

# Human senescent fibroblasts trigger progressive lung fibrosis in mice

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## ABSTRACT

Cell senescence has recently emerged as a potentially relevant pathogenic mechanism in fibrosing interstitial lung diseases (f-ILDs), particularly in idiopathic pulmonary fibrosis. We hypothesized that senescent human fibroblasts may suffice to trigger a progressive fibrogenic reaction in the lung. To address this, senescent human lung fibroblasts, or their secretome (SASP), were instilled into the lungs of immunodeficient mice. We found that: (1) human senescent fibroblasts engraft in the lungs of immunodeficient mice and trigger progressive lung fibrosis associated to increasing levels of mouse senescent cells, whereas non-senescent fibroblasts do not trigger fibrosis; (2) the SASP of human senescent fibroblasts is pro-senescence and pro-fibrotic both *in vitro* when added to mouse recipient cells and *in vivo* when delivered into the lungs of mice, whereas the conditioned medium (CM) from non-senescent fibroblasts lacks these activities; and, (3) navitoclax, nintedanib and pirfenidone ameliorate lung fibrosis induced by senescent human fibroblasts in mice, albeit only navitoclax displayed senolytic activity. We conclude that human senescent fibroblasts, through their bioactive secretome, trigger a progressive fibrogenic reaction in the lungs of immunodeficient mice that includes the induction of paracrine senescence in the cells of the host, supporting the concept that senescent cells actively contribute to disease progression in patients with f-ILDs.

## INTRODUCTION

Fibrosing interstitial lung diseases (f-ILDs) constitute a complex and heterogeneous group of diseases characterized by non-resolving pulmonary fibrosis [1]. Idiopathic pulmonary fibrosis (IPF) is the most frequent and representative f-ILD [2, 3]. The pathogenesis of f-ILD is complex and still incompletely understood but cell senescence has recently emerged as a potentially relevant pathogenic player [4–7]. Cell senescence is an adaptation of cells to circumstances of unreparable cellular damage [8, 9]. The entry in senescence involves a profound rewiring of cellular biology that is largely irreversible, with a permanent exit from the cell cycle (in the case of proliferating cells), the acquisition of stable epigenetic changes, the expansion of the lysosomal compartment and a vigorous Senescence Associated Secretory Phenotype (SASP) [10]. The SASP includes multiple pro-inflammatory and tissue remodelling mediators that can foster a fibrogenic cascade and propagate the senescent phenotype to the surrounding cells [11]. In support of the pathogenic role of the SASP in age-related diseases, senescent cells cause systemic frailty [12] and organ deterioration [13, 14] when transplanted to healthy animals. Moreover, lung fibroblasts isolated from patients with IPF display features of cellular senescence [15]. Senescence-related processes are among the top upregulated transcriptional signatures in IPF alveolar type 2 (AT2) cells [16]. However, the contribution of senescent lung fibroblasts in the initiation and progression of lung fibrosis seen in f-ILD is poorly defined.

We hypothesized that xeno-transplantation of senescent human fibroblasts into the lungs of immunodeficient mice may trigger a fibrogenic reaction. Here we: (1) explored this hypothesis *in vivo*; (2) investigated the potential underlying biological mechanisms *in vitro*; and (3) studied the effects of one experimental senolytic compound (navitoclax) and two anti-fibrotic drugs currently used in the treatment of IPF in humans (nintedanib and pirfenidone), both *in vivo* and *in vitro*.

## RESULTS

### Senescent human lung fibroblasts induce progressive lung fibrosis in mice

As a cellular model, we used human lung fibroblasts (IMR90), which is a normal diploid non-immortalized cell line. Senescence was induced *in vitro* by treatment with  $\gamma$ -radiation as confirmed by detection of senescence-associated beta-galactosidase (SABG) activity, which is a common marker of cellular senescence (Figure 1A). We will refer to these cells as

SEN-IMR90. To determine if the xeno-transplantation of senescent human cells in the mouse lungs would initiate a fibrogenic reaction, IMR90, SEN-IMR90, or PBS as control, were intratracheally instilled in the lungs of immunodeficient mice (Figure 1B). After 21 days, collagen deposition, as assessed by Masson's trichrome (MT) staining, was increased in the lungs from animals xeno-transplanted with SEN-IMR90 cells, but not with IMR90 (or PBS) (Figure 1C). Likewise, the lungs of animals xeno-transplanted with SEN-IMR90 cells had a significantly higher modified Ashcroft score (Figure 1D), higher level of hydroxyproline content, and higher levels of murine *Col6a3* expression (Figure 1E), three well-established markers of fibrosis [17], compared to the IMR90 group. Collectively, these results indicate that human senescent IMR90 cells mediate pro-fibrotic effects in mouse lungs.

### Senescent human lung fibroblasts engraft successfully in mouse lungs

Based on the pro-fibrotic effects induced by SEN-IMR90 reported above, we explored if these cells can successfully engraft within the lung parenchyma of immunodeficient mice after instillation. We found that human senescent cells, as detected by staining with antibodies against human CDKN1A/p21<sup>Cip1/Waf1</sup>, a senescence marker, and Anti-Human Nuclear Antigen (HuNu), can be localized in the interstitium of mouse lungs 3 hours post-transplantation, and are still detectable 48 hours after instillation (Figure 2A). Similar observations were made when we measured human *CDKN1A/p21<sup>Cip1/Waf1</sup>* mRNA levels (Figure 2B). We verified senescence in the engrafted cells by serial sectioning and staining with antibodies against human CDKN1A/p21<sup>Cip1/Waf1</sup> and Phospho-H2AX ( $\gamma$ -H2AX) as a marker for DNA damage and genomic instability (Figure 2C).

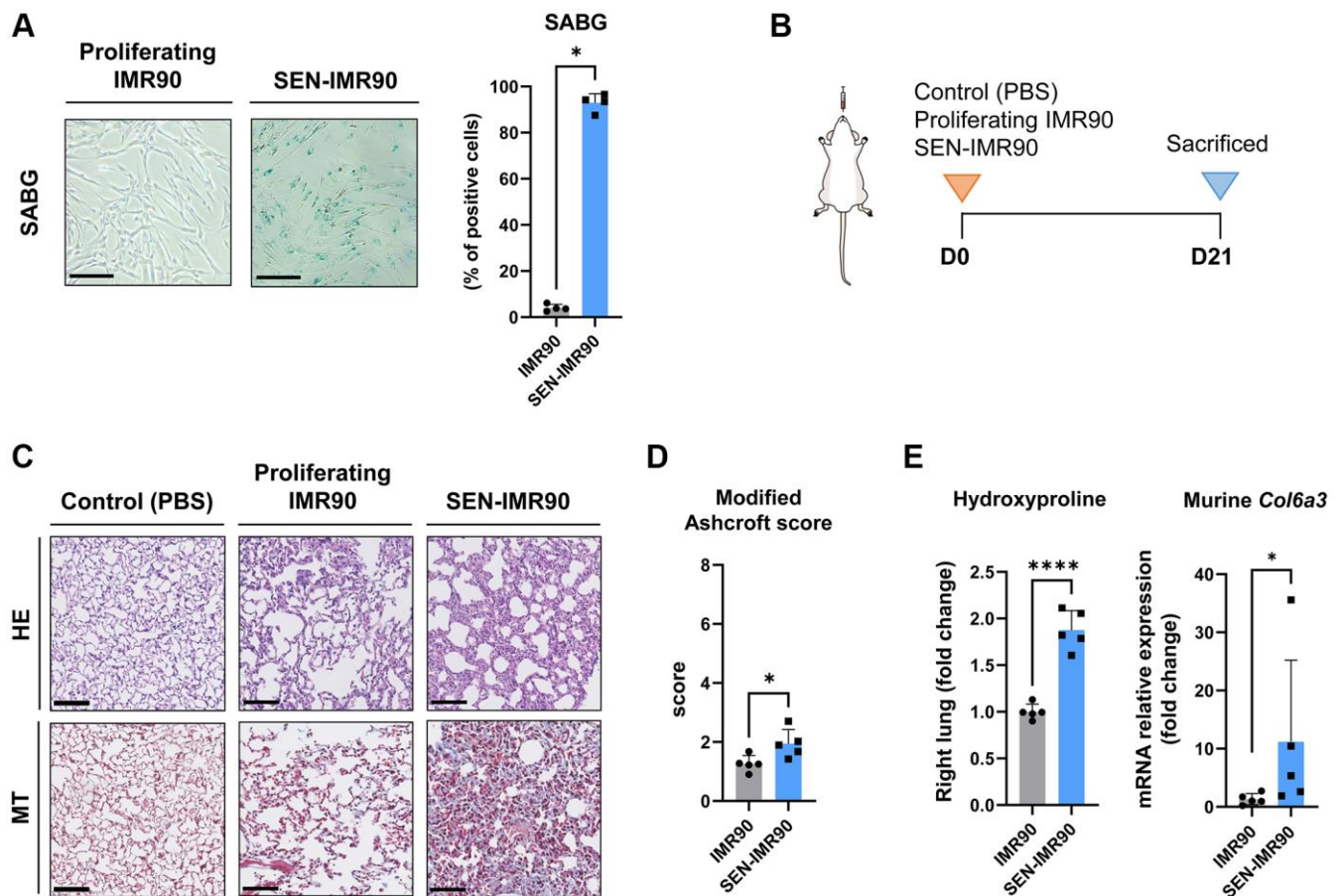
### Progression of murine lung fibrosis initiated by senescent human lung fibroblasts

To explore how fibrosis evolves over time, we monitored the dynamics of lung fibrosis at different time points after SEN-IMR90 transplant, using PBS as a negative control (Figure 3A). We observed that, over time (up until 2 months after transplant), there was a progressive increase in several fibrotic markers including hydroxyproline content (Figure 3B), and mRNA levels of murine *Colla2* (Figure 3C) and the murine senescent marker *Cdkn1a/p21<sup>Cip1/Waf1</sup>* (Figure 3D). This suggests that fibrosis initiated by senescent human cells evolves into murine senescence and fibrosis at late stages of disease. At the latest endpoint analyzed (two months post challenge), fibrosis content was compared to mice treated with a single dose of bleomycin [18, 19]. The levels of

hydroxyproline content (Supplementary Figure 1A) and murine *Col1a2* (Supplementary Figure 1B) and *Cdkn1a/p21<sup>Cip1/Waf1</sup>* (Supplementary Figure 1C) mRNAs were similar in SEN-IMR90 transplanted animals compared to those treated with bleomycin. The increase in murine senescence was confirmed by immunohistochemistry, which showed high levels of p21-positive cells 1 month after instillation of human SEN-IMR90 cells (Figure 3E). This contrasted with the progressive elimination of the engrafted human SEN-IMR90 cells, which were essentially undetectable by qRT-PCR (Figure 2B) or by immunohistochemistry (Figure 3E) 1 month after intratracheal instillation. Therefore, senescent human lung fibroblasts initiate a process of fibrosis that is progressive and associated to an increase in senescent murine cells.

### The secretome of senescent human lung fibroblasts is profibrotic both *in vitro* and *in vivo*

Based on the observations above, we explored the profibrogenic potential of the secretome of human senescent IMR90 lung fibroblasts (SASP), both *in vitro* and *in vivo*. As shown in Figure 4A, we found that the levels of many SASP mediators measured using a multiplex antibody-based commercial assay, including RANTES (Regulated upon Activation, Normal T Cell Expressed and Presumably Secreted), cathepsin D, C-X-C motif chemokine ligand 1 (CXCL1), interleukin (IL)-1 $\alpha$ , interferon gamma-induced protein 10 (IP-10), monocyte chemoattractant protein 3 (MCP-3), macrophage colony stimulating factor (M-CSF), monokine induced by gamma interferon (MIG), and macrophage



**Figure 1. Senescent human lung fibroblasts induce lung fibrosis in mice.** (A) IMR90 lung fibroblasts were exposed to  $\gamma$ -irradiation (20 Gy). Fourteen days later, senescence was confirmed by SABG staining (scale bar, 100  $\mu$ m). (B) Immunodeficient (nude) mice were randomized to receive intratracheal instillation of proliferating human lung fibroblasts (IMR90) or senescent IMR90 (SEN-IMR90). PBS was used as a negative control. (C) Representative images of lung sections stained with Hematoxylin Eosin (HE) and Masson's Trichrome (MT) from mice injected with IMR90 cells, SEN-IMR90 cells or PBS at 21 days post-injection. Scale bar 100  $\mu$ m. (D) Modified Ashcroft score of MT staining in sections from mice injected with IMR90 or SEN-IMR90 cells at 21 days post-injection;  $n = 5$ . These data are part of a larger experiment presented in Triana-Martinez F, et al. [27]. (E) Hydroxyproline content in the right lung tissue of mice injected with SEN-IMR90 compared with IMR90 group at 21 days post-injection;  $n = 5$  (left panel). These data are part of a larger experiment presented in Triana-Martinez F, et al. [27]. Relative expression of the mRNA coding for *Col6a3* relative to *Actin-b* levels in lung cell extracts from mice injected with IMR90 and SEN-IMR90 cells at 21 days post-injection;  $n = 5$  (right panel). All values are expressed as fold change relative to IMR90 group. Statistical significance was assessed by *U*-Mann Whitney test: \* $p < 0.05$ , \*\*\*\* $p < 0.0001$ . For further explanations, see text.

inflammatory protein 1-alpha (MIP-1 $\alpha$ ), were significantly higher (and that of platelet derived growth factor-AA, lower) in conditioned medium (CM) from SEN-IMR90 cells (SEN-CM) compared to CM from non-senescent IMR90 cells (NS-CM), with no significant differences in IL-6 or plasminogen activator inhibitor-1 (PAI-1) levels.

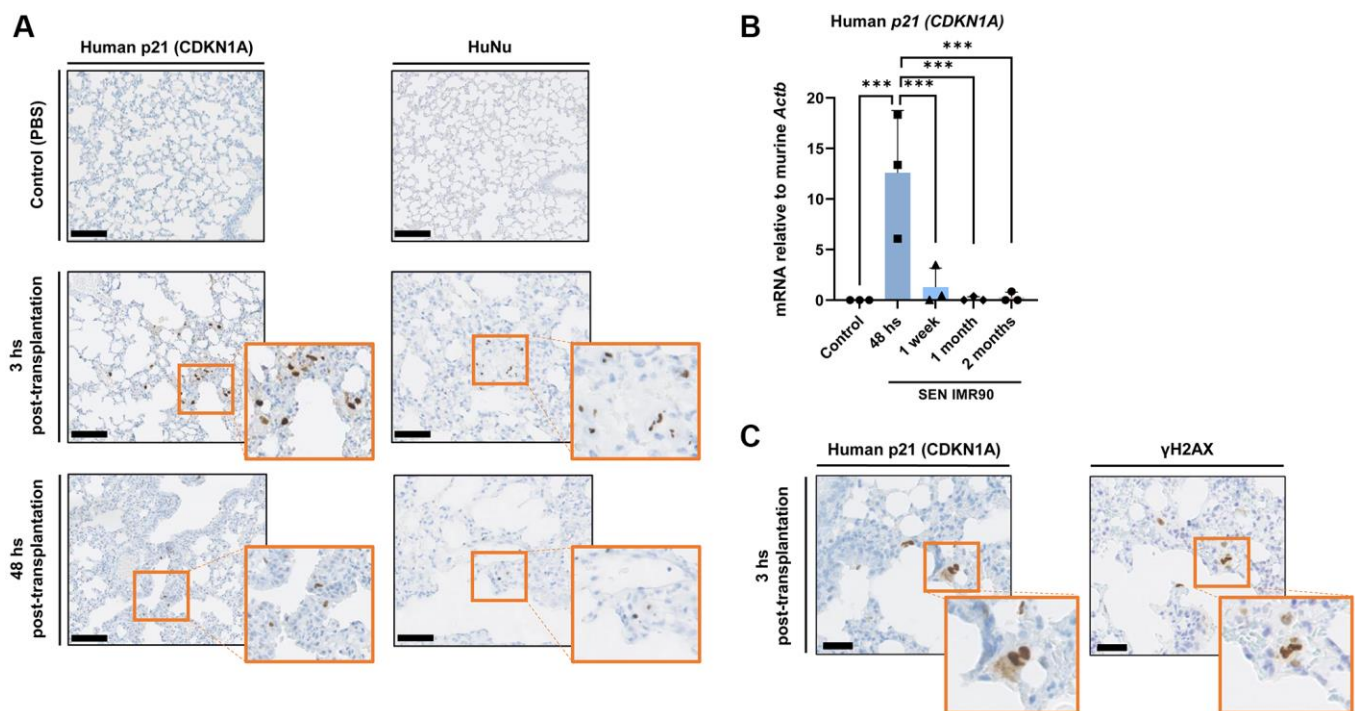
Then, to explore if human SASP factors released by SEN-IMR90 induce senescence in mouse cells, we incubated mouse embryo fibroblasts (MEFs) with CM derived from 6-day cultured SEN-IMR90 and confirmed that it does indeed trigger senescence in MEFs, as indicated by enhanced SABG staining (Figure 4B), and transcriptional upregulation of the profibrotic SASP components *IL-6*, *Tgf- $\beta$*  and *Colla2* (Figure 4C). In addition, we detected by immunofluorescence the expression of CDKN1A/p21<sup>Cip1/Waf1</sup> and the myofibroblast activation marker alpha smooth muscle actin ( $\alpha$ -SMA) (Figure 4D). The amount of double positive  $\alpha$ -SMA/p21 cells in MEFs treated with SEN-CM was significantly higher compared to those treated with NS-CM (Figure 4E). Finally, we observed that the instillation of SEN-CM to immunodeficient

mice (Figure 4F) triggered a fibrogenic response 21 days after treatment, as indicated by augmented hydroxyproline levels (Figure 4G) and MT staining (Figure 4H).

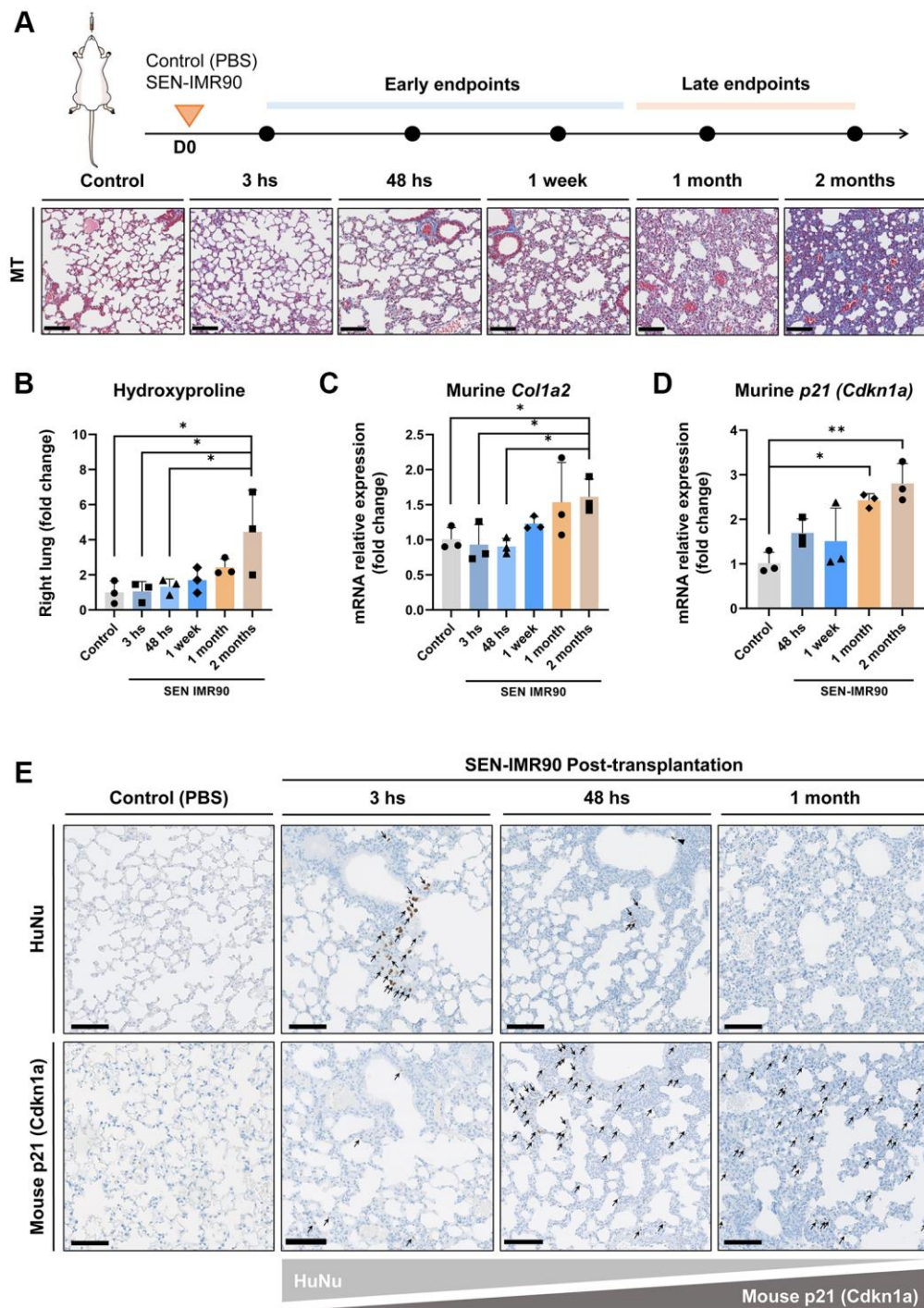
### Effect of antifibrotic and senolytic drugs

Twenty-one days after the instillation of SEN-IMR90 cells (or proliferating IMR90 cells), mice were randomized to receive nintedanib, pirfenidone, navitoclax or vehicle for two weeks (Figure 5A). We observed that hydroxyproline levels (Figure 5B) were significantly lower in mice treated with any of these drugs vs. vehicle.

Since pirfenidone and nintedanib were able to rescue the fibrotic phenotype in mouse lungs initiated by senescent human fibroblasts, we then investigated if these agents had senolytic activity in mouse cells *in vitro*. To this end, we induced senescence in MEFs with bleomycin for 7 days and senescent (and non-senescent) MEFs were then exposed to increasing concentrations of nintedanib, pirfenidone, navitoclax or vehicle for 72 hours. We observed that, in line with its



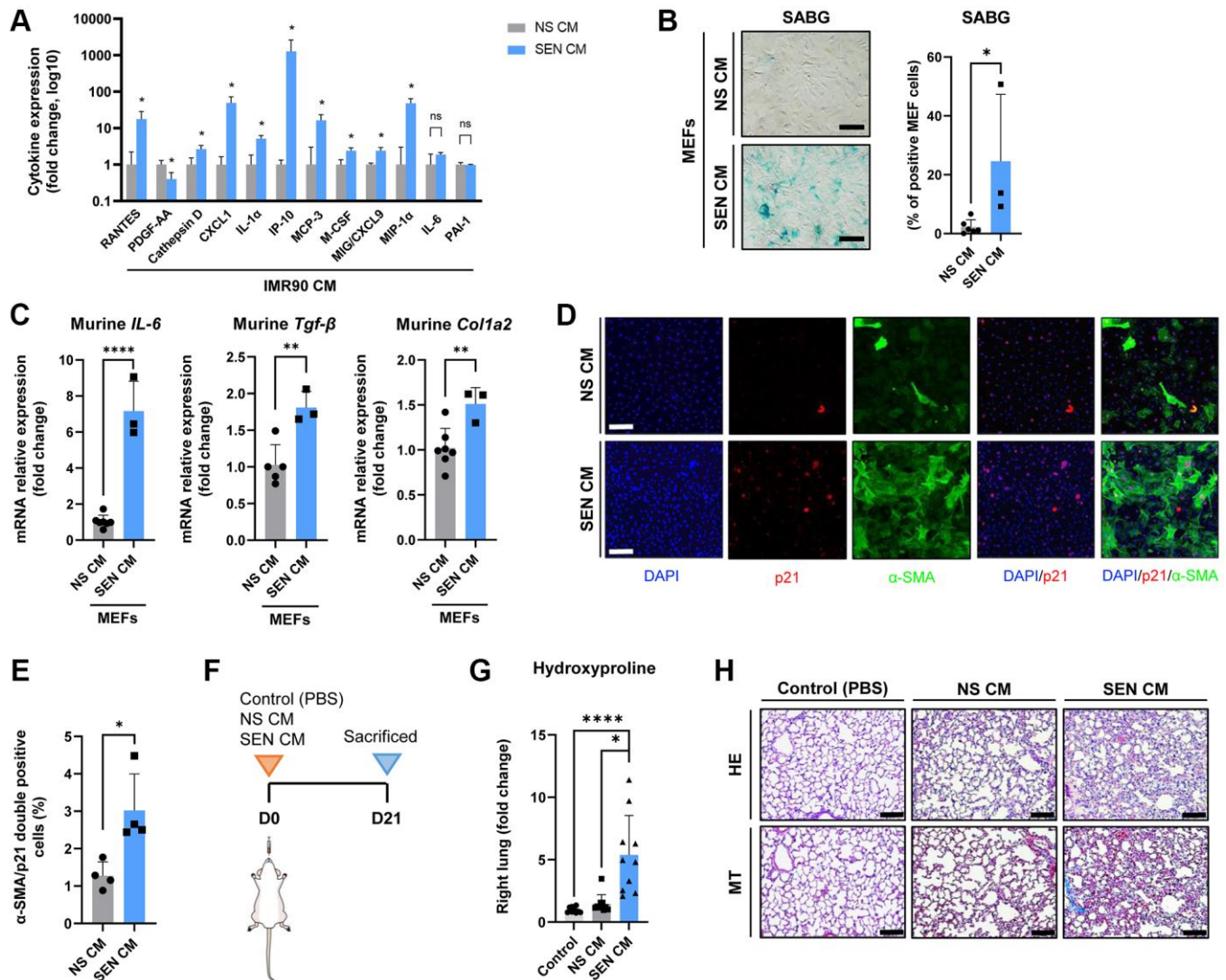
**Figure 2. Senescent human lung fibroblasts infiltration and engraftment in mice lungs.** (A) Images of lung sections showing IHC staining for *CDKN1A/p21<sup>Cip1/Waf1</sup>* (left panel) and HuNu (right panel) from mice injected with SEN-IMR90 or control. Engraftment of senescent cells (arrows) in mice sacrificed after 3 and 48 hours post-transplantation. Positive cells were confirmed using histology at low magnification (20 $\times$ , scale bar 100  $\mu$ m), and high magnification (40 $\times$ , orange box). (B) RT-PCR expression of *CDKN1A/p21<sup>Cip1/Waf1</sup>* was measured relative to murine *Actin-b* to demonstrate the engraftment of SEN-IMR90 in the lungs of nude mice after 48 hours post-transplantation, and their presence at later endpoints compared to control;  $n = 3$  each group. Statistical significance was assessed by the one-way ANOVA with Tukey test: \*\*\* $p < 0.001$ . (C) Serial section and staining with antibodies against human *CDKN1A/p21<sup>Cip1/Waf1</sup>* and Phospho-H2AX ( $\gamma$ -H2AX) to demonstrate senescence state of the engrafted cells. For further explanations, see text.



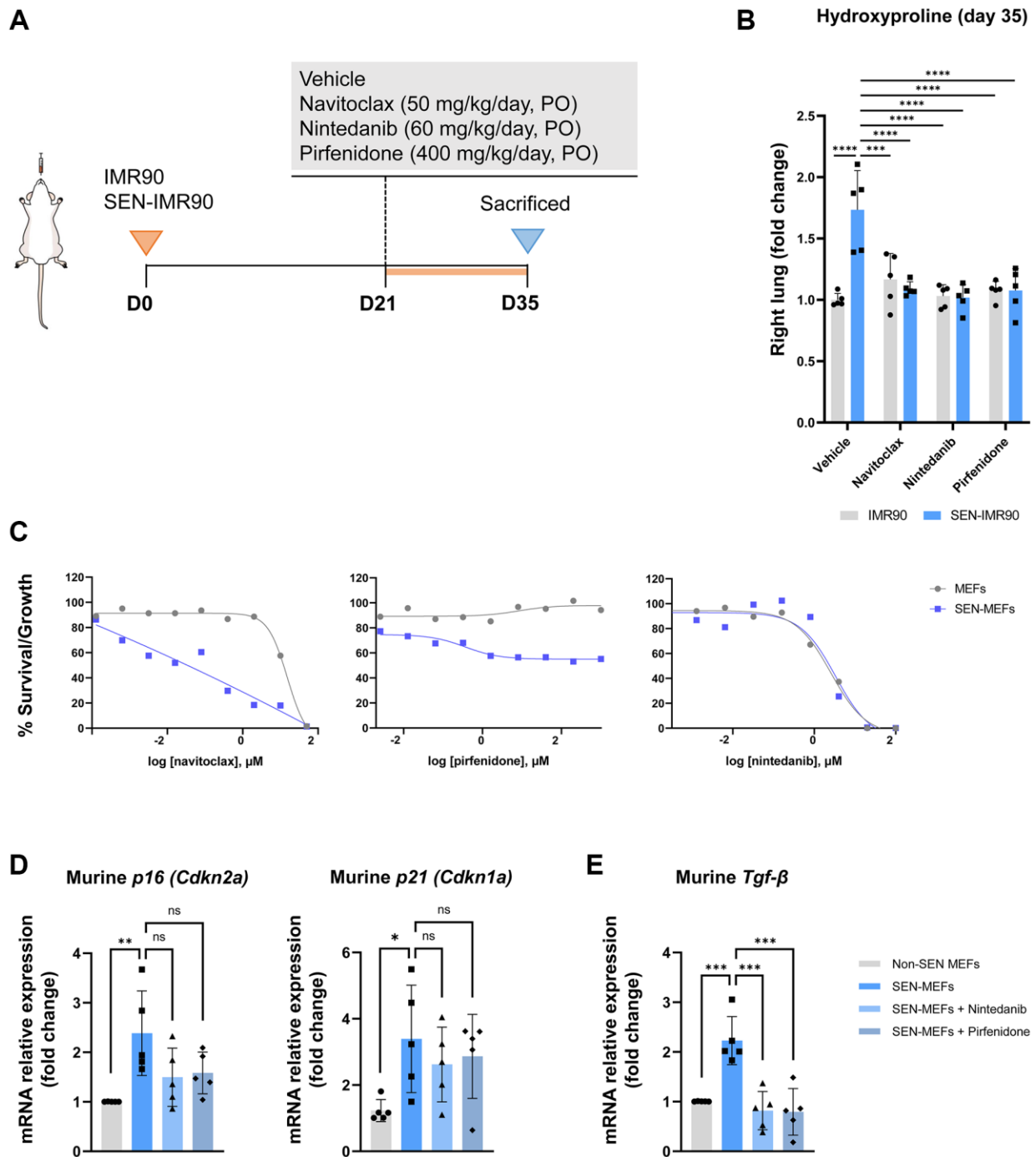
**Figure 3. Lung fibrosis induced by senescent human lung fibroblasts is progressive.** (A) Diagram showing the experimental plan to evaluate the dynamics of the development of pulmonary fibrosis in nude mice, as well as representative images of lung sections stained with Masson's Trichrome (MT) (20 $\times$ , scale bar 100  $\mu$ m). These animals received intratracheal instillation of irradiated SEN-IMR90, compared with PBS-exposed mice at early endpoints, and bleomycin-challenged mice (single injection, dose of 3 UI/kg) at late endpoints;  $n = 3$  each group. (B) Hydroxyproline content in the right lung of mice injected with SEN-IMR90 compared with control (PBS). (C) Relative expression of the mRNA coding for murine *Col1a2* in the lungs of the same mice as in panel B. (D) Relative expression of the mRNA coding for murine *Cdkn1a*/p21<sup>Cip1/Waf1</sup> in the same mice as in panel B. For panels B, C and D,  $n = 3$  for each experimental group and statistical significance was assessed by the one-way ANOVA with Tukey test: \*\* $p < 0.01$ ; \* $p < 0.05$ . For panels B, C and D, the group labelled SEN-IMR90 (2 months) is the same group labelled SEN-IMR90 in Supplementary Figure 1A–1C, and the data are the same. (E) Images of lung sections showing positive cells using IHC staining for HuNu and mouse p21 (Cdkn1a/p21<sup>Cip1/Waf1</sup>) from mice injected with SEN-IMR90 or control. Engraftment of senescent cells (arrows) in mice sacrificed after different time points (3 hours, 48 hours, and 1 month), showing their dramatic reduction after 48 hours post-transplantation, and the gradually increase of mouse p21 over time (20 $\times$ , scale bar 100  $\mu$ m).

well documented senolytic properties, navitoclax induced apoptosis with higher efficiency in senescent MEFs compared to non-senescent MEFs (Figure 5C). In

contrast, nintedanib or pirfenidone were not pro-apoptotic (pirfenidone) or were equally pro-apoptotic over senescent and non-senescent MEFs (nintedanib)



**Figure 4. The secretome of senescent human lung fibroblasts as mediator of murine lung fibrosis.** (A) Diagram showing cytokine concentrations in conditioned media (CM) corresponding to 0.5 million cells collected from irradiated SEN-IMR90 (SEN-CM) compared with proliferating IMR90-derived CM (NS-CM) as control, were quantified by using human cytokine arrays;  $n = 4$  each group, independent experiments. Statistical significance was assessed by the two-tailed Student's test:  $*p < 0.05$ . (B) Mouse Embryo Fibroblasts (MEF) incubated with SEN-CM or NS-CM as control for 6 days. Senescence was confirmed by SABG staining (scale bar, 100  $\mu\text{m}$ ). (C) Transcriptional upregulation of the profibrotic secretome components (*IL-6*, *Tgf- $\beta$*  and *Col1a2*) was confirmed by RT-PCR at 2 days post-exposure to the indicated CM;  $n = 6$  SEN-CM and  $n = 3$  NS-CM group, independent experiments. Statistical significance was assessed by the two-tailed Student's test:  $*p < 0.05$ . (D) Representative immunofluorescence images showing double staining of *Cdkn1a/p21<sup>Cip1/Waf1</sup>* (red), and  $\alpha$ -smooth muscle actin ( $\alpha$ -SMA) (green) (10 $\times$ , scale bar, 100  $\mu\text{m}$ ). (E) Quantification of the average number of  $\alpha$ -SMA/p21 double positive cells as observed in the images shown in panel D using ImageJ. Quantification was performed from 4 experiments with  $>25$  cells quantified for each condition. Statistical significance was assessed by the two-tailed Student's *t*-test:  $*p < 0.05$ . (F) Diagram showing the experimental plan to evaluate the effect of the secretome in nude mice lung. These animals were intratracheally delivered SEN-CM or NS-CM, normalized by the number of cells (corresponding to  $5 \times 10^5$  cells each group), using PBS as negative control;  $n = 10$  each group. (G) Hydroxyproline content in the right lung tissues of mice injected with SEN-CM or NS-CM, compared with control;  $n = 10$  each group. (H) Representative images of lung sections of nude mice 21 days after instillation of SEN-CM or NS-CM, or PBS as negative control, stained with Hematoxylin Eosin (HE) and Masson's Trichrome (MT) (40 $\times$ , scale bar 100  $\mu\text{m}$ ) showing that SEN-CM initiated a cascade of the events that induced mild fibrosis. Statistical significance was assessed by the one-way ANOVA with Tukey test:  $***p < 0.001$ ;  $*p < 0.05$ . For further explanations, see text.



**Figure 5. Effects of antifibrotic and senolytic drugs.** (A) Scheme showing the experimental design to assess the effect of antifibrotic or senolytic drugs. Nude mice were randomized after 21 days post-injection of irradiated SEN-IMR90 cells or IMR90 cells as negative control, to either the two approved antifibrotics drugs (nintedanib or pirfenidone), a senolytic drug (navitoclax), or vehicle, for two weeks. (B) Hydroxyproline content in the right lung tissues of mice treated with navitoclax, nintedanib or pirfenidone, compared with control;  $n = 5$  each group. Statistical significance was assessed by the one-way ANOVA with Tukey test: \*\*\*\* $p < 0.0001$ ; \*\*\*\*\* $p < 0.00001$ . (C) Senolytic activity of navitoclax (left panel), pirfenidone (middle panel) or nintedanib (right panel). Diagram showing the senolytic activity of these agents after exposure of senescent MEFs (SEN-MEFs) or non-senescent MEFs (NS-MEFs) to increasing concentration of navitoclax, pirfenidone, nintedanib or vehicle for 72 hours, as confirmed by relative expression of the mRNA coding for murine senescence markers (*Cdkn2a*/p16<sup>INK4a</sup> and *Cdkn1a*/p21<sup>Cip1/Waf1</sup>), measured relative to *Actin-b* levels in lung cell extracts of nintedanib or pirfenidone group compared to control (D);  $n = 5$  each group, independent experiments. Statistical significance was assessed by the one-way ANOVA with Tukey test: \*\* $p < 0.01$ ; \* $p < 0.05$ . (E) Relative expression of the mRNA coding for *Tgf-β* (transforming growth factor-β) was measured relative to *Actin-b* levels in SEN-MEFs treated with pirfenidone or nintedanib, compared with control;  $n = 5$  each group, independent experiments. Statistical significance was assessed by the one-way ANOVA with Tukey test: \*\*\* $p < 0.001$ . For further explanations, see text.

(Figure 5C) [20]. In keeping with this observation, neither nintedanib or pirfenidone reduced the levels of the senescent markers *Cdkn2a/p16<sup>INK4a</sup>* and *Cdkn1a/p21<sup>Cip1/Waf1</sup>* in senescent MEFs (Figure 5D). Thus, the anti-fibrotic effect of pirfenidone and nintedanib seems unrelated to promoting selective apoptosis of senescent cells in the lung. As anticipated, both drugs decreased the expression of *Tgf-β* in senescent MEFs treated with these molecules (Figure 5E), reinforcing the widely accepted concept that these drugs operate mainly by reducing TGF-β.

## DISCUSSION

The main results of this study are that: (1) human senescent fibroblasts engraft successfully in the lungs of immunodeficient mice and trigger progressive lung fibrosis; (2) the secretome of senescent human lung fibroblasts (SASP) is profibrotic both *in vitro* and *in vivo*; and, finally, (3) navitoclax (an experimental senolytic compound) and two anti-fibrotic drugs currently used in the treatment of IPF in humans (nintedanib and pirfenidone) ameliorate lung fibrosis induced by senescent human fibroblasts *in vivo*, albeit only navitoclax displayed clear direct senolytic activity.

### Previous studies

Existing animal models of f-ILDs do not fully recapitulate the complex pathobiology of human interstitial diseases, thus limiting their use to explore potential candidate therapeutic drugs. For instance, spontaneous resolution of tissue injury is a major drawback of the bleomycin-induced lung fibrosis model, which currently is the most widely employed animal model of IPF [21–26]. On the other hand, cell senescence has been recently identified as a potentially relevant pathogenic mechanism in IPF [15, 16]. Accordingly, we sought to better evaluate the contribution made by senescent human lung fibroblasts toward progressive tissue remodelling observed in fibrotic lung disease. Here we have developed a novel f-ILD experimental model where progressive pulmonary fibrosis in immunodeficient mice is triggered by the intratracheal xeno-transplantation of human senescent fibroblasts. We propose that this new model of fibrosis initiated by human senescent cells in mice may open a window of opportunity for a better understanding of the pathobiological mechanisms underlying f-ILDs in humans, as well as for the preclinical investigation of potential anti-fibrotic candidate drugs [16, 27–31].

### Interpretation of novel findings

The results of this study support the tenet that senescent cells integrating the lung parenchyma are sufficient to

recapitulate key pathologic features of f-ILDs and enable the propagation of the senescent phenotype (secondary senescence) at later stages of disease. In particular, we showed that transplanted human senescent cells initiate a process that evolves into murine fibrosis at later stages of disease. This observation reinforces the concept that the senescent secretome causally accounts for the ability of senescent cells to initiate and feed the fibrogenic cascade in the lung [7, 32, 33]. In keeping with this observation, we demonstrate that SASP factors derived from human senescent fibroblasts induce a molecular signature of senescence in mouse recipient cells *in vitro*, while mirroring the pro-fibrotic effect of transplanted human senescent cells in inducing fibrotic damage in the mouse lung. Thus, collectively, our *in vitro* and *in vivo* results pinpoint towards the secretome of senescent human fibroblasts, which is a known source of factors implicated in proliferation and tissue rearrangement in lung fibrosis [7, 32–34], as a mechanism by which these cells mediate a humanized fibrotic pulmonary disorder in mice.

On the other hand, and in keeping with the notion that senolytic drugs may expand the available toolbox to treat pulmonary fibrosis [32], we also showed that the prototypical senolytic drug, navitoclax, reduces the burden of collagen deposition in our experimental model. This is apparently in contrast with the mechanisms of action of two drugs currently used in the clinic to treat IPF (pirfenidone and nintedanib) which also ameliorate lung fibrosis independently of their ability to promote apoptotic death of senescent cells. In agreement with previous reports [35, 36], we show that pirfenidone and nintedanib can reduce the expression of the pro-fibrotic factor TGF-β. Based on these findings, we propose that these two drugs act as “senomorphic” agents by modifying the composition of the SASP [37].

### Potential limitations

We are aware that the method of senescence induction impacts the repertoire of SASP factors expressed, which further adds to the complexity of the SASP which may also be both cell type and microenvironment specific [38]. Our study shows the contribution of the secretory profile of human senescent fibroblasts triggered by γ-radiation, since ionizing radiation-induced senescence is known to cause tissue fibrosis [39, 40]. The diverse biological roles of other senescence inducers (DNA-damaging chemotherapeutic agents, oncogene-induced senescence, or replicative senescence) or cell types in lung fibrosis using this system still need to be investigated, particularly *in vivo*.

Furthermore, we used immunodeficient mice to facilitate the engraftment of human cells in the lung, but



we recognize that this may represent a limitation of our model because these mice are unable to mount adaptive immune responses. T cells can be implicated in the pathogenesis of pulmonary fibrosis, can participate in the surveillance of senescent lung cells and may even be involved in the therapeutic pathways of antifibrotic molecules [41]. In this sense, the interaction of the immune response with senescent cells cannot be widely studied with this new model. However, we also showed that our experimental model can be a useful preclinical platform to evaluate antifibrotic therapies, as testified by the fact that two approved drugs for IPF (pirfenidone and nintedanib) significantly improved the fibrotic phenotype initiated by human senescent cells.

## CONCLUSIONS

Our results indicate that human senescent fibroblasts trigger a progressive diffuse mild fibrogenic reaction in the lung of immunodeficient mice through their bioactive secretome. These observations support that accumulation of senescent cells may contribute to fibrotic lung disease in patients with f-ILDs, particularly IPF.

## METHODS

Methods are summarized below and presented in detail in the online supplement.

### *In vivo* humanized mouse model

Human lung fibroblasts (IMR90) were purchased from the American Type Culture Collection (ATCC, Manassas, VA, USA) and grown in Dulbecco's Modified Eagle Medium (DMEM) (Gibco®) supplemented with 10% foetal bovine serum and 100 U/mL penicillin/streptomycin, hereinafter referred to as DM10 media, and maintained at 37°C, 5% CO<sub>2</sub>. To induce cell senescence, when cells reached 50% confluence, proliferating IMR90 cells were exposed to ionizing  $\gamma$ -radiation (20 Gy). Senescence induction was assessed by monitoring senescence-associated  $\beta$ -galactosidase (SABG) activity using a protocol adapted from Dimri et al. [42]. Then, normal proliferating IMR90 or  $\gamma$ -irradiated senescent human fibroblasts IMR90 (SEN-IMR90) ( $5 \times 10^5$  cells each group) were washed and resuspended in PBS, and then instilled intratracheally to the lung of 6–8 weeks old male nude athymic (*nu/nu*) mice (Envigo Laboratory). At the time points indicated in each experiment, mice were euthanized, and lungs were removed for analysis. The degree of pulmonary fibrosis induced was determined semi-quantitatively by Masson's trichrome (MT) staining (AR17311-2, Dako – Agilent), and quantitatively by the concentration of lung hydroxyproline, the modified Ashcroft score and the

collagen mRNA levels, determined using real-time quantitative PCR (RT-qPCR) (PowerUp™ SYBR® Green Master Mix, Applied Biosystems, Foster City, CA, USA) [17, 43].

### Instillation of IMR90-derived conditioned medium into mouse lung

IMR90 and SEN-IMR90 cells were cultured in DM10 medium, at 37°C and under hypoxic conditions (5% CO<sub>2</sub>). After 14 days, medium was removed and exchanged with 25 ml fresh culture medium, and then collected 24 hrs later. Levels of cytokines, chemokines, and growth factors in conditioned medium (CM) from proliferating IMR90 and SEN-IMR90 were quantitated using Human Cytokine 48-Plex Discovery Assay (Eve Technologies Corporation, Canada). CM corresponding to 0.5 million IMR90 cells was transferred to mice as detailed in the Supplementary Materials, and PBS was used as negative control (see details in the Supplementary Materials); 3 weeks later lungs were removed and analysed.

### *In vitro* experiments

Mouse embryonic fibroblasts (MEFs) were isolated from 13.5-day C57BL/6J mouse embryos and grown in DM10 media at 37°C and under hypoxic conditions (5% CO<sub>2</sub>), as detailed in the Supplementary Materials. Senescence was induced by either: (1) paracrine induction by CM from IMR90 control or SEN-IMR90, as detailed in the Supplementary Materials; or (2) addition of 100  $\mu$ M Bleomycin sulphate (A10152, Sigma-Aldrich) to cell culture for 7 days. The degree of senescence induction was monitored by SABG activity as above [42], gene expression analysis (RT-PCR), or immunofluorescence (against p21) as detailed in the Supplementary Materials.

### Effect of antifibrotics and senolytic treatments

#### *In vivo*

Twenty-one days after instillation of IMR90 or SEN-IMR90 in 6–8 weeks old male nude athymic (*nu/nu*) mice, treatment with navitoclax (50 mg/kg, oral gavage, 14 days), nintedanib (60 mg/kg, oral gavage, twice per day, 14 days; Boehringer Ingelheim), or pirfenidone (400 mg/kg, oral gavage, twice per day, 14 days; Roche), or vehicle was started, and mice were euthanized thereafter. The degree of pulmonary fibrosis was determined quantitatively by the concentration of lung hydroxyproline.

#### *In vitro*

To determine the induction of apoptosis after treatment with navitoclax, pirfenidone or nintedanib in control and senescent MEFs, we used CellTiter-Glo® Luminescent Cell Viability Assay (Promega, Madison, WI, USA) or

CellTiter-Blue® Cell Viability Reagent (Promega) as detailed in the Supplementary Materials.

### Statistical analysis

Results are presented as mean  $\pm$  SD. Groups were compared using non-parametric Mann-Whitney *U* test, unpaired Student's *t* tests or one-way ANOVA with Bonferroni post hoc tests, as appropriate, using Prism 9 software (GraphPad Prism Software, San Diego, CA, USA). A *p* value < 0.05 was considered significant in all cases.

### AUTHOR CONTRIBUTIONS

F.H.-G., F.P. and M.S. performed and designed all the experiments; N.P. and M.A., and D.M. collaborated with histological and immunohistochemical studies and analyses of mice lungs. V.R. performed additional analysis of drugs senolytic activity; J.A.L., K.M. and M.I.M.M. helped with experiments; A.A., J.S. collaborated with M.S., F.H.-G., and F.P. in coordinating the work, designing the experiments, and interpreting the results; F.H.-G., F.P., A.A., and M.S. wrote the manuscript.

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### CONFLICTS OF INTEREST

M.S. is a shareholder of Senolytic Therapeutics, Life Biosciences, Rejuveron Senescence Therapeutics and Altos Labs, and is an advisor of Rejuveron Senescence Therapeutics and Altos Labs. The funders had no role in the study design, data collection and analysis, decision to publish, or manuscript preparation. F.H.-G. and J.S. report honoraria for lectures, presentations, or educational events, and support for attending meetings and/or travel received from Roche and Boehringer Ingelheim (disclosures made outside the submitted work). F.P., N.P., V.R., J.A.L.-D., K.M., M.A., M.I.M.M., D.M., A.A., R.F. do not have conflicts of interest to declare in relation to this study.

### ETHICAL STATEMENT

All mouse procedures were performed in compliance with guidelines established by the Barcelona Science Park's

Committee on Animal Care and under the corresponding approved ethics protocol (CEEA-PCB 10884).

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## SUPPLEMENTARY MATERIALS

### Supplementary Materials and Methods

#### *In vivo* humanized mouse model

##### *Induction of senescence in cultured human lung fibroblasts*

Mouse embryonic fibroblasts (MEFs) were isolated from a 13.5-day C57BL/6J mouse embryo. Cells were grown in Dulbecco's Modified Eagle Medium (DMEM) (Gibco®) supplemented with 10% fetal bovine serum and 100 U/mL penicillin/streptomycin, hereinafter referred to as DM10 media, and maintained at 37°C, 5% CO<sub>2</sub>. Passaging of cells was performed by enzymatic detachment using 0.05% Trypsin-EDTA (Gibco®) on cells for 5 minutes followed by inactivation in DM10 media and centrifugation at 180xg for 5 min. Supernatant was aspirated to remove dead cells and debris and pellet were resuspended in fresh DM10 media. For *in vivo* experiments, cells were washed and resuspended in PBS before inoculation.

For senescence induction, proliferating IMR90 cells were exposed to ionizing  $\gamma$ -radiation (20 Gy) when they reached 50% confluence. Senescence was induced in MEFs by treatment with 100  $\mu$ M Bleomycin sulphate (Sigma Aldrich, A10152) for 7 days. For the assessment of optimal senescence induction, senescence-associated  $\beta$ -galactosidase (SABG) activity was monitored using a protocol adapted from Dimri et al. [1]. Briefly, cells were fixed (5 mM EGTA, 2 mM MgCl<sub>2</sub>, 0.2% (w/v) glutaraldehyde in 0.1M phosphate buffer (pH 7.3)) for 10 min and washed twice with PBS. Staining was performed by incubating cells with an X-gal staining solution (40 mM citric acid, 5 mM potassium hexacyanoferrate (II), 5 mM potassium hexacyanoferrate (III), 150 mM sodium chloride and 2 mM magnesium chloride in 0.1M phosphate buffer (pH 7.3) adjusted to pH 6.0) for 14–18 hrs at 37°C in a non-CO<sub>2</sub> regulated incubator. Images were acquired by brightfield microscopy using an inverted microscope (Olympus CKX41) equipped with a digital camera (Olympus DP20).

##### *Instillation of IMR90 fibroblasts into mouse lung*

Normal proliferating IMR90 (IMR90) or  $\gamma$ -irradiated senescent human fibroblasts IMR90 (SEN-IMR90) ( $5 \times 10^5$  cells each group) were delivered to the lung of six-to-eight weeks old male nude athymic (*nu/nu*) mice (Envigo Laboratory). At the time points indicated in each experiment, mice were euthanized by cervical dislocation and lungs were removed for further analysis. All mouse procedures were performed in compliance with guidelines established by the Barcelona Science Park's Committee on Animal Care and under the

corresponding approved ethics protocol (CEEA-PCB 10884).

##### *Instillation of IMR90-derived conditioned medium into mouse lung*

Normal proliferating IMR90 or SEN-IMR90 were cultured in DM10 medium under standard conditions. Twenty-four hours before medium collection, cells were incubated in serum-free DM10 medium. Conditioned medium (CM) was collected and filtered (0.45  $\mu$ m) prior to buffer exchange procedure (PD MidiTrap™ G-10 column; GE Life Sciences). CM was then lyophilized and stored at –80°C. CM from IMR90 and SEN-IMR90, normalized by the number of cells, was resuspended in sterile PBS on the same day of the experiment. CM derived from IMR90 or SEN-IMR90 corresponding to  $5 \times 10^5$  cells was delivered intratracheally to six-to-eight weeks old male immunodeficient mice (Envigo Laboratory). PBS was used as negative control. Three weeks after intratracheal instillation, lungs were removed and analyzed.

##### *Senolytic and antifibrotic treatments in vivo*

NS-IMR90 or SEN-IMR90 were instilled in the lung of six-to-eight weeks old male nude athymic (*nu/nu*) mice as described above. Twenty-one days after instillation, mice were treated with navitoclax (100 mg/kg, oral gavage, 14 days; ABT-263, Selleckchem, ref.S1001), nintedanib (50 mg/kg, oral gavage, twice per day, 14 days; Ofev®, Boehringer Ingelheim), or pirfenidone (400 mg/kg, oral gavage, twice per day, 14 days; Esbriet®, Roche), or vehicle. At the end of the treatment, mice were sacrificed.

##### *Histologic assessment of mouse pulmonary fibrosis*

Left lung tissue was fixed in 10% neutral buffered formalin solution for 24 hours, transferred into tissue cassettes and washed with PBS for at least 24 hours. Tissues were then sent to the Histopathology Facility of the Institute for Research in Biomedicine (IRB, Barcelona) for standard histological procedures. 4–5  $\mu$ m tissue sections were obtained and stained with Hematoxylin Eosin and Masson's Trichrome (AR17311-2, Dako-Agilent) for the histological analysis. Samples were first examined in blind, and then in an unblinded fashion for confirmation, by a pathologist. Semiquantitative histological scoring of fibrosis in Masson's Trichrome stained sections was determined at 20–40 $\times$  using the following scale: 1,  $\times$ 1; 2,  $\times$ 2; 3,  $\times$ 3 increase the thickening of alveolar walls; 4,  $>$  $\times$ 3 thickening of alveolar walls and focal areas of single fibrotic masses. In the case of ambiguous scoring, the intervening number was given.

Immunohistochemistry for anti-Human p21<sup>WAF1/Cip1</sup> (M7202, Agilent), for anti-Mouse p21 clone HUGO 291H/B5 (CNIO) and for anti-Human Nucleoli [HuNu] (Merck, MAB4383) was performed using a Ventana discovery XT at 1:50–1:250, ready to use (RTU) and 1:300, respectively, for 60 min, for anti-phospho H2A Histone Family Member X [gH2AX] (9718, Cellsignaling) using a Leica BOND RX at 1:750 for 120 min and manually for anti-Human Nucleoli [HuNu, NM95] (ab190710, Abcam) at 1:100 and incubated overnight at 4°C. Antibodies were diluted with EnVision FLEX Antibody Diluent (K800621, Dako-Agilent). Antigen retrieval for p21<sup>WAF1/Cip1</sup> and p21 clone HUGO 291H/B5 was performed with Cell Conditioning 1 (CC1) buffer (6414575001, Roche) and with Ultra Cell Conditioning 2 (CC2) buffer (5279798001, Roche) for HuNu MAB4383. The recombinant Anti-IgG1 + IgG2a + IgG3 antibody [M204-3] (ab133469, Abcam) at 1:500 for 32min was used to enhance specific labelling of p21<sup>WAF1/Cip1</sup> and HuNu MAB4383. Secondary antibodies used was OmniMap™ anti-Rb HRP (760-4311, Roche) or the OmniMap anti-Rat HRP (760-4457, Roche). Blocking was done with Casein (ref: 760-219, Roche). Antigen-antibody complexes were revealed with ChromoMap DAB Kit (760-159, Roche). Antigen retrieval for gH2AX was performed with BOND Epitope Retrieval 2 – ER2 (AR9640, Leica) buffer for 20 min and with citrate buffer pH6 at 121°C for 20 min for anti-HuNu NM95. Quenching of endogenous peroxidase was performed by 10 min of incubation with Peroxidase-Blocking Solution (S2023, Dako-Agilent). Unspecific unions were blocked using 5% of goat normal serum (16210064, Life technology) with 2.5% BSA (10735078001, Sigma) for 60min. Blocking of unspecific endogenous mouse Ig staining was also performed using Mouse on mouse (M.O.M) Immunodetection Kit – (BMK-2202, Vector Laboratories) incubated 60 min at room temperature. Secondary antibody was a polyclonal Goat Anti-Mouse 1:100 (P0447, Dako, Agilent) incubated for 30 min or the BrightVision poly HRP-Anti-Rabbit IgG, incubated for 45 min (DPVR-110HRP, ImmunoLogic). Antigen-antibody complexes were revealed with 3-3'-diaminobenzidine (K3468, Dako) or with the DAB (Polymer) (RE7230-CE, Leica), with the same time exposure per antibody. Sections were counterstained with Hematoxylin (Dako, S202084) and mounted with Mounting Medium, Toluene-Free (CS705, Dako) using

a Dako CoverStainer. Specificity of staining was confirmed by the mouse IgG1, kappa monoclonal [MOPC-21] (ab18443, Abcam), the rabbit IgG (NBP2-24891, Novus biotech) or the rat IgG (6-001-F, R&D Systems) isotype controls.

### Image acquisition

Brightfield images were acquired with a NanoZoomer-2.0 HT C9600 digital scanner (Hamamatsu Photonics, France) equipped with a 20× objective. All images were visualized with a gamma correction set at 1.8 in the image control panel of the NDP view 2 U12388-01 software (Hamamatsu, Photonics, France).

### Hydroxyproline assay

Right lung lobes were surgically dissected, weighed, placed into 1.5-mL sterile tubes, and snap-frozen. Frozen lung samples were grinded using a liquid-nitrogen filled mortar and pestle. On the day of the assay, thawed and resuspended in 1 mL of distilled water. Tissues were homogenized using a microsample homogenizer (Precellys). 200 µL of 12N hydrogen chloride was added to 200 µL of homogenized tissues. Samples were placed into a preheated oven set to 120°C and incubated overnight. Biochemical quantification of hydroxyproline was performed using a hydroxyproline assay kit (Amsbio®).

### RNA extraction and quantitative Real Time-PCR

Total RNA from lung tissue was extracted by mechanical disruption in 1000 µL of TRIzol® reagent (Invitrogen) using a microsample homogenizer (Precellys) according to manufacturer's instructions. RNA concentration was determined using NanoDrop® ND-1000 UV-Vis spectrophotometer at 260nm wavelength. For extraction of total RNA from cells, 6-well plates were scraped in 1000 µL of TRIzol® reagent (Invitrogen). cDNA was synthesized using the SuperScript™ III Reverse Transcriptase (Thermo Fisher, Waltham, MA, USA). mRNA expression analysis was performed using real-time quantitative PCR (RT-qPCR) (PowerUp™ SYBR® Green Master Mix, Applied Biosystems, Foster City, CA, USA) run on a CFX96™ Real-Time PCR Detection system (Bio-Rad). Relative gene expression was determined using  $\Delta\Delta C_t$  method by measuring RT-qPCR signal relative to signal of housekeeping gene (*Actb*). The RT-qPCR primers used in this study are listed below:

Target	Species	Forward primer	Reverse primer
Actin b	Mouse	ATGGAGGGGAATACAGCCC	TTCTTTGCAGCTCCTTCGTT
Gapdh	Mouse	AACTTTGGCATTGTGGAAGG	ACACATTGGGGGTAGGAACA
Col6a3	Mouse	ACTGGAACCACGGAAGTTCA	GTCACCTCCAACATCGAGGC
Col1a2	Mouse	AGGTCTTCCTGGAGCTGATG	ACCCACAGGGCCTTCTTTAC
Cdkn1a	Mouse	GTGGGTCTGACTCCAGCCC	CCTTCTCGTGAGACGCTTAC

IL-6	Mouse	ACTCACCTCTTCAGAACGA	CCATCTTTGGAAGGTTTCAG
Tgf- $\beta$	Mouse	CGGAGAGCCCTGGATACCA	ACTTCCAACCCAGGTCCTTC
Pai-1	Mouse	CCAACATCTTGGATGCTGAA	GCCAGGGTTGCACTAAACAT
Cdkn2a	Mouse	TACCCCGATTACAGGTGAT	TTGAGCAGAAGAGCTGCTACGT
$\alpha$ -SMA	Mouse	CCCAAAGCTAACCGGGAGAAG	CCAGAATCCAACACGATGCC
CDKN1A	Human	TGTCCGTCAGAACCCATGC	AAAGTCGAAGTTCCATCGC

## ***In vitro* experiments**

### ***Cell culture-conditioned medium experiment***

IMR90 cells were cultured in DM10 medium, at 37°C and under hypoxic conditions (5% CO<sub>2</sub>). For senescence induction, cells were irradiated with 10 Gy and cultured 14 more days. Control cells were cultured in parallel with 3–4 $\times$  sub-passaging steps in between. After 14 days, medium was removed and exchanged to 25 ml fresh culture medium for senescent and for control cells. CM was collected 24 hrs later and used for the medium transfer experiments. MEFs were generated as described previously. Three to four different MEF clones are representing different biological replicates. MEFs were cultured in DM10 medium, at 37°C and under hypoxic conditions (5% CO<sub>2</sub>). MEFs were seeded reaching 30–40% confluency prior the first medium transfer for paracrine senescence induction. Next day CM from IMR90 control or IMR90 senescent cells was sterile filtered with a 0.2 $\mu$ m filter and transferred to the MEFs. Two days later, a second CM transfer was conducted the same way. Cells were grown for another 3 days and then harvested for the according readout, which is gene expression analysis (RT-PCR), SABG staining and immunofluorescence staining.

### ***Immunofluorescence staining in vitro***

MEFs grown in a black 96well plate were fixed with 4% PFA in PBS for 10 min at RT and followed by 2 PBS washing steps. Then, the cells were permeabilized by 0.2% Triton incubation for 5 min at RT followed by 2 PBS washing steps. Anti-smooth muscle actin antibody and (A5228, Sigma) anti-p21 (ab188224, Abcam) were added and cells were transferred to 37°C shaking platform for 30 min. After 2 PBS washing steps MEFs were incubated with the according secondary antibodies (Thermo Fisher, Waltham, MA, USA) and DAPI for another 30 min, shaking at 37°C. Cells were then washed twice with PBS before adding 50% glycerol and

storage at 4°C until microscopy was performed. Microscopy was performed within 1 week post staining using an LSM680 ZEISS confocal microscope (10x objective) taking 2–3 independent field acquisitions per 96 well of each condition. Positive cells were quantified using ImageJ software and represented in percentage.

### ***Senolytic assay in vitro***

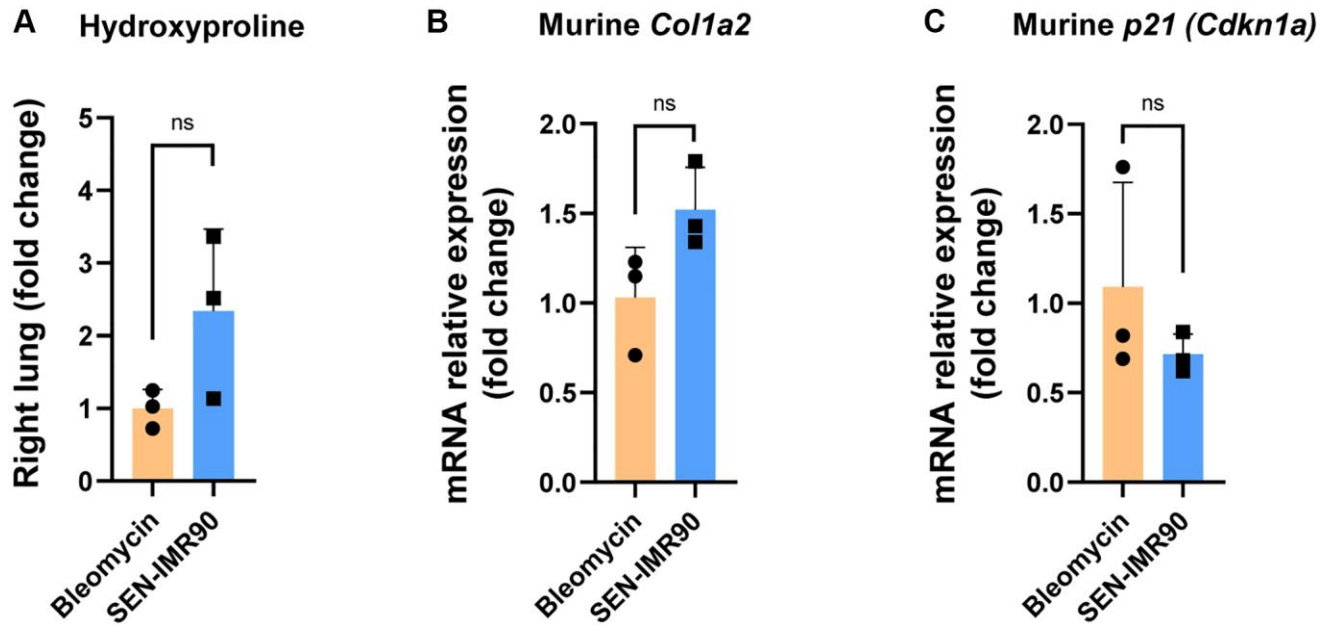
For senescence induction, MEFs cells were treated with 100 $\mu$ M bleomycin sulfate (Sigma Aldrich, A10152) for 7 days. To determine the induction of apoptosis after the treatment with navitoclax, pirfenidone and nintedanib, control and senescent cells were seeded in flat-bottom-clear 96-well plates at a density of 6,000–8,000 and 4,000–6,000 cells per well, respectively, at 37°C in a 5% CO<sub>2</sub> humidified atmosphere. The following day cells were treated with serial dilutions of navitoclax, pirfenidone or nintedanib in 0.2% FBS-containing media. Vehicle treatments were used as control for the entire duration of the assay. Viability was assessed 48–72 hs after treatment upon by means of CellTiter-Glo<sup>®</sup> Luminescent Cell Viability Assay (Promega, Madison, WI, USA) or CellTiter-Blue<sup>®</sup> Cell Viability Reagent (Promega). Raw data were acquired by measuring luminescence in a VICTOR Multilabel Plate Reader (Pelkin Elmer) or fluorescence at an excitation/emission wavelength of 560 nm/590 nm in an Infinite 200 PRO Multimode Spectrophotometer (TECAN).

## **Supplementary Reference**

1. Dimri GP, Lee X, Basile G, Acosta M, Scott G, Roskelley C, Medrano EE, Linskens M, Rubelj I, and Pereira-Smith O. A biomarker that identifies senescent human cells in culture and in aging skin in vivo. *Proc Natl Acad Sci U S A*. 1995; 92:9363–67. <https://doi.org/10.1073/pnas.92.20.9363> PMID:[7568133](https://pubmed.ncbi.nlm.nih.gov/7568133/)



## Supplementary Figure



**Supplementary Figure 1. Lung fibrosis induced by senescent human lung fibroblasts recapitulates other features of the bleomycin-induced lung fibrosis model.** (A) Lung fibrosis was ascertained by hydroxyproline assay, and (B) mRNA expression of *Col1a2* and (C) *Cdkn1a/p21<sup>Cip1/Waf1</sup>* in lungs of SEN-IMR90 injected mice compared to bleomycin treated mice, 2 months after treatment. For each experimental group,  $n = 3$ . The group labelled SEN-IMR90 is the same group labelled SEN-IMR90 (2 months) in Figure 3B, 3C, 3D and the absolute data are the same. Statistical significance was assessed by two-tailed Student's *t*-test. Abbreviation: NS: not significant. For further explanations, see text.