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Airborne particulate matter (PM_{2.5}) triggers ocular hypertension and glaucoma through pyroptosis

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Abstract

Background: Particulate matter (PM) is strongly linked to human health and has detrimental effects on the eye. Studies have, however, focused on the ocular surface, with limited research on the impact of PM_{2.5} on intraocular pressure (IOP).

Methods: To investigate the impact of PM_{2.5} on IOP and the associated mechanism, C57BL/6 mouse eyes were topically exposed to a PM_{2.5} suspension for 3 months, and human trabecular meshwork (HTM) cells were subjected to various PM_{2.5} concentrations in vitro. Cell viability, NLRP3/caspase-1, IL-1 β , and GSDMD expression, reactive oxygen species (ROS) production and cell contractility were measured by western blot, ELISA, cell counting kit-8, ROS assay kit or a cell contractility assay. ROS scavenger N-acetyl-L-cysteine (NAC) and caspase-1 inhibitor VX-765 were used to intervene in PM_{2.5}-induced damages.

Results: The results revealed that the IOP increased gradually after PM_{2.5} exposure, and upregulations of the NLRP3 inflammasome, caspase-1, IL-1 β , and GSDMD protein levels were observed in outflow tissues. PM_{2.5} exposure decreased HTM cell viability and affected contraction. Furthermore, elevated ROS levels were observed as well as an activation of the NLRP3 inflammasome and downstream inflammatory factors caspase-1 and IL-1 β . NAC improved HTM cell viability, inhibited the activation of the NLRP3 inflammasome axis, and HTM cell contraction by scavenging ROS. VX-765 showed similar protection against the PM_{2.5} induced adverse effects.

Conclusion: This study provides novel evidence that PM_{2.5} has a direct toxic effect on intraocular tissues and may contribute to the initiation and development of ocular hypertension and glaucoma. This occurs as a result of increased oxidative stress and the subsequent induction of NLRP3 inflammasome mediated pyroptosis in trabecular meshwork cells.

Keywords: PM_{2.5}, NLRP3, Intraocular pressure, Human trabecular meshwork cell, Pyroptosis

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Highlights

- PM_{2.5} exposure increases intraocular pressure accompanied by intraocular tissue pyroptosis
- PM_{2.5} triggers oxidative stress which activates pyroptosis in human trabecular meshwork cells
- NLRP3 inflammasome mediates PM_{2.5} induced trabecular meshwork pyroptosis

Introduction

Epidemiological and experimental studies suggest that particulate matter (PM), especially PM_{2.5} (aerodynamic diameter $\leq 2.5 \mu\text{m}$), is strongly associated with respiratory, cardiovascular, metabolic, and even emotional disorders [1]. Although the eyes are in direct contact with the external environment, studies on the impact of PM_{2.5} on ocular health remain scarce [2]. According to previous studies, PM causes ophthalmic diseases such as conjunctivitis, keratitis, and dry eye syndrome [3–6]. However, these PM-associated studies focused on the ocular surface, with little consideration of the particles' potential to penetrate the human cornea and affect tissues within the eye, and their involvement in the initiation and development of the intraocular diseases.

The recent increase in air pollution may be an important cause of, for example, glaucoma [7]. An epidemiological study, for the first time, presented a relationship between long-term air pollution and intraocular pressure (IOP) elevation [7]. Other studies have demonstrated the association between PM_{2.5} exposure and glaucoma, without any link to IOP elevation, but through neurotoxicity and vascular dysfunction in the retina [8, 9]. Therefore, conducting investigations to elucidate the link between PM_{2.5} and glaucoma as well as the associated underlying mechanisms is of relevance.

The production and outflow of aqueous humor dynamically maintain IOP. Outflow resistance is reported in the juxtacanalicular tissue region, where adjacent trabecular meshwork (TM) and Schlemm's canal endothelial cells regulate aqueous humor outflow [10]. Cellular or extracellular stimuli such as oxidative stress can damage the TM, causing inflammation and cell death, thereby causing IOP elevation and, eventually, glaucoma [11, 12].

The current study aimed to investigate the association between PM_{2.5} and ocular hypertension and to elucidate the mechanisms underlying this relationship. This study was incentivized by the unintentional discovery that topical application of fluorescent mock PM_{2.5} to the eye caused deposition in outflow tissues including the iris, ciliary body, and TM. These tissues are vital for regulating aqueous humor outflow resistance and IOP [10]. The current findings suggest that, apart from its

involvement in ocular surface disease, PM_{2.5} pollution also affects tissues inside the eye, possibly participating in the development of intraocular diseases such as glaucoma.

Results

The size distribution of the PM

The dynamic light-scattering of the PM suspension was analyzed and the result revealed PM with sizes ranging between 712 nm and 1280 nm. 99.95% of the particles were less than 1280 nm (Fig. 1a).

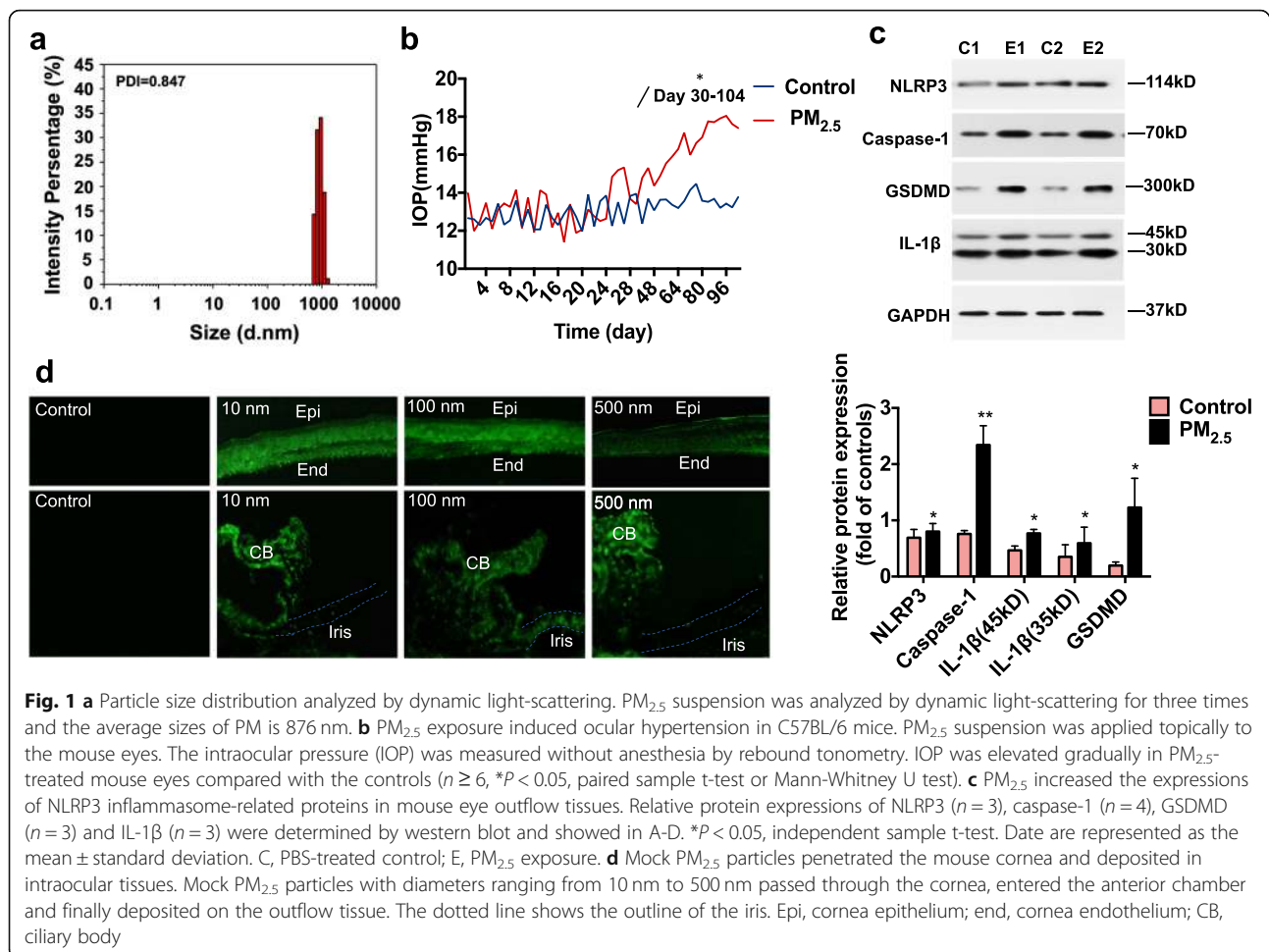
PM_{2.5} exposure induced ocular hypertension in C57BL/6 mice

Elevated IOP values were observed in mouse eyes exposed to PM_{2.5} from day 25 to 104. During this period, the average IOP elevation was 2.6 mmHg, with a maximum of 4.7 mmHg observed on day 92 ($n = 7$, $P < 0.05$, Fig. 1b). From day 1 to day 24, IOP values for both eyes (treated and controls) were similar. On day 25, IOP values of PM_{2.5}-treated eyes significantly increased relative to those of phosphate-buffered saline (PBS)-treated eyes (14.8 ± 0.6 mmHg, $n = 10$ vs 12.3 ± 0.8 mmHg, $n = 7$, $P < 0.05$). Elevated IOP was sustained for 3 days, followed by 2 days of normalization. Thereafter, IOP values increased again and remained consistently higher compared to the values of control eyes. IOP elevation was steadily significant from day 30 onwards, with IOP values of 14.9 ± 1.4 mmHg and 12.4 ± 1.1 mmHg for PM_{2.5}-treated eyes and PBS-treated eyes, respectively (PBS treated, $n = 10$; PM_{2.5}-treated, $n = 10$; $P < 0.05$). Our data demonstrate that PM_{2.5} exposure adversely affected IOP, thus representing a risk factor for glaucoma.

Characteristic features of pyroptosis in outflow tissues following PM_{2.5} exposure

To understand the underlying mechanism of PM_{2.5}-related ocular hypertension, we measured the levels of proteins associated with the classic pyroptosis pathway (NLRP3/caspase-1/IL-1 β /GSDMD) in aqueous humor outflow tissue (Fig. 1c). NLRP3 expression increased 1.2-fold in outflow tissue of PM_{2.5}-treated eyes compared to that of control eyes ($n = 3$, $P < 0.05$). Furthermore, levels of the NLRP3 downstream proteins caspase-1 and GSDMD increased 3.1- and 6.3-fold, respectively ($n = 4$, $P < 0.05$). In addition, total IL-1 β was 1.7 times higher in outflow tissue from PM_{2.5}-treated eyes compared to contralateral controls, whereas cleaved IL-1 β increased 5.5-fold ($n = 3$, $P < 0.05$). These observations suggested that PM_{2.5} caused tissue injury via pyroptosis of cells in outflow tissue of the eye.

To understand PM_{2.5} entry into the anterior chamber, fluorescent mock PM_{2.5} tracers were topically applied to the eye. Particles with diameters from 10 to 500 nm



passed through the cornea into the anterior chamber and were mainly deposited in outflow tissue, with ciliary body deposition being the most pronounced (Fig. 1d). Fluorescent tracers distributions revealed that some PM_{2.5} crossed the cornea and entered the eye, thereby affecting intraocular function.

PM_{2.5} exposure triggered pyroptosis in HTM cells

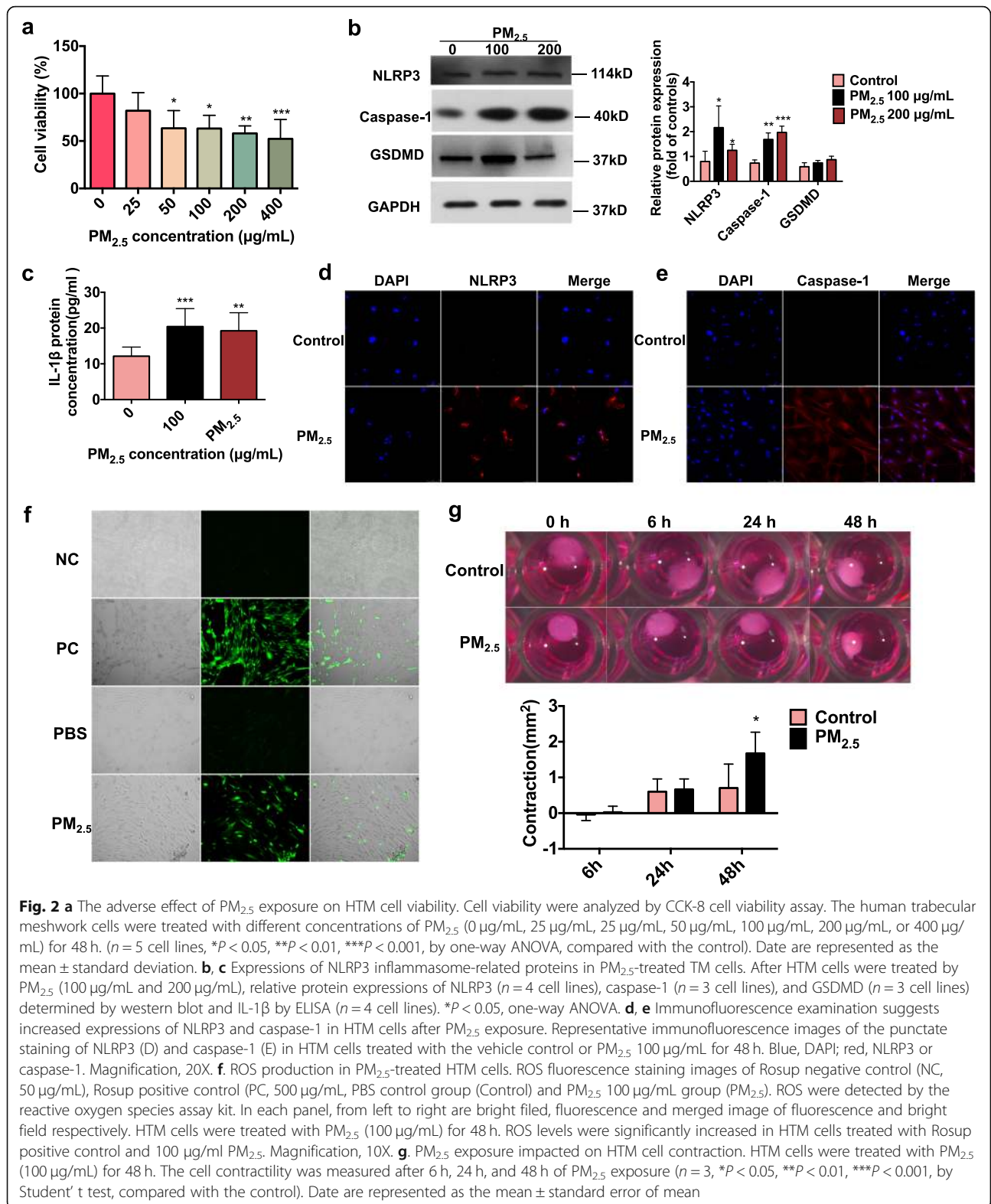
To further verify the relationship between PM_{2.5} and pyroptosis in order to explain the effect of PM_{2.5} on IOP, we utilized HTM cells as an in vitro model. As previously mentioned, HTM cells are vital components of outflow tissue, which is the main site of outflow resistance and a key IOP determinant. We initially found that PM_{2.5} was toxic to HTM cells even at 25 $\mu\text{g}/\text{mL}$. When HTM cells were exposed to different PM_{2.5} concentrations for 48 h, a concentration-dependent decrease of cell viability was observed. HTM cell viability was reduced by 18, 37, 37, 42, and 48% after treatment with concentrations of 25, 50, 100, 200, and 400 $\mu\text{g}/\text{mL}$ PM_{2.5}, respectively ($n = 5$ cell lines, $P \leq 0.05$, vs. controls groups, Fig. 2a). Similarly, HTM cell viability was

reduced by 8, 17, 25, 23, and 37% after treatment with concentrations of 25, 50, 100, 200, and 400 $\mu\text{g}/\text{mL}$ PM_{2.5} from Shanghai, respectively ($n = 5$ cell lines, $P \leq 0.05$, vs. controls groups, Fig. S1A).

We further observed that NLRP3 protein levels were upregulated 2.7- and 1.6-fold after HTM cells were treated with 100 and 200 $\mu\text{g}/\text{mL}$ of PM_{2.5}, respectively, for 48 h ($n = 4$ cell lines, $P < 0.05$, Fig. 2b). Quantitative real-time polymerase chain reaction (RT-qPCR) results revealed that 48 h of PM_{2.5} exposure increased the relative expression of NLRP3 mRNA 1.3- and 1.4-fold in 100 $\mu\text{g}/\text{mL}$ and 200 $\mu\text{g}/\text{mL}$ PM_{2.5}-treated HTM cells ($n = 3$ cell lines, $P < 0.05$, Fig. S2A), respectively.

Moreover, the protein expression of caspase-1 increased by factors of 2.3 and 2.6 in PM_{2.5}-treated HTM cells compared to controls (100 $\mu\text{g}/\text{mL}$ and 200 $\mu\text{g}/\text{mL}$, $n = 3$ cell lines, $P < 0.01$, Fig. 2b). However, the relative expression of caspase-1 mRNA decreased 0.8-fold in PM_{2.5}-treated cells (100 $\mu\text{g}/\text{mL}$ or 200 $\mu\text{g}/\text{mL}$, $n = 3$ cell lines, $P < 0.05$, Fig. S2A).

Activation of caspase-1 facilitates cleavage of GSDMD, release of the N-terminal GSDMD fragment, and



maturation of IL-1β, leading to inflammatory cell death [13]. Notably, in control cell supernatant, the average protein concentration of IL-1β was 12.13 ± 2.56 pg/mL,

whereas in PM_{2.5}-treated cell culture supernatant, IL-1β levels of 20.42 ± 5.02 and 19.23 ± 5.06 pg/mL were observed after 100 and 200 µg/mL PM_{2.5} treatment,

respectively ($n = 8$ cell lines, $P < 0.05$, Fig. 2b). IL-1 β mRNA expression was also higher in PM_{2.5}-treated HTM cells compared with controls by factors of 15.0 and 15.4 for treatment concentrations of 100 and 200 $\mu\text{g}/\text{mL}$, respectively ($n = 6$ cell lines, $P < 0.05$, Fig. S2A). The protein expression of GSDMD increased by factors of 1.4 and 1.2 in PM_{2.5}-treated HTM cells compared to controls (100 $\mu\text{g}/\text{mL}$ and 200 $\mu\text{g}/\text{mL}$, $n = 3$ cell lines, $P > 0.05$, Fig. 2b).

To further characterize PM_{2.5}-induced pyroptosis of HTM cells, we performed immunofluorescence staining for NLRP3 and caspase-1 after 100 $\mu\text{g}/\text{mL}$ PM_{2.5} exposure for 48 h. As shown in Fig. 2d and e, NLRP3 and caspase-1 protein levels increased in PM_{2.5}-exposed HTM cells. PM samples from a different location (Shanghai) were also used to test the above-mentioned molecules, which confirmed increased expressions of NLRP3 and caspase-1 by 1.3-fold and 1.0-fold in 100 $\mu\text{g}/\text{mL}$ PM_{2.5}-treated HTM cells and 2.8-fold and 2.4-fold in 200 $\mu\text{g}/\text{mL}$ PM_{2.5}-treated HTM cells, respectively ($n = 3$ cell lines, $P < 0.05$, Fig. S1B). GSDMD was upregulated by 1.4-fold, but was not statistically significant which may be due to the small sample sizes ($n = 3$ cell lines, $P > 0.05$, Fig. S1B).

Taken together, these observations revealed the cellular and molecular mechanisms through which PM_{2.5} increased IOP. PM_{2.5} caused cellular toxicity and pyroptosis by activating NLRP3, caspase-1, and GSDMD. This suggests that pyroptosis is an important mediator of ocular cell damage and the decreased cell viability induced by PM_{2.5}.

Increased ROS production and enhanced HTM cell contraction following PM_{2.5} exposure

Prior studies demonstrated that ROS elevation is essential for inflammasome activation [4, 14]. Therefore, in order to verify the role of ROS in PM_{2.5}-induced pyroptosis, we measured the changes in intracellular ROS levels in HTM cells after treatment with PM_{2.5} from different locations (100 $\mu\text{g}/\text{mL}$) for 48 h. Representative micrographs of the DCF fluoresce-labeled cells indicate that the PM_{2.5} from different locations elevated ROS production in HTM cells (Fig. 2f, S1C). Furthermore, the contractility of PM_{2.5}-treated HTM cells increased 2.4-fold compared to controls after 48 h of treatment ($n = 3$, $P < 0.05$, Fig. 2g), probably due to the increase in ROS levels. Combined with the results presented in Fig. 2, it can be inferred that PM_{2.5}-related ROS production likely reduced cell viability and caused pyroptosis. However, further pharmacological experiments are required to test this hypothesis.

Role of ROS and caspase-1 in PM_{2.5}-induced HTM cell pyroptosis

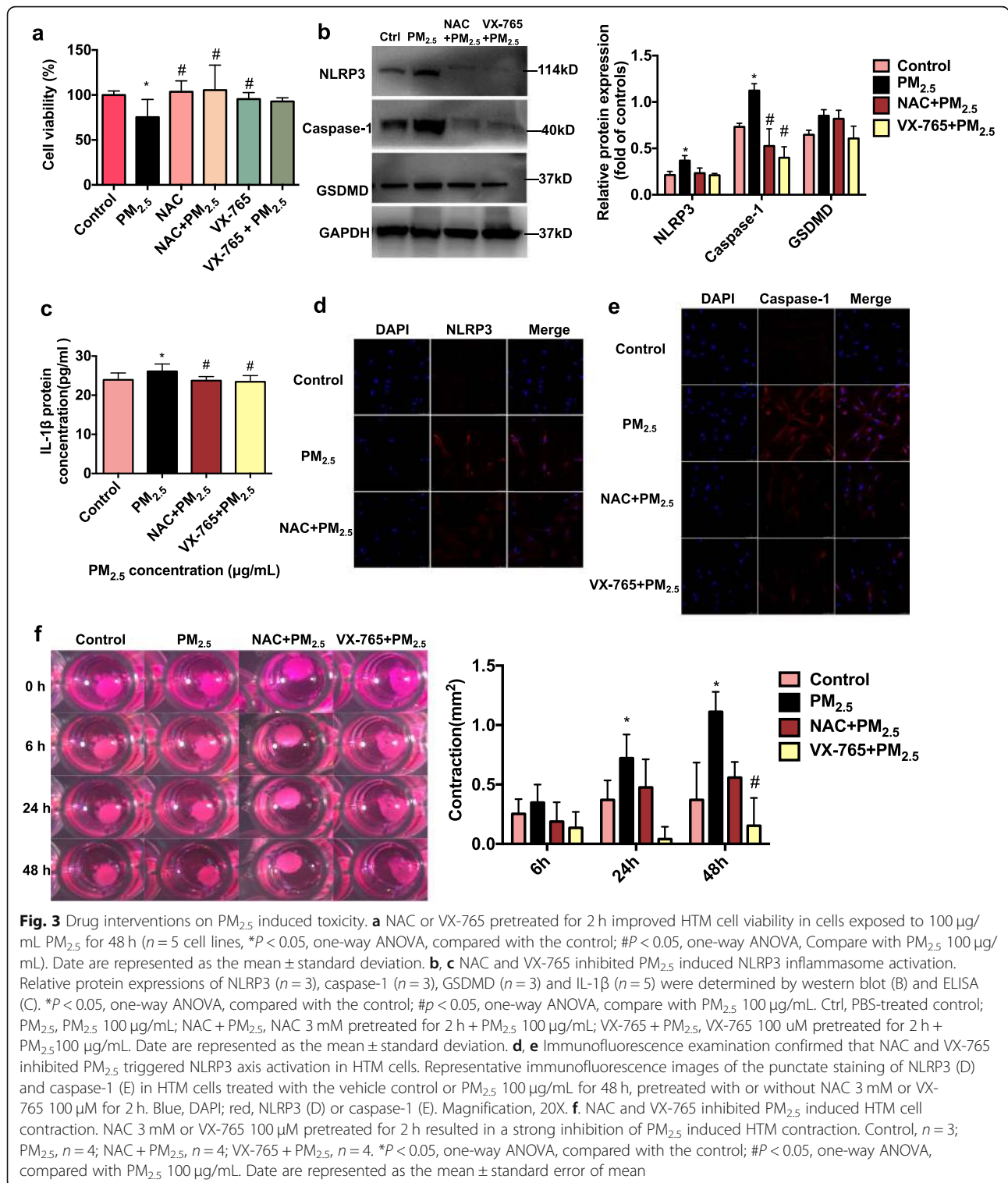
To further assess if PM_{2.5}-induced pyroptosis of HTM cells is triggered by ROS and dependent on caspase-1, we conducted ROS and caspase-1 inhibitory experiments. The ROS scavenger N-acetyl-L-cysteine (NAC) and the caspase-1 selective inhibitor belnacasan (VX-765) were tested for their potential in preventing/alleviating PM_{2.5}-induced cell damage. Figure 3a reveals that PM_{2.5} exposure (100 $\mu\text{g}/\text{mL}$) for 48 h lowered HTM cell viability by 25% ($n = 5$ cell lines, $P < 0.05$), while pretreatment with NAC (3 mM) markedly increased cell viability by 40% ($n = 5$, $P < 0.05$). Furthermore, NAC (3 mM) pretreatment reduced ROS production in HTM cells treated with PM_{2.5} from different locations (Fig. S1C, S2B). Similarly, after VX-765 (100 μM) pretreatment, the viability of HTM cells exposed to PM_{2.5} increased by 23% ($n = 5$, $P > 0.05$). Cell viability was similar in HTM cells treated with NAC, VX-765, and the drug vehicle (Fig. 3a, Fig. S2C). Consistent with these results, NAC and VX-765 significantly inhibited PM_{2.5}-induced HTM cell contraction (Fig. 3f). These results suggest that the ROS scavenger and caspase-1 inhibitor may help prevent PM_{2.5}-induced cell damage.

Our results further demonstrated that NAC and VX-765 significantly downregulated the expressions of the NLRP3, caspase-1, GSDMD, and IL-1 β (Fig. 3b and c). Figure 3d and e showed that NAC inhibited NLRP3 and caspase-1 expressions in PM_{2.5}-exposed HTM cells, and a distinct decrease in caspase-1 was observed in the HTM cells treated with VX-765 after PM_{2.5} exposure (Fig. 3e).

Discussion

This is the first study to demonstrate that PM_{2.5} exposure leads to ocular hypertension and glaucoma by inducing cell pyroptosis and inflammation in intraocular tissues responsible for controlling IOP. Reducing ROS production or inhibiting caspase-1 prevented PM_{2.5}-induced inflammation and pyroptosis of HTM cells.

Various eye diseases such as dry eye syndrome, conjunctivitis, and keratitis are attributed to air pollution, especially PM_{2.5} pollution, according to epidemiological and experimental studies [15–18]. PM_{2.5} exposure is also linked to high blood pressure [19], a condition with a pathological mechanism resembling that of high IOP [20, 21]. For example, oxidative stress promotes vascular aging and damage, which contributes to hypertension [22]. Similarly, oxidative stress induces TM cell damage, which impairs aqueous humor drainage, thereby causing ocular hypertension [23]. However, PM_{2.5} has not yet been reported to induce ocular hypertension. Jama et al., however, suggested that environmental black carbon exposure may represent a risk factor for increased IOP in



individuals susceptible to other biological oxidative stressors [7]. Consistent with previous studies [7–9], we observed that topical application of PM_{2.5} suspension (1 mg/mL) can cause ocular hypertension in mice (Fig. 1b). To the best of our knowledge, the relationship between

PM_{2.5} and IOP-related disease was unproven in vivo, and this study is the first to report the association between PM_{2.5} and IOP changes in mice.

The classical pyroptosis pathway is central to PM_{2.5}-induced injury. Previous studies have reported PM_{2.5}-

induced NLRP3 inflammasome activation and ocular injury in vivo and in vitro [16, 24–32]. The NLRP3 inflammasome is a vital component of sterile- and infection-triggered inflammation as well as of the immune responses to various diseases [33]. Classical pyroptosis is mediated by the NLRP3 inflammasome, and caspase-1 activation promotes the cleavage of pro-IL-1 β and GSDMD [34]. Previous studies have revealed that NLRP3 inflammasome activation greatly contributes to cardiovascular, neurological, and lung disease development [26–32]. In addition, ROS has been reported to activate the NLRP3 inflammasome in environment-induced dry eye syndrome and conjunctivitis, and a significant increase in inflammatory factors, such as IL-1 β , was observed [15–18]. However, whether the NLRP3 inflammasome participates in PM-induced injury in outflow tissues has been poorly studied. Consistent with previous studies, our results revealed an increase in NLRP3 protein levels in outflow tissues of PM_{2.5} topically-treated mouse eyes (Fig. 1c). We further demonstrated the PM_{2.5}-induced upregulation of the caspase-1, which indicates NLRP3 inflammasome activation. Increased IL-1 β protein and cleaved IL-1 β were also observed (Fig. 1c). In addition, most of the PM used in the experiments was less than 2.5 μ m in size by DLS detection (Fig. 1a) which was consistent with fluorescent PM_{2.5} tracer experiments that some PM_{2.5} could cross the cornea and entered the eye (Fig. 1d). These findings suggest that the NLRP3/caspase-1/IL-1 β axis is active in PM_{2.5}-induced ocular hypertension. Since TM tissue is vital for the regulation of IOP, and their damage is closely associated with increased aqueous outflow resistance and IOP elevation [35], we suggest that HTM cells undergo pyroptosis during PM_{2.5}-induced ocular hypertension.

TM cells are involved in IOP regulation, and PM_{2.5} reduced the viability of the HTM cells by triggering cell pyroptosis. It has been shown that the malfunction of HTM induced by oxidative stress or vascular damage increases IOP [36]. PM_{2.5} carries toxic components, such as metal, nitrate, sulfate and polycyclic aromatic hydrocarbons (PAHs), and a biological fraction (such as endotoxin), and is known to induce cell death in various ways, including autophagy and pyroptosis [37, 38]. In our study, PM_{2.5}-induced pyroptosis and cell dysfunction were observed in HTM cells. The PM_{2.5} used in our study is rich in PAHs and was collected from Lanzhou [39]. When HTM cells were exposed to different PM_{2.5} concentrations (25–400 μ g/mL) for 48 h, cell viability was reduced in a concentration-dependent manner (Fig. 2a, Fig. S1A). PM_{2.5} toxicity was also observed in PM_{2.5}-exposed or diesel exhaust particle-treated human umbilical vein endothelial cells, (HUVECs), hippocampal neuron

cells, bronchial epithelium cells, cornea, and conjunctiva human cell lines [4, 25, 38, 40–42]. According to the published studies [38, 43], PM_{2.5} concentrations from 50 μ g/mL to 10 mg/mL have been used in other cells types, therefore, we selected 100 μ g/mL and 200 μ g/mL of PM_{2.5} to investigate the effect of PM_{2.5} in in vitro toxicology experiments. We further found that PM_{2.5} exposure increased the expressions of NLRP3, caspase-1, IL-1 β , and GSDMD proteins in HTM cells (Fig. 2b–e, Fig. S1B). These findings were consistent with the observations in the in vivo experiments (Fig. 1) and were further verified in the next rescue experiments.

ROS production can trigger NLRP3 inflammasome-associated proteins production and inflammatory responses [4], and high ROS levels are detrimental to HTM cells and have been linked to ocular hypertension in glaucoma studies [35, 44]. In our study, a significant elevation of ROS was observed in HTM cells treated with 100 μ g/mL PM_{2.5} for 48 h (Fig. 2f, Fig. S1C), along with enhanced HTM cell contractility (Fig. 2g). HTM contractility is an important regulator of conductivity and decreases cell permeability and aqueous humor outflow by reducing the size of intercellular spaces, thereby promoting IOP elevation. In contrast, cell relaxation has the opposite effects [45]. The current observations indicate that oxidative stress damage induces HTM cell dysfunction through NLRP3-mediated pyroptosis.

NAC is a well-known ROS scavenger that decreases ROS production [46], while VX-765 is an effective selective inhibitor of caspase-1 with potent anti-inflammatory activity through inhibition of IL-1 β and IL-18 release [47]. Previous studies also reported that NAC can decrease ROS levels and downregulated NLRP3 expression in HUVECs treated with cooking oil fume-derived PM_{2.5} [25]. Nicotine-induced atherosclerosis via ROS/NLRP3-mediated endothelial cell pyroptosis is also prevented by NAC and VX-765 [48]. In our study, NAC and VX-765 were used to further verify the PM_{2.5}-related HTM cell injury mechanism. PM_{2.5} at a concentration of 100 μ g/mL was employed for further mechanistic experiments. Consistent with other studies, our results revealed that NAC (3 mM) or VX-765 (100 μ M) pretreated for 2 h improved HTM cell viability following PM_{2.5} exposure (Fig. 3a). NAC pretreatment efficiently reduced ROS levels and HTM contraction associated with PM_{2.5} exposure (Fig. S1C, S2B, Fig. 3f), inhibiting NLRP3, caspase-1, IL-1 β , and GSDMD activation (Fig. 3b–e). VX-765 pretreatment also resulted in the relaxation of HTM cells following PM_{2.5}-induced contraction (Fig. 3f) and inhibited caspase-1-mediated pyroptosis (Fig. 3b–e). Hence, these results indicate that the PM_{2.5} exposure elevated oxidative stress, which partially enhanced HTM cell contraction and contributed

to an increase in IOP and activation of NLRP3/caspase-1/IL-1 β signaling. These in vitro observations supported the results from experiments in mice where elevated IOP was observed following PM_{2.5} exposure, in parallel to oxidative stress damage and NLRP3/caspase-1-mediated pyroptosis. Together with the results in Fig. 2, our data suggest that PM_{2.5}-induced ROS production further triggered HTM cell pyroptosis, which is the underlying mechanism of ocular hypertension. NAC and VX-765 might be viable protective compounds against PM_{2.5}-induced ocular hypertension.

This study used glass fibre filters (without resin binding) to collect PM samples. Glass filters are non-hygroscopic and are extremely difficult to ash by chemicals or heat thus are increasingly used for PM sampling. A previous study reported a loss of filter mass after PM extraction and the blank filters control caused an increase in cytokine release from macrophages. In our study, we did not observe any visible breakage or particle dissociation from the glass fibre filters during the PM extraction process. There was a slight change in the filter weights before PM collection and after a standard extractions procedure used in our study (Supplement Table S1). Nevertheless, the glass fibre toxicity on TM cells was evaluated (Fig. S3A). Assuming 5–30% glass fibre contamination, a dose-response relationship was established which showed little cytotoxicity. NLRP3 and caspase-1 expression in TM cells was not affected by 30% glass fibre contamination (Fig. S3B). This is consistent with what has been reported using human mesothelial cells, in that, NLRP3 inflammasome cannot be activated by glass beads [49].

Topical applications of PM_{2.5} (1 mg/mL) caused corneal injury according to our previous study as well as other in vivo studies [43, 50, 51]. The concentration was 5–10 times higher than the concentration used for in vitro experiments. It was estimated that less than 5% of the drops can enter the anterior chamber [52]. Assuming 5% of the drops enter the anterior chamber, each application of PM_{2.5} delivers 0.1 μ g PM_{2.5} into the anterior chamber. And in this study, PM_{2.5} was applied three times a day for 3 months.

There are several limitations to our study. We were not able to collect sufficient PM_{2.5} samples for fine particulate matter components analysis. At the same location, our group analysed PM_{2.5} components and found elevated amounts of PAHs [39]. Future studies should investigate the effect of major PM_{2.5} components and the role of each one in the pathological process of PM_{2.5}-induced ocular hypertension. Furthermore, we were not able to calculate the exposure concentration from the particle deposition rate for the study period. The PM_{2.5} level ranges from 46 to 163 μ g/m³ (average level 42.8 μ g/m³, Fig. S4), which is markedly higher than

the nation's air quality standards (35 μ g/m³) according to the Environment Protection Agency. However, the PM_{2.5} exposure concentrations used in this study are consistent with the common concentration range in the literature [50, 53, 54], which has been calculated based on the particle deposition rate [54]. For cell culture experiments, cytotoxicity was observed with 500 μ g/mL PM_{2.5} [54] and 1 mg/mL PM_{2.5}, which caused ocular surface damage in rats [50]. Material and assay interference was not measured in the study. However, in each experiment, a stringent control is included and we showed, using cell viability as an example, that materials used had little interference on the results. HTM cells were treated with culture medium, medium plus PBS, and medium plus DMSO and PBS, no difference in cell viability was observed (Fig. S2C).

Conclusion

This study provides novel evidence that PM_{2.5} has a direct toxic effect on the intraocular tissues and may contribute to the initiation and development of ocular hypertension and glaucoma. This occurs as a result of increased oxidative stress and the subsequent induction of NLRP3 inflammasome mediated pyroptosis in TM cells.

Materials and methods

Airborne particulate matter (PM_{2.5}) collection

Atmospheric PM_{2.5} samples were obtained from May 2016 to Dec 2018 in Lanzhou, China, and from Jan 2018 to Dec 2019 in Shanghai, China, according to our previously described methods [39]. PM samples were gathered on glass fibre filters (APFF04700, Millipore, USA) by a flow air particle sampler (TH-150C, Wuhan Tianhong Instrument Factory, Wuhan, China) at a constant flow rate of 100 L/min. The filter membranes were then cut into 1 cm \times 1 cm squares and extracted three times using an ultrasonic extractor at 100 W for 15 min in deionized water. Next, each sample suspension was filtered using 12 gauze layers, and dried by a vacuum freeze-drying machine (Labconco, Kansas, USA). PM_{2.5} samples were stored at -80° C, followed by resuspension of the resulting pellets in phosphate-buffered saline (PBS) before use.

Dynamic light-scattering (DLS)

The size distribution of the PM_{2.5} suspended in pure water was analyzed using a Nano-Zetasizer (Malvern Instruments Ltd., Worcestershire, UK) based on the dynamic light-scattering measurement technique, and the sample was first ultrasonicated with water (100 W, 15 min), then 1 ml sample was put into the measuring instrument start to start the measurement.

Animals and topical exposure of PM_{2.5} suspension

C57BL/6 mice (3–4 weeks old) were purchased from the Shanghai Sippr-BK Laboratory Animal Co. Ltd. Mice were housed in clear cages loosely covered with air filters and containing a corncob pad as bedding. After a week of acclimatization, mice were exposed to 1 mg/mL PM_{2.5} suspension thrice daily (3 × 2 μL drops) for 3 months, with PBS applied topically to the contralateral eye as a control. After 3 months, the mice were sacrificed and outflow tissues were isolated and collected for western blot analysis. All experiments complied with the Association for Research in Vision and Ophthalmology Statement for the use of animals for ophthalmic and vision research and were performed under the guidance of the Animal Care and Use Committee of Fudan University (Shanghai, China).

PM_{2.5} distribution

The fluorescent mock PM_{2.5} particles prepared from SiO₂ (diameter: 10–500 nm) were a gift from Prof. Yonghui Deng at Fudan University. Particles were dissolved in PBS and sonication was performed to disperse SiO₂ particles. Then, the fluorescent particles in PBS were topically applied to the eyes of mice. Frozen sections were prepared, and particle distributions were observed under a confocal fluorescence microscope (Leica, Shanghai, China).

Mice IOP measurements

The IOP for both eyes was measured without anaesthesia by rebound tonometry (TonoLab; ICare, Espoo, Finland). IOP was measured three times, and the average was used as the final value.

Western blotting

Mouse outflow tissue and HTM cells were lysed using a RIPA solution (Beyotime, Shanghai, China), and the protein concentration was determined using the BCA method (Beyotime). Approximately 5–20 μg of protein were loaded onto gels and separated by SDS-PAGE (10% or 12% acrylamide). Proteins were then transferred onto polyvinylidene fluoride membranes (PVDF, 0.45 μm;

Millipore, Shanghai, China) by electrophoresis. Membranes were blocked with 5% non-fat dry milk for 2 h at room temperature and probed with a primary antibody (dilutions of the primary antibodies are presented in Table 1), followed by incubation with peroxidase-conjugated secondary antibodies. Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) was used as a loading control.

HTM cell culture and PM_{2.5} treatment

HTM cells were purchased from ScienCell Research Laboratories (Shanghai, China). HTM cells were incubated in Trabecular Meshwork Cell Medium (ScienCell, Cat. No. 6591) containing 2% foetal bovine serum (FBS, Cat. No. 0010), 1% HTM growth supplement (TMC GS, Cat. No. 6592), and 1% penicillin/streptomycin (P/S, Cat. No. 0503) at 37 °C and 5% CO₂.

For the experiments, HTM cells were seeded at a concentration of 5 × 10⁵ cells/well in 6-well plates and 5 × 10³ or 5 × 10⁶ cells/well in 96-well plates. After cell attachment, the culture medium was replaced with a fresh medium containing a PM_{2.5} suspension or an equal volume of medium as a control. NAC (Sigma, Shanghai, China) 3 mM or VX-765 (Selleck, Shanghai, China) 100 μM were used. NAC and VX-765 were first dissolved in DMSO (10,010,023, Gibco, Shanghai, China) and stored at –20 °C. For rescue experiments, NA and VX-765 were diluted with culture medium (NAC: 3 mM; VX-765: 100 μM). All experiments were performed at least three times.

Cell viability test

The viability of PM_{2.5}-treated HTM cells was tested using a cell counting kit-8 (CCK-8) assay (Dojindo, Kumamoto, Japan) according to the manufacturer's instructions. Briefly, 100 μL of HTM suspensions (5000 cells/well) were added into the wells of a 96-well plate. After cell attachment, the culture medium was replaced with different concentrations of PM_{2.5} suspension (25, 50, 100, 200, and 400 μg/mL) diluted with fresh medium or an equal volume of medium as a control. After a 48-h incubation, CCK-8 solution (1:10) was added to each

Table 1 Primary antibodies used in the study

Antibody	Source	Catalog No	Type of Ab	Dilution	MW
NLRP3	Abcam	ab214185	Rabbit polyclonal	1:1000 (WB) 1:200 (IF)	114 kD
Caspase-1	Abcam	ab207802	Rabbit polyclonal	1:1000 (WB)	35, 70 kD
Caspase-1	Proteintech	22,915–1-AP	Rabbit polyclonal	1:200 (IF)	
IL-1β	Abcam	ab9722	Rabbit polyclonal	1:1000 (WB)	30, 45 kD
GSDMD	Abcam	ab215203	Rabbit monoclonal	1:1000 (WB)	35, 250 kD
GAPDH	Abcam	ab8245	Mouse monoclonal	1:1000 (WB)	36 kD

CST Cell Signaling Technology, IF immunofluorescence, MW molecular weight, WB western blotting

well of the plate and cultured at 37 °C for 2 h. Then, the optical density (OD) was measured at 450 nm using a microplate reader (Tecan, Männedorf, Switzerland), and cell viability was reported as a percentage of the OD values from unexposed control cells (100%). In NAC or VX-765 rescue experiments, NAC (3 mM) or VX-765 (100 µM) was applied 2 h before PM_{2.5} exposure. And cell viability of HTM cells treated with culture medium, medium plus PBS, and medium plus DMSO and PBS were tested to find out whether the materials we used had interference on the results.

Contractility assay and treatments

Collagen gels were prepared in 96-well plates from a collagen solution (1.85 mg/mL, Cell Biolabs, Beijing, China) by following the manufacturer's instructions. Briefly, HTM (4×10^6 cells/mL medium) was added onto the collagen gel and incubated for 1 h at 37 °C with 5% CO₂. After collagen polymerization, the culture medium was added to each collagen gel lattice. Following a 48-h incubation, the edge of the gel was gently detached using a pipette tip. The gel area was then imaged using a Fluorescent Stereomicroscope (M165 FC, Leica) every hour for 15 h to determine the time required for cessation of the "natural contraction" of the gel by HTM cells. Drugs (NAC (3 mM) or VX-765 (100 µM)) were added to the medium, and images were captured at 6, 24, and 48 h. The gel area was calculated using the Fluorescent Stereomicroscope (M165 FC, Leica, China).

ROS production detection

The intracellular ROS levels were detected using the Reactive Oxygen Species Assay Kit (ROS Assay Kit), following the manufacturer's instructions (Beyotime). After 100 µg/mL PM_{2.5} exposure for 48 h, HTM cells were washed with DMEM/F12 without FBS and then treated with 2',7'-dichlorodihydrofluorescein diacetate (DCFH-DA, 1:1000, diluted with DMEM/F12) at 37 °C for 20 min. After washing three times with the medium without FBS, the DCF fluorescence distribution of cells was detected. Rosup 50 µg/mL and 500 µg/mL were employed as the negative and positive controls, respectively. The DCF fluorescence distribution of cells was observed under a fluorescence microscope (ZEISS, Shanghai, China).

RNA isolation and RT-qPCR

HTM cells were exposed to PM_{2.5} (100 µg/mL and 200 µg/mL) for 48 h, while control cells were treated with PBS. Total RNA was then extracted using an RNeasy Mini Kit (Qiagen, Valencia, CA, USA), and RNA concentration was measured using a NanoDrop 2000 spectrophotometer (Thermo Scientific, Wilmington, DE, USA). mRNA expression was measured using the SYBR

Green quantitative real-time PCR kit (Takara, Osaka, Japan) according to the manufacturer's instructions. Samples were amplified in a ViiA 7 Real-Time PCR System (Life Technologies, Pleasanton, CA, USA), and mRNA expression was normalized to β-actin (the house-keeping gene). Expression was estimated using the comparative CT method ($2^{-\Delta\Delta CT}$) of relative quantification using the ViiA 7 Software (Life Technologies). Three independent experiments were conducted. The primer sequences used for the RT-qPCR are presented in Table 2.

Enzyme-linked immunosorbent assay (ELISA)

The IL-1β protein levels in the HTM cell culture medium were quantified using the human IL-1β ELISA kit (Abcam, Boston, MA, USA) following the manufacturer's instructions. Briefly, 100 µL of each standard and sample were added into appropriate wells and cover well and incubated for 2.5 h at room temperature. Then, the solution was discarded, washed four times, 100 µL of 1× Biotinylated IL-1 beta Detection Antibody was added to each well and incubated for 1 h at room temperature with gentle shaking. Next, the solution was discarded, washed four times, 100 µL of 1× HRP-Streptavidin solution was added to each well and incubated for 45 min at room temperature with gentle shaking. After that, the solution was discarded, washed four times, 100 µL of TMB One-Step Substrate Reagent was added to each well and incubated for 30 min at room temperature in the dark with gentle shaking. Finally, 50 µL of Stop Solution was added to each well and the OD was immediately read at 450 nm using a microplate reader (Tecan).

Immunofluorescence

Immunofluorescence analysis was performed using specific primary antibodies against NLRP3 (Abcam) and caspase-1 (ProteinTech, Shanghai, China). HTM cells grown on coverslips were fixed in 4% paraformaldehyde for 30 min at room temperature and washed three times with PBS. Then, the cells were treated with 0.1% Triton X-100 (Biotech Well, Shanghai, China) in PBS for 10 min and once again washed three times with PBS. This

Table 2 Primers used for RT-PCR

Gene	Primer
NLRP3	Forward: 5'-GCACTTGCTGGACCATCCTC-3'
	Reverse: 5'-GTCCAGTGCACACGATCCAG-3'
Caspase-1	Forward: 5'-AAGACCCGAGCTTTGATTGACTC-3'
	Reverse: 5'-AAATCTCTGCCACTTTTGTTC-3'
IL-1β	Forward: 5'-TATTACAGTGGCAATGAGG-3'
	Reverse: 5'-ATGAAGGGAAAGAAGTG-3'
β-actin	Forward: 5'-CCCTGGACTTCGAGCAAGAG - 3'
	Reverse: 5'-TCACACTTCATGATGGAGTTG-3'

was followed by blocking in PBS containing 0.5% bovine serum albumin (BSA, Roche, Shanghai, China) for 1 h at room temperature in a humidified chamber. The coverslips were then incubated with a primary antibody (dilutions of the primary antibodies are presented in Table 1) diluted in PBS containing 0.5% BSA overnight at 4 °C in a humidified chamber. The cells were then washed three times with PBS and incubated with Alexa Fluor[®]555 anti-rabbit IgG (H + L) (1:200; donkey polyclonal; Beyotime) for 1 h at room temperature. After further washing with PBS, coverslips were stained with DAPI and stored at 4 °C in the dark before being viewed under a confocal fluorescence microscope (Sp8, Leica).

Statistics

The results are presented as the mean \pm standard deviation (SD) or mean \pm standard error of mean (SEM). Data were analyzed using SPSS 21.0 (IBM, Chicago, IL, USA). For normally distributed data, the paired t-test, independent t-test, or the Student's t-test were used for two-level comparisons, while one-way analysis of variance (ANOVA) was used for ≥ 3 -level comparison. The Mann-Whitney U test was used for two-level comparisons, while the Kruskal-Wallis H test was used for ≥ 3 -level comparisons of non-normally distributed data. In all cases, differences were considered significant at $P < 0.05$.

Abbreviations

PM: Particulate matter; IOP: intraocular pressure; HTM: Human trabecular meshwork; ROS: Reactive oxygen species; TM: Trabecular meshwork; SC: Schlemm's canal; NAC: N-Acetyl-L-cysteine; VX-765: Belnacasan; CCK-8: Cell counting kit-8

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12989-021-00403-4>.

Additional file 1: Fig. S1. A. Cytotoxicity of the PM_{2.5} on HTM cell. For the PM_{2.5} collected in Shanghai between Jan 2018 to Dec 2019, cell viability was analyzed by CCK-8 assay. The human trabecular meshwork cells were treated with different concentrations of Shanghai PM_{2.5} (0 $\mu\text{g}/\text{mL}$, 25 $\mu\text{g}/\text{mL}$, 50 $\mu\text{g}/\text{mL}$, 100 $\mu\text{g}/\text{mL}$, 200 $\mu\text{g}/\text{mL}$, or 400 $\mu\text{g}/\text{mL}$) from Shanghai for 48 h ($n = 5$ cell lines, $*P < 0.05$, $**P < 0.01$, $***P < 0.001$, by one-way ANOVA, compared with the control). Data are represented as the mean \pm standard deviation. **B.** Expressions of NLRP3 inflammasome-related proteins exposed to PM_{2.5} collected in Shanghai. After HTM cells were treated by PM_{2.5} 100 $\mu\text{g}/\text{mL}$ or 200 $\mu\text{g}/\text{mL}$, relative protein expressions of NLRP3 ($n = 3$), caspase-1 ($n = 3$), and GSDMD ($n = 3$) determined by western blot ($*P < 0.05$, one-way ANOVA, compared with the control). Data are represented as the mean \pm standard deviation. **C.** ROS production in HTM cells exposed to the PM_{2.5} collected in Shanghai. Representative ROS fluorescence staining images of Rosup positive control (PC, 500 $\mu\text{g}/\text{mL}$), PBS control group (control), PM_{2.5} 100 $\mu\text{g}/\text{mL}$ group (PM_{2.5}) and NAC + PM_{2.5} group. ROS levels were detected by Reactive Oxygen Species Assay Kit. In each panel, from left to right are bright field, fluorescence, and merged images of fluorescence and bright field. HTMs were treated with Shanghai PM_{2.5} 100 $\mu\text{g}/\text{mL}$ for 48 h, pretreated with or without NAC 3 mM for 2 h. ROS levels were significantly increased in HTMs treated with Rosup positive control and 100 $\mu\text{g}/\text{mL}$ PM_{2.5}. Magnification, 5X. **Fig. S2. A.** Expressions of NLRP3 inflammasome-related

mRNA in PM_{2.5}-treated TM cells. qPCR showed mRNA expression of NLRP3 increased significantly after HTMs treated by PM_{2.5} 100 $\mu\text{g}/\text{mL}$ or 200 $\mu\text{g}/\text{mL}$ ($n = 3$ cell lines, $*P < 0.05$, one-way ANOVA, compared with the control). The mRNA expression of caspase-1 decreased after HTMs treated by PM_{2.5} 100 $\mu\text{g}/\text{mL}$ or 200 $\mu\text{g}/\text{mL}$ ($n = 3$ cell lines, $*P < 0.05$, one-way ANOVA, compared with the control). The mRNA expression of IL-1 β increased significantly after HTMs treated by PM_{2.5} 100 $\mu\text{g}/\text{mL}$ or 200 $\mu\text{g}/\text{mL}$ ($n = 6$ cell lines, $*P < 0.05$, one-way ANOVA, compared with the control). **B.** NAC inhibited ROS elevation in Lanzhou PM_{2.5}-treated HTM cells. Representative ROS fluorescence staining images of Rosup negative control (NC, 50 $\mu\text{g}/\text{mL}$), Rosup positive control (PC, 500 $\mu\text{g}/\text{mL}$), PBS control group (control), PM_{2.5} 100 $\mu\text{g}/\text{mL}$ group (PM_{2.5}) and NAC + PM_{2.5} group. ROS levels were detected by Reactive Oxygen Species Assay Kit. In each panel, from left to right are bright field, fluorescence, and merged images of fluorescence and bright field. HTMs were treated with PM_{2.5} 100 $\mu\text{g}/\text{mL}$ for 48 h, pretreated with or without NAC 3 mM for 2 h. ROS levels were significantly increased in HTMs treated with Rosup positive control and 100 $\mu\text{g}/\text{mL}$ PM_{2.5}. Magnification, 10X. **C.** Cytotoxicity of the PBS and DMSO on HTM cell. Cell viability was analyzed by CCK-8 assay. The human trabecular meshwork cells were treated with culture medium (control), medium plus PBS, medium plus DMSO and PBS, respectively according to our used dose in the study ($n = 5$ cell lines, $p > 0.05$, by one-way ANOVA, compared with the control). Data are represented as the mean \pm standard deviation. **Fig. S3. A.** Effects of fiber-exposure on HTM cell viability. Cell viability were analyzed by cell viability assay (CCK-8). The human trabecular meshwork cells were treated with different concentrations of fiber (0 $\mu\text{g}/\text{mL}$, 5 $\mu\text{g}/\text{mL}$, 20 $\mu\text{g}/\text{mL}$, 30 $\mu\text{g}/\text{mL}$) for 48 h. ($n = 6$ cell lines, $*P < 0.05$, by one-way ANOVA, compared with the control). Data are represented as the mean \pm standard deviation. **B.** NLRP3 inflammasome-related protein expressions in glass fibre treated HTM cells. After HTM cells were treated by 30 $\mu\text{g}/\text{mL}$ fibers obtained from glass fibre filters that were used for PM collection. Relative protein expressions of NLRP3 and caspase-1 were determined by western blot ($n = 3$ cell lines, $P > 0.05$, independent t-tests, compared with the control), which did not differ significantly from control cells. Data are represented as the mean \pm standard deviation. **Fig. S4. A.** Air quality index (AQI) in Lanzhou between May 2016 and Dec 2018. The PM_{2.5} level ranges from 46 to 163 $\mu\text{g}/\text{m}^3$ with an average level of 42.8 $\mu\text{g}/\text{m}^3$ from May 2016 to Dec 2018 in Lanzhou. **B.** Air pollutants in Lanzhou between May 2016 and Dec 2018. The figure shows the concentration of PM_{2.5} ($\mu\text{g}/\text{m}^3$), PM₁₀ ($\mu\text{g}/\text{m}^3$), SO₂ ($\mu\text{g}/\text{m}^3$), CO (mg/m^3), NO₂ ($\mu\text{g}/\text{m}^3$), O₃ ($\mu\text{g}/\text{m}^3$) in the air from May 2016 to Dec 2018 in Lanzhou. **Table 1** Weight of glass fiber filters before and after PM extraction.

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Authors' contributions

Y.R, Y.L and H.S designed and supervised all the experiment work. L.L and C.X (contributed equally) acquired and analyzed the data used in the present study with assistance from L.N, B.L, J.Z, M.S., C.H, J.N. All authors contributed to the writing of the manuscript. And all authors read and approved the final manuscript.

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Availability of data and materials

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

Declarations**Ethics approval and consent to participate**

All experiments complied with the Association for Research in Vision and Ophthalmology Statement on the use of animals in ophthalmic and vision research. The animal study was approved by the Animal Care and Use Committee of Eye and ENT Hospital of Fudan University (Shanghai, China).

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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