

# The effects of BMMSC treatment on lung degeneration in elderly macaques

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## Research

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# Abstract

**Background:** Age-related degeneration of lung tissues increases the risk of lung injury and exacerbates lung disease. It is also the main risk factor for chronic lung diseases (such as COPD, idiopathic pulmonary fibrosis, cancer, etc.).

**Methods:** we performed systematic screening, evaluation of elderly macaque model. A senile multiple organ dysfunction model was used to explore whether BMMSCS could improve degeneration of lung tissues in an elderly macaque model.

**Results:** Using model evaluation tests, we found that the average alveolar area, Mean linear intercept (MLI) and fibrosis area in the elderly macaque models were significantly larger than in young rhesus monkeys ( $P < 0.05$ ), and the capillary density around the alveoli was significantly lower than in young macaque models ( $P < 0.05$ ). Intravenous infusion of BMMSCS reduced the degree of pulmonary fibrosis in elderly macaque, increased the density of capillaries around the alveoli ( $P < 0.05$ ), and the number of type II alveolar epithelium in elderly macaque ( $P < 0.05$ ). BMMSCS infusion reduced lung tissue ROS level, systemic and lung tissue inflammation level and Treg cell ratio in elderly macaque model ( $P < 0.05$ ). Indirect co-cultivation revealed that BMMSCS reduced the expression of senescence-related genes, ROS levels, apoptosis rate of aging type II alveolar epithelial cells (A549 cells) and promoted their proliferation ( $P < 0.05$ ).

**Conclusions:** BMMSC treatment can improve age-related degeneration of macaque lung tissue

## Introduction

The aging population in China is one of the main problems currently being faced. By 2050, it is expected that the percentage of the population aged 65 years or above will represent about 20% of the population<sup>[1]</sup>. With age, the respiratory system undergoes various structural changes including a gradual increase in aging-related deterioration in lung tissue, alveolar enlargement, alveolar wall destruction, reduced gas exchange surface area, increased airway obstruction or occlusion, decreased pulmonary vascular density, increased collagen deposition and decreased elastin, etc<sup>[2-4]</sup>. Immune system disorders, including non-specific inflammation and suppressed immune responses, etc<sup>[5]</sup> functional changes, such as loss of elastic recoil, increased residual volume and barriers to gas exchange<sup>[2, 6]</sup> increases the susceptibility of the lungs to injury among the elderly and development of chronic lung diseases such as COPD, idiopathic pulmonary fibrosis and cancer<sup>[7]</sup>. Therefore, improving and delaying aging-related lung degenerative changes should be investigated.

Mesenchymal stem cells are multipotent stem cells characterized by low immunogenicity, self-renewal ability, and multi-directional differentiation potential. Because of the significance of immune regulation in anti-apoptosis, angiogenesis, migration, differentiation to target organs, support of the growth and

differentiation of local stem cells and progenitor cells, anti-scarring, and chemical gravitational have been used in the treatment of various diseases<sup>[8]</sup>.

Numerous studies have suggested that mesenchymal stem cells can repair lung injury and effectively treat acute and/or chronic lung diseases<sup>[9-11]</sup>. However, the majority of these studies have been performed on rodents, and very few studies on primates exist. Moreover, the efficacy of mesenchymal stem cells has not been extensively studied in age-related lung tissue degeneration.

Macaque is one of the most popular non-human primates, known to possess many biological characteristics similar to humans. As a model organism, the macaque is physiologically similar to humans, their genomes have a 93% average sequence identity with humans, hence serve as an ideal model to studies of human health and disease.

Besides, macaques as a model animal, have numerous advantages including controllability of environmental factors, ease of scale, and it is widely used in basic and applied research in biomedicine<sup>[12, 13]</sup>.

In our previous studies, we reported the correlation between aging and bone marrow mesenchymal stem cells in macaque as well as its anti-aging ability in 293T cells *in vitro*<sup>[14]</sup>. Therefore, in this study, macaque was used as the experimental animal to determine the effects of bone marrow mesenchymal stem cell(BMMSCS) on age-related degeneration both *in vivo* and *in vitro*.

## Materials And Methods

### Animal and cell sources

The macaques used in this experiment were obtained from the Kunming Institute of Zoology of Chinese Academy of Sciences [SCXK (Yunnan) K2017-0003]. In total, we used 30 animals, aged between 2-26 years old and weighing 2.2-12 kg. Among them, 25 female macaques, aged 6-26 years old. 5 male macaques, aged 2-3 years old. The animals were housed in the Experimental Animal Center of the 920th Hospital of the Chinese People's Liberation Army Joint Logistics Support Force, experimental animal license number: SYXK (Military) 2017-0051, the research program was passed by the experimental animal council of the 920th Hospital of Joint Logistics Support Force.

The bone marrow used in our experiments was self-made; A549 cells were purchased from the Wuhan Sevier company with identification certificate, and were subcultured in a culture flask with DMEM medium containing 10% fetal bovine serum, and placed the culture flask in an incubator containing 5% carbon dioxide at 37°C in our laboratory.

### Main Reagents and antibodies

FBS and penicillin - streptomycin solution were purchased from Servicebio; DMEM / F12 media were purchased from Hyclone; 0.25% pancreatin-0.04% EDTA were purchased from Invitrogene; 30% hydrogen

peroxide solution was purchased from Solarbio; Cell senescence  $\beta$ -galactosidase staining kit, apoptosis detection kit, active oxygen detection kit, cell cycle and apoptosis detection kit were purchased from Beyotime Biotechnology; Anti-proSP-C antibody (AB3786) was purchased from Sigma; Mouse anti-human CD45(555555), mouse anti-human CD73(344007), Cell staining buffer, True-Nuclear Transcription Factor Buffer Set, Alexa Fluor® 647 anti-human FoxP3(320113), FITC anti-human CD4(317408), PE anti-human CD25 antibody(356103), PE Mouse IgG1  $\kappa$  Isotype Ctrl and Alexa Fluor® 647 Mouse IgG1  $\kappa$  Isotype Ctrl(400135) were purchased from Biolegend; Monkey interleukin 1 $\beta$  (IL-1 $\beta$ ) ELISA kit, monkey interleukin-17A (IL-17A) ELISA kit, human tumor necrosis factor alpha (TNF- $\alpha$ ) ELISA kit were purchased from MeiMian; Fluorescent secondary antibody HRP, fluorescent secondary antibody CY3 and primers were purchased from Servicebio; GoScript™ Reverse Transcription System, GoTaq® qPCR Master Mix were purchased from Promega Corporation; TNF alpha Antibody(7B8A11), IL-10 Antibody(20850-1-AP) and CEBPB Antibody(2B6E10) were purchased from Proteintech group; Adipogenic differentiation medium, osteogenic differentiation medium, and chondrogenic differentiation medium were purchased from Guangzhou Saiye Biological Technology Co., Ltd.; <sup>18</sup>F-FDG was provided by the First Affiliated Hospital of Kunming Medical University

## **Experimental protocols**

### **Screening and evaluation of senile lung degeneration macaque models**

Female macaque aged 22-26 years old were used as the elderly model group, while female young macaque aged 6-8 years old were used as the young control group, with 5 animals in each group. About 5ml of peripheral blood was collected from cynomolgus monkeys, and centrifuged to obtain serum. These sera samples were used for ELISA tests to quantify TNF- $\alpha$  and IL-1 $\beta$  levels. Next, the macaque were anesthetized with 3% pentobarbital sodium 5ml / kg, and sacrificed to obtain lung tissues. Some the tissue samples were used for size, morphological and texture analysis, while the remaining were subjected to Hematoxylin-eosin staining. The degree of pulmonary fibrosis in the two groups was determined by Masson's Trichrome stain. Capillary density was examined by immunohistochemistry.

### **Preparation and identification of macaque bone marrow mesenchymal stem cells**

Two 2-3 years old macaques randomly were selected and anesthetized with intramuscular injection of 3% pentobarbital sodium at 1 ml / kg. They were placed on the operating table in a supine position, and bone marrow aspiration was performed above and behind the anterior superior iliac spine. This was done using a 20 ml syringe containing 5 ml of heparin sodium saline (100 U / ml). Briefly, 5 ml of bone marrow were drawn. Thereafter, red blood cells were removed by addition of 0.38% ammonium chloride to the bone marrow samples, followed by centrifugation to prepare single cells. The cells were resuspended in DMEM / F12 medium containing 20% fetal bovine serum and seeded in a 175 cm<sup>2</sup> cell flask. The medium was changed 5 days later, and at intervals of 3 days thereafter. When the adherent cells reached became 80% confluent, they were passaged. The P4 generation of cells was used to perform morphological examination and characterization of growth patterns, expression of surface antigens CD29, CD45, CD73,

CD90 and CD184 was determined by flow cytometry. In order to confirm their differentiation ability in vitro. Meanwhile, the P4 generation of cells are transferred into different special induction mediums to induce them to differentiate into osteocytes, chondrocytes, fat cells. The proliferation ability of cells was determined by CCK-8 assay.

### **Cell processing and injection**

When the fusion degree of the cultured the P4 generation BMMSCS reaches over 80%, the cells are digested and washed, and the cells were diluted with 0.9% sterile sodium at a concentration of  $2 \times 10^6$  cells/ml by cytometer. After the macaques were fixed, the BMMSCSs were infused in the femoral vein at a cell dose of  $1 \times 10^7$  cells/kg per macaque, once every other day, for a total of 3 infusions. On the other hand, the macaques in the control group and the model group received equal volumes of 0.9% sterile sodium at the same time.

### **Observation of the histological structure of lung tissues of macaque after BMMSCS infusion**

Changes in lungs were imaged with PET-CT before BMMSCS treatment and at 90 days and 180 days after treatment. Each macaque was fasted for more than 6 hours, and anesthetized with 3% sodium pentobarbital at a dose of 1 ml/kg, and then blood glucose was measured. After completing the above operations, <sup>18</sup>F-FDG was injected through the brachial vein at a dose of 3.70~4.44 MBq/kg, and the scan was performed after 60 minutes of rest. Use GE Discovery™ PET/CT Elite to scan the whole body. CT adopts conventional whole-body spiral scanning, tube voltage 120 kV, tube current 240 mA, pitch 0.561, rotation speed 0.5 s/week, layer thickness 3.75 mm, interval 3.75 mm, interval moment 512X512. After collecting image data, import the image data into the PET/CT AW4.6 post-processing workstation to calculate SUV value and Hounsfield unit.

At 180 days after BMMSCS treatment, macaque were killed by anesthesia, lung tissue was collected, observed and photographed. The right lung was excised, fixed with 4% paraformaldehyde solution, embedded in paraffin and sectioned. The tissue sections were used for Hematoxylin-eosin staining, We took the left peripheral lung tissues of 5 animals to make slices respectively, each slice was 0.5cm apart, and a total of 5 slices were cut for Hematoxylin-eosin staining. Then each slice randomly selected 3 fields of 400× view to observed lung tissue structure (i.e., the change of general lung anatomy, alveolar size, inflammation, alveolar septum thickness, pigmentation). Let the entire sliced tissue fill the entire field of view. Use Image-Pro Plus 6.0 software to calculate the area of each field of view (mm<sup>2</sup>) and the area of Hematoxylin-eosin staining positive area (lung parenchyma) (mm<sup>2</sup>) using the 400-fold scale as the standard. The average alveolar area (mm<sup>2</sup>) = (field of view area mm<sup>2</sup> - Area of lung parenchyma mm<sup>2</sup>) / number of alveoli. In addition, draw a cross line at the center of each field of view of each slice and extend to the edge, then calculate the number of alveolar intervals passed by the cross line, measure the total length of the cross line (mm), and calculate the average lining interval (Mm) = total length of crosshair [mm] / number of alveolar intervals.

The tissue sections were also used for Masson's Trichrome stain. The sectioning method is the same as Hematoxylin-eosin staining. Observe under the microscope, at least two 100x fields of view are randomly selected for each slice to take pictures. When taking pictures, try to fill the entire field of view with tissue, and the background light of each photo should be as consistent as possible. Use Image-Pro Plus 6.0 software to select the measurement area by adjusting the threshold, then measure the area of blue collagen fibers. Finally calculate the area percentage of collagen fibers.

The tissue sections were also used for Immunohistochemistry. The sectioning method is the same as Hematoxylin-eosin staining. After capturing the image, import the image to Densito Quant in Quant Center. Then set the dark brown, brown, light brown, and blue nuclei of the slices to be strong positive, moderate positive, weak positive, and negative respectively. The next step is to identify and calculate the area of strong positive, moderate positive, weak positive and negative, and the percentage of positive. Finally, the number of positive cells in each slice and hematoxylin-eosin staining intensity are converted into numerical values for H-score scoring.

### **Analysis of the effect of BMMSCS on type II alveolar epithelial cells**

A549 cells were placed in a complete medium containing 200 $\mu$ mol / L, 400 $\mu$ mol / L, 600 $\mu$ mol / L, 800 $\mu$ mol / L, 1000 $\mu$ mol / L, 1200 $\mu$ mol / L hydrogen peroxide. PCR assay was performed on the cells to determine expression of P53 gene while  $\beta$ -galactosidase staining was carried out to determine the optimal concentration of hydrogen peroxide. The expression of *TERT*, *TCAB1*, *P53*, *P21* was quantified by RT-PCR. The Experimental procedures for RT-PCR follow the programs of GoScript™ Reverse Transcription System and GoTaq® qPCR Master Mix from Promega Corporation. The expression of these genes reflect the level of senescence of A549 cells at the specific hydrogen peroxide concentration. Once a senescence model of type II alveolar epithelial cell was established, the senescent cells were seeded in the lower chamber of a transwell with a pore size of 0.4  $\mu$ m, and an equal proportion of BMMSCS cells were seeded in the upper chamber of the transwell chamber. After 48 hours of co-cultivation, the expression of P53, P21, TCAB1 of A549 cells was determined using RT-PCR, method as above.

Meanwhile the rate of apoptosis of A549 cells was determined by flow cytometry according to the Annexin V Alexa Fluor488/PI manual of 4ABIO. ROS levels and cell cycle progression were compared between the model and treatment group according to the Reactive Oxygen Species Assay Kit manual of beyotime. In addition, Immunohistochemistry was performed to detect proSPC as markers of type II alveolar epithelial cells. After staining, 3 fields were randomly selected, and each field was counted 200 cells, and then the percentage of type II alveolar epithelial cells to the total number of cells was calculated.

### **Analysis of the effect of BMMSCS treatment on ROS, inflammatory factors and VEGF in elderly macaques**

Serum was prepared from the peripheral blood of macaque obtained at 0, 30, 60, and 90 days after BMMSCS treatment. The level of inflammatory factors IL-1 $\beta$ , IL-17A, and TNF- $\alpha$  in the peripheral blood

was detected by ELISA.

After BMMSCS treatment, left lung tissues were used for ROS staining. After finishing ROS staining, put sections under the same laser intensity for the same time exposure and capture images. Then use the Densito Quant in the Quant Center to set dark red, brown red, light red, and blue nuclei as strong positive, moderate positive, weak positive and negative respectively and perform the H-score scoring. The protein level of proinflammatory factors IL-6, TNF- $\alpha$ , IL-1 $\beta$  and anti-inflammatory factor IL-10 in the lung tissue was detected by Western blot. Expression of VEGF in lung tissue after BMMSCS treatment was also determined by Western blot. Then use ImageJ to analyze the gray values of all western blot images obtained, and compare the gray value of the internal control band with the gray value of the target protein band. Analysis of the effect of BMMSCS treatment on the ratio of peripheral blood Treg cells and FOXP4 in lung tissues of elderly macaque

Lymphocytes were isolated from blood samples obtained from the animals at 0, 30, 60, and 90 days after BMMSCS treatment. Changes in Treg cell ratio in peripheral blood were detected by flow cytometry; Treg cells were labeled with FOXP4, and the changes in FOXP4 content were analyzed by immunohistochemistry.

### **Statistical analysis**

Statistical analyses were performed using SPSS 21.0 statistical software. Measurement data are expressed as mean  $\pm$  standard deviation ( $\bar{X} \pm s$ ). The means of three or more than three groups were analyzed by one-way ANOVA (One-Way ANOVA).

## **Results**

### **Differences in appearance and lung tissue structure between the young and elderly macaques**

Elderly macaques were found to have a dull coat that turned white, especially around the head and face. Besides, their skin was loose and dry, and the face red (Fig. 1).

The lung tissue of the young control group and the elderly model group were soft, butterfly-shaped, flexible, and pale red. However, when compared with the young control group, the lung size of the elderly model macaque was significantly larger than that of the young control (Fig. 2).

The young control group showed clear lung structures, thin and smooth alveolar walls, no thickening of the alveolar space, no exudate, and only a small amount of inflammatory cellular infiltrate around the blood vessels. In the elderly model group, although the alveolar wall thickness was uniform and the alveoli clean, there was no exudate, the alveolar cavity became irregularly enlarged to form the pulmonary bullae and there was visible pigmentation (as indicated by the black arrow). The elderly model group showed severe inflammation compared with the young control group. The average area and MLI were significantly increased in the elderly model group compared to the young control group (Fig 3), ( $P < 0.0001$ ).

Masson's Trichrome stain colored the collagen fibers blue. The collagen area in the lungs of the elderly model group was significantly increased, compared with the young control group (Fig .4),(P <0.01).

Immunohistochemical staining to label the vascular endothelial cells with CD31 revealed that the lung tissue cells nuclear were stained blue, and the surface markers of vascular endothelial cells CD31 were stained brown. In the elderly group, the expression of CD31 in the lungs was significantly reduced compared with the young control group (Fig .5), (P=0.0001).

### Cultivation and identification of BMMSCS

Macaque BMMSCSs were isolated from bone marrow aspirates and cultured by adherent culture screening. A few fusiform adherent cells were observed under an inverted phase-contrast microscope after 3-4 days. The cell fusion rate had reached about 80% after 9 days. BMMSCSs of passage 3 to 5 showed uniform morphology, they grew densely spiral and were isolated (Fig .6A).

To confirm the purity of the cultured cells, the immunophenotypes of the P4 generation of juvenile macaque BMMSCS were analyzed by flow cytometry. A panel of surface antigens was analyzed. The results showed that the BMMSCSs were positive for CD29, CD45, CD73, CD90 and CD184 at percentage rates of  $96.35 \pm 0.62$ ,  $0.16 \pm 0.12$ ,  $95.22 \pm 0.37$ ,  $96.25 \pm 1.71$ ,  $93.53 \pm 2.76$ , respectively (Fig .6B,Table 1).

**Table 1 Flow cytometry analysis of surface antigens**

Surface antigen	n	Cell positive rate
CD29	3	96.35±0.62
CD45	3	00.16±0.12
CD73	3	95.22±0.37
CD90	3	96.25±1.71
CD184	3	93.53±2.76

n is the number of repeated experiments

The proliferation assay showed that the BMMSCSs took on an "S" shape, the cells remained latent for the first 1-2 days, and entered a logarithmic proliferation phase on 3 to 7 days, where the cells grew vigorously and had the best vitality. On the 8th day, they entered the plateau phase which was characterized by a reduction in proliferation (Fig .6C).

The P4 generation of young macaque BMMSCS was used to determine the differentiation ability and proliferation *in vitro*. The duration of the differentiation experiment was 14-21 days. The cells were cultured in osteogenic induction medium and allowed to aggregate, form nodules, and accumulate calcium deposits. Alizarin red stain was used to detect the precipitated calcium deposits which were an indication of differentiation. Intracellular lipid droplets were stained with oil red O, and red-stained lipid

droplets were found in the cells. Proteoglycans were stained with Alcian blue and appeared as smears (Fig .6D).

### The Changes in lung tissue structure after BMMSCS treatment

PET-CT examination was performed on the macaques, before treatment, at 90 and 180 days after treatment. The results showed that the texture of the lungs of the macaque before treatment was grid-like, ground-glass opacity, honeycomb-shaped, with the peripheral, subpleural, and lower lung lobes as the main features and emphysema was obvious. The average Hounsfield unit was significantly decreased before treatment compared with the control group ( $P < 0.01$ ). Before treatment, HRCT showed irregular thickening of the leaflet intervals, and the small blood vessels in the leaflet became obvious due to the thickening of the wall in the treatment group. PET showed that the  $^{18}\text{F}$ -FDG uptake quantified as the glucose uptake in the lungs decreased after treatment. At 90 and 180 days, following treatment, CT showed that the lungs' texture was clear, and both hilar were normal. The Hounsfield unit was higher than before treatment. The average Hounsfield units were  $(685 \pm 12.53)$  and  $(705 \pm 18.53)$ , respectively (Table 2, Fig .7).

**Table 2 Changes in PET-CT in elderly macaque lung after BMMSCS treatment**

	n	Hounsfield Unit	SUV max
Control	5	$672 \pm 12.52$	$0.4 \pm 0.09$
Prior treatment	5	$853 \pm 25.32$ \$	$0.7 \pm 0.06$ \$
90 days after treatment	5	$685 \pm 12.53$ *	$0.5 \pm 0.08$ *
180 days after treatment	5	$705 \pm 18.53$ *	$0.3 \pm 0.07$ **

n is for the number of animals analyzed, \$  $P < 0.05$  when compared with the control group, \*  $P < 0.05$  when compared with prior treatment, \*\*  $P < 0.01$  when compared with the prior treatment

Examination of material from the lung tissue after 180 days of treatment, showed that the lung tissue appeared dark white and red without embolism. However, there were no significant changes observed in both the treatment group compared with the model group (Fig .8).

HE results showed that the control group had clear lung structures, thin and smooth alveolar walls, no thickening of the alveolar spaces, no exudates, and only a small amount of inflammatory cellular infiltration around the blood vessels. The alveolar cavity showed irregular enlargement and the formation of pulmonary bullae the alveoli were clean with no exudates seen, a small amount of inflammatory cell infiltration was observed around the blood vessels with pigmentation in the model group macaques and the treatment group macaques. Although the inflammation score was not statistically different in those groups, the treatment group had lower inflammation than the model group. The average alveolar area and alveolar lining interval (MLI) of the model group and the treatment group were both significantly

increased compared with the control group, ( $P < 0.05$ ); however, there was no significant difference between the model group and the treatment group (Fig .9), ( $P > 0.05$ ).

Masson's Trichrome stain showed blue collagen. The collagen area of the treatment group was significantly reduced ( $P < 0.05$ ) when compared with the model group (Fig .10).

To determine the changes in capillary density around the alveoli after cell transplantation, immunohistochemistry was performed using CD31 as a marker of vascular endothelial cells. The nucleus stained blue and capillaries with CD31 surface markers were stained brown. The content of CD31 around the alveoli was significantly increased in the treatment group, compared with the model group (Fig .11), ( $P < 0.0001$ ).

### **Effect of BMSCS on senile type 2 alveolar epithelial cells**

Type 2 alveolar epithelium plays a significant role in lung aging. In the elderly, the quantity and quality of type 2 alveolar epithelial cells are significantly reduced [15]. In this study, the effect of BMSCS on lung structure was observed using type 2 alveolar epithelial cells to explore the specific effect on lung cells. Hydrogen peroxide was used to establish an aging model of A549 cells. Different concentrations of hydrogen peroxide were found to induce different degrees of aging in A549 cells. Following SA- $\beta$ -gal staining, we found that when the hydrogen peroxide concentration was 600 $\mu$ m/L and 800 $\mu$ m/L, the senescence rate of A549 cells was the highest. In addition, 1000 $\mu$ mol / L and 1200 $\mu$ mol / L were not considered as most of the cells were apoptotic and deformed (Fig 12).

RT-PCR was used to detect the expression of P53 gene in A549 cells after induction of aging. The increase in P53 expression was most significant ( $P < 0.001$ ,  $P < 0.0001$ ) at 600 $\mu$ mol / L, 800 $\mu$ mol / L hydrogen peroxide concentration (Fig 13A). Therefore, 600 $\mu$ mol/L was chosen as the optimal concentration to induce senescence of A549 cells.

RT-PCR was used to further compare the expression of P53, P21, TERT, TCAB1 before and after induction of aging using hydrogen peroxide at a concentration of 600  $\mu$ mol / L. At 6h, after induction, the changes in P53 and P21 expression were significantly increased ( $P < 0.0001$ ,  $P < 0.001$ ); however, TERT and TCAB1 were significantly decreased ( $P < 0.001$ ,  $P < 0.01$ ) 6 hours after induction (Fig .13B). At 24h, 48h, and 72h after changing the medium, TCAB1 did not show any significant change ( $P > 0.05$ ), while P21 was significantly increased ( $P < 0.0001$ ). Even though the expression of P53 was significantly decreased after induction ( $P < 0.05$ ), it remained higher than before induction (Fig .13C).

Following the indirect co-culture of A549 cells aging model with BMSCS for 48H, the lower layer of A549 cell was collected to explore the effect of BMSCS on the A549 aging model by RT-PCR. The expression levels of P53 and P21 were found to be significantly decreased ( $P < 0.001$ ,  $P < 0.01$ ) in the treatment group compared with the model group. However, the expression level of TCAB1 increased significantly ( $P < 0.05$ ) (Fig .14A).

The ROS level, apoptosis ratio, and cell cycle of A549 cells were detected by flow cytometry after indirect co-culture. The ROS level and apoptosis ratio of the treatment group were found to be significantly reduced compared with the model group (Fig .14B, Fig .14C), ( $P < 0.0001$ ,  $P < 0.001$ ). Proliferation in the treatment group accelerated to the G2 phase (Fig .14D), ( $P < 0.01$ ).

In vitro experiments, revealed the effects of BMSCS on the aging of the A549 cell model. To verify these effects in vivo, proSPC was used as a marker for type  $\alpha$  alveolar epithelial cells. The results showed that type  $\alpha$  alveolar epithelial cells were round or oval and scattered in the alveolar wall. The number of type  $\alpha$  alveolar epithelial cells in the model group was significantly reduced compared with the control group ( $P < 0.001$ ). However, in the treatment group, type  $\alpha$  alveolar epithelial cells were significantly increased compared with the model group (Fig .15), ( $P < 0.01$ ).

### **Changes in VEGF expression level in lung tissue**

There were observed changes in the density of capillaries around the alveoli, hence there was a need to check on the level of VEGF in the lungs. Western blot analysis of the lung tissue showed that the VEGF in the model group decreased significantly compared with the control group ( $P < 0.05$ ). Besides, after BMSCS treatment, the VEGF level in the treatment group was significantly higher compared with the model group (Fig .16) ( $P < 0.05$ ).

### **Changes in the level of ROS and inflammatory factors after BMSCS treatment**

Extensive experiments in a wide range of organisms from yeast to primates have revealed that the nine hallmarks of aging are stem cell failure, changes in intercellular communication, genomic instability and telomere wear, epigenetic changes, loss of protein homeostasis, nutrition changes, mitochondrial dysfunction and cellular senescence<sup>[16]</sup>. There are still many unresolved issues on the main causes and impacts of these events. However, emerging research suggests that the causes and commonalities of these events are related to the immune system. Inflammatory aging is characterized by elevated levels of immune cell infiltration and elevated levels of pro-inflammatory cytokines and chemokines in the tissue microenvironment and circulatory system<sup>[16]</sup>. Under normal physiological conditions, ROS in the cells is constantly generated and eliminated. Therefore, maintaining appropriate levels of ROS in the cells plays an important role in the stability of cell functions. However, in the state of aging, the level of ROS may also be elevated due to mitochondrial stress and damage and persistent inflammation<sup>[17]</sup>. High levels of ROS not only increase damage to the cells but also stimulates immune cells to produce more pro-inflammatory factors to form a vicious circle<sup>[18]</sup>. The immune regulation and damage repair functions of mesenchymal stem cells are very critical. Studies have reported that MSCs control inflammation and ROS production through paracrine and mitochondrial transfer between MSCs and aging cells<sup>[19,20]</sup>. Therefore, this study proposed that mesenchymal stem cells altered inflammation and ROS levels in elderly macaques thus affecting lung degeneration. The frozen section of lung tissue was used to detect the level of ROS. Using the inverted fluorescent microscope, the lung cells' nucleus was stained blue, and the

red fluorescence was distributed in the cytoplasm. Compared with the model group, the ROS level of the treatment group was significantly reduced (Fig .17),(P <0.01).

To explain the regulatory effect of BMMSCS on aging-related inflammation, the levels of IL-1 $\beta$ , IL-17A, and TNF- $\alpha$  were detected by ELISA. IL-1 $\beta$  was found to be significantly decreased in blood serum (P <0.05) compared with the model group at 30 and 60 days after BMMSCS treatment and returned to the original levels after 90 days. Besides, TNF- $\alpha$  was found to be significantly decreased (TNF- $\alpha$ ) after 30 days (P <0.05), and returned to original levels after 60 days, and remained unchanged. There was no significant change in IL-17A levels (Fig .18A).

The changes in the levels of inflammatory factors including IL-1 $\beta$ , IL-6, TNF- $\alpha$ , and IL-10 in the lung after BMMSCS treatment were determined by western blot analysis. The levels of IL-1 $\beta$ , IL-6, TNF- $\alpha$  in the treatment group were significantly lower than in the model group. The level of IL-10 in model group was significantly lower than in the control group (P <0.05), but was significantly increased after BMMSCS treatment (Fig .18B), (P <0.05).

### **Effect of BMMSCS treatment on immune regulatory cells**

As described above, there were observed changes in the expression levels of inflammatory factors in peripheral blood and lung tissue after BMMSCS treatment. Treg cells are a class of cells with immune-regulatory functions and also play a vital role in the regulation of inflammation. The ratio of Treg cells in the peripheral blood was measured by flow cytometry. The results showed that the Treg ratio in macaque peripheral blood decreased significantly at 30 days after BMMSCS treatment (P <0.01), and reached its minimum at 60 days after treatment (P <0.0001) compared with the model group. However, there was no significant change in the ratio of Treg cells at 60 and 90 days (Fig .19), (P> 0.05).

To determine whether the changes in Treg cells in the periphery and lung tissue were consistent, the Treg cell surface marker FOXP4 was used and detection was performed by immunohistochemistry. The content of FOXP4 in the lung tissue of the model group was found to be significantly higher compared with the control group (P < 0.01). Besides, the content of FOXP4 in the treatment group was significantly lower than that in the model group (P <0.0001), and the control group (Fig .20).

## **Discussion**

Changes in lung tissue structure associated with aging, are characterized by enlarged alveoli, damaged alveolar walls, decreased gas exchange surface area, increased airway obstruction or occlusion, decreased pulmonary vascular density, deepened fibrosis, and decreased elastin content [2]. In this study, we observed these phenomena through pathological analysis.

As a common type of stem cells, MSC can be affected by chemotaxis and accumulate at the injured site, and secrete some cytokines (such as KGF, HGF) and some RNA-rich microvesicles. This paracrine effect can delay cell senescence and apoptosis, repair lung tissue and resist fibrosis. Meanwhile, mitochondrial

transfer and communication between MSC and neighboring cells can also play a role in the repair of such damage. Therefore, the lung, as the first organ through which MSC pass in the body, may benefit more<sup>[21-23]</sup>.

Therefore, in this study, we focused on observing the positive effect of BMMSCS on lung tissue degeneration. The changes in the lungs were detected by PET-CT, which revealed a significant increase in x-ray transmittance (Hounsfield unit) in lung tissue after BMMSCS treatment. However, a significant decrease in SUV max values and fibrosis were observed. Masson's Trichrome stain revealed that collagen deposition in the treatment group was significantly reduced compared to the model group. Immunohistochemistry analysis showed that alveolar vascular density in the treatment group was significantly increased compared to the model group. However, Hematoxylin-eosin staining showed no significant reduction in alveolar size, which was inconsistent with the results of PET-CT and results reported in other previous studies on mesenchymal stem cells in the treatment of COPD<sup>[24-26]</sup>. Breathing is a dynamic process, and changes in the tissues around the alveoli determine the lung's inspiratory function and Alveolar size. Due to changes in the air in the lungs during inspiration and expiration, PET-CT or Hematoxylin-eosin staining may not sufficiently evaluate whether BMMSCS can repair the deterioration of alveolar size during actual respiration in elderly lung tissue. Therefore, to assess the beneficial effects of BMMSCS, observation should be performed from multiple directions.

SA- $\beta$ -gal is a hydrolase enzyme that catalyzes the hydrolysis of  $\beta$ -galactosides into monosaccharides only in senescent cells. P53 and its downstream gene P21 and TCAB1, have been associated with the control of cellular aging<sup>[14, 16]</sup>.

In this study, type  $\square$  alveolar epithelium cells (A549 cells) were used, and aging was induced by hydrogen peroxide. Following co-culture with BMMSCS,  $\beta$ -galactoside staining, RT-PCR, and flow cytometry were used to determine the changes in  $\beta$ -galactosidase, P53, P21, and TCAB1 expression levels as well as the oxidative stress levels, apoptosis, and proliferation in A549 cells. The results confirmed that BMMSCS can reverse the aging-related characteristics of type  $\square$  alveolar epithelial cells (A549 cells). These results were consistent with those reported in our previous study on the effect of bone marrow mesenchymal stem cells on the 293T cell senescence model<sup>[14]</sup>. Besides, we used proSPC to label type  $\square$  alveolar epithelial cells, and determine the number by immunofluorescence. The results showed that the number of type  $\square$  alveolar epithelial cells in the treatment group was significantly higher than that in the model group. To some extent, these results were consistent with the *in vitro* experiments.

High inflammation and oxidative stress are important factors driving degeneration of tissues and organs during aging<sup>[7, 16, 27]</sup>. In this study, we found high level of age-related inflammation and oxidative stress in elderly macaque. Besides, the expression of inflammatory factors TNF- $\alpha$ , IL-1 $\beta$ , and IL-17A in the peripheral blood was analyzed after 30 $\square$ 60 and 90 days after BMMSCS treatment. After 180 days histological and ROS level analysis was performed and the results showed that the serum IL-1 $\beta$  level was significantly reduced 30 days after BMMSCS treatment, but the level returned to normal at 60 and 90 days. However, western blot analysis of the lung tissue, showed that IL-1 $\beta$  remained relatively lower than

in the model group after 6 months of treatment. Furthermore, TNF- $\alpha$  expression levels appeared to be inconsistent in the lungs and peripheral blood. The content of TNF- $\alpha$  in the peripheral blood decreased significantly after 30 days of cell transfusion and returned to the levels before treatment at 60 and 90 days. However, after 6 months, western blot analysis of the lung tissue, found that TNF- $\alpha$  expression levels were significantly lower than in the model group. This inconsistency between the circulation and lung tissue expression levels differed from that reported in previous studies on rodent lung injury models [28, 29]. Besides, our results show that IL-6 was downregulated in the lung tissue compared to the model group after BMSCS treatment. Therefore, mesenchymal stem cells, such as BMSCS, have a regenerative effect in the aging lung tissue of macaque.

Treg cells play an important role in the control of the immune response. Specifically, they play a central role in immune homeostasis and in preventing autoimmunity. They are produced by the thymus and lymph and exported to the entire body, where they inhibit the activation and proliferation of potential self-reactive T cells, thereby regulating the body's immunity and inflammation. Some studies have found that the number and function of Treg cells change significantly in the aging body. Zhao et al reported that the ratio of Treg cells in the CD4 + cells of peripheral blood in elderly mice, was significantly increased and the function was significantly reduced [30]. IL-10 is a multi-functional cytokine, which regulates the growth and differentiation of cells, participates in inflammatory reactions and immune responses, and is recognized as an immunomodulatory cytokine used by almost all innate immune cells [31]. Current studies have shown that the number and function of Treg cells have a regulatory effect on CD4 + and CD25- cells to activate the release of IL-10. Aging affects the ability of CD4 (+), CD25 (+), FOXP4 (+) T cells to regulate the production of IL-10 [32]. Besides, mesenchymal stem cells, are powerful immune regulatory cells, with not only a regulatory effect on Treg cells but also strongly influence IL-10 production. Some studies have reported that mesenchymal stem cells can increase the production of IL-10 by communicating with macrophages, B lymphocytes, and dendritic cells or by secreting PGE2, IDO, IL-6, and HO-1 [33]. In this study, CD4, CD25, and FOXP4 were used as markers to detect the changes in the ratio of Treg cells in peripheral blood within 90 days after mesenchymal stem cell treatment. Treg cells in peripheral blood were found to continuously decrease from 0 to 60 days after BMSCS treatment, and the low levels were maintained from 60 to 90 days. These results were inconsistent with those reported in previous studies on the regulation of Treg cells by mesenchymal stem cells under extremely high inflammatory conditions [33]. Therefore, it was not clear whether BMSCS regulated the entry of Treg cells into the tissues, hence reducing the Treg ratio in peripheral blood. Therefore, FOXP4 was used as a marker to further explore the level of Treg cells in the lung tissue. However, the findings from the lung tissues were consistent with those in the peripheral blood, but the FOXP4 levels in the lung tissue of the treatment group were significantly reduced. Because there are no studies showing that MSCs play a major role in immune regulation in healthy individuals, and their effects have only been demonstrated in cell-based therapy and disease scenarios. Therefore, regarding this phenomenon that we found, it may need to be further studied.

The content of IL-10 in the lung tissue was detected by western blot analysis. The results showed that after 6 months of BMMSCS treatment, the content of IL-10 in the lung tissue of macaque was significantly higher than that of the model group. These findings were consistent with the short-term observation in a previous mesenchymal stem cell treatment model of lung injury<sup>[34]</sup>. However, the decrease in the ratio of Treg cells was inconsistent with the phenomenon of elevated IL-10, and this shows that BMMSCS treatment can promote the secretion of IL-10 through other mechanisms not related to Treg.

In summary, this study reveals for the first time that BMMSCS delays aging-related lung degeneration in the elderly macaque. This study had limitations. Due to experimental limitations, the impact of BMMSCS on lung function was not explored in this study. Besides, the mechanisms by which BMMSCS improved lung degeneration was also not explored.

## Conclusions

1. Obtaining elderly model of pulmonary degenerative macaque shows that the alveolar cavity is enlarged, the structure of the lung is disordered, the pigmentation is increased, the degree of fibrosis is increased, and the capillary density is decreased.
2. BMMSCS can reduce the degree of pulmonary fibrosis in elderly macaques, reduce the level of inflammation in the lung and peripheral blood, increase the expression of VEGF in lung tissue, increase the density of capillaries around the alveoli, and reduce the content of Treg cells in peripheral blood and lung tissue.
3. BMMSCS can reduce the expression of type  $\alpha$  alveolar epithelial aging-related genes, reduce its apoptosis and oxidative stress levels, and promote proliferation. BMMSCS can increase the number of type  $\alpha$  alveolar epithelium in lung tissue.

## Declarations

- **Ethics approval and consent to participate**

Experimental protocols were approved by the Experimental Animal Ethics Committee of the 920th Hospital of the PLA Joint Logistics Support Force.

- **Consent for publication**

Not applicable.

- **Availability of data and material**

All data generated or analysed during this study are included in this published article.

- **Competing interests**

The authors declare that they have no competing interests.

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### **Authors' contributions**

YKY,XQZ,YL and YYW made substantial contributions to study conception and design, data acquisition, or data analysis and interpretation.

YKY and RGP agree to be accountable for all aspects of the work and ensure that questions related to the accuracy or integrity of any part of the work will be appropriately investigated and resolved.

XHP and XQZ have given final approval of this version of the manuscript for publication.

XQZ, XHP, HYH, YKY and RQP have been involved in drafting the manuscript or revising it critically for important intellectual content.

All authors read and approved the final manuscript.

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## **Abbreviations**

Abbreviations	Full name
BMMSCs	Bone marrow mesenchymal stem cells
FOXP3	Recombinant Forkhead Box P3
IL	Interleukin
P53	Tumor Protein 53
P21	Tumor Protein 21
TERT	telomerase reverse transcriptase
MSCs	Mesenchymalstemcells
CCK-8	Cellcountingkit-8
<sup>18</sup> F-FDG	β-2-[ 18 F]-Fluoro-2-deoxy-D-glucose

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## Figures

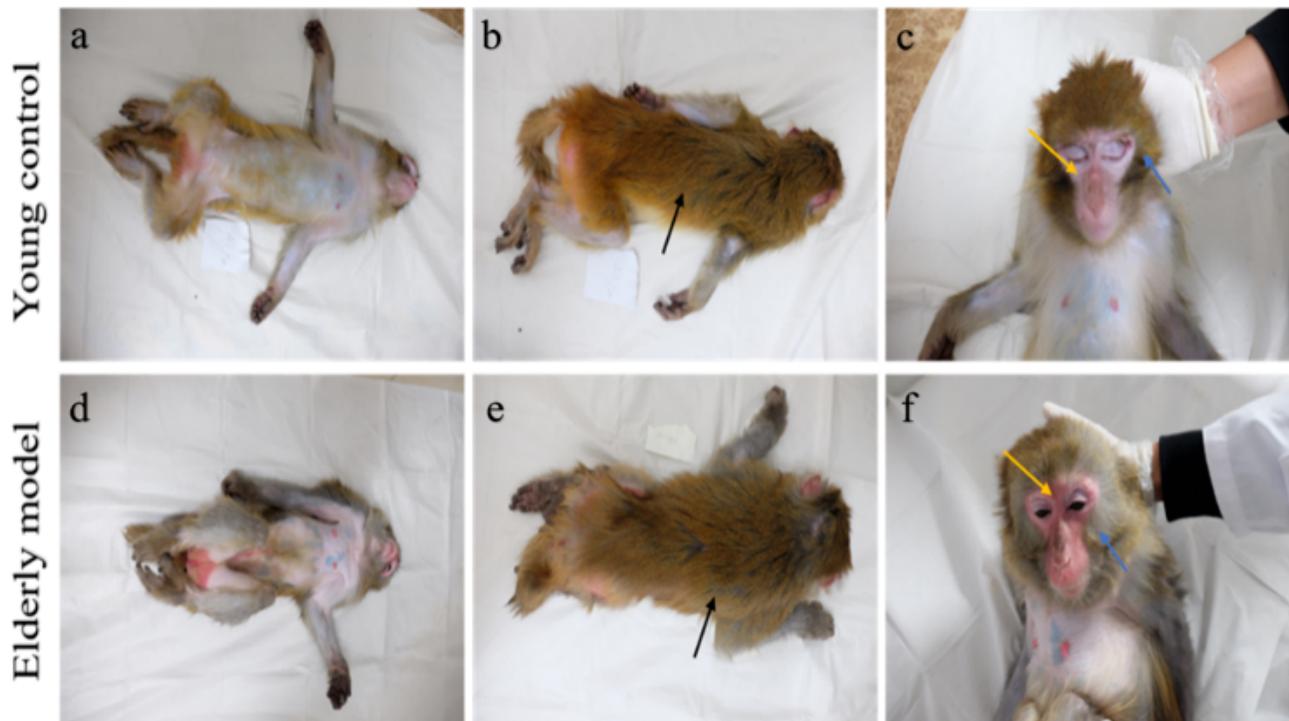
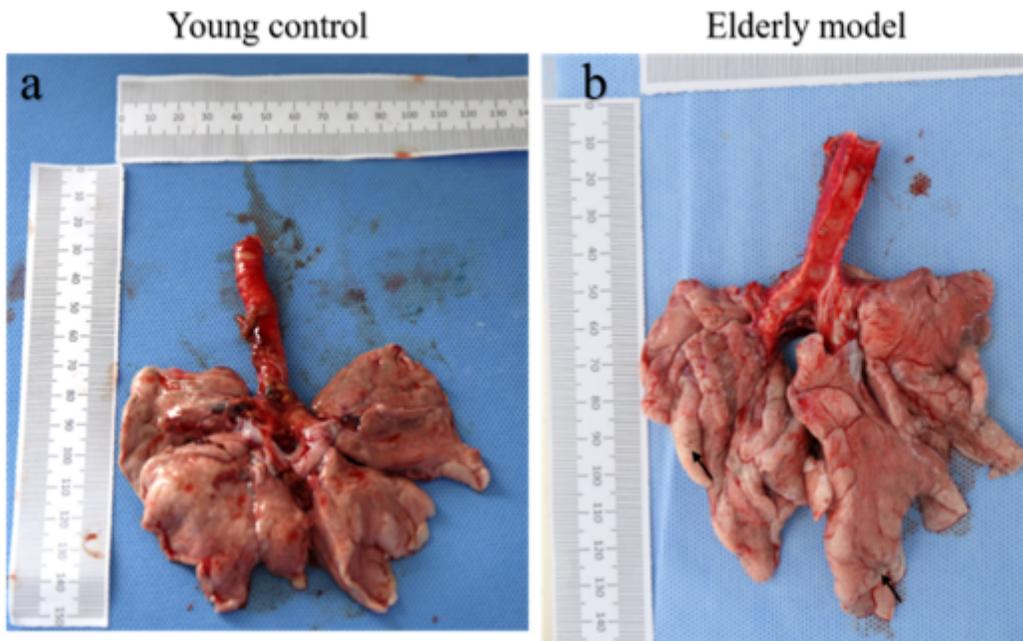


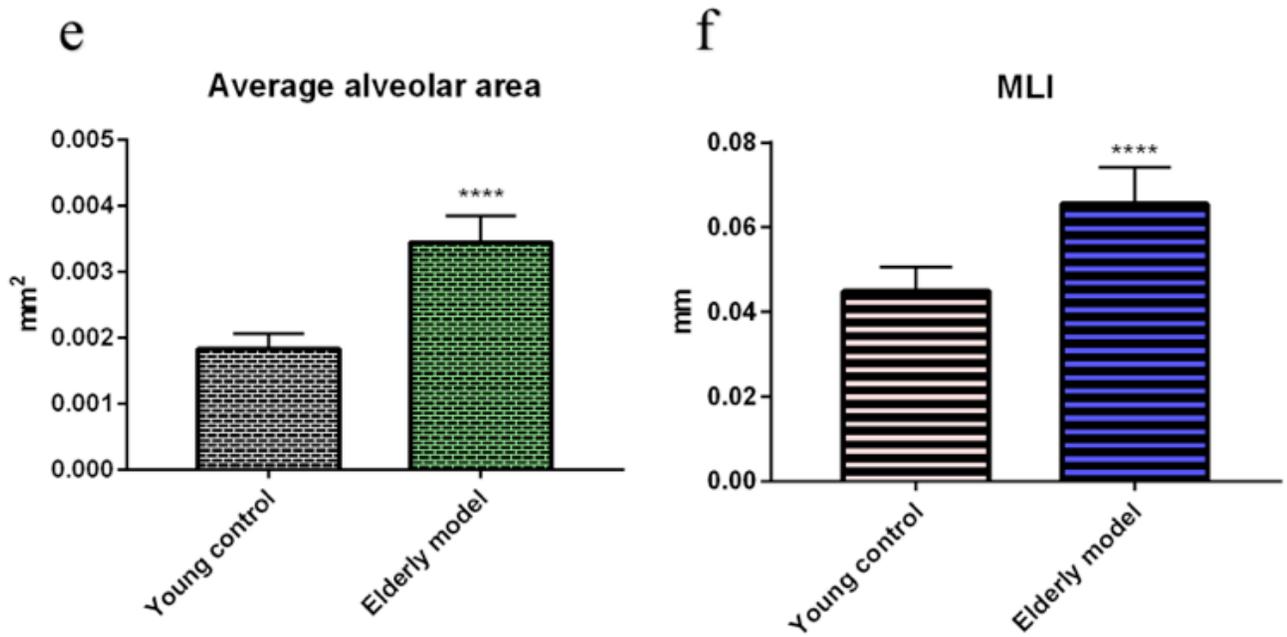
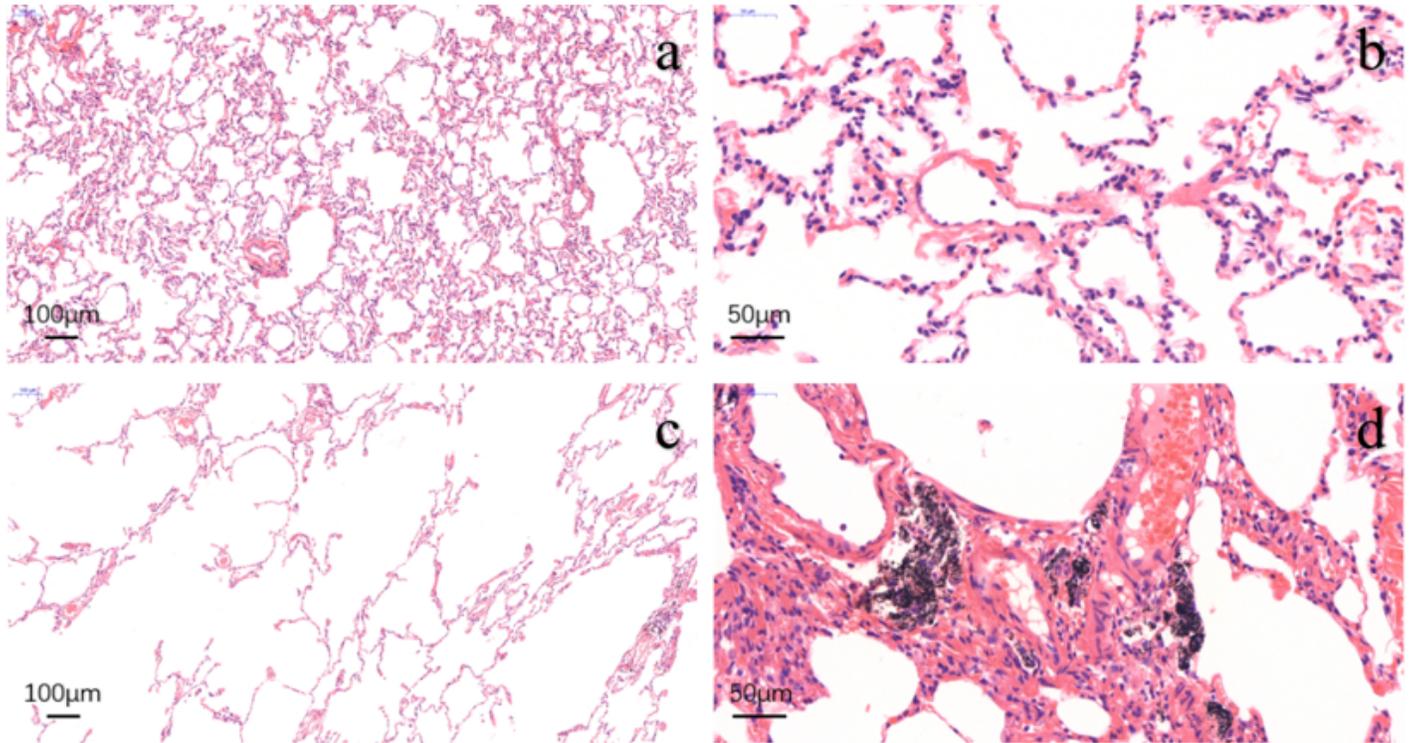
Figure 1

age-related change in appearance between the young control group and the elderly model group (a, b, c are the front, back and face of the macaques in the elderly model group respectively, d, e, f are the front, back and face of the macaque in young control respectively)



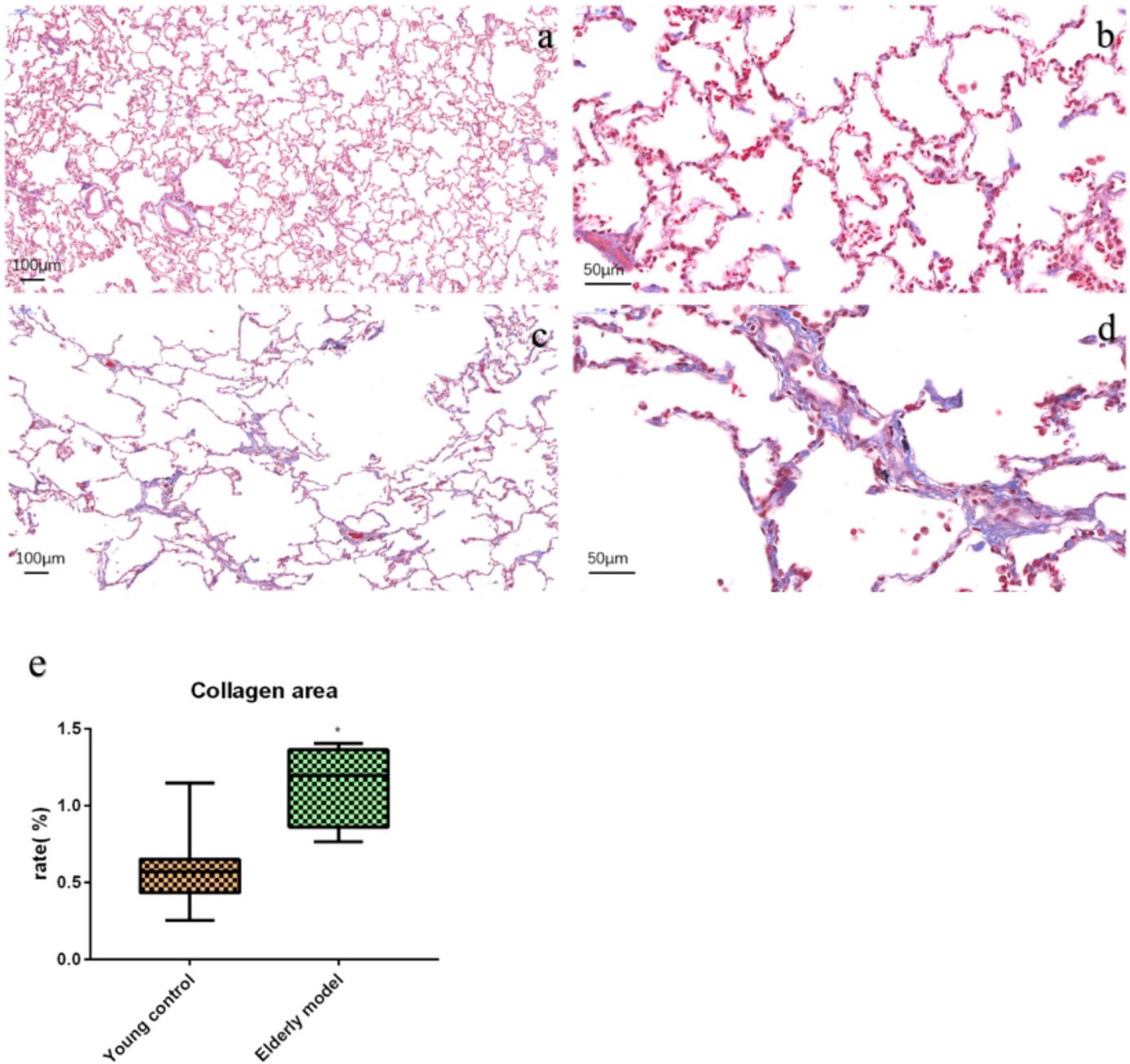
**Figure 2**

Comparison of lung size between the young control group and the elderly model group.(a is the isolated lung of the macaques in young control group, b is the isolated lung tissue of the elderly group of macaque.)



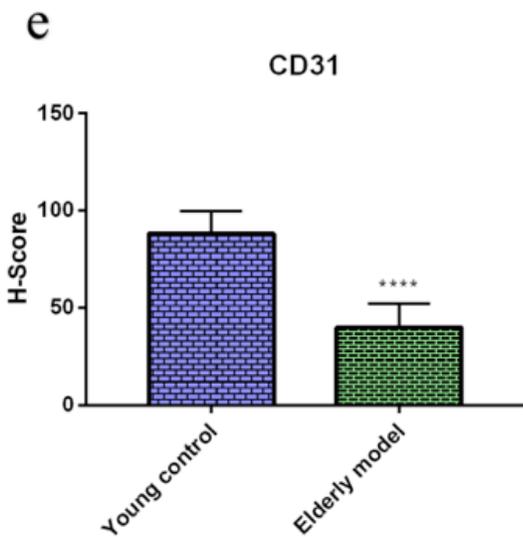
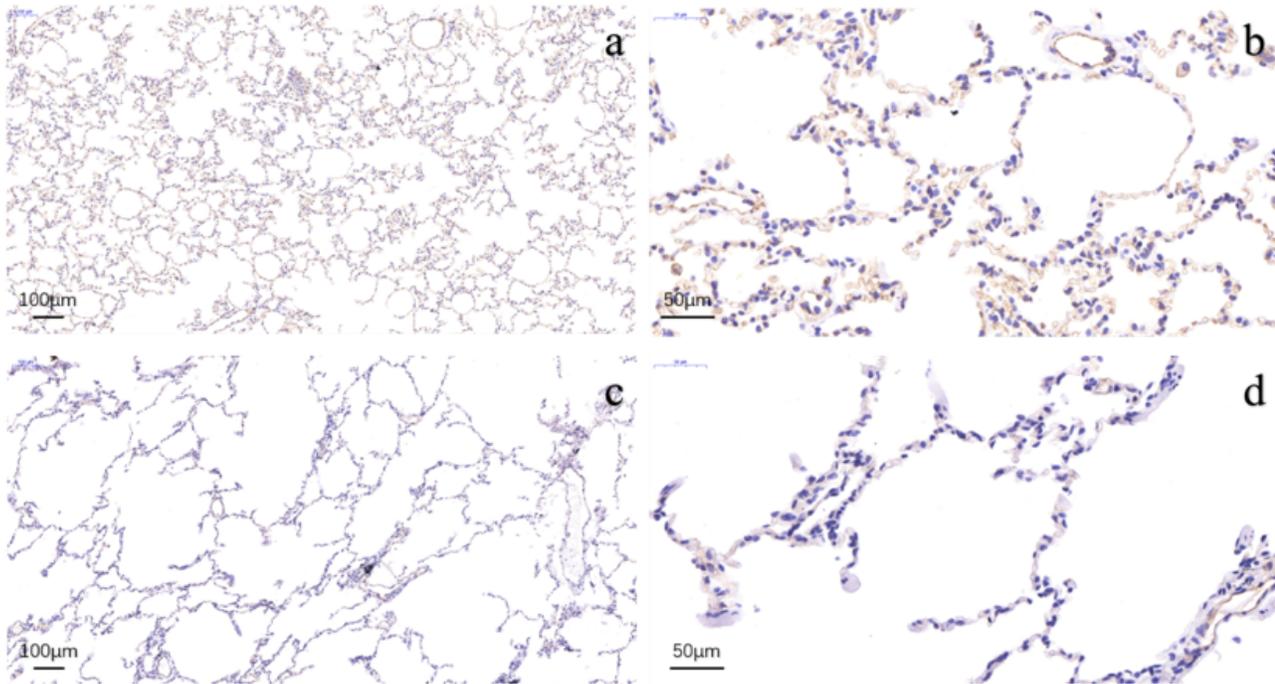
**Figure 3**

Comparison of the lung tissue structure between the young control group and the elderly model group (a, b are HE stained pictures of lung tissue of macaques in the young control group. c, d are HE stained pictures of lung tissue of macaques in the elderly model group. e is the statistical graph of the average alveolar area. f is statistical of the average lining interval. n=5, n is for the number of animals analyzed, \*\*\*\* P < 0.0001 when compared with the young control group)



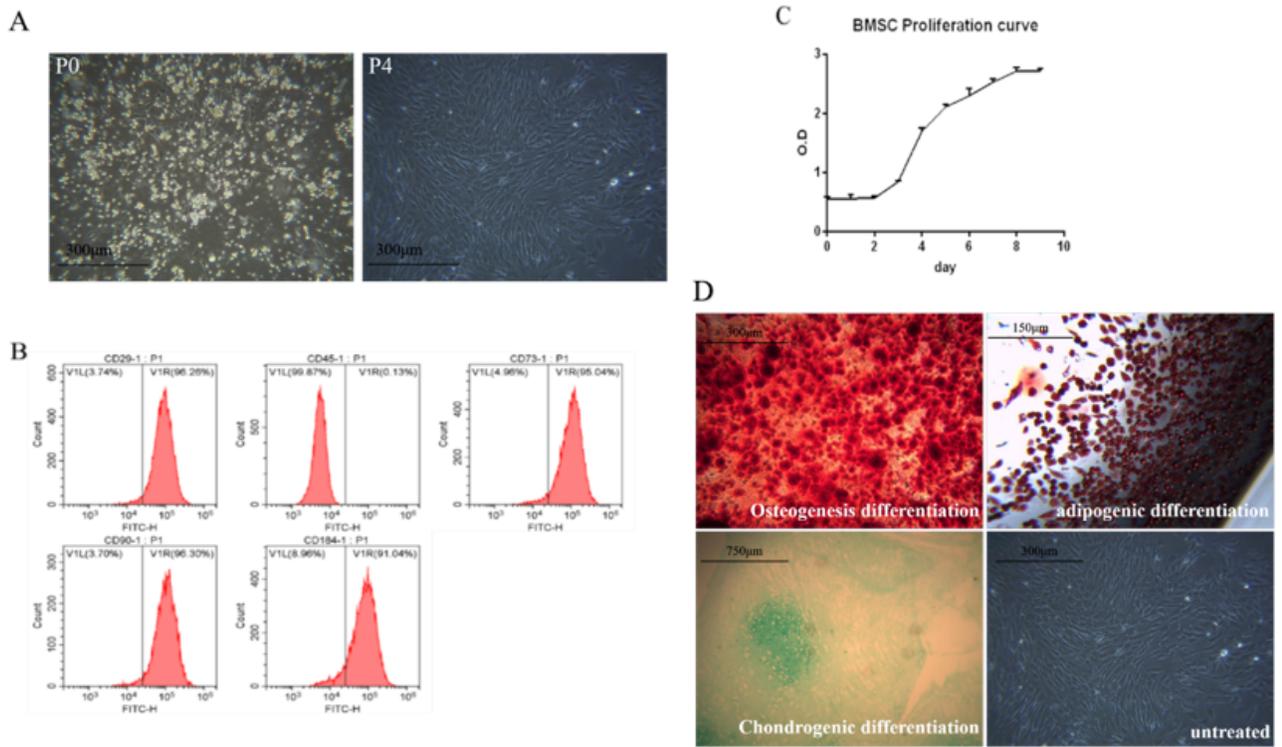
**Figure 4**

Comparison of the collagen area between the young control group and the elderly model group ((a, b are Masson's Trichrome stained pictures of lung tissue of macaques in the young control group. c, d are Masson's Trichrome stained pictures of lung tissue of macaques in the elderly model group. e is the rate of collagen area statistics graph. n=5, n is for the number of animals analyzed, \*\* P < 0.01 when compared with the young control group)



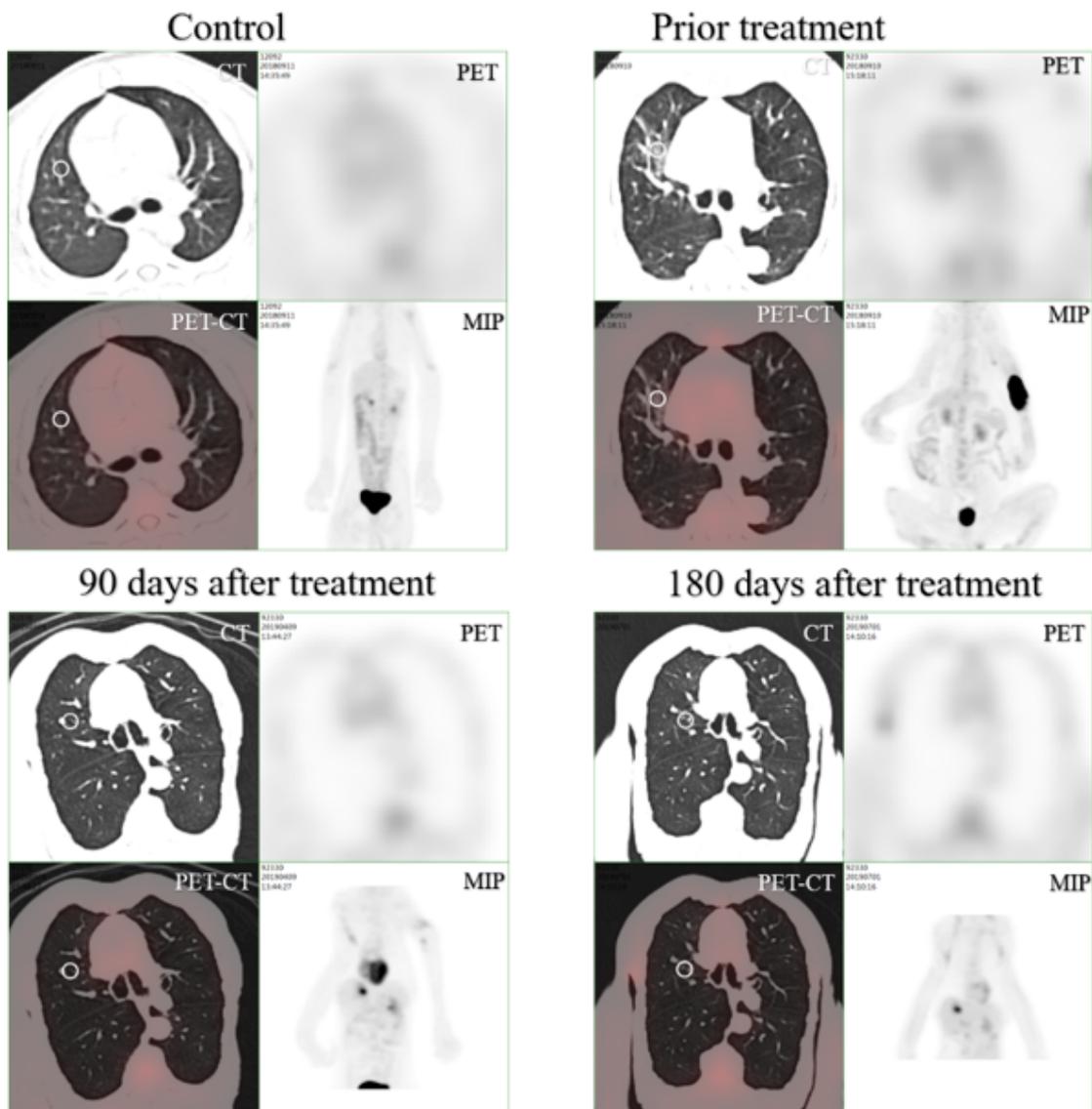
**Figure 5**

Comparison of CD31 expression around the alveoli between the young control group and the elderly model group((a, b are immunohistochemistry pictures of lung tissue of macaques in the young control group. d, e are immunohistochemistry pictures of lung tissue of macaques in the elderly model group. e is the H-Score statistical graph of the above picture. n=5, n is for the number of animals analyzed, \*\*\*\* P <0.0001 when compared with the young control group)



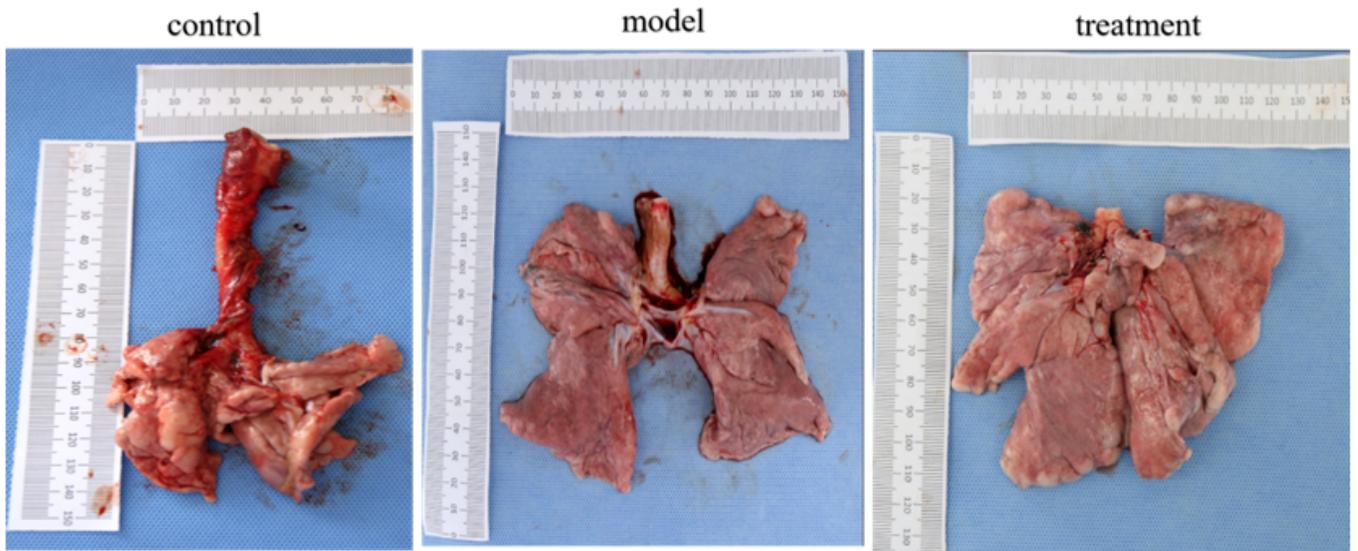
**Figure 6**

Cultivation and identification of BMMSCS a. BMMSCS form b. BMMSCS flow analysis c. BMMSCS proliferation curve d. BMMSCS differentiation



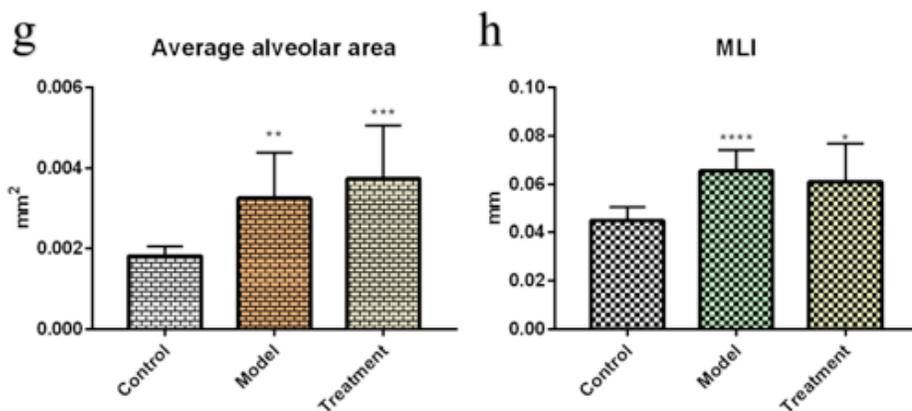
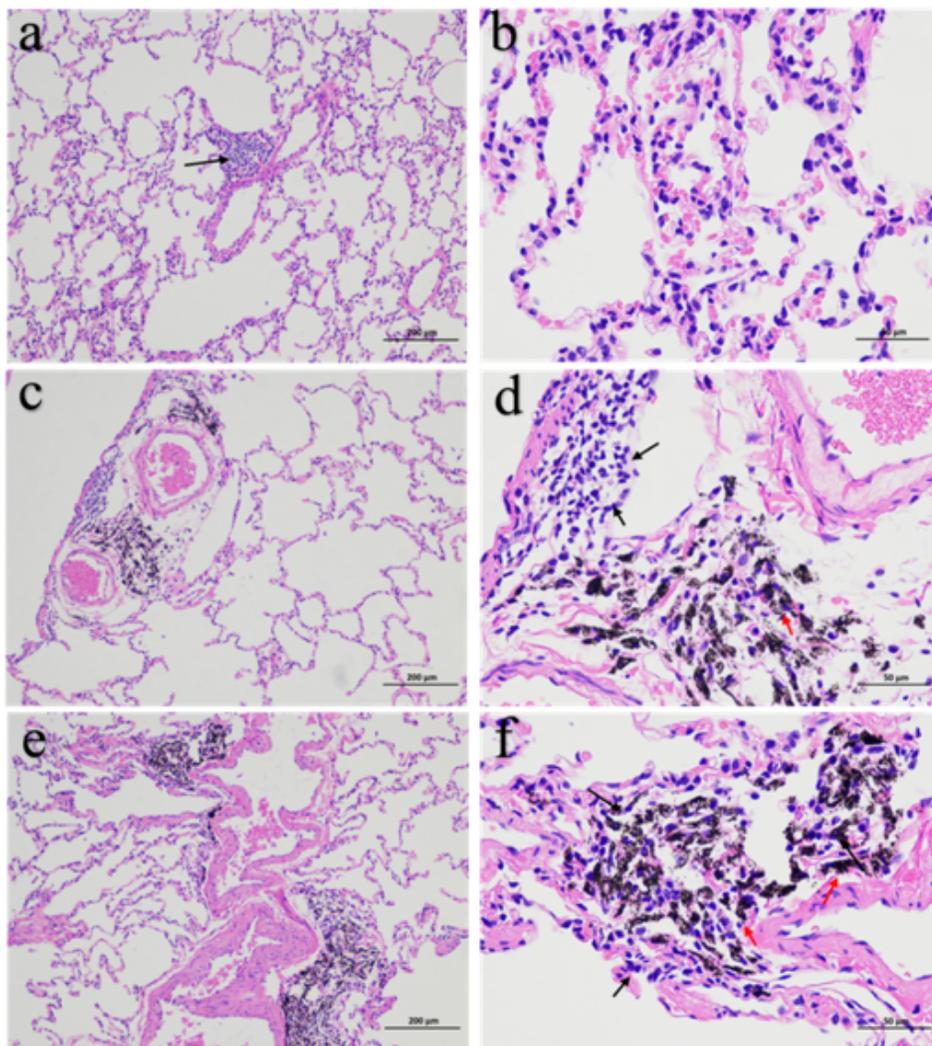
**Figure 7**

Changes in PET-CT in elderly macaque lung after BMMSCS treatment (FDG dosage is 0.2 mCi/kg)



**Figure 8**

Gross anatomical changes of lung tissue in elderly macaque after BMMSCS treatment



**Figure 9**

Changes in the structure of lung tissue after BMMSCS treatment by Hematoxylin-eosin staining (a, b are the Hematoxylin-eosin staining pictures of the lung tissue of the control group of macaques. c, d are the Hematoxylin-eosin staining pictures of the lung tissue of the model group of macaques. e, f are the Hematoxylin-eosin staining pictures of the lung tissue of the treatment group of macaques. g is the statistical graph of the average alveolar area. h is statistical graph of the average lining interval. n=5, n=5,

n is for the number of animals analyzed, \* P <0.05 when compared with the control group, \*\* P <0.01 when compared with the control group, \*\*\* P <0.001 when compared with the control group).

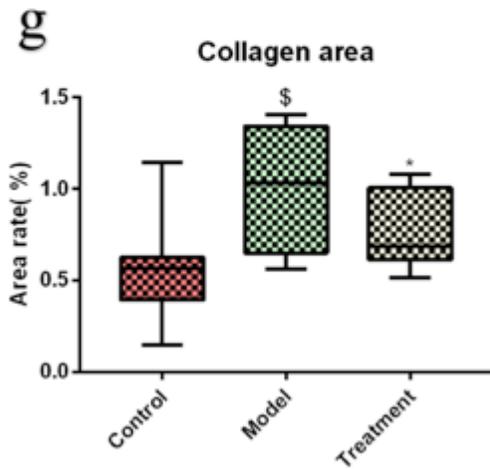
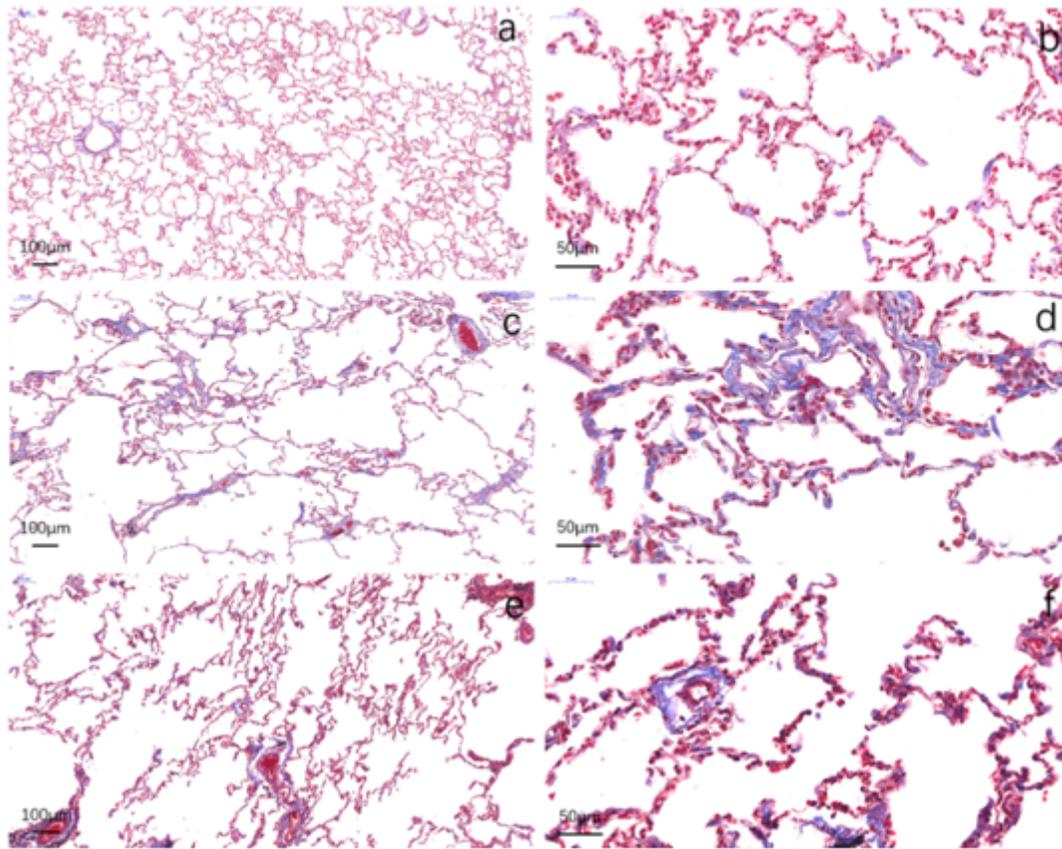


Figure 10

Changes in collagen area in lung tissue after BMMSCS treatment (a, b are the Masson's Trichrome stain pictures of the lung tissue of the control group of macaques. c, d are the Masson's Trichrome stain pictures of the lung tissue of the model group of macaques. e, f are the Masson's Trichrome stain pictures of the lung tissue of the treatment group of macaques. g is the the rate of collagen area statistics graph.

n=5, n is for the number of animals analyzed, \$ P <0.05 when compared with the control group, \*P<0.05 when compared with the model group).

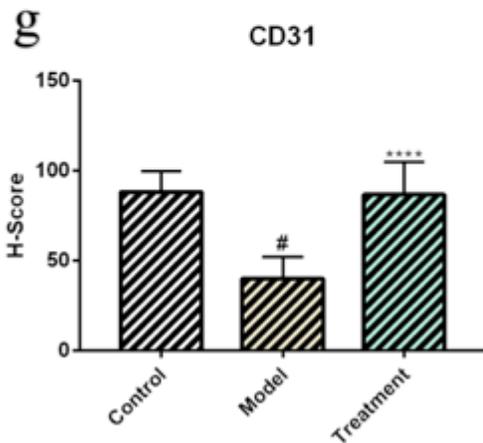
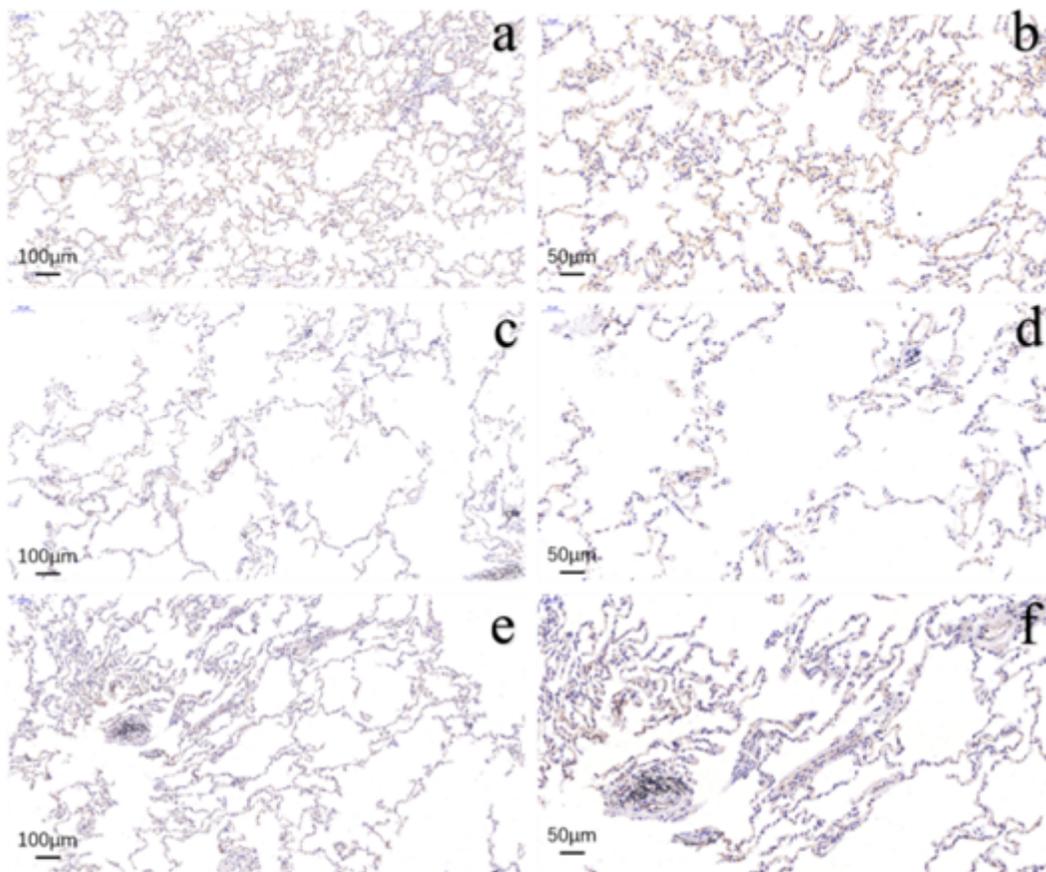
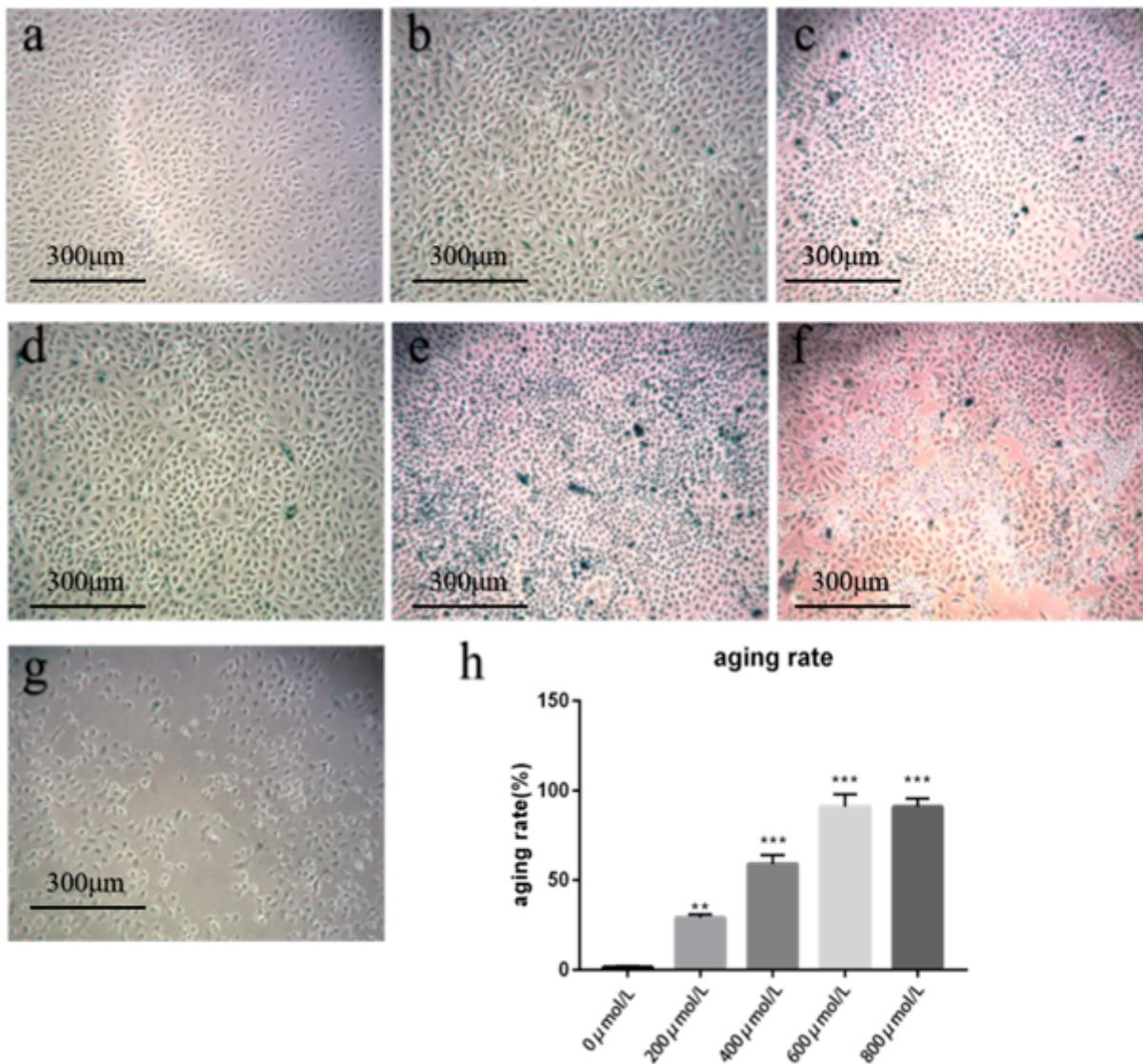


Figure 11

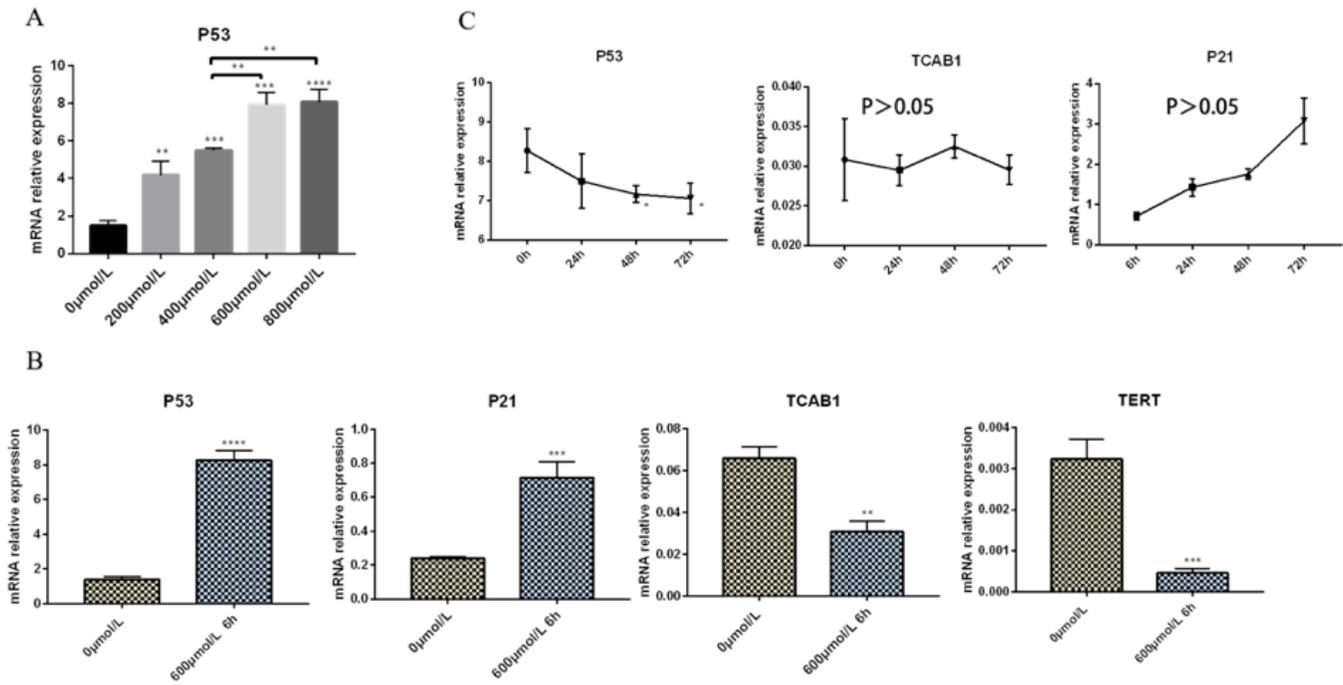
Changes in CD31 surface markers around the alveoli after BMMSCS treatment (a, b are the immunohistochemistry pictures of the lung tissue of the control group of macaques. c, d are the immunohistochemistry pictures of the lung tissue of the model group of macaques. e, f are the immunohistochemistry pictures of the lung tissue of the treatment group of macaques. g is the H-Score

statistical graph of the above pictures. n=5, n is for the number of animals analyzed, #P <0.001 when compared with the control group, \*\*\*\*P<0.0001 when compared with the model group)



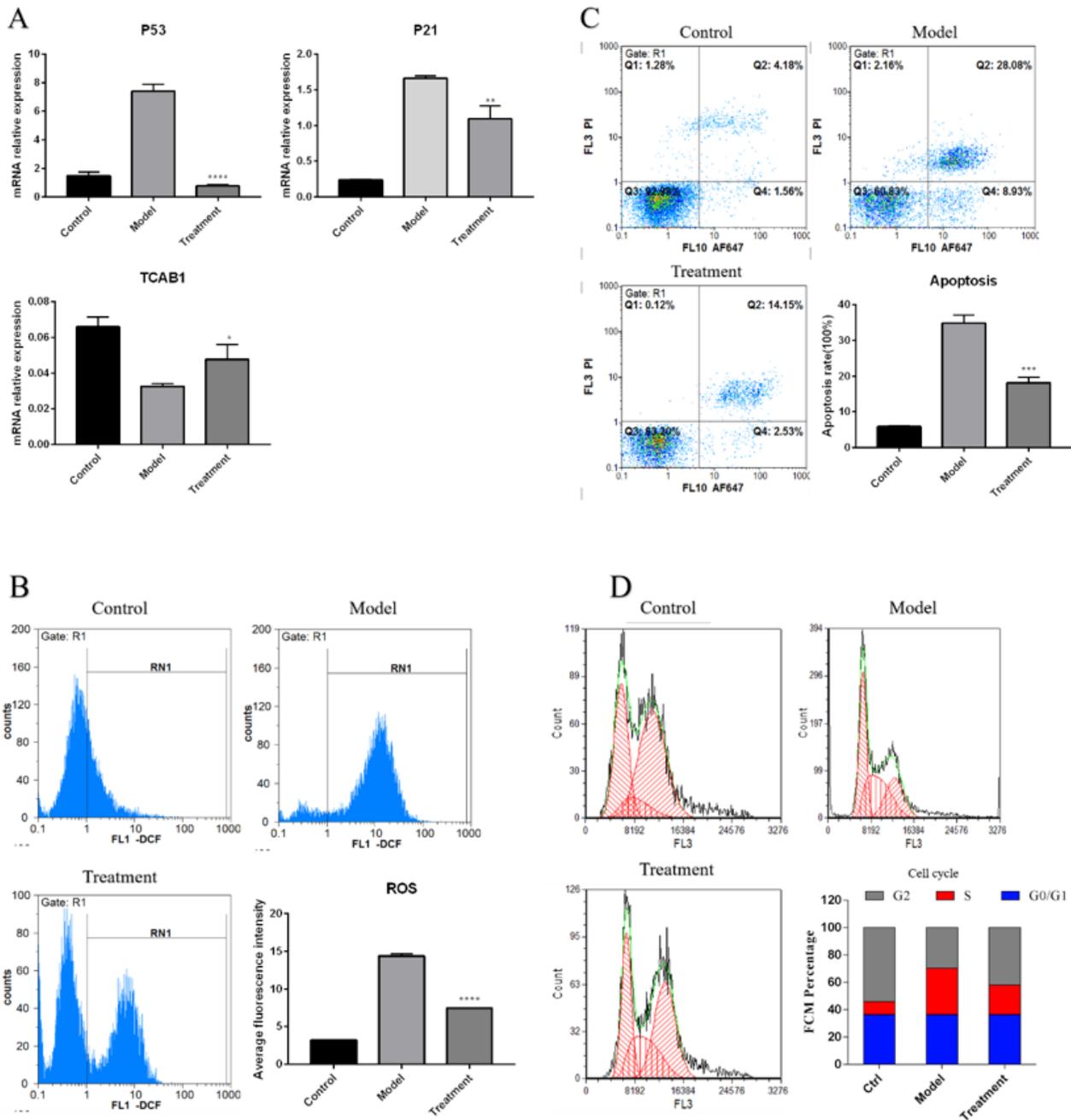
**Figure 12**

Hydrogen peroxide-induced senescence of A549 cells and β-galactosidase staining (100 ×) a, b, c, d, e, f, g are the SA-β-gal staining of BMMSCS after 6 hours of induction at 0 μmol/L, 200 μmol/L, 400 μmol/L, 600 μmol/L, 800 μmol/L, 1000 μmol/L and 1200 μmol/L hydrogen peroxide concentration, respectively. h is the percentage chart of senescent cells. n=3, n is the number of repeated experiments, \*\*P <0.01 when compared with the 0 μmol/L group, \*\*\*\*P <0.0001 when compared with the 0 μmol/L group)



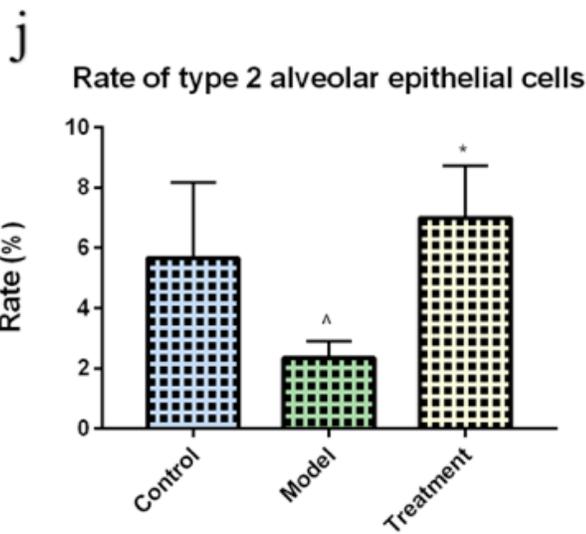
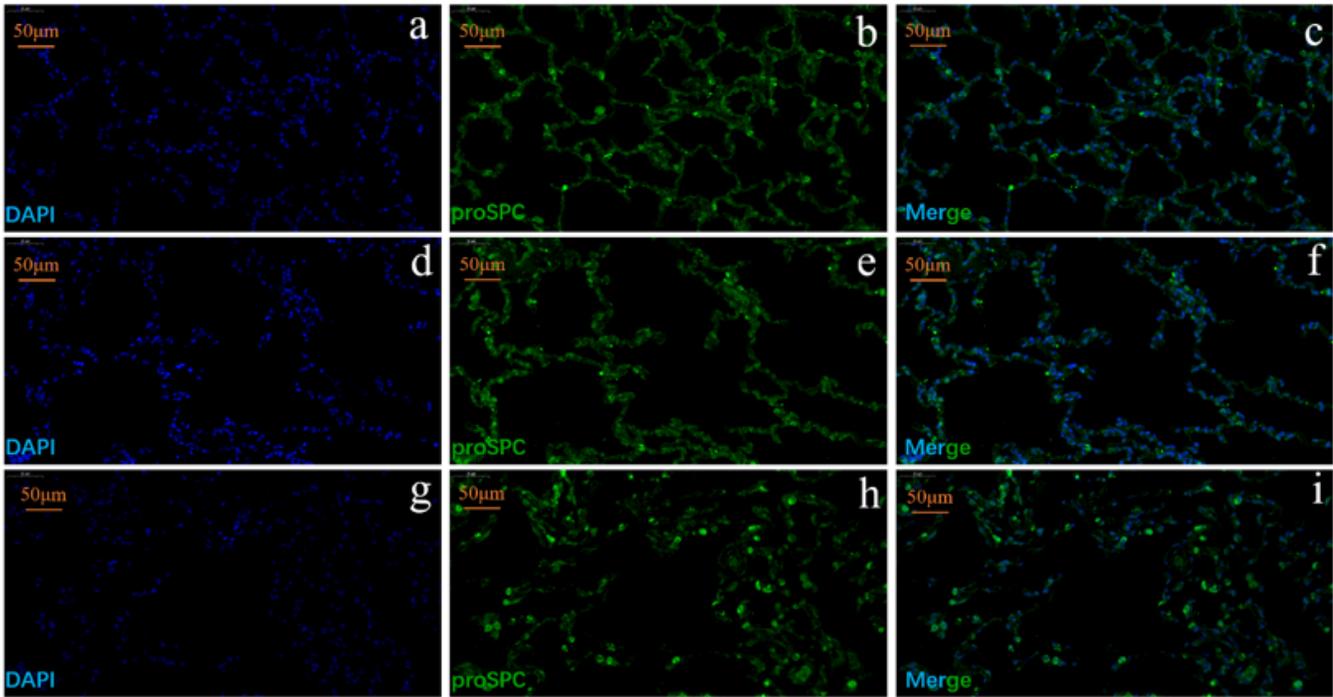
**Figure 13**

Identification of A549 Aging Model a. qPCR P53 expression After induction (n=3 is the number of repeated experiments, #P>0.05, \*\*P<0.01 when compared with the 0μmol/L group, \*\*\* P<0.001 when compared with the 0μmol/L group, \*\*\*\*P<0.0001 when compared with the 0μmol/L group) b. qPCR validation confirming the aging characteristics of A549 cells aging model (n=3 is the number of repeated experiments, \*\*P<0.01 when compared with the 0μmol/L group, \*\*\*P<0.001 when compared with the 0μmol/L group, \*\*\*\* P<0.01 when compared with the 0μmol/L group) c. qPCR validation confirming the aging stability of the A549 cells aging model (n=3 is the number of repeated experiments, \*P<0.05 when compared with the 0h group)



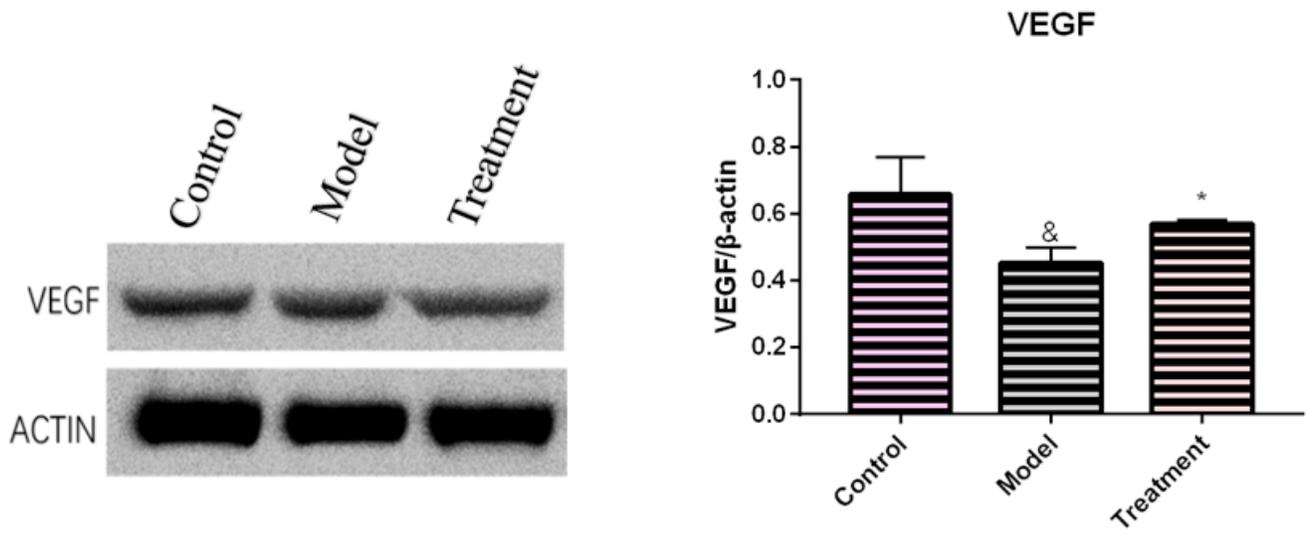
**Figure 14**

Effects of BMMSCS on the aging model of type II alveolar epithelial cells a. qPCR detection of aging-related gene expression after co-culture (n=3, n is the number of repeated experiments, \* P<0.05 when compared with the model group, \*\* P<0.01 when compared with the model group, \* \*\*\*P<0.0001 when compared with the model group) b. Changes in cell ROS level after co-culture (n=3, n is the number of repeated experiments, \*\*\*P<0.001 when compared with the model group, \*\*\*\* P<0.0001 when compared with the model group) c. Changes in cell apoptosis rate after co-culture (n=3, n is the number of repeated experiments, \*\*\*P<0.001 compared with the model group, \*\*\*\* P<0.0001 compared with the model group) d. Changes in cell proliferation (n=3, n is the number of repeated experiments)



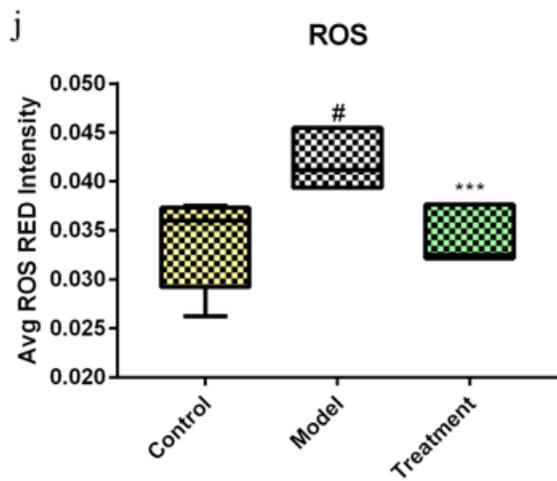
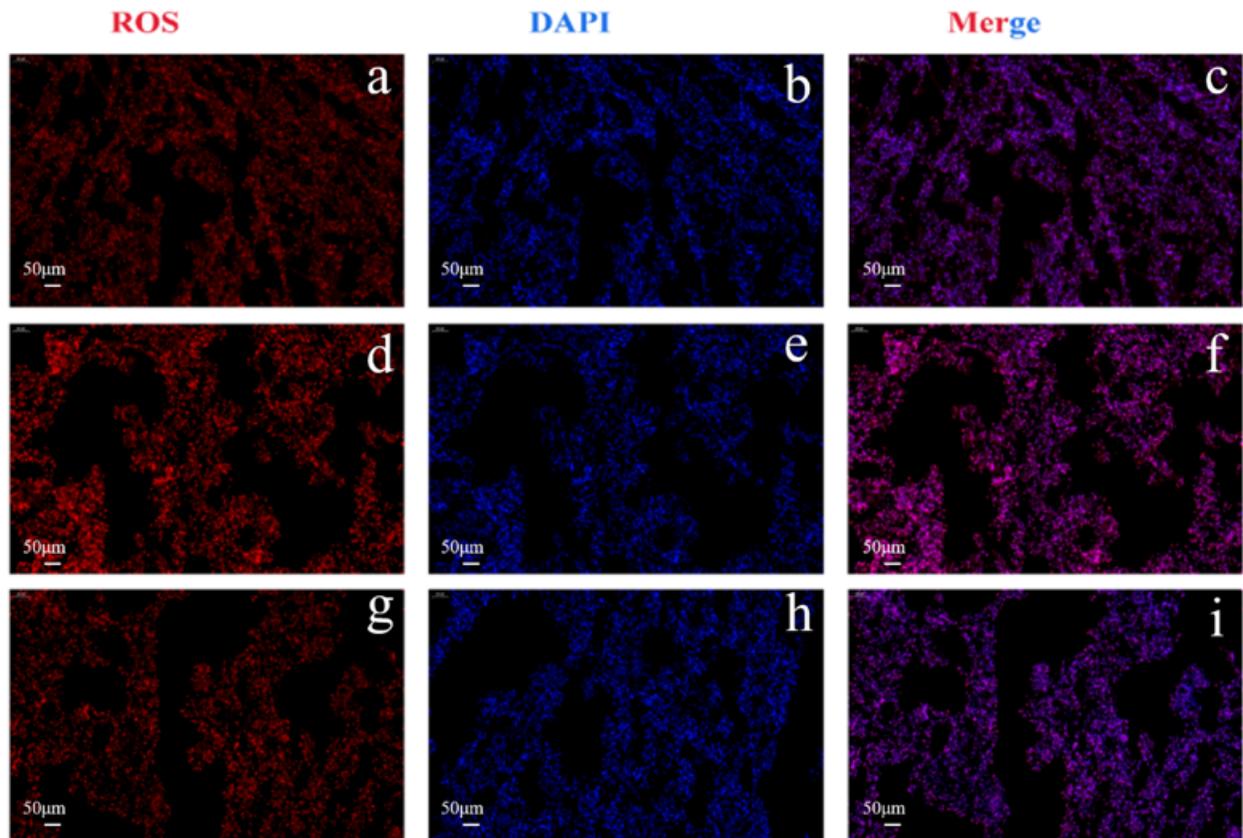
**Figure 15**

Effect of BMMSCS on the number of type 2 alveolar epithelial cells in the lung tissue (a, b, c are the Immunofluorescence pictures of the lung tissue of the control group of macaques. d, e, f are the Immunofluorescence pictures of the lung tissue of the model group of macaques. g, h, i are the Immunofluorescence pictures of the lung tissue of the treatment group of macaques. n=5, n is for the number of animals analyzed, \*\*P<0.01 when compared with the model group, ^P<0.001 compared with the control group)



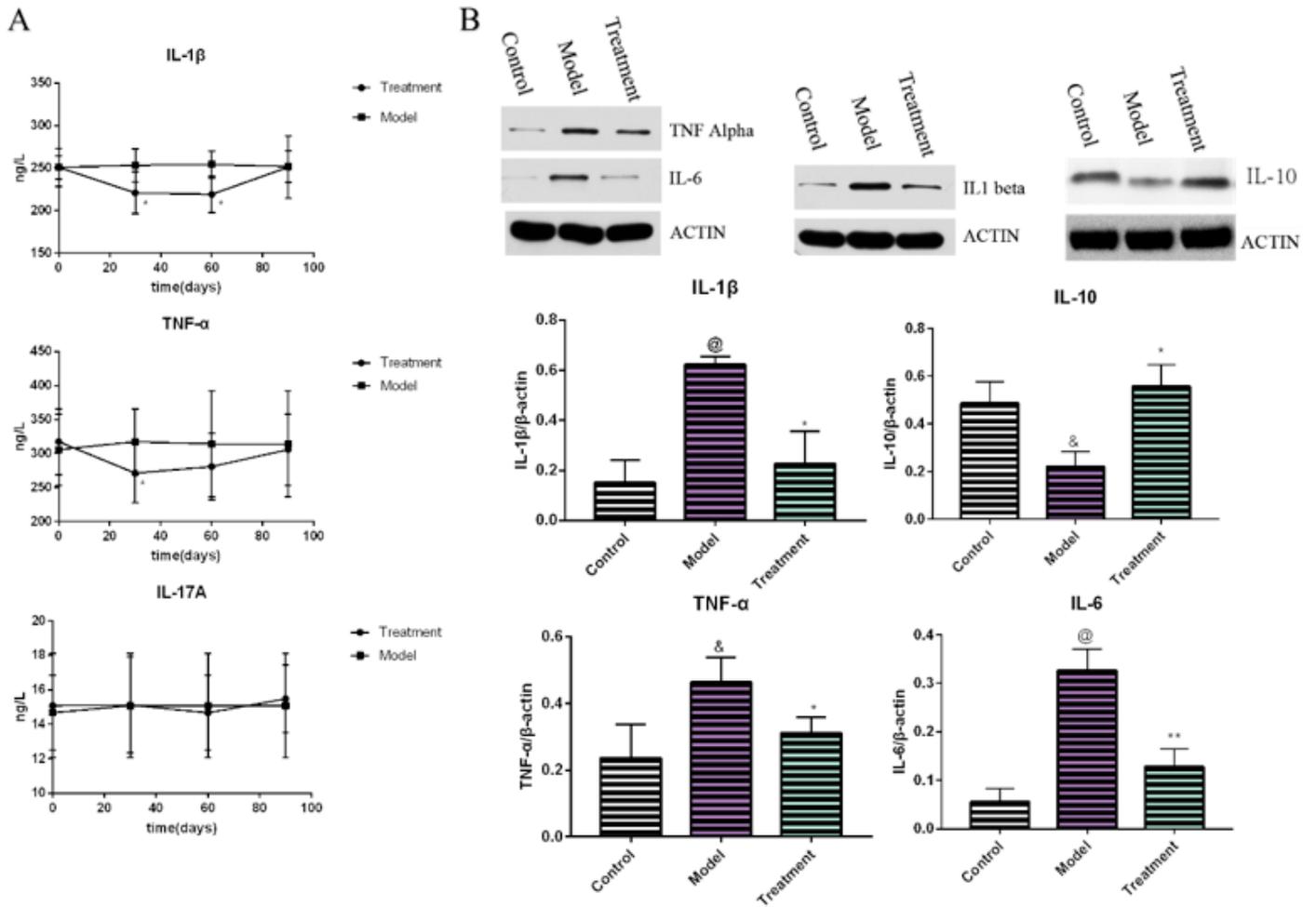
**Figure 16**

Changes in VEGF level in the lung tissue after BMMSCS treatment (n=5, n is for the number of animals analyzed. &P <0.05 compared with the control group, \*P<0.05 compared with the model group)



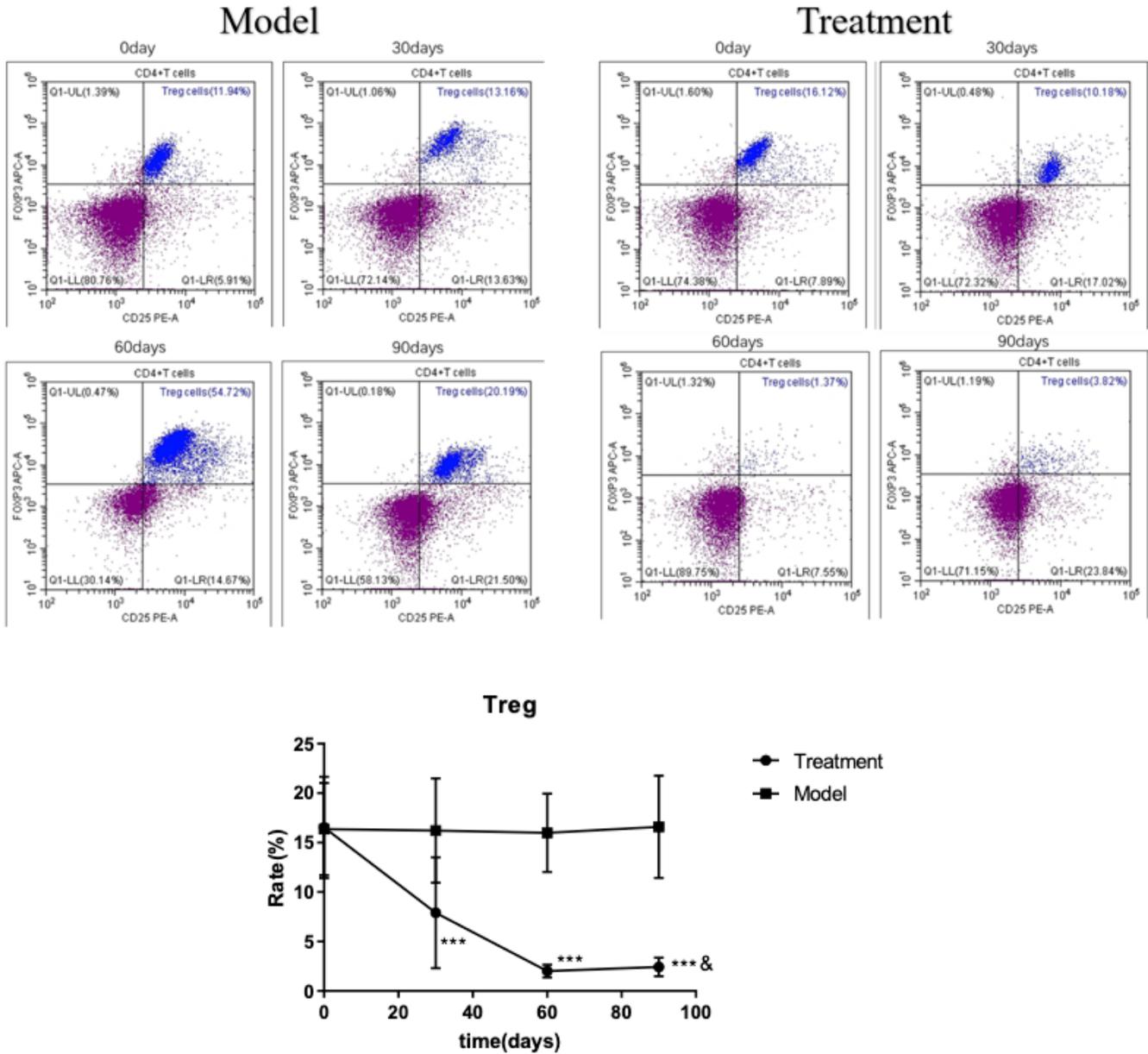
**Figure 17**

Changes in ROS level after BMMSCS treatment (200 ×) ((a, b, c are the ROS staining pictures of the lung tissue of the control group of macaques. d, e, f are the ROS staining pictures of the lung tissue of the model group of macaques. j, h, i are the ROS staining pictures of the lung tissue of the treatment group of macaques. J is the H-Score statistical graph of the above pictures. n=5, n is for the number of animals analyzed, #P <0.001 compared with the control group, \*\*\*P <0.001 compared with the model group)



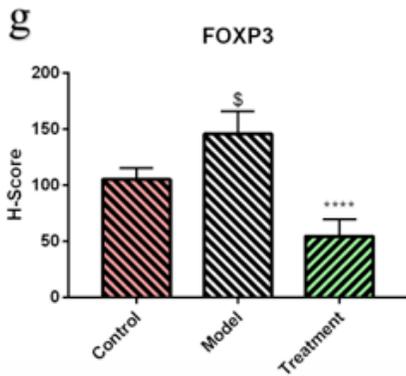
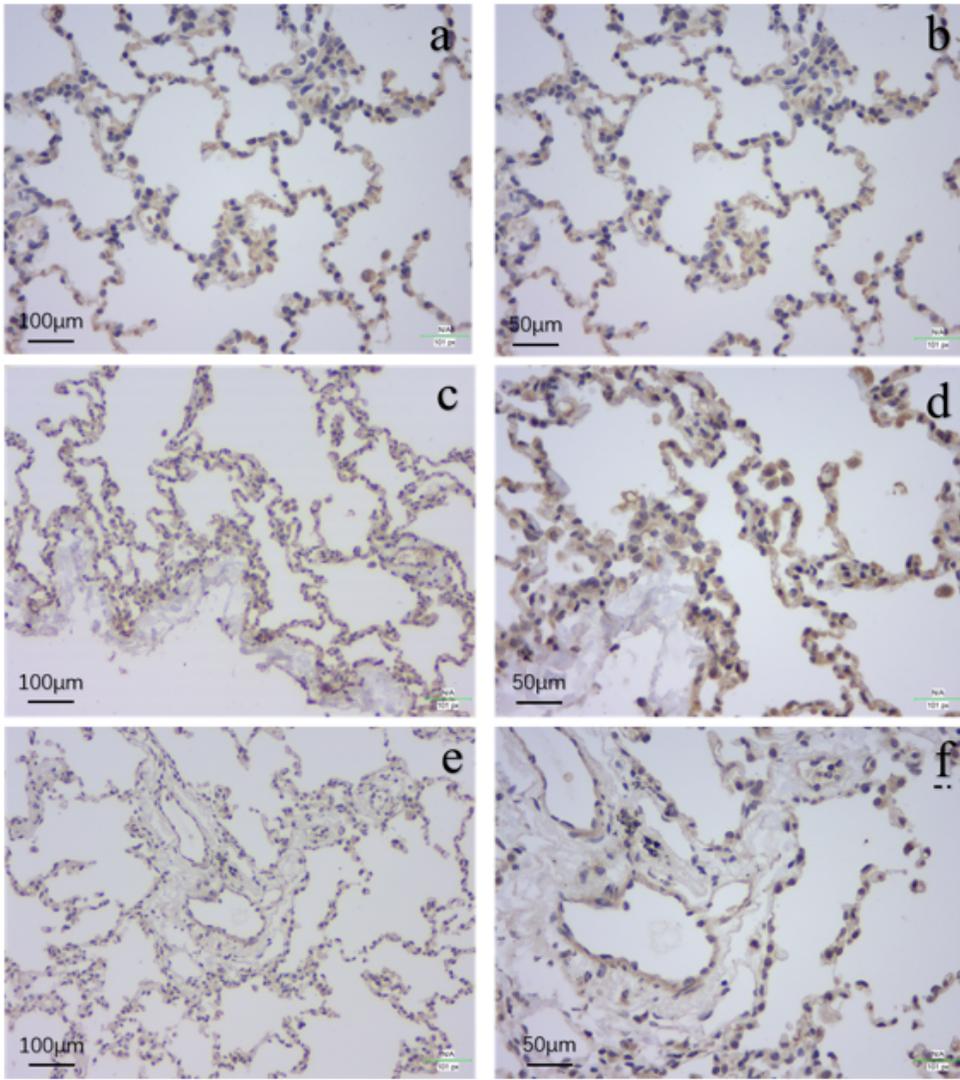
**Figure 18**

Changes in the inflammatory factors in the lung tissue after BMMSCS treatment a. Changes in the level of inflammatory factors in peripheral blood after cell therapy (n=5, n is for the number of animals analyzed \*P <0.05 compared with the model) b. Changes in inflammatory factors in lung tissue after BMMSCS treatment (n=5, n is for the number of animals analyzed, \*P <0.05 compared with the model group, \*\* P <0.01 compared with the model group, & P <0.05 compared with the control group, @P <0.001 compared with the control group).



**Figure 19**

Changes in Treg cell ratio in peripheral blood after BMMSCS treatment (FOXP4 antibody, CD25 antibody and CD4 antibody were used to co-label Treg cells,, n=5, n is for the number of animals analyzed. \*\*\*P <0.001 compared with the model group, & P<0.05 compared with 60 days after treatment)



**Figure 20**

Changes in FOXP4 content in the lung tissue after BMMSCS treatment (a, b are the immunohistochemistry pictures of the lung tissue of the control group of macaques. c, d are the immunohistochemistry pictures of the lung tissue of the model group of macaques. e, f are the immunohistochemistry pictures of the lung tissue of the treatment group of macaques. g is the H-Score

statistical graph of the above pictures. n=5, n is for the number of animals analyzed. \*\*\*\*P <0.0001 compared with the model group, \$ \*P <0.001 compared with the control group).