was performed using anti-SMAD2/3 and IgG antibodies followed by PCR analysis. Data presented as mean  $\pm$  SD. \*\*\*\*p < .001, \*\*\*p < .001, \*\*p < .01, and \*p < .05 between indicated groups.

mitochondrial structure and mtROS accumulation. eHSP90 $\alpha$  activates TGF- $\beta$  signaling to enhance the transcription of senescence genes (Figure 9). We found that eHSP90 $\alpha$  blockade by 1G6-D7 decreased age-related lung fibrosis.

Our study showed that the level of serum HSP90 $\alpha$  was upregulated in BLM-injured mice and associated with disease severity, which is consistent with published data.<sup>4</sup> Administration of rHSP90a had a pro-fibrotic effect, as shown in our previous study<sup>5</sup> and Martin's study,<sup>4</sup> which highlights the importance of eHSP90 $\alpha$  in pulmonary fibrosis. Previous data demonstrated that eHSP90 $\alpha$  might activate ERK signaling. However, whether eHSP90 $\alpha$  mediates the senescence phenotype in pulmonary fibrosis is unknown. Interestingly, the level of HSP90 $\alpha$  was correlated with the levels of SASP factors, such as IL-6, TGF $\beta$ 1, and PAI1. Furthermore, the expression of p16 and p21 was upregulated from day 7, suggesting that senescent cells contribute to the inflammatory and fibrotic phases.

HSP90 has been shown to affect cellular senescence. Destabilizing the CDK6 interaction with HSP90 promotes premature cellular senescence.<sup>41</sup> A recent study showed that HSP90 $\beta$  interacts with MDM2 to suppress p53-dependent senescence.<sup>42</sup> Moreover, HSP90 inhibitors have been identified as novel senolytics.<sup>7,43</sup> In contrast, our in vitro experiments showed that eHSP90 $\alpha$ , but not intracellular HSP90 $\alpha$ , was upregulated in the secretome of etoposide-exposed fibroblasts. We also found that



**FIGURE 7** Blockade of eHSP90 $\alpha$  by 1G6-D7 attenuates cellular senescence and lung fibrosis in vivo. (A) Outline the design of therapeutic dosing of 1G6-D7 in mice with established fibrosis following bleomycin-induced lung injury. (B) Representative images show hematoxylin and eosin (H&E) staining of lung sections from the indicated groups of mice. (C) Twenty-one-day body weight was compared with baseline body weight. Lung resistance (D) and compliance (E) were measured by a pulmonary function test. (F) Hydroxyproline (HYP) contents in lung tissues from mice. Representative western blots (G) and quantitative analysis (H–K) show the amounts of p16, p21,  $\alpha$ -SMA, fibronectin and  $\beta$ -actin in lung homogenates from mice treated as shown in (A). Data presented as mean ± SD. \*\*\*\*p < .0001, \*\*\*p < .001, \*\*p < .01, and \*p < .05 between indicated groups. Representative western blots (L) and quantitative analysis (M, N) show the amounts of Psmad2, Psmad3 in lung homogenates. (O) Representative images show  $\alpha$ -SMA, fibronectin, and p16 staining of lung sections from the indicated groups of mice. (P) Immunofluorescence show  $\gamma$ H2AX staining.



**FIGURE 8** Blockade of eHSP90 $\alpha$  by 1G6-D7 accelerates resolution of bleomycin-induced lung fibrosis in aged mice. (A) Serum level of HSP90 $\alpha$  was measured by ELISA on 2-month-old and 18-month-old mice. Representative images show H&E (B), Masson's trichrome (C), and  $\alpha$ -SMA(D) staining of lung sections from the indicated groups of mice. (E–G) Representative immunoblots of  $\alpha$ -SMA, fibronectin, and  $\beta$ -actin in whole-lung homogenates. Data presented as mean  $\pm$  SD. \*\*\*\*p < .0001 and \*p < .05 between indicated groups.



**Fibroblast senescence** 

**FIGURE 9** Schematic illustration of mechanisms of eHSP90 $\alpha$  in cellular senescence.

eHSP90α stimulated a fibroblast senescence phenotype that included, for example, increased p16 and p21 expression and SA-β-gal activity. Using a model of BLM-induced lung fibrosis, we demonstrated that treatment with the anti-HSP90a antibody 1G6D7 significantly decreased cellular senescence. Thus, HSP90α functional diversity may depend on cell type and disease. Several studies have confirmed that the effect of eHSP90 $\alpha$  is mediated by lowdensity lipoprotein receptor-related protein 1 (LRP1).44-46 Intracellular HSP90ß may stabilize the TGF-BRII receptor to promote smad2/3 translocation into the nucleus.<sup>47</sup> However, the receptor to which eHSP90a binds to activate downstream pathways remains unknown. The SASP is an important pathological feature of senescence released through nuclear factor  $\kappa B$  (NF- $\kappa B$ ) signaling activation.<sup>48</sup> In this study, eHSP90a also augmented SASP factor expression, which promotes senescence. Overall, our findings highlight the importance of eHSP90 $\alpha$  in regulating fibroblast senescence.

Several studies have demonstrated the importance of TGF $\beta$  signaling in pulmonary fibrosis.<sup>17,49</sup> We showed that eHSP90 $\alpha$  activated TGF $\beta$  signaling in a manner dependent on mitochondrion-derived ROS accumulation. TGF $\beta$  signaling activation triggers translocation of the SMAD complex into the nucleus to promote target gene transcription.

SMADs have been implicated in mediating p21 transcription.<sup>22–24</sup> Our data showed that SMAD2/3, but not SMAD4, binds a specific sequence in the promoters of senescence markers. Furthermore, eHSP90 $\alpha$  promoted p53 and p21 transcription by enhancing the binding of SMAD2/3 to their promoters. As p53 also acts as a transcription factor to control p21 transcription, we demonstrated that eHSP90 $\alpha$  induces p21 transcription in direct and indirect ways. The accumulation of cytosolic p53, in particular, promotes mitochondrial dysfunction.<sup>50</sup>

Our previous experiments evaluated the preventive effect of the anti-HSP90a antibody 1G6-D7 in a BLMinduced injury model.<sup>5</sup> To accurately assess anti-fibrotic efficacy, the intervention should inhibit fibrosis without impacting early inflammation.<sup>51</sup> In this study, our therapeutic model employed 1G6-D7 administration from day 8 to day 20, which led to improved lung function and decreased collagen deposition. While aging is a major risk factor for IPF and cellular senescence has been proposed as a typical feature,<sup>52</sup> we observed a normal lung structure with the absence of inflammation or fibrosis in 18-month-old mice. In contrast, some studies found significant alveolar changes with age.<sup>53,54</sup> Previous research<sup>55</sup> demonstrated that old mice (above 18-month-old) are more sensitive to bleomycin injury which is consistent with our results. A recent study performed an analysis on human lungs from donors age between 16 and 76 years. They found damage of alveolar epithelial cell and an increasing proportion of fibroblasts with age.<sup>56</sup> Relationship between fibroblast proliferation and cellular senescence should be further explored. Interestingly, 18-month-old mice presented high levels of circulating HSP90 $\alpha$  than 2-month-old mice, suggesting eHSP90 $\alpha$  might be involved in injury response. Additionally, eHSP90a blockade significantly reduced lung fibrosis in the old model mice. The effect of 1G6-D7 on BLM-injured old mice highlights its capacity to reduce age-related fibrosis and cellular procession.

In conclusion, the present study has demonstrated that eHSP90 $\alpha$  causes mitochondrial dysfunction, activating mtROS-dependent TGF-beta signaling to induce fibroblast senescence. eHSP90 $\alpha$  blockade reduced lung fibrosis in both young and old mode mice. This study adds to the literature and identifies eHSP90 $\alpha$  as an essential contributor to fibroblast senescence in pulmonary fibrosis, suggesting an anti-hsp90a antibody as a potential treatment for age-related fibrotic disease.

#### AUTHOR CONTRIBUTIONS

Wenshan Zhong, Weimou Chen, and Yuanyuan Liu designed and performed experiments, analyzed data, interpreted the results, and wrote the manuscript. Jinming Zhang and Ye Lu performed experiments and analyzed data. Xuan Wan, Yujie Qiao, Haohua Huang, and Zhaojin Zeng performed experiments. Xiaojing Meng, Wei Li, Haijin Zhao, and Mengchen Zou interpreted the results and edited the manuscript. Hangming Dong and Shaoxi Cai designed and supervised the study, interpreted the results, wrote, and edited the manuscript.

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#### DISCLOSURES

The authors declare no conflicts of interest.

#### DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available in the methods and/or Supporting Information of this article.

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#### SUPPORTING INFORMATION

Additional supporting information can be found online in the Supporting Information section at the end of this article.

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