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SPECIAL SECTION: MULTISYSTEMS PHARMACOLOGY AND THERAPEUTIC APPLICATION OF ADIPOKINES AND NEUROPEPTIDES IN OBESITY—ARTICLE

Adiponectin in cigarette smoke–induced epithelial-mesenchymal transition



Francesca Panico¹ , Giuseppe Spaziano² , Ersilia Nigro^{2,3}, Davida Mirra^{2,*} ,
Renata Esposito², Eleonora Caiazzo² , Aurora Daniele^{3,4}, Bruno D'Agostino²

¹ Science of Health Department, School of Medicine, University “Magna Graecia” of Catanzaro, Catanzaro, Italy

² Department of Environmental, Biological and Pharmaceutical Sciences and Technologies, University of Campania “Luigi Vanvitelli”, Caserta, Italy

³ CEINGE Biotechnologie Avanzate “Franco Salvatore” S.c.a r.l., Naples, Italy

⁴ Department of Molecular Medicine and Medical Biotechnology, University of Naples Federico II, Naples, Italy

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ABSTRACT

Cigarette smoke (CS) contributes to a wide range of chronic respiratory diseases by inducing epithelial-mesenchymal transition (EMT), which drives tissue remodeling, fibrosis, and tumor progression. In this scenario, adiponectin (Acrp30) has emerged as a potential EMT-negative regulator, inhibiting migration and invasion in non-small cell lung cancer cell lines. This study aimed to investigate ability of Acrp30 to prevent CS-induced EMT and profibrotic remodeling in normal and adenocarcinoma lung cells, through AdipoRon, a synthetic Acrp30 receptor agonist. BEAS-2B and A549 cells were used to preliminarily assess the optimal dose and timing for CS and AdipoRon exposure for subsequent evaluations by thiazolyl blue tetrazolium bromide assay. Expression of epithelial (E-cadherin), mesenchymal (vimentin), and profibrotic (α -smooth muscle actin, transforming growth factor- β , total collagen deposition, and collagen type I α 1 chain) markers was examined through immunofluorescence, reverse transcription polymerase chain reaction, and Sirius Red staining. Wound-healing assay was used to assess CS and Acrp30 ability to influence cell migration. Reactive oxygen species levels were measured by 2',7'-dichlorofluorescein diacetate assay. Acrp30 prevents CS-induced dysregulation of E-cadherin and vimentin at both mRNA and protein levels, preserving the epithelial-mesenchymal balance in both cell lines. Furthermore, CS-induced increase in profibrotic marker expression was significantly attenuated by AdipoRon pretreatment in both BEAS-2B and A549 cells. The CS-induced increase in migratory capacity was significantly reversed by Acrp30 in both cell lines, confirming its ability to prevent EMT. Finally, Acrp30 also markedly reduced CS-induced reactive oxygen species overproduction in BEAS-2B and A549 cells, indicating a role in modulating oxidative stress-driven epithelial plasticity. Thus, Acrp30 effectively preserves epithelial markers, limits the mesenchymal shift and profibrotic remodeling, and reduces migratory capacity by mitigating oxidative stress. These results support Acrp30 as a promising therapeutic candidate for preventing airway remodeling and disease progression in smoking-related conditions.

Significance Statement: Cigarette smoke (CS) induces epithelial-mesenchymal transition and fibrosis in both healthy and cancerous lung epithelial cells. Adiponectin pretreatment prevents CS-induced epithelial-mesenchymal transition and profibrotic remodeling in both cell lines; adiponectin can prevent airway remodeling and disease progression in CS conditions.

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* Address correspondence to: Dr Davida Mirra, Department of Environmental, Biological and Pharmaceutical Sciences and Technologies, University of Campania “Luigi Vanvitelli”, Caserta, Italy. E-mail: davida.mirra@unicampania.it

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F.P. and G.S. contributed equally to this work.

1. Introduction

1.1. Cigarette smoke–induced lung injury and epithelial–mesenchymal transition

Cigarette smoke (CS) is a mixture of many different chemical compounds and represents a widely recognized primary risk factor for several lung conditions, including chronic obstructive pulmonary disease (COPD) and lung cancer.^{1–3} CS exposure can lead to chronic pulmonary injury and dysfunction through different complex mechanisms, including DNA damage, epigenetic alteration, oxidative stress, and alteration in inflammatory pathways.^{4–11} The bronchial and alveolar epithelium are the first line of defense against inhaled toxicants such as those contained in CS,^{12–16} which can lead to significant alterations of epithelial integrity and cell function, ultimately promoting epithelial–mesenchymal transition (EMT), a crucial event in tissue remodeling, tumor progression, and metastasis.^{17–20} At the molecular level, the EMT process is defined by the downregulation of epithelial junction proteins such as E-cadherin as well as the upregulation of mesenchymal markers such as vimentin.²¹ EMT not only contributes to tissue remodeling and fibrosis but is also involved in carcinogenesis and metastasis.^{22,23} Indeed, in lung cancer, EMT drives the invasion process through desmoplastic stroma formation and contributes to the conversion of cancer cells into invasive malignant phenotypes.^{24,25} Additionally, in the small airway of smokers with COPD who develop lung cancer, EMT plays a key role in airway remodeling and cancer progression.²⁶

1.2. Oxidative stress as a driver of CS-induced EMT

Oxidative stress is a critical driver of EMT induction. Although the exact signaling pathways through which reactive oxygen species (ROS) promote this process remain unclear, ROS can modulate cellular signaling to activate EMT-associated pathways and drive the acquisition of a cancerous phenotype.²⁷ CS, in addition to direct oxidative injury, promotes the trafficking and persistence of inflammatory cells in the lung. These immune cells, particularly neutrophils and macrophages, release a range of chemotactic and cytotoxic mediators, such as proteolytic enzymes and ROS, contributing to extracellular matrix degradation and epithelial barrier dysfunction.^{28–30} Currently, bronchodilators and anti-inflammatory agents, commonly drugs used for the treatment of smoke-related lung injury,³¹ are not able to act on EMT, highlighting the urgent need to identify new therapeutic targets.

1.3. Adiponectin as a potential regulator of epithelial plasticity and tumor progression

Adiponectin (Acrp30), an adipokine produced by adipose tissues and well established as a pleiotropic mediator,³² plays a critical role in the lung microenvironment by reducing local inflammation and inhibiting alveolar macrophage activation.³³ This adipokine exerts its biological functions through specific receptors, adiponectin receptor 1 (AdipoR1), AdipoR2, and T-cadherin. Dysregulation of the Acrp30 system has been reported in several malignancies, including non–small cell lung carcinoma (NSCLC). Previously, we observed reduced levels of total adiponectin and T-cadherin as well as an increase in high molecular weight oligomers and AdipoR1 in NSCLC lung tissue, suggesting a dysfunctional Acrp30 axis.³⁴ Emerging evidence indicates that Acrp30 may inhibit migration and invasion by suppressing the EMT process in NSCLC cells.³⁵ However, the study by Cui et al³⁵ focused on Acrp30 effects in established NSCLC cells under basal conditions, without addressing the contribution of CS, which is the

major etiological factor in lung carcinogenesis and a key inducer of EMT-related alterations in airway epithelium. In patients with nasopharyngeal carcinoma, an inverse correlation has been observed between serum Acrp30 level and tumor stage, recurrence and metastasis, and poor metastasis-free survival.³⁶ Furthermore, circulating Acrp30 levels are reduced in smokers, suggesting a potential link between Acrp30 deficiency and increased susceptibility to smoke-induced lung conditions.³⁷

1.4. Study aim

The potential of Acrp30 in counteracting CS-induced EMT onset in healthy epithelium as well as in tumor progression has not been fully elucidated. Therefore, the aim of this study was to investigate the protective effects of Acrp30 against CS-induced EMT, to our knowledge, here addressed for the first time, in healthy and cancerous lung epithelial cells, through AdipoRon, a synthetic Acrp30 agonist that activates the AMP-activated protein kinase and peroxisome proliferator-activated receptor- α pathways, reducing oxidative stress and inflammation.³⁸

2. Materials and methods

2.1. Cell culture and treatment

The normal human bronchial epithelial cell line BEAS-2B was purchased from American Type Culture Collection (ATCC CCL-185) and cultured in RPMI-1640 Medium (Gibco) supplemented with 10% FBS (Gibco) and 1% penicillin/streptomycin (Gibco) at 37 °C and in 5% CO₂/95% humidified air. The human lung adenocarcinoma cell line A549 was purchased from ATCC (ATCC CCL-185) and cultured in Dulbecco's modified Eagle's medium/nutrient mixture F-12 (Gibco) supplemented with 10% FBS (Gibco) and 1% penicillin/streptomycin (Gibco) at 37 °C and in 5% CO₂/95% humidified air. Both cell lines were grown in 75 cm² tissue culture flasks. Confluence was reached after 3 to 4 days in culture, and all experiments were performed on cells between passage 15 and 25. The AdipoRon compound (Merck) was solubilized in DMSO to produce a stock solution (Sigma-Aldrich) and subsequently diluted to working concentration. Three independent experiments for each evaluation were performed in both cell lines.

2.2. Preparation of CS extract

The cigarette smoke extract (CSE) was prepared by bubbling 10 Red Marlboro cigarettes (Phillip Morris), as previously reported.³⁹ The obtained extract was considered as 100% CSE. The CSE was sterile-filtered through a 0.20- μ m pore filter, aliquoted, and stored at –80 °C to reduce variability and ensure reproducibility across all experiments. Before use, CSE was diluted with RMPI or Dulbecco's modified Eagle's medium to the indicated concentration.

2.3. Thiazolyl blue tetrazolium bromide assay

BEAS-2B or A549 cells were used to determine the timing and optimal dose for CSE and AdipoRon exposure for subsequent analyses. The cells were seeded at a density of 1×10^4 per well in 96-well plates and allowed to attach overnight at 37 °C in 5% CO₂/95% humidified air. The cells were then exposed to increasing CSE concentrations, ranging from 1% to 20% for 24 hours. In another set of experiments, cells were exposed to increasing AdipoRon concentrations ranging from 5 to 50 μ g/mL for 24 and 48 hours. Cellular viability was determined by the thiazolyl blue tetrazolium bromide assay (M5655, Sigma-Aldrich). Briefly, 25 μ L of thiazolyl blue tetrazolium bromide (2 mg/mL) was added to each well and

incubated for 4 hours to perform the proliferation assay. After this time, DMSO was added to measure the absorbance. The cellular viability was calculated as the percentage of absorbance at 570 nm of untreated cells using a microplate reader (GloMax Explorer, Promega).

2.4. ROS determination

To evaluate the AdipoRon potential to counteract CSE effects on the redox balance of BEAS-2B or A549 cells, ROS species were measured with 2',7'-dichlorodihydrofluorescein diacetate assay. The cells were seeded at a density of 1×10^4 per well in 96-well plates and allowed to attach overnight at 37 °C in 5% CO₂/95% humidified air. The cells were then pretreated or not for 2 hours with AdipoRon (10 µg/mL) or 1 mM N-acetylcysteine (NAC), used as an antioxidant control, and then exposed to 2% or 10% CSE for 24 hours. H₂O₂ was used as a positive oxidative stress control in the presence or absence of NAC. After treatments, 200 µL of 50 µM 2',7'-dichlorodihydrofluorescein diacetate (Merck) in Hanks' balanced salt solution (Sigma-Aldrich) were added in each well and incubated for 1 hour at 37 °C. Then, the cells were subsequently washed 3 times with PBS to remove the extracellular fluorescent compounds and minimize background signal. Fluorescence was measured at an excitation wavelength of 540 nm and an emission wavelength of 590 nm using a microplate reader (GloMax Explorer, Promega). The relative ROS production was normalized to the fluorescence intensity of untreated control cells.

2.5. Reverse transcriptase and quantitative real-time polymerase chain reaction

mRNA basal expression levels of AdipoR1 and AdipoR2 were preliminarily evaluated in both cell lines. Moreover, to evaluate the ability of AdipoRon to counteract CSE effects on fibrosis and EMT, the expression of transforming growth factor- β (TGF- β), epithelial (E-cadherin), and mesenchymal (vimentin) markers was examined through reverse transcription polymerase chain reaction (RT-PCR). Notably, 1×10^6 cells per well BEAS-2B and A549 were seeded in 6-well plates and allowed to attach overnight at 37 °C in 5% CO₂/95% humidified air. Both cell lines were pretreated or not for 2 hours with AdipoRon (AR) and then exposed to 2% or 10% CSE for 24 hours. Cells were then lysed and total RNA was extracted using the TRIzol method. Then, 2 µg of total RNA was subjected to RT-PCR using the High-Capacity cDNA Reverse Transcription Kit (Thermo Fisher Scientific), according to the manufacturer's instructions. The thermocycling conditions were set as follows: 25 °C for 10 minutes, 37 °C for 120 minutes, 85 °C for 5 minutes, and 4 °C hold. RT-PCR was performed using the TaqMan Universal PCR Master Mix (Thermo Fisher Scientific) according to the manufacturer's instructions and QuantumStudio3 Real-Time PCR Systems. The thermocycling conditions were set as follows: 50 °C for 2 minutes, 95 °C for 10 minutes, 95 °C for 15 seconds, and 60 °C for 1 minutes. Data from all RT-PCR experiments and mRNA expression levels were normalized to the endogenous housekeeping gene control applying the comparative $2^{-\Delta\Delta Ct}$ method, where $\Delta Ct = Ct_{mRNA} - Ct_{housekeeping\ mRNA}$, whereas the relative quantification of differences in expression was conducted with $\Delta\Delta Ct = \Delta Ct_{TREATMENT} - \Delta Ct_{CTR}$.

2.6. Immunofluorescence staining

To further investigate AdipoRon's ability to prevent CSE effects on EMT and profibrotic remodeling, expression of E-cadherin, vimentin, α -smooth muscle actin (α -SMA), and collagen type I $\alpha 1$ chain (COL1A1) was evaluated through immunofluorescence

staining. BEAS-2B and A549 cells were seeded on 12-mm coverslips at a density of 5×10^4 per well in 24-well plates and allowed to attach overnight at 37 °C in 5% CO₂/95% humidified air. Both cell lines were pretreated or not for 2 hours with AdipoRon and then exposed to 2% or 10% CSE for 24 hours. After treatments, cells were washed 3 times in PBS, fixed in 4% formaldehyde at room temperature for 20 minutes, followed by permeabilization with 1% Triton X-100 in PBS for 10 minutes. Cells were then washed 3 times with PBS and stained with an antivimentin /E-cadherin/COL1A1 (ABclonal; Elabscience) and anti- α -SMA (Sigma-Aldrich) antibodies followed by donkey anti-rabbit IgG 555/488 (Alexa Fluor) and donkey anti-mouse IgG 555 (Alexa Fluor) secondary antibodies. After staining, the coverslips were rinsed in PBS and mounted on glass slides. Images were acquired with an Apotome.2 microscope (Zeiss) at 20 $\times$ magnification and analyzed with ImageJ software (National Institutes of Health, NIH).

2.7. Sirius Red staining

Total collagen deposition was assessed in BEAS-2B and A549 cells by Sirius Red staining. Cells were seeded at a density of 1×10^4 per well in 96-well plates and allowed to attach overnight at 37 °C in 5% CO₂/95% humidified air. Both cell lines were pretreated or not for 2 hours with AdipoRon and then exposed to 2% or 10% CSE for 24 hours. After treatments, cells were washed once with PBS and then fixed with 4% formaldehyde for 20 minutes at room temperature. Following fixation, cells were washed 3 times with PBS and incubated with 50 µL of 0.1% Sirius Red (Direct Red 80; Sigma-Aldrich) prepared in 1% acetic acid for 60 minutes at room temperature. Cells were then washed with 0.1 M HCl and then 0.1 M sodium hydroxide (NaOH) solution was added to each well to eluate the bound dye from the cell-associated collagens. Representative images of stained wells were acquired before dye elution. Subsequently, 100 µL of the eluted solution was transferred to a new 96-well plate, and absorbance was measured at 540 nm using a microplate reader (GloMax Explorer, Promega). The relative collagen content was normalized to the absorbance of untreated control cells.

2.8. Wound-healing assay

To further explore AdipoRon's potential in counteracting CSE-induced EMT, cell migration was examined through the wound-healing assay. BEAS-2B and A549 cells were seeded at a density of 2×10^5 cells/well into a 12-well plate and allowed to attach overnight at 37 °C in 5% CO₂/95% humidified air. The monolayer was scratched using a sterile tip and washed with PBS to remove detached cells. Then the cells were pretreated or not for 2 hours with AdipoRon and then exposed to 2% or 10% CSE for 24 hours. Both cell lines were photographed at 4, 12, and 24 hours post-wounding. The relative closure area of the wound at the chosen time points was normalized to the initial wound area of untreated cells.

2.9. Statistical analysis

The normal distribution of data was assessed before statistical analysis using appropriate tests. Specifically, normal distribution was evaluated using the Shapiro-Wilk test. Variables showing normal distribution were analyzed with parametric tests. The one-way ANOVA followed by Tukey multiple comparison test were used for statistical analyses using the GraphPad software (version 9.0) (GraphPad Software). Differences were considered statistically significant at $P < .05$. Data are presented as the mean $\pm$ SD.

Only statistically significant differences relevant to the purpose of this study are displayed in the graphs.

3. Results

3.1. CS effects on cellular viability

To determine the optimal CSE exposure for downstream analyses, BEAS-2B and A549 cells were treated with different concentrations of CSE for 24 hours. Both cell lines showed a dose-dependent decrease in viability, with the most pronounced reduction observed at 5% CSE for BEAS-2B ($P < .0001$) and 10% CSE for A549 ($P < .0001$) (Fig. 1, A and C). From these concentrations onward, no further decrease was observed in either cell line. In BEAS-2B, evident morphological alteration, including cell rounding and partial detachment, became significant starting from 5% CSE (Fig. 1B), whereas significant morphological changes in A549 cells were evident only upon exposure to the highest concentration of CSE (Fig. 1D). Based on these results, 2% CSE for BEAS-2B and 10% CSE for A549 were selected as working concentrations for subsequent experiments, representing the doses at which the most pronounced functional impairments were observed, without altering cell morphology.

3.2. AR effects on cellular viability

To establish the exact dose and exposure time of AR for downstream analyses, BEAS-2B and A549 cells were treated with increasing concentrations of AR ranging from 5 to 50 $\mu\text{g/mL}$ for 24 or 48 hours (Fig. 2). After 24 hours, BEAS-2B viability was unaffected at 5 and 10 $\mu\text{g/mL}$, whereas 50 $\mu\text{g/mL}$ induced a significant ~40% reduction ($P < .01$) (Fig. 2A). After 48 hours, BEAS-2B cells exhibited a reduction in viability across both AR-treated and corresponding vehicle-treated groups, with the most pronounced effect observed at 50 $\mu\text{g/mL}$ ($P < .0001$) (Fig. 2B). In contrast, A549 cells did not show significant changes in viability at any AR concentration or time point (Fig. 2, C and D). Based on these findings, 10 $\mu\text{g/mL}$ AR was selected as the optimal and noncytotoxic working condition for both cell lines, whereas the 2-hour pretreatment was established in accordance with previous reports^{40–42} for subsequent experiments.

3.3. AR pretreatment modulation of CSE-induced EMT markers dysregulation

Basal expression levels of AdipoR1 and AdipoR2 were preliminarily evaluated in both cell lines. BEAS-2B cells showed

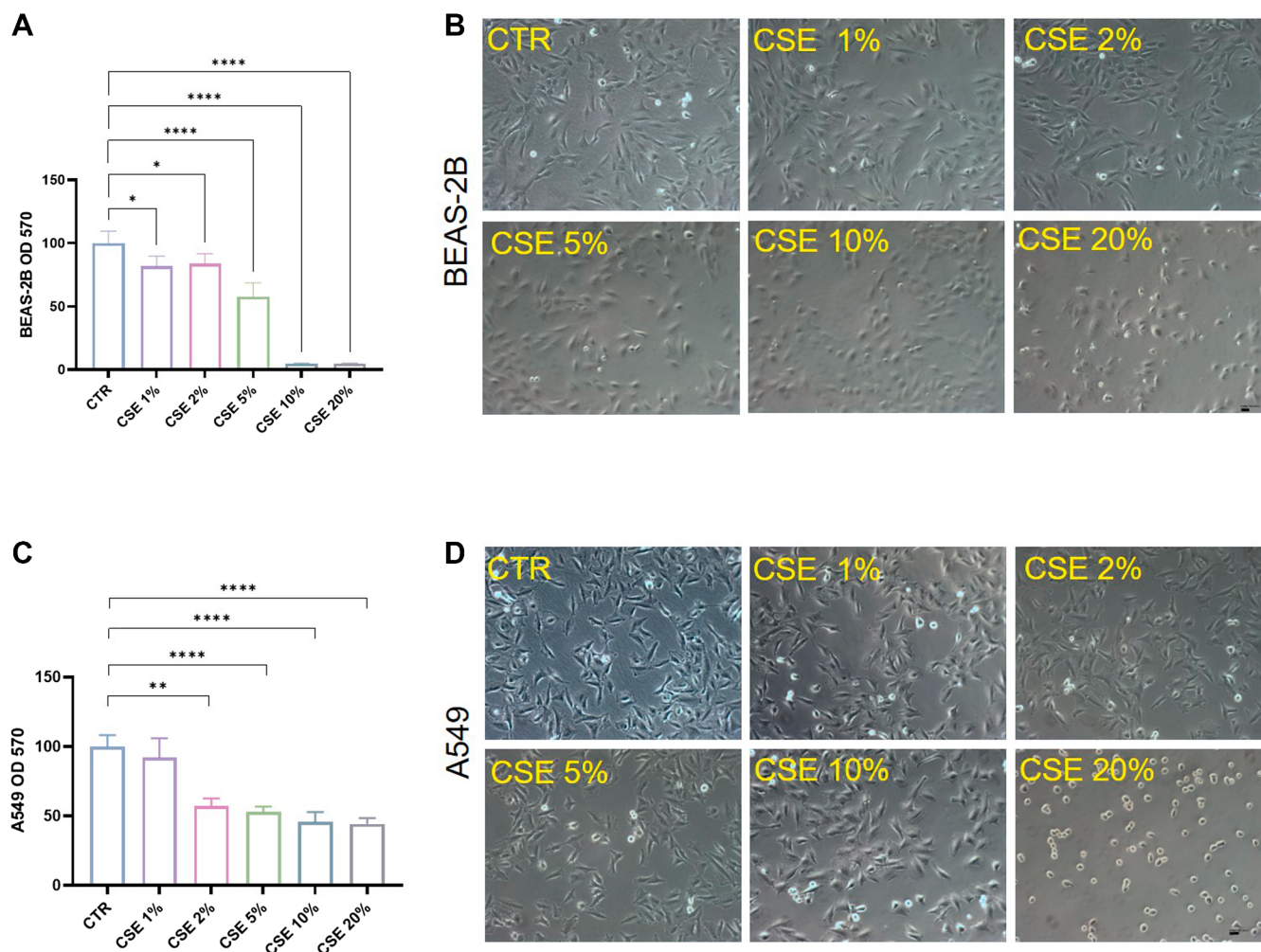


Fig. 1. BEAS-2B and A549 cells were used to assess the exact dose of CSE exposure for downstream analyses. Both cell lines were exposed to different CSE concentrations ranging from 1% to 20% for 24 hours. Following treatments, cellular viability was determined by the thiazolyl blue tetrazolium bromide assay. The relative cellular viability was calculated as the percentage of absorbance at 570 nm of untreated cells (A and B). Representative images of BEAS-2B (C) and A549 (D) cells treated with increasing concentrations of CSE. Scale bar, 100 μm . Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, **** $P < .0001$.

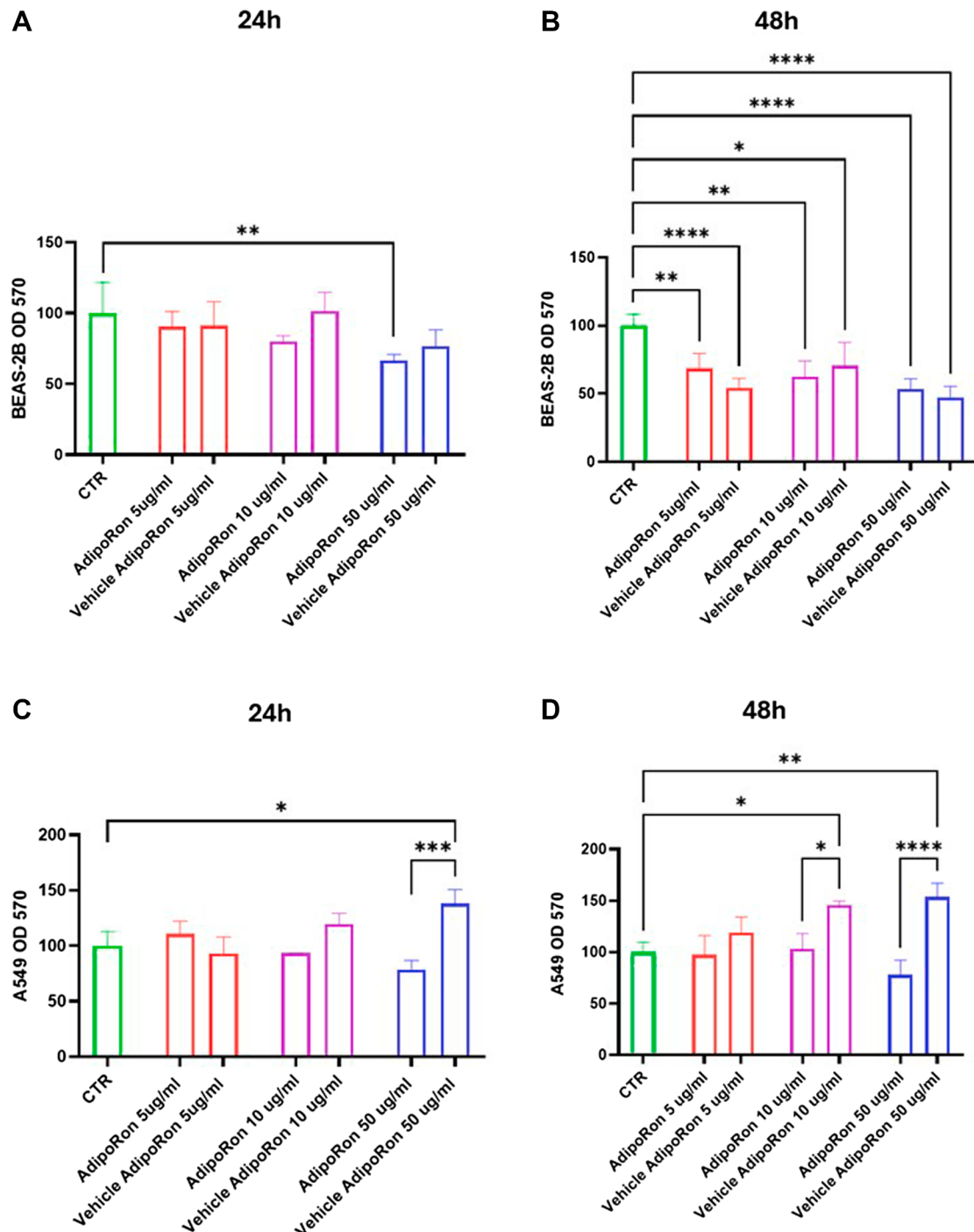


Fig. 2. BEAS-2B and A549 cells were used to assess the exact dose and timing of AR exposure for downstream analyses. Both cell lines were exposed to different AR concentrations ranging from 5 to 50 $\mu\text{g}/\text{mL}$ for 24 hours (A and C) and 48 hours (B and D). Following treatments, cellular viability was determined by the thiazolyl blue tetrazolium bromide assay. The relative cellular viability was calculated as the percentage of absorbance at 570 nm of untreated cells. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$, **** $P < .0001$.

significantly higher expression of both receptors compared with A549 cells ($P < .0001$) (Supplemental Fig. 1). To assess the potential AR ability to counteract CSE-induced EMT, the mRNA expression levels of mesenchymal (vimentin) and epithelial (E-cadherin) markers were analyzed by RT-PCR in both cell lines. CSE exposure led to a marked upregulation of vimentin ($P < .05$ in both cell lines) (Fig. 3, A and C) and a concomitant downregulation of E-cadherin ($P < .05$ in BEAS-2B and $P < .001$ in A549) (Fig. 3, B and D) mRNA expression levels compared with the control (CTR) group.

Although AR alone did not significantly affect the mRNA expression levels of vimentin and E-cadherin, AR pretreatment significantly prevented CSE effects on vimentin and E-cadherin levels ($P < .01$ and $P < .05$ in both cell lines, respectively) by restoring the expression of these markers toward CTR levels (Fig. 3). To further validate these findings, the expression of the above-mentioned markers was detected also by immunofluorescence staining (Fig. 4, A and C; Fig. 5, A and C). Consistent with the RT-PCR results, CSE increased vimentin levels ($P < .01$ in BEAS-2B; $P < .05$ in A549)

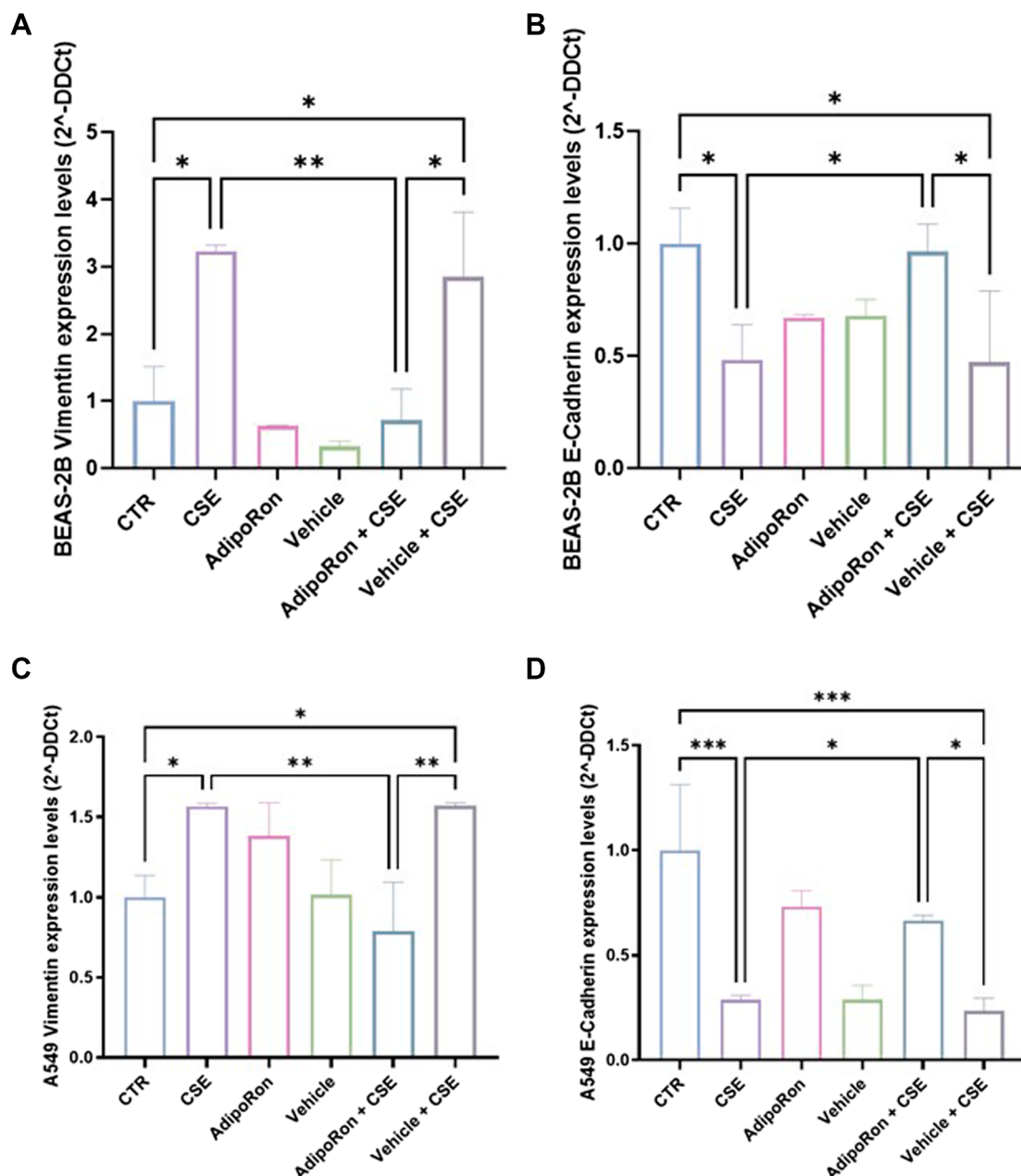


Fig. 3. RT-PCR analysis of EMT marker expression in BEAS-2B and A549 cells. The mRNA expression levels of the mesenchymal marker vimentin (A and C) and the epithelial marker E-cadherin (B and D) were evaluated to assess AR ability to mitigate CSE effects on EMT. Cells were pretreated or not for 2 hours with AR and then exposed to CSE for 24 hours before performing RT-PCR. Data from all RT-PCR experiments and mRNA expression levels were analyzed by normalizing to the endogenous housekeeping gene control applying the comparative $2^{-\Delta\Delta C_t}$ method. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$.

(Fig. 4A; Fig. 5A) and markedly reduced E-cadherin ($P < .001$ in BEAS-2B; $P < .0001$ in A549) (Fig. 4C; Fig. 5C) at the protein level. AR pretreatment mitigated these changes, preserving the epithelial ($P < .05$ in both cell lines) and mesenchymal ($P < .01$ in BEAS-2B; $P < .05$ in A549) balance. Fluorescence imaging showed that CSE ability to promote the transition of epithelial cells toward a mesenchymal phenotype and AR protective role in preventing these alterations (Fig. 4, B and D; Fig. 5, B and D).

3.4. Role of AR in CSE-induced profibrotic response

To investigate AR and CSE effects on profibrotic pathways, the expression of the profibrotic mediators α -SMA and TGF- β , total

collagen deposition, and collagen type I protein was evaluated in both cell lines. As shown in Fig. 6, A and C, α -SMA expression was markedly enhanced following CSE exposure in both cell lines ($P < .01$ vs CTR). This response was significantly blunted by AR pretreatment, which reduced α -SMA levels in BEAS-2B and A549 cells ($P < .0001$ and $P < .05$, respectively) (Fig. 6, A and C). Figure 6, B and D show representative immunofluorescence images of labeled α -SMA. RT-PCR analysis revealed a significant increase in TGF- β mRNA expression following CSE exposure compared with CTR cells ($P < .001$ in both cell lines) (Fig. 6, E and F), indicating the activation of signaling pathways involved in both EMT progression and early fibrotic remodeling. AR pretreatment significantly attenuated this response ($P < .001$ in A549 and $P < .01$ in BEAS-2B),

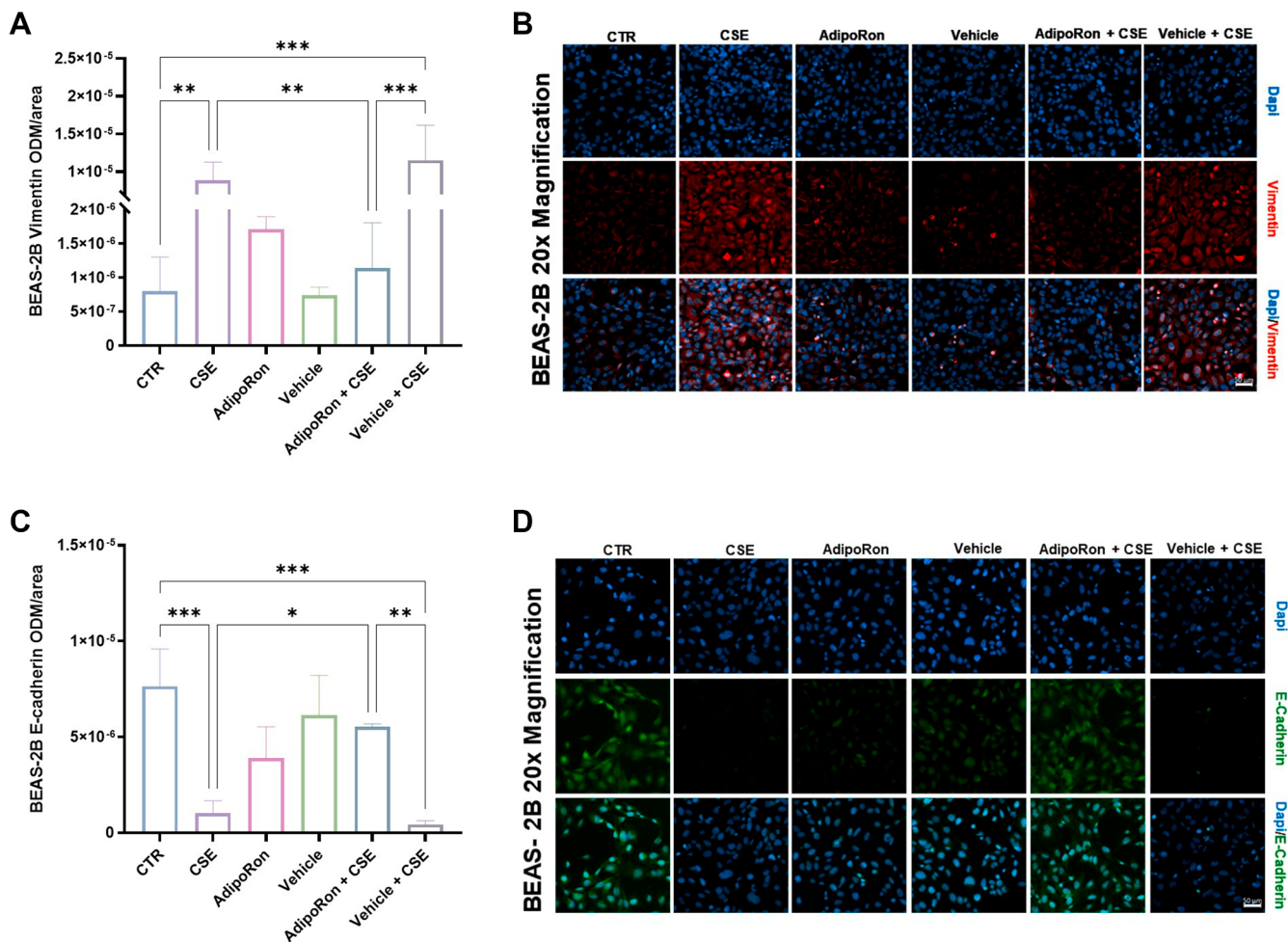


Fig. 4. Immunofluorescence staining of EMT marker expression in BEAS-2B. The expression of the mesenchymal marker vimentin (A) and the epithelial marker E-cadherin (C) was evaluated to assess AR ability to mitigate CSE effects on EMT. Cells were pretreated or not for 2 hours with AR and then exposed to CSE for 24 hours before performing immunofluorescence staining. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$. Representative images of labeled vimentin (B) and E-cadherin (D) proteins in BEAS-2B cells. Scale bar, 50 μ m. Cell fluorescence images were acquired by the Apotome.2 microscope (Zeiss) and analyzed with ImageJ software (NIH).

restoring TGF- β expression toward CTR levels (Fig. 6, E and F). Collagen deposition was then quantified by Sirius Red staining. CSE exposure significantly increased total collagen content compared with untreated cells ($P < .05$ in BEAS-2B and $P < .001$ in A549), while AR pretreatment markedly reduced collagen accumulation ($P < .0001$ in both cell lines) (Fig. 7, A and C), reflecting its ability to attenuate the CSE-induced profibrotic effect. Representative Sirius Red images show collagen content in both cell lines, indicating more intense staining in CSE-treated cells and a substantial reduction following AR exposure (Fig. 6, B and D). COL1A1 protein levels, assessed by immunofluorescence staining, followed a similar trend. Notably, CSE exposure significantly enhanced COL1A1 expression compared with CTR cells ($P < .0001$ in A549 and $P < .01$ in BEAS-2B), whereas AR pretreatment significantly prevented this increase ($P < .001$ in both cell lines) (Fig. 7, E and G). Fluorescence images show these findings, in Fig. 7, F and H.

3.5. AR pretreatment effects on CSE-induced enhancement of cell migratory capacity

BEAS-2B and A549 migration and the wound closure phenomenon were monitored at 4, 12, and 24 hours postscratch. The relative closure area of the wound at the chosen time points was

normalized to the initial wound area of untreated cells, as shown in Fig. 8, A and B. No statistically significant differences were observed in cell migration between experimental groups at 4 (Fig. 8A) and 12 hours (data not shown) for both cell lines. However, at 24 hours, BEAS-2B (Fig. 8A) and A549 (Fig. 8B) cells exposed to CSE showed a significantly reduced scratch area compared with the CTR group ($P < .001$ and $P < .05$, respectively). In contrast, pretreatment with AR led to a significantly residual wound area at 24 hours in both cell lines ($P < .0001$, respectively), reflecting a reduction in cell migratory potential and a slower wound healing process. Figure 8, C and D display the scratch-wound closure over time, highlighting the temporal progression of wound healing. Figure 8, E and F display representative micrographs of scratch areas at 4 and 24 hours.

3.6. AR pretreatment impact on CSE-induced oxidative unbalance

To evaluate whether AR and CSE effects on redox status, intracellular ROS levels were measured in BEAS-2B and A549 cells. CSE exposure markedly increased ROS production ($P < .05$ in BEAS-2B and $P < .001$ for A549) (Fig. 9), compared with untreated controls, confirming its ability to induce oxidative stress. Pretreatment with AR significantly attenuated these effects, demonstrating ROS-

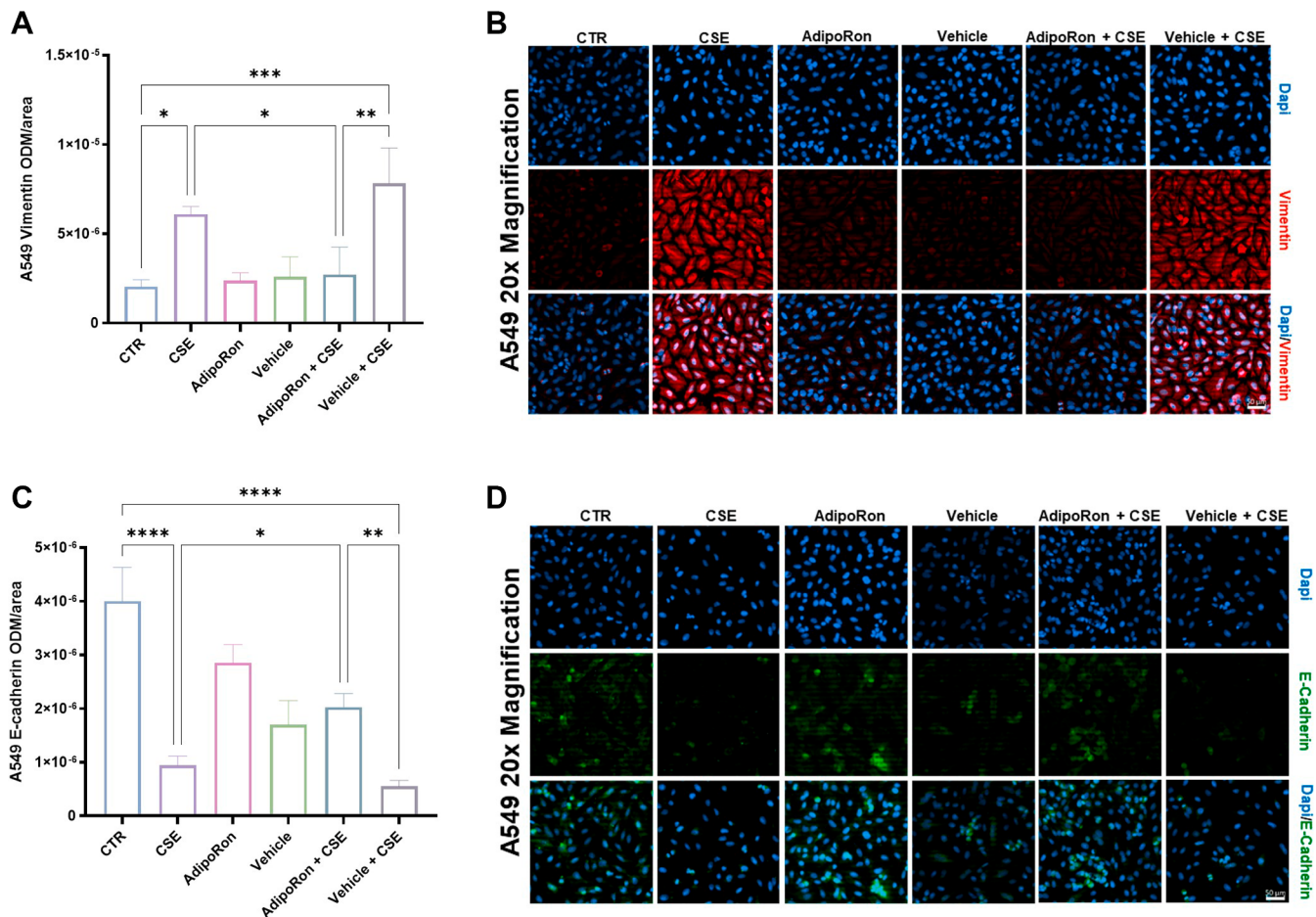


Fig. 5. Immunofluorescence staining of EMT marker expression in A549 cells. The expression of the mesenchymal marker vimentin (A) and the epithelial marker E-cadherin (B) was evaluated to assess AR ability to mitigate CSE effects on EMT. Cells were pretreated or not for 2 hours with AR and then exposed to CSE for 24 hours before performing immunofluorescence staining. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$, **** $P < .0001$. Representative images of labeled vimentin (B) and E-cadherin (D) proteins in A549 cells. Scale bar, 50 μ m. Cell fluorescence images were acquired by the Apotome.2 microscope (Zeiss) and analyzed with ImageJ software (NIH).

scavenging activity by restoring redox balance in BEAS-2B ($P < .0001$) (Fig. 9A) and A549 cells ($P < .05$) (Fig. 9B). Similarly, treatment with the antioxidant control NAC significantly attenuated CSE-induced ROS generation ($P < .0001$ for both cell lines) (Fig. 9) and effectively counteracted H₂O₂-induced oxidative stress in BEAS-2B ($P < .0001$) and A549 ($P < .05$) (Fig. 9). Overall, these findings support the antioxidant activity of AR against CSE-induced oxidative imbalance.

4. Discussion

4.1. Acrp30 counteracts CS-induced EMT and fibrotic remodeling

This study highlights the role of Acrp30 as a negative modulator of EMT and profibrotic response driven by CS, restoring epithelial marker expression and reducing migratory capacity. A growing body of evidence indicates that CS-induced pathological remodeling of the respiratory epithelium is largely driven by EMT, a dynamic process implicated in airway structural changes and tumor progression.^{17–20} In this context, Acrp30 has been identified as a negative modulator of EMT.³⁵ Moreover, reduced systemic Acrp30 levels have been observed in current smokers compared with never smokers.³⁷ However, its role in counteracting CS-induced EMT in both normal and malignant lung epithelial cells remains poorly understood. Exposure to CSE significantly reduced

the viability of both healthy bronchial epithelial cells and lung adenocarcinoma cell models, accompanied by morphological alterations such as cell rounding and detachment, indicative of cellular stress. Consistent with our findings, different studies reported a similar CSE cytotoxicity of pulmonary cells confirming its ability to compromise epithelial cell integrity.^{18,43,44} EMT has been reported as an active process in CS-associated COPD, where reticular basement membrane hypervascularity and fibrotic remodeling are closely related to ongoing EMT in pulmonary lesions.⁴⁵ CS-induced EMT leads to endothelial cells and epithelial barrier dysfunction, interferes with tissue repair, ultimately leading to airway fibrosis.³¹ In accordance with these evidence, in our study, normal bronchial epithelial cells exposed to CSE displayed a marked downregulation of E-cadherin, a transmembrane protein implicated in epithelial cellular adhesion, and a significant upregulation of vimentin and α -SMA, which are characteristic markers of fibrogenic and mesenchymal cells, confirming EMT profibrotic remodeling induction by CSE. Interestingly, we observed similar EMT-related alterations in the lung adenocarcinoma cell line, supporting CS role as a driving force of lung cancer progression and metastasis.⁴⁶ TGF- β is widely recognized as a central mediator of EMT and fibrotic remodeling, promoting fibroblast activation, extracellular matrix deposition, and epithelial plasticity.⁴⁷ We further demonstrated that CSE exposure significantly increased TGF- β mRNA expression levels in both models, supporting the

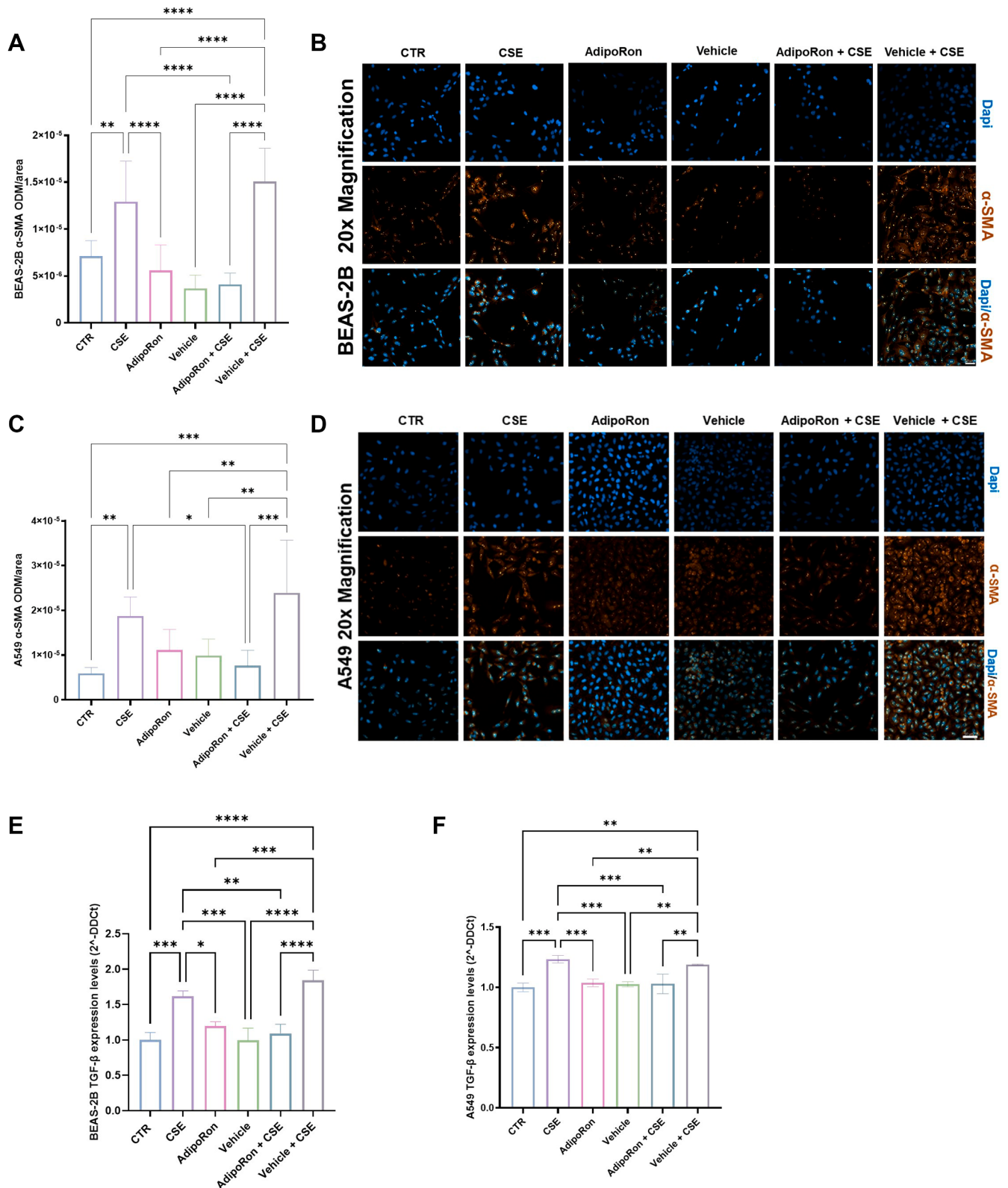


Fig. 6. BEAS-2B and A549 cells were used to further explore AR potential in counteracting CSE-induced effects on the expression of the profibrotic mediators α -SMA (A and C) and TGF- β (E and F) through immunofluorescence staining and RT-PCR, respectively. Cells were pretreated or not for 2 hours with AR and then exposed to CSE for 24 hours before performing immunofluorescence staining and RT-PCR. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$, **** $P < .0001$. Representative images of labeled α -SMA in BEAS-2B (B) and A549 (D) cells. Scale bar, 50 μ m. Cell fluorescence images were acquired by the Apotome.2 microscope (Zeiss) and analyzed with ImageJ software (NIH).

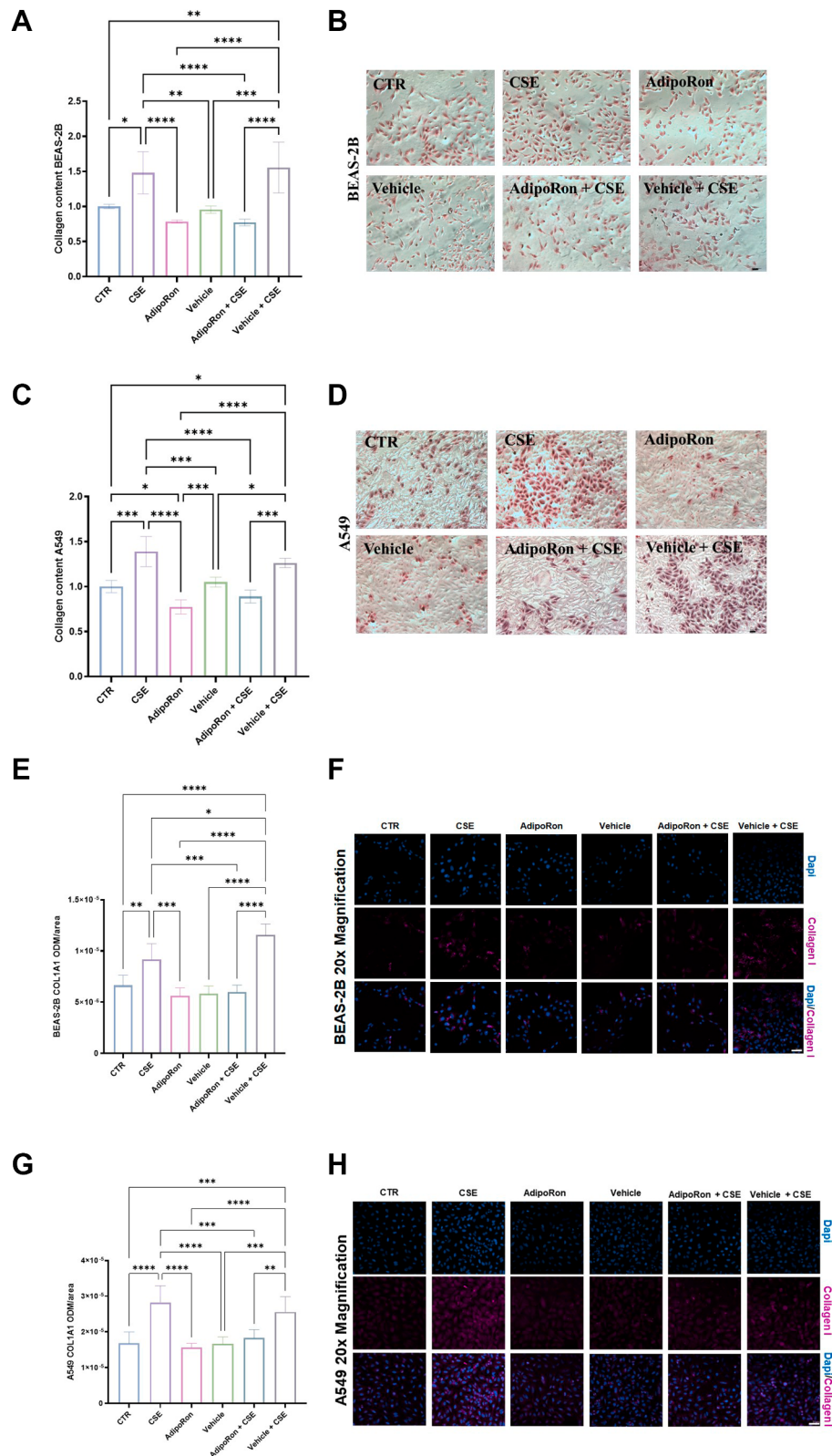


Fig. 7. To investigate AR and CSE effects on profibrotic pathways, the expression of total collagen deposition (A and C) and collagen type I protein (E and G) was evaluated in both cell lines through Sirius Red staining and immunofluorescence staining, respectively. Cells were pretreated or not for 2 hours with AR and then exposed to CSE for 24 hours before performing Sirius Red staining and immunofluorescence staining. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$, **** $P < .0001$. Representative images of labeled total collagen in BEAS-2B (B) and A549 (D) cells. Scale bar, 100 μ m. Representative images of labeled COL1A1 in BEAS-2B (F) and A549 (H) cells. Scale bar, 50 μ m. Cell fluorescence images were acquired by the Apotome.2 microscope (Zeiss) and analyzed with ImageJ software (NIH).

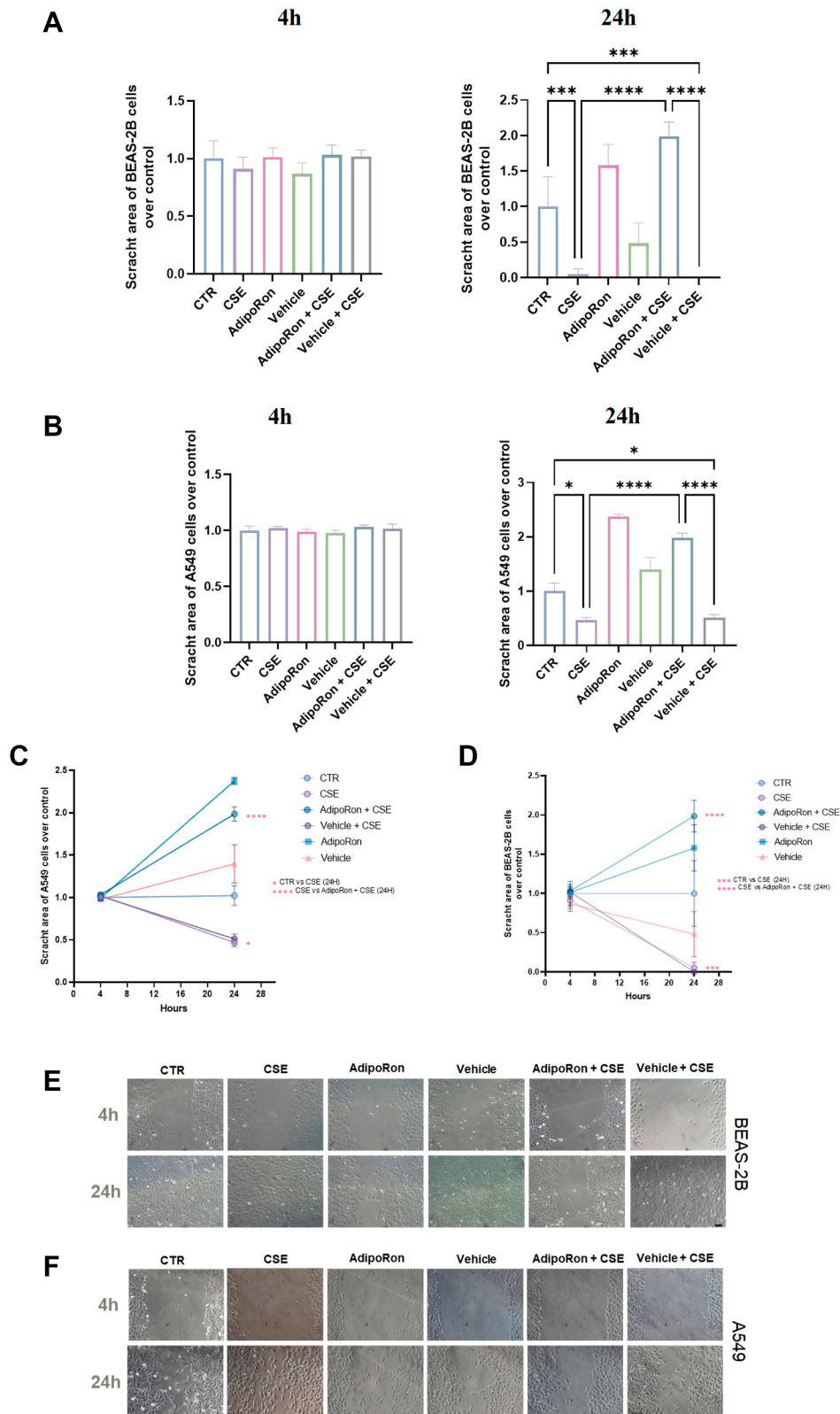


Fig. 8. BEAS-2B and A549 cells were used to further explore AR potential in counteracting CSE-induced effects on cell migration through the wound-healing assay. The monolayer was scratched using a tip and washed with PBS to remove detached cells. Then the cells were pretreated or not for 2 hours with AR and then exposed to CSE for 24 hours. The relative closure area of the wound at the chosen time points was normalized to the initial wound area of untreated BEAS-2B (A) and A549 cells (B). Scratch-wound closure over time of BEAS-2B (C) and A549 (D). Representative images of BEAS-2B (E) and A549 (F) scratch profiles in all groups at 4 and 24 hours. Scale bar, 100 μ m. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$, **** $P < .0001$.

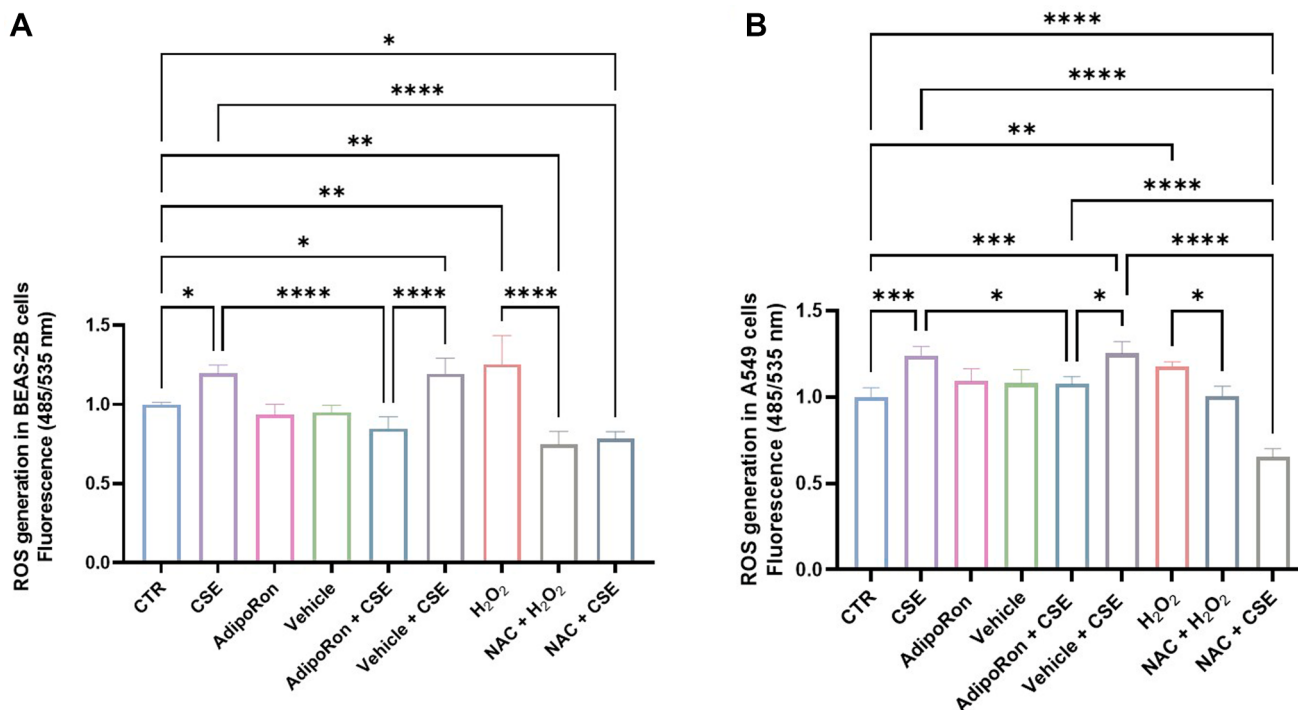


Fig. 9. BEAS-2B (A) and A549 cells (B) were used to assess AR and CSE effects on redox balance. The cells were pretreated or not for 2 hours with AR 10 $\mu\text{g}/\text{mL}$ and then exposed to 2% or 10% CSE for 24 hours. Following treatments, ROS generation was determined by the 2',7'-dichlorodihydrofluorescein diacetate fluorometric assay. The relative ROS production was normalized to the fluorescence intensity of untreated cells. Data are representative of 3 independent experiments, and values are expressed as mean $\pm$ SD. The statistical test used in these analyses was one-way ANOVA followed by Tukey multiple comparison test. * $P < .05$, ** $P < .01$, *** $P < .001$, **** $P < .0001$.

activation of signaling pathways involved not only in EMT progression but also in fibrotic remodeling. Importantly, the activation of these profibrotic pathways was accompanied by a significant increase in total collagen and COL1A1, indicating that CSE-induced EMT is associated not only with phenotypic epithelial plasticity but also with the establishment of a fibrotic microenvironment. AdipoRon pretreatment effectively mitigated these CSE-induced changes, reducing TGF- β expression, preserving the epithelial phenotype, attenuating the mesenchymal shift, and limiting collagen accumulation, suggesting that its protective activity extends beyond EMT modulation and includes the suppression of downstream fibrotic remodeling. Different authors reported that AdipoRon suppresses pro-EMT transcription factors, restoring E-cadherin and vimentin expression levels and attenuating EMT and fibrotic remodeling.^{48–51} It has been documented that AdipoR1 knockdown in breast cancer cells leads to enhanced invasion and EMT, suggesting that AdipoR1 and consequently Acrp30 could exert an inhibitory role on EMT and invasive processes.⁵² Consistently, previous studies have shown that AdipoRon suppresses both phosphorylated and total nuclear factor κB , a key driver of mesenchymal switch, and EMT-related transcription factors counteracting phenotypic transition and cell invasion processes.⁵³ Furthermore, Li et al⁵⁴ demonstrated that AdipoRon alleviates hypertension-induced EMT and renal fibrosis, thereby supporting it as a broad EMT inhibitor across different epithelial models.

4.2. Acrp30 prevents CS-induced cell migration and invasive phenotype

Cell migration is a fundamental process involved in physiological as well as pathological conditions, including immune responses, tissue regeneration, and tumor progression. Evaluating

cell migration is a crucial functional marker of a mesenchymal phenotype, especially in the context of EMT. Indeed, during EMT epithelial cells lose their intercellular contact, and acquire increased motility and invasiveness.⁵⁵ In our models, exposure to CSE was found to enhance the migratory capacity of both normal human bronchial epithelial cells and lung adenocarcinoma cells, providing functional evidence of CS ability to promote a mesenchymal-like phenotype characterized by increased invasive potential. These findings are consistent with previous studies where CS derived, via redox-dependent modifications, the transition toward a mesenchymal phenotype with enhanced migratory capacity.^{46,56} Similarly, CS exposure has been reported to promote anchorage-independent growth, as well as migration, invasion, and morphological alterations in mammary epithelial and breast cancer cells.⁵⁷ Interestingly, in our study, AdipoRon pretreatment reduced the cells' migratory potential and induced a slower wound healing process in both normal and cancerous cell lines. In agreement with our results, literature data showed Acrp30 ability to suppress proliferation, migration, invasion, and colony formation in thyroid cancer cells, NSCLC cells, and in healthy human colorectal cell lines, highlighting its broad impact on tumor cell behavior.^{35,58–60}

4.3. Acrp30 protects against CS-induced effects by mitigating oxidative stress

It is fundamental to consider that ROS play an important role as intracellular signaling molecules involved in various pathological conditions, including EMT activation.^{61,62} Chronic exposure to CS perpetuates oxidative damage and inflammatory responses in pulmonary tissues, resulting in ongoing tissue injury and aberrant repair processes that contribute to airway structural changes.^{31,63–65} In line with this evidence, our data showed that

CSE altered the redox balance of normal BEAS-2B and adenocarcinoma A549 cells, leading to a significant increase in ROS levels. The oxidative stress induced by CSE may thus account, at least in part, for the EMT-related alterations observed in our models. This aligns with previous reports documenting that CSE-induced EMT has been associated with oxidative stress-mediated mechanisms. CSE exposure leads to excessive production of ROS, either directly from smoke components or indirectly through activation of intracellular oxidant-generating systems such as NADPH oxidases.⁶⁶ It has been demonstrated that a redox imbalance can trigger redox-sensitive transcriptional reprogramming characteristic of EMT, ultimately leading to alteration in E-cadherin and vimentin markers expression.^{61,67} Importantly, we documented that AdipoRon pretreatment confers a protective effect against CSE-induced oxidative stress, attenuating ROS accumulation. Notably, the comparable antioxidant effects observed with NAC and AdipoRon further support the hypothesis that the protective action of AR is, at least in part, mediated through attenuation of ROS-dependent signaling pathways. AdipoRon capacity to mitigate oxidative damage is probably linked to its ability to regulate antioxidant enzymes activity through Acpr30 receptor activation. Indeed, different studies showed that Acpr30 stimulates the increase of superoxide dismutase, catalase, and glutathione peroxidase activity,^{68–70} thereby preserving redox homeostasis and preventing ROS-dependent EMT induction.^{48–50} As documented in literature by Park et al,⁷¹ overexpression of AdipoR2 reduces oxidative stress by increasing peroxisome proliferator-activated receptor- α activity and catalase expression, exerting protective effects against inflammation and fibrotic progression. This mechanism could explain the observed preservation of epithelial integrity following AdipoRon pretreatment, highlighting Acpr30 potential as a regulator of oxidative stress-mediated epithelial plasticity.

4.4. Conclusions

Collectively, these findings highlighted the protective role of AR in counteracting CS-induced EMT onset and fibrosis, by mitigating oxidative stress, preserving epithelial-mesenchymal balance, and reducing migratory capacity in healthy epithelium as well as in tumor progression. One limitation of this study is the lack of investigation into additional molecular mechanisms and the relative contribution of each receptor by which AR could inhibit CS-induced EMT and fibrosis. Nevertheless, these results suggest that AR may offer promising therapeutic potential in preventing airway remodeling and disease progression in smoking-related conditions.

Abbreviations

Acpr30, adiponectin; AdipoR1, adiponectin receptor 1; AR, AdipoRon; ATCC, American Type Culture Collection; COL1A1, collagen type I $\alpha 1$ chain; COPD, chronic obstructive pulmonary disease; CS, cigarette smoke; CSE, cigarette smoke extract; EMT, epithelial-mesenchymal transition; NAC, N-acetylcysteine; NSCLC, non-small cell lung carcinoma; ROS, reactive oxygen species; RT-PCR, reverse transcription polymerase chain reaction; α -SMA, α -smooth muscle actin; TGF- β , transforming growth factor- β .

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Conflict of interest

The authors declare no conflicts of interest.

Data availability

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

CrediT author contribution statement

Francesca Panico: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Writing – Original draft. **Giuseppe Spaziano:** Conceptualization, Formal analysis, Investigation, Methodology, Writing – Original draft. **Ersilia Nigro:** Data curation, Formal analysis, Investigation, Methodology, Validation. **Davida Mirra:** Conceptualization, Data curation, Project administration, Software, Supervision, Validation, Writing – Review and Editing. **Renata Esposito:** Data curation, Formal analysis, Investigation, Validation, Visualization. **Eleonora Caiazzo:** Formal analysis, Investigation. **Aurora Daniele:** Supervision, Validation, Writing – Review and Editing. **Bruno D'Agostino:** Conceptualization, Funding acquisition, Supervision, Validation, Writing – Review and Editing.

Supplemental material

This article has supplemental material available at jpet.aspetjournals.org.

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