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## Pulmonary health effects of wintertime particulate matter from California and China following repeated exposure and cessation

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### Abstract

Epidemiological studies show strong associations between fine particulate matter ( $PM_{2.5}$ ) air pollution and adverse pulmonary effects. In the present study, wintertime  $PM_{2.5}$  samples were collected from three geographically similar regions—Sacramento, California, USA; Jinan, Shandong, China; and Taiyuan, Shanxi, China—and extracted to form  $PM_{CA}$ ,  $PM_{SD}$ , and  $PM_{SX}$ , respectively, for comparison in a BALB/c mouse model. Each of four groups was oropharyngeally administered Milli-Q water vehicle control ( $50 \ \mu$  L) or one type of PM extract ( $20 \ \mu$ g/50  $\mu$ L) five times over two weeks. Mice were necropsied on post-exposure days 1, 2, and 4 and examined using bronchoalveolar lavage (BAL), histopathology, and assessments of cytokine/ chemokine mRNA and protein expression. Chemical analysis demonstrated all three extracts contained black carbon, but  $PM_{SX}$  contained more sulfates and polycyclic aromatic hydrocarbons (PAHs) associated with significantly greater neutrophil numbers and greater alveolar/bronchiolar inflammation on post-exposure days 1 and 4. On day 4,  $PM_{SX}$ -exposed mice also exhibited significant increases in interleukin-1 beta, tumor necrosis factor-alpha, and chemokine C-X-C motif ligands-3 and –5 mRNA, and monocyte chemoattractant protein-1 protein. These combined

Appendix A. Supplementary data

Supplementary material related to this article can be found, in the online version, at doi:https://doi.org/10.1016/j.toxlet.2021.10.014.

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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

findings suggest greater sulfate and PAH content contributed to a more intense and progressive inflammatory response with repeated  $PM_{SX}$  compared to  $PM_{CA}$  or  $PM_{SD}$  exposure.

#### **GRAPHICAL ABSTRACT**



#### Keywords

PM<sub>2.5</sub>; Lung; Sulfate; PAHs; Time-lag; Inflammation

#### 1. Introduction

The physicochemical characteristics of particulate matter (PM) can be highly variable due to different emission sources, weather conditions, and interactions with other solid/liquid and organic/inorganic particles suspended in the atmosphere (Kelly and Fussell, 2012). PM originating from different sources, locations, and times can exhibit unique levels of toxicity associated with respiratory effects, such as asthma and COPD (Lee et al., 2014). PM size can also greatly influence health outcomes. Although coarse PM (particles with a diameter between 10 and 2.5  $\mu$ m; PM<sub>10-2.5</sub>) is known to irritate the eyes and nose, ambient fine PM (particles with an aerodynamic diameter <2.5  $\mu$ m; PM<sub>2.5</sub>) has a greater potential for harmful health effects due to its ability to penetrate beyond the air passages of the upper respiratory tract and deposit in the distal lung and alveoli where gas exchange occurs (EPA, 2017; Ma et al., 2015). With clearance from the respiratory tract via mucociliary transport, lymphatic drainage, or systemic blood circulation, PM<sub>2.5</sub> can also bind to cells and tissues to affect extra-pulmonary organs (Kim et al., 2015; Phalen, 2004). For these and other reasons, with more than 4.2 million premature deaths worldwide attributed to PM<sub>2.5</sub> exposure in 2016 alone, PM<sub>2.5</sub> is a growing universal health concern.

In the present study, PM samples were collected from cities in California (Sacramento) and China (Jinan and Taiyuan) to compare the biological effects of  $PM_{2.5}$  from different geographic locations. The cities were chosen due to their heavy urbanization, industrial and agricultural economies, moderately dry and sunny winters, and long histories of relatively high  $PM_{2.5}$  levels, especially during the winter season, when meteorological inversions are

more frequent and longer in duration (Liu et al., 2010; Xia et al., 2013). California's Central Valley has increased levels of air pollution due to its geologic bowl shape leading to the retention and accumulation of PM from vehicular exhaust, local farming, and industrial processes (Plummer et al., 2015). Several studies have shown respiratory-related emergency department visits increase with increasing  $PM_{2.5}$  concentrations (Anderson et al., 2012; Cadelis et al., 2014), and active asthma prevalence in Sacramento County individuals age 18–64 years was 12 % compared to 7.6 % for age-matched individuals state-wide. Active asthma was defined as having been diagnosed by a professional and afflicted by an asthma attack within the preceding year (CDPH, 2016).

The capital cities of Jinan and Taiyuan in the Chinese provinces of Shandong and Shanxi, respectively, represent regions of strong economic growth dominated by abundant production and industrial emissions. Shandong Province is one of the most prosperous and heavily industrialized provinces in the country, with significant coal mining, oil refineries, and developed metallurgical and mechanical sectors (Liu et al., 2018). In 2011, Jinan has experienced an annual average PM<sub>2.5</sub> concentration of 149  $\mu$ g/m<sup>3</sup>, one of the highest reported for cities worldwide (Yang et al., 2012). Similarly, Shanxi Province is China's largest coal-producing region, with Taiyuan as one of the most polluted cities in China. From 2009 to 2010, the annual average PM<sub>2.5</sub> concentration in Taiyuan was 220  $\mu$ g/m<sup>3</sup>, more than 20 times the World Health Organization's 10  $\mu$ g/m<sup>3</sup> air quality guideline for mean annual PM<sub>2.5</sub> emissions (He et al., 2015; Stanway, 2017; WHO, 2018; Xia et al., 2013).

A recent study (Yuan et al., 2020) from our laboratory demonstrated a single acute exposure to extracts of  $PM_{2.5}$  collected from California, Shandong, and Shanxi ( $PM_{CA}$ ,  $PM_{SD}$ , and  $PM_{SX}$  extracts, respectively) could cause lung inflammation and induce toxic responses in human immortalized U937 macrophages. The current study furthers our previous work by exploring pulmonary inflammatory responses to repeated oropharyngeal aspiration (OPA) of  $PM_{CA}$ ,  $PM_{SD}$ , or  $PM_{SX}$  over two weeks, in BALB/c mice. Although comparative analyses of pulmonary inflammatory responses to PM from California and China have been reported previously (Sun et al., 2017; Yuan et al., 2020; Zhang et al., 2018), the present study is the first to assess the differential time-lag effects of repeated, low-dose exposure to PM from Sacramento, Jinan, and Taiyuan. The findings of this study are discussed in relation to previous (Sun et al., 2017; Yuan et al., 2020; Zhang et al., 2018) to illustrate further how different exposure paradigms influence observed pulmonary responses.

#### 2. Materials and methods

The methods in the present study have been reported previously (Sun et al., 2017; Yuan et al., 2020; Zhang et al., 2018) and restated herein for reader accessibility.

#### 2.1. Ambient PM<sub>2.5</sub> collection

Ambient winter PM<sub>2.5</sub> collection occurred in Sacramento, California; Jinan, Shandong; and Taiyuan, Shanxi. The California sampling site was located in downtown Sacramento, at the northeast corner of T and 13th streets, on the rooftop of a two-story building (N38°34', W121°29'). Downtown Jinan's collection site was located on the rooftop of a three-story primary school, Wang She Ren (N36°40', E117°09'). The sampling site in downtown

Taiyuan was situated on the rooftop of the five-story College of Environmental Science and Resources building at Shanxi University (N37°47', E112°34'). All three sampling locations are proximal to major freeways and surrounded by a combination of residential, commercial, and industrial sites.

In Sacramento, a high-volume sampler system (Tisch Environmental Inc., TE-6070V2.5-HVS) was used to collect  $PM_{2.5}$  for seven days. The system had a  $PM_{2.5}$  size-selective head (Tisch Environmental Inc., TE-6001) that ran at a flow rate of 40 cubic feet per minute (cfm) and was fitted with Teflon-coated borosilicate glass microfiber filters (Pall Corporation, TX40H120WW). The filters were pre-cleaned by repeated sonication in a mixture of Milli-Q water, dichloromethane, and hexane prior to sampling (Bein and Wexler, 2015).

In Jinan and Taiyuan,  $PM_{2.5}$  was collected for one day using a high-volume particle collector (Thermo Anderson, HVAIR100) operated at a flow rate of 40 cfm. The shorter collection times in China were due to higher density and heavier loading of PM relative to the sampling site in California. The sampler was fitted with a  $PM_{2.5}$  size-selective opening and 90-mm diameter quartz microfiber filters (Whatman, WHA1851090). During the 24 h before PM collection, the quartz filters were pre-heated at 450 °C for the removal of any potential endotoxin from the filters.

#### 2.2. PM sample extraction

Following particle sample collection in Sacramento and Jinan, all filters were post-weighed to determine the collected PM mass (post-weight – pre-weight = PM mass). Filters were placed in Milli-Q water and sonicated for 1 h to obtain PM extracts. The sonicated PM extracts were passed through 0.2- $\mu$ m pore size syringe filters, and the collected solution (approximately 100 mL) was lyophilized and stored at 80 °C. A more detailed description of the PM extraction methods was provided by Bein and Wexler(2015).

Filters from Taiyuan were post-weighed to determine the collected PM mass, subsequently cut into segments and placed in a 250-mL conical flask with 30 mL Milli-Q water, and sonicated for 30 min (3 cycles of 10 min each) to yield a PM extract. The extract was filtered through six layers of sterile gauze, lyophilized to powder, and stored at -80 °C. For all extraction procedures, field blanks were included to ensure no extractable compounds from filters were present in the suspension to produce an effect beyond that noted with Milli-Q water suspension alone.

Lyophilized PM extracts from Sacramento, Jinan, and Taiyuan were weighed, resuspended in Milli-Q water, and sonicated for 20 min to derive stock PM extract samples ( $PM_{CA}$ ,  $PM_{SD}$ , and  $PM_{SX}$ , respectively) of the same concentration (1 µg/mL; Fig. 1). All stock PM samples were frozen at –20 °C until needed.

#### 2.3. Characterization of the hydrodynamic particle size

Dynamic light scattering (DLS) was used to determine the hydrodynamic particle size distribution of each stock extract sample following 20 min of sonication to insure adequate dispersion of particles (Sun et al., 2017).

# 2.4. Chemical characterization of stock extract samples via high-resolution time-of-flight aerosol mass spectrometry (HR-AMS)

Chemical characterization of all three PM extracts was performed at the University of California, Davis. Frozen stock PM extracts were thawed to 4 °C, sonicated for 20 min for thorough PM dispersal, and diluted by Milli-Q water to  $0.1 \mu g/mL$ . Subsequently, 1 mL of each diluted stock sample was individually passed through a constant output aerosol generator with an inert argon carrier gas, and a silica gel diffuser where the sample became atomized. HR-AMS was used to determine the chemical composition of each extract sample. Chemical species such as nitrate, sulfate, chloride, ammonium, and organic compounds were quantified (Canagaratna et al., 2007). The HR-AMS spectra of organic compounds were further analyzed to determine PAH contents and associated elemental (hydrogen, carbon, oxygen, and nitrogen) composition and average degree of oxidation as detailed by Aiken et al. (2008), Ghio and Devlin (2001), Sun et al. (2017), Zhang et al. (2018), Yuan et al. (2020) and Dzepina et al. (2007).

#### 2.5. Animals and exposure

The UC Davis Institutional Animal Care and Use Committee approved all procedures and animal housing practices. All methods in this study were carried out following relevant guidelines and regulations. As BALB/c mice exhibit strong T-helper (Th)-2-mediated responses that facilitate the detection of asthmatic and other inflammatory responses driven by PM (Li et al., 2009), 63 male, six-week-old BALB/c mice were purchased (Envigo, Hayward, CA) and randomly assigned to one of four groups for exposure to 1) vehicle control (Milli-Q water; n = 9), 2) PM<sub>CA</sub> (n = 18), 3) PM<sub>SD</sub> (n = 18); or 4) PM<sub>SX</sub> (n = 18). All mice were acclimated for 2 weeks before the start of the exposure study, maintained on a 12-h light/dark cycle, and housed three per cage using sterile laboratory bedding and access to food and water ad libitum.

Immediately before exposure, stock extracts of  $PM_{CA}$ ,  $PM_{SD}$ , and  $PM_{SX}$  were thawed, sonicated for 20 min, and diluted with Milli-Q water to a concentration of 20 µg/50 µL. Exposure by OPA occurred five times, once per day on days 1, 4, 7, 10, and 14 (Fig. 2). During exposure, mice were sedated by inhaling isoflurane in oxygen at a 3:1 ratio (Plummer et al., 2015). Each mouse was given 50 µL of Milli-Q water, or 20 µg of  $PM_{CA}$ ,  $PM_{SD}$ , or  $PM_{SX}$  in 50 µL Milli-Q water via OPA such that the exposure volume was consistent across all groups. A single dose of this nature has been used previously (Sun et al., 2017) to examine PM effects. With high ambient PM levels noted in each of the sampled cities (in particular China), a total PM dose of 100 µg administered over two weeks could approach levels experienced by inhalation of ambient air over the same period.

#### 2.6. Bronchoalveolar lavage fluid (BALF)

Mice were euthanized for necropsy at 1, 2, and 4 day(s) after the final OPA exposure (study days 15, 16, and 18, respectively; Fig. 2) with a 0.2-mL intraperitoneal injection of Beuthanasia-D pentobarbital solution (65 mg/kg body weight; Nembutal, Cardinal Health, Sacramento, CA). In the present study, the term "necropsy" refers to the post-mortem examination and collection of biological tissues from study animals. Each mouse was cannulated intratracheally, and the left mainstem bronchus was clamped, while the right

lung was lavaged with two aliquots (0.6 mL/aliquot) of Dulbecco's Phosphate Buffered Saline (PBS; Sigma-Aldrich, St Louis, MO). The collected BALF was centrifuged at  $500 \times$  g at 4 °C for 15 min to pellet the cells. BALF supernatant was decanted, frozen in liquid nitrogen, and stored at -80 °C. Cell pellets were resuspended in 500 mL PBS to determine total cell numbers and viability using a hemocytometer and 0.4 % Trypan Blue solution (Sigma-Aldrich).

BALF cytospin slides were prepared using a Shandon Cytospin (Thermo Shandon, Inc., Pittsburg, PA). The slides were then stained with DippKwik Differential Stain (American MasterTech, Lodi, CA) to determine the proportions of BAL macrophages, neutrophils, eosinophils, and lymphocytes in cell counts (500 cells/animal) performed using brightfield microscopy. The experimental protocol is shown in Fig. 3.

#### 2.7. Lung tissue collection

The left lungs were inflated-fixed at 30 cm of pressure with 4% paraformaldehyde for one hour, stored in 4% paraformaldehyde for 48 h, and subsequently transferred into 70 % ethanol for later tissue processing for histopathology. The right lung lobes of each mouse were placed in two cryovials—one vial for the cranial and middle lobes—and the other for the caudal and accessory lobes. All vials were stored at -80 °C until further use.

#### 2.8. Semi-quantitative lung histopathology

Following placement of the left lung in 70 % ethanol, four transverse slices (levels) were prepared as a method to sample uniformly histological changes throughout the entire lobe. These tissue slices were dehydrated, embedded in paraffin, and sectioned at a thickness of 5 µm. The sections were placed on glass slides, stained with Harris hematoxylin and eosin (H&E; American MasterTech, Lodi, CA), and coverslipped.

Four transverse slices were stained for each mouse (n = 9/control or 18/PM group). All slides were examined independently by two blinded observers (WY and SCV) for the presence of inflammation, cellular infiltrates, and cellular/tissue remodeling (i.e., desquamation and squamous metaplasia of airway epithelial cells, septal wall thickening) in alveolar ducts and airways using a semi-quantitative scoring scale. Previously published rubrics (Silva et al., 2013) were used to rank the severity (absent to marked; 0-3) and extent (0: no changes, 1: less than one-third of the slide, 2: one-half of the slide, 3: two-thirds of the slide) of the observed alveolar and bronchiolar inflammation. Alveolar inflammation involves the lung parenchyma, which includes the alveolar airspaces and pulmonary interstitium (i.e., the collection of support tissues within the lung that includes the alveolar epithelium, pulmonary capillary endothelium, basement membrane, and perivascular and perilymphatic tissues). With alveolar inflammation, focal accumulations of inflammatory cells composed of macrophages and/or polymorphonuclear cells (PMNs), e.g., neutrophils and eosinophils, can fill the alveolar airspace. Bronchiolar inflammation involves the airways, which can include the influx of PMNs and/or phagocytes (e.g., macrophages) in the airway wall and contiguous alveoli, as well as signs of airway cell damage (e.g., sloughing). For each mouse, the final score for a given parameter (e.g., alveolar inflammation) was the averaged product of the severity and extent scores for the four transverse slices. The use

of the products resulted in scores ranging from 0 to 9, thus increasing the probability of finding significant (p < 0.05) differences between groups (Rutgers et al., 2010). Detailed semi-quantitative scoring guidelines are shown in Supplementary Table S1.

#### 2.9. Gene expression analysis

As ribonucleic acids (RNAs) are the initial products of gene expression, messenger RNA (mRNA) was isolated from the right caudal and accessory lung lobes using TRI Reagent (Sigma-Aldrich) and a Quick-RNA Miniprep kit (Zymo Research, Irvine, CA). The mRNA was converted to complementary deoxyribonucleic acid (cDNA) using a highcapacity cDNA Reverse Transcription Kit (Applied Biosystems, Indianapolis, IN). Genespecific mouse primers (0.2 µM; IDT, Coralville, IA), cDNA (2 µL/reaction), and SYBR Green (Applied Biosystems) DNA-binding stain were used for quantitative polymerase chain reaction (qPCR) measurements. Expression of genes for inflammatory cytokines, interleukin-1 beta  $(IL-1\beta)$  and tumor necrosis factor-alpha (TNF-a), and neutrophil chemokines, chemokine (C-X-C Motif) ligands-3 and -5 (CXCL-3 and CXCL-5) were examined. Expression was assessed using the -Ct method and standardized to the expression of elongation factor 1-alpha 1 (*EEF1a1*) "housekeeping genes" (Castaneda et al., 2018). Housekeeping genes are 1) typically constitutive (i.e., continuously expressed); 2) required for maintenance of basic cellular functions essential for the existence of a cell irrespective of its role in tissue or the organism; and 3) expressed in all cells of an organism under normal and pathophysiological conditions (Pampel, 2017). The *EEF1a1* gene is one of the alpha subunit forms of the elongation factor 1 complex that interacts with aminoacylated transfer RNA (tRNA) and delivers it to the A site of the ribosome during the elongation phase of protein synthesis. It is constitutively expressed in all tissues in mice and humans, eukaryotes in general (Gentile et al., 2016). The use of EEF1a1 genes is an accepted method for qPCR gene expression analysis of mouse lung tissue (Castaneda et al., 2018).

Mouse gene primers were designed using Primer3 primer design software (Untergasser et al., 2012). Primers used in this study are detailed in Supplementary Table S2.

#### 2.10. Enzyme-Linked immunosorbent assay (ELISA)

ELISAs (Biolegend, San Diego, CA) were performed on the homogenized cranial and middle lobes of the right lung to analyze concentrations of specific proteins, including TNF- $\alpha$ ; monocyte chemoattractant protein-1 (MCP-1), a chemoattractant responsible for monocyte and macrophage recruitment; CXCL-1, seen in inflammation or wound healing; and IL-1 $\beta$  and IL-6, both mediators of inflammatory responses. The cranial and middle lobes from control- and PM-exposed animals and standards from the R&D Systems ELISA kits (1000 µg/mL to 7.8 µg/mL) were prepared and examined in duplicate in 96-well plates using a SpectroMax plate reader (Molecular Devices, Sunnyvale, CA). Duplicate readings were averaged. All concentrations were normalized to total lung protein and reported in pg of specific protein per mg of lung tissue.

#### 2.11. Statistical analysis

No data points were excluded before statistical analysis. All statistical tests were performed using GraphPad PRISM 8.0 software. A value of p < 0.05 was considered statistically

significant. For each measured endpoint, a Shapiro-Wilk test was first used to detect normality, and a one-way analysis of variance (ANOVA) and *post hoc* Tukey's test were performed to determine differences due to treatment. All data in the present document are expressed as the mean  $\pm$  standard error of the mean (SEM).

#### 3. Results

#### 3.1. DLS particle size analysis

All stock PM extracts exhibited unimodal hydrodynamic size distributions.  $PM_{CA}$  ranged between 100–600 nm, while  $PM_{SD}$  and  $PM_{SX}$  ranged between 100 nm–1800 nm (Fig. 4). Based on the hydrodynamic diameters measured by DLS, all PM samples were within the  $PM_{2.5}$  size range in their hydrodynamic measured diameters (Fig. 4).

#### 3.2. Chemical composition of PM extracts

Organic compounds comprised the largest fraction of the total solute mass in each extract sample. While the PM<sub>SX</sub> extract was the darkest (Fig. 1), suggesting it had the highest black carbon content, PMSD was also found to contain carbon. Organic compounds accounted for 54 %, 57 %, and 45 % of the PM<sub>CA</sub>, PM<sub>SD</sub>, and PM<sub>SX</sub> mass, respectively (Fig. 5a-c). These results based on the color and chemical analysis of each PM extract suspension suggests the forms of carbon are different, at least with respect to the PM<sub>SX</sub> and PM<sub>SD</sub> extracts. Within the organic fraction, carbon and oxygen accounted for at least 87 % of the mass (Fig. 5d-f). For other compounds measured by HR-AMS, nitrate and sulfate fractions were highly variable, while those of chloride and ammonium were relatively similar among the California and China PM extracts. Nitrate was more abundant in California PM than China PM, while sulfate was most prevalent in PM<sub>SX</sub> versus the other two PM extracts (Fig. 5a-c). The sulfate and nitrate measured in all three PM samples were principally inorganic, identified by the presence of anions in both the ammonium sulfate and ammonium nitrate. However, based on ion balance analysis and the abundance of sulfur-containing organic ions in the HR-AMS spectrum, it is likely that PM<sub>SX</sub> contained more organic sulfate than the other two PM extracts, perhaps a reflection of sulfur-containing coal combusted locally. Of the PAH levels analyzed by HR-AMS, the PAH/Organics ratios were 0.0017, 0.004, and 0.0024 of the PM<sub>CA</sub>, PM<sub>SD</sub>, and PM<sub>SX</sub>, respectively. The percentage of dissolved PM mass varied between samples due to the different organic fractions. Therefore, based on the calculated enhancement factors relative to PM<sub>CA</sub>, PM<sub>SD</sub> and PM<sub>SX</sub> had 2.48 and 11.50 times higher levels than PM<sub>CA</sub>, respectively (Table 1).

#### 3.3. Cell differentials in BALF

The BALF cell analyses were based on a total of 9 control mice (n = 3/time-point) and 54 PM-exposed mice (n = 6/group/time-point). A total of 9 control mice pooled from post-OPA days 1, 2, and 4 were used for comparisons since there were no statistical differences among controls over time. Recovery time did not appear to play a role in the responses observed in BALF as there were no statistically significant differences between necropsy days (Fig. 6a-b). No significant (p < 0.05) exposure-related differences were noted for total cell counts of mice exposed to the vehicle control versus PM<sub>CA</sub>, PM<sub>SD</sub>, or PM<sub>SX</sub> irrespective of the recovery time post-OPA (Fig. 6a). In contrast, significantly increased neutrophil numbers

were observed on post-OPA days 1 and 4 in all PM-exposed groups relative to controls (p < 0.05 for all comparisons), and on day 4 in PM<sub>SX</sub>-exposed mice relative to their PM<sub>CA</sub>- and PM<sub>SD</sub>-exposed counterparts (p = 0.0326 and p = 0.0196, respectively; Fig. 6b). No statistically significant treatment-related effects were observed on day 2 (Fig. 6b), as neutrophil numbers in all PM-exposed groups dropped temporarily to near-control levels. However, the elevated neutrophil numbers observed in PM- versus control-exposed mice on day 4 is not unlike results of previous studies (Noel et al., 2016; Smith et al., 2006) in which incomplete lung recovery—including elevated BALF neutrophils and other measures of inflammation— has been reported in mice and rats exposed sub-acutely to combustion-derived PM and examined up to 10 days post exposure.

#### 3.4. Histological analysis

There were no significant (p < 0.05) differences in bronchiolar inflammation scores on post-OPA day 1 (Fig. 6c). However, on day 2, mice exposed to PM<sub>SX</sub> exhibited significantly (p = 0.0179) more bronchiolar inflammation than controls (Fig. 6c). By day 4, both groups given Chinese PM extracts exhibited more severe bronchiolar inflammation than controls (p < 0.04 for PM<sub>SD</sub>; p < 0.01 for PM<sub>SX</sub>; Fig. 6c), with neutrophilic influxes into the peribronchiolar and alveolar regions of the lungs that were not observed in other groups (Fig. 7a-d). Mice exposed to PM<sub>SX</sub> were found to have visible black particles—likely due to coal combustion—in macrophages throughout the alveolar regions of the lungs.

In contrast to the bronchiolar, perivascular, and subpleural regions, the alveoli appeared to be most affected by PM. On post-OPA days 1 and 2, significantly greater alveolar inflammation was observed in groups administered  $PM_{SD}$  (p = 0.0206 and 0.0361, respectively) or  $PM_{SX}$ (p = 0.0057 and 0.0172, respectively) versus the vehicle control (Fig. 6d). Over these two days, no significant intra-group differences in alveolar inflammation were observed within the PM<sub>SD</sub> or PM<sub>SX</sub> groups. By day 4, all three PM-exposed groups exhibited higher scores for alveolar inflammation than controls (Fig. 6d; p < 0.01, p < 0.01, and p < 0.01, p < 0.01, and p < 0.01, 0.0001 for PM<sub>CA</sub>, PM<sub>SD</sub>, and PM<sub>SX</sub>, respectively). Mice exposed to PM<sub>SD</sub> and PM<sub>SX</sub> had average scores of 1.8 and 2.4, respectively, for alveolar inflammation, with notable changes in alveolar wall thickening, and numerous free macrophages distributed throughout the parenchyma (Fig. 7e-f), while control mice had an average score of 0.7, with thin-walled alveolar septa and relatively few lumenal macrophages (Figs. 6d and 7e). Although intergroup differences were not statistically significant among the PM-exposed mice, those administered  $PM_{SX}$  appeared to demonstrate the greatest effects (Fig. 7e-f) with cellular debris and numerous foamy, particle-laden macrophages aggregated in the alveolar airspaces (Fig. 7f). Trends toward greater but not statistically significant bronchiolar and alveolar inflammation in the PM-exposed groups over time (Fig. 6c-d) support the aforementioned findings of elevated BALF neutrophils in these groups relative to controls on post-OPA day 4 (Fig. 6b), and the likelihood of unresolved and/or late-peaking inflammation with repeated exposure to combustion-derived PM.

#### 3.5. qPCR analysis of gene expression

PM-induced recruitment of neutrophils into the lungs can be directed by numerous chemotactic mediators, including *IL-1\beta*, *TNF-a*, *CXCL-3*, and *CXCL-5* (Maueroder et

al., 2015). Results demonstrated expression of *IL-1β* and *TNF-a* genes was significantly increased in PM<sub>SX</sub>- versus control-exposed mice on days 2 (p = 0.0261 and p = 0.0415, respectively) and 4 (p < 0.0001 and p = 0.0062, respectively; Fig. 8a-b) post OPA. Similar increases were observed for post-OPA day 4 *IL-1β* in PM<sub>CA</sub>-exposed mice relative to controls (p = 0.0049) and in PM<sub>SX</sub>-exposed mice relative to their PM<sub>SD</sub>-exposed counterparts (p = 0.0152; Fig. 8a-b). Gene expression of *CXCL-3* and *CXCL-5* was also significantly elevated in PM-versus control-exposed mice, with PM<sub>SX</sub> producing increases in one or both chemokines at all post-OPA time points (p < 0.05); PM<sub>CA</sub> producing increases in *CXCL-3* (p = 0.0351) and *CXCL-5* (p < 0.0001) on post-OPA day 4 alone; and PM<sub>SD</sub> producing an increase in *CXCL-5* (p = 0.0233) on post-OPA day 4 alone (Fig. 8c-d). *CXCL-3* and/or –5 mRNA expressions increased in all PM-versus control-exposed groups on post-OPA day 4.

#### 3.6. Quantification of cytokines by ELISA

Although six cytokines were examined, concentrations of three proteins (TNF- $\alpha$ , CXCL-1, and IL-6) were too low to quantify reliably with limits of detection at 5.16, 66.521, 8.301 ng/mL, respectively. Of the two remaining cytokines, IL-1 $\beta$  (data not shown) and MCP-1 (Fig. 9), only protein levels of the latter varied significantly among the PM- and control-exposed groups, with PM<sub>SX</sub> producing increases relative to control on post-OPA days 2 and 4 (p < 0.05 and p < 0.0001, respectively); and PM<sub>CA</sub> and PM<sub>SD</sub> producing similar increases on post-OPA day 4 alone (p < 0.05 and p < 0.05, respectively).

#### 4. Discussion

Global PM emissions are of major concern, with the duration of PM exposure and physicochemical characteristics of PM (e.g., diameter and composition) among the primary factors influencing adverse health effects. Of the various PM size fractions, PM<sub>2.5</sub> is most associated with morbidity and premature mortality (Rajak and Chattopadhyay, 2019) after long-term exposure, and, because it contributes to 7 million deaths each year (WHO, 2019), PM<sub>2.5</sub> is often described as one of the leading risk factors for premature mortality. Though mortality was not examined in the present study, cumulative results add to the existing body of literature regarding 1) the impacts of repeated seasonal PM<sub>2.5</sub> exposure on pulmonary morbidity, and 2) the variability of PM<sub>2.5</sub> toxicity related to chemical composition.

The prior two studies performed in our laboratory (Sun et al., 2017; Yuan et al., 2020), with similar PM extracts administered in single-exposure paradigms, showed striking differences in findings compared to the current study with repeated PM exposure of different duration and frequency. In the present study, many of the statistically significant responses were observed on post-OPA day 4, with BAL (Fig. 6b), histopathology (Fig. 6c-d), gene expression (Fig. 8), and ELISA (Fig. 9) analyses. Less severe inflammatory responses were noted on days 1 and 2. The cumulative findings suggest repeated exposure to  $PM_{CA}$ ,  $PM_{SD}$ , or  $PM_{SX}$  was associated with significant and persistent inflammation relative to the sham controls. These results are in striking contrast to those from our previous study (Yuan et al., 2020), which measured the inflammatory effects of a single 50-µL OPA exposure to Milli-Q water or a  $PM_{CA}$ ,  $PM_{SD}$ , or  $PM_{SX}$  stock (1 µg/µL) extract on post-OPA days 1, 2

and 4. In that study, the three PM extracts produced the most severe inflammatory responses on post-OPA day 1, with all measured biological parameters returning to control levels by days 2 and 4. Given the identical animal model and PM extract composition for both studies, the drivers for different findings in the two studies are likely to be the 1) longer exposure duration, which stimulated inflammation pathways that are initiated subacutely; 2) intermittent, repeated exposure frequency that prevented resolution of the inflammatory response; 3) lower dose rate which could have prevented the strong acute inflammation response; and 4) higher total dose of PM delivered (100 µg versus 50 µg) in the present versus the previous (Yuan et al., 2020) study.

Several reviews have concluded the chemical components of PM can play an important role in post-exposure health effects (Chen and Lippmann, 2009; Rohr and Wyzga, 2012). Therefore, to enact the proper exposure controls (e.g., protective equipment, concentration limits), it is critical to determine which PM components, or combinations thereof, are most harmful to human health (Wyzga and Rohr, 2015). In the present study, repeated exposure to  $PM_{SX}$  frequently produced the greatest differences relative to control (Figs. 6,8, and 9). The combined findings for BALF neutrophil numbers and lung histopathology provided some confirmation that the chemical composition of  $PM_{SX}$  may contribute to greater toxicity than that observed for  $PM_{CA}$  and  $PM_{SD}$  administered on an equal PM mass basis. Differences between the PM-exposed groups were most likely due to differing chemical compositions influenced by regional emission sources such as vehicular exhaust, agricultural and industrial operations in the Sacramento region, or coal production in the Shandong and Shanxi provinces (Plummer et al., 2015; Xia et al., 2013).

Although we were unable to definitively identify the precise chemical constituent(s) that produced the observed biological effects, it appears that organosulfates and polyaromatic hydrocarbons (PAHs) likely contributed to the differential induction of biological effects observed in the present study (Table 1). Sulfate levels measured at 2%, 14 %, and 26 % in  $PM_{CA}$ ,  $PM_{SD}$ , and  $PM_{SX}$ , respectively (Fig. 5a-c). Several studies (Chen and Lippmann, 2009; Krewski et al., 2005) have indicated sulfate-associated particles (i.e., fossil fuel combustion products) are among the most toxic in terms of cardiopulmonary and cardiovascular disease effects and lung cancer mortality. Particulate air pollution, characterized by high secondary aerosol concentrations, including organosulfates, has been a serious environmental problem during recent winters in China (Huang et al., 2018; Wang et al., 2017). Thus, higher fractions of organosulfates are plausible for  $PM_{SX}$  and  $PM_{SD}$  compared to  $PM_{CA}$ . However, in the present study, speciation of sulfates was not possible; therefore, future exposure studies could benefit from quantifying organic and inorganic sulfates as a means to understand better the health risks they pose.

All three PM extracts used in the current study were found to activate the aryl hydrocarbon receptor (AhR), as well as *CYP1A1* mRNA expression in human HepG2 cells (Yuan et al., 2020). These effects are most likely due to PAHs in the carbonaceous fraction of each PM extract. In the current research, compared to PM<sub>CA</sub>, the PAH levels were 2.48 and 11.5 higher in PM<sub>SD</sub> and PM<sub>SX</sub>, respectively (Table 1). PAHs have received considerable attention due to their potential toxic, carcinogenic, and mutagenic effects. Coal combustion is known to produce PAHs (Shen et al., 2010), and in 2012, China's coal consumption was

2.75 billion tons, approximately one-half of the global coal consumption. Shanxi province is one of China's largest coal-producing centers (Yang and Teng, 2018).

Repeated exposure to  $PM_{SX}$  produced a higher BALF neutrophil count than  $PM_{CA}$  and  $PM_{SD}$  on post-OPA day 4 (Fig. 6b). Repeated exposure to PM has been shown to increase BAL neutrophils that may be causally associated with lung injury (Maueroder et al., 2015; Pardo et al., 2016), with the higher numbers of neutrophils entering the lungs typically reflective of the severity of the inflammatory and cellular response. Neutrophil influx was clearly observed in our study, with neutrophils present in the lung interstitium and among epithelial cells (Fig. 7a-d). Treatment groups in which neutrophil numbers in BALF attained a level of statistical significance also exhibited significantly greater histological inflammatory scores in the airways and alveoli on day 4 post OPA (Fig. 6c-d). Neutrophils are generally present in these two areas of the lungs during periods of inflammation.

In the present study, mice exposed to  $PM_{SX}$  exhibited significantly increased mRNA for *TNF-a* and *IL-1β*, cytokines involved in pulmonary inflammatory reactions (Li et al., 2017), on post-OPA days 2 and 4 (Fig. 8a-b), as well as *CXCL-3* and *CXCL-5* neutrophil chemokines (Bhatia et al., 2012; Narayanan et al., 2005) on post-OPA day 4 (Fig. 8c-d) when compared to control-,  $PM_{CA}$ -, and  $PM_{SD}$ -exposed groups. MCP-1 protein was increased in all PM-exposed groups relative to controls on day 4 post OPA, but  $PM_{SX}$  produced the longest-lasting response that was also stronger than that for  $PM_{SD}$  (Fig. 9). MCP-1 is a chemoattractant responsible for monocyte and macrophage recruitment. However, growing evidence shows MCP-1 may also be involved in attracting neutrophils (Balamayooran et al., 2011). When considered cumulatively, the concomitant elevation of cytokine (*TNF-a* and *IL-1β*) and chemokine (*CXCL-3*, *CXCL-5*) mRNA, MCP-1 protein, and BALF neutrophils on post-OPA day 4 suggest multiple chemical mediators may be driving high neutrophil numbers even after 4 days of recovery.

Findings from the present study provide further evidence for potential mechanisms by which PM extracts from different regions of the world promote time-lagged effects. Future studies could use later post-OPA time points to compare persistent pulmonary health effects of repeated exposure to extracts of  $PM_{2.5}$  and test the impact of elevated PM-sulfur on the degree of inflammation. These future studies would significantly add to the existing knowledge to correlate PM composition to pulmonary responses, inform future regulations on source-specific emissions, and better protect and benefit human health.

#### 5. Conclusions

Consistent with universal time-lag effects of PM on health, the results in the current study showed exposure to  $PM_{CA}$ ,  $PM_{SD}$ , or  $PM_{SX}$  produced greater toxicity on post-OPA day 4, compared to days 1 and 2. These results suggest sulfate and PAH content may have influenced the observed toxicity for  $PM_{CA}$ ,  $PM_{SD}$ , and  $PM_{SX}$ . Mice exposed to  $PM_{SX}$  demonstrated the greatest inflammatory responses, as evidenced by statistically significant (p < 0.05) increases in pulmonary neutrophil numbers and pro-inflammatory cytokines and chemokines that may be due to higher sulfate and PAH content, compared to that of  $PM_{CA}$  and  $PM_{SD}$ . In summary, the current study provides compelling evidence that PM extracts

from different regions promote adverse time-lagged health effects influenced by chemical composition. These findings further highlight the need to develop source-specific regulations to protect human health.

#### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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#### HIGHLIGHTS

- Time-lagged effects were evaluated in mice following repeated exposures to one of three PM extract samples.
- All three PM extracts induced progressive inflammatory responses over time.
- These time-lagged effects were influenced by the chemical composition and potentially the PAH content of each PM extract.



#### Fig. 1. PM samples at equal mass concentrations.

The image shows stock extracts of PM from Sacramento, California (left); Jinan, Shandong (middle); and Taiyuan, Shanxi (right) at a concentration of 1  $\mu$ g/ $\mu$ L (1 mg/mL) suspended in nanopure water Milli-Q water. The PM sample from Taiyuan is strikingly darker in appearance.



#### Fig. 2. Exposure design timeline.

The experimental schedules for four different mouse treatment groups are shown, with the x-axis representing the exposure timeline (days). The aspirates were 50  $\mu$ L Milli-Q water (control; white triangles), or a 20 $\mu$ g/50  $\mu$ L PM extract from Sacramento, California (gray triangles); Jinan, Shandong (striped triangles); or Taiyuan, Shanxi (black triangles). Each triangle in the figure represents a single exposure on days 1, 4, 7, 10, or 14. Each "X" represents day 1, 2, or 4 post OPA (shown as days 15, 16, or 18, respectively), when the euthanasias and collection of biological samples occurred. A total of 3 controls and 6 PM-exposed mice were euthanized on each necropsy day.



#### Fig. 3. Experimental protocol.

Four different treatment groups are shown along with their respective oropharyngeal aspiration exposures and post-necropsy processing.



Fig. 4. Hydrodynamic particle size distributions of wintertime PM extracts from Sacramento, California ( $PM_{CA}$ ); Jinan, Shandong ( $PM_{SD}$ ); and Taiyuan, Shanxi ( $PM_{SX}$ ). The graph shows the results from dynamic light scattering analyses. Particle density (intensity %) is shown on the y-axis as a function of particle diameter (size on the x-axis) in nanometers.



Fig. 5. Chemical composition of wintertime PM extracts from Sacramento, California; Jinan, Shandong; and Taiyuan, Shanxi.

Characterization was performed using high-resolution time-of-flight aerosol mass spectrometry. Charts (a-c) in the top row illustrate bulk composition, including organic matter. Charts (d-f) in the bottom row show the average elemental composition of the organic matter. Abbreviations: C- carbon; H – hydrogen; N – nitrogen; O – oxygen.



Fig. 6. Oropharyngeal aspiration of particulate matter (PM) extracts produced elevated BALF neutrophil numbers and tissue inflammation in mice up to four days post exposure. Animals were exposed on five separate days over two weeks to Milli-Q water (Control), or a PM extract ( $20 \ \mu g/50 \ \mu L$ ) from Sacramento, CA; Jinan, Shandong; or Taiyuan, Shanxi (PM<sub>CA</sub>, PM<sub>SD</sub> or PM<sub>SX</sub> respectively). N = 9 for controls, and 18/PM group (6/group/day). Each mouse received a 50- $\mu$ L aspirate on the exposure days (total PM dose = 0 or 100

Each mouse received a 50-µL aspirate on the exposure days (total PM dose = 0 or 100 µg/mouse), and necropsies occurred on day 1, 2, or 4 post exposure. Panels a-b show counts of total bronchoalveolar lavage fluid (BALF) cells and neutrophils, respectively. Panels c-d show semi-quantitative histopathology scores from the bronchiolar and alveolar lung regions, respectively. Final histopathology scores were averaged from the products of the severity and extent scores in each region of the lungs. Data from different time points in each panel were analyzed separately via one-way ANOVAs and presented as the mean  $\pm$  standard error of the mean. Asterisks (\*) indicate significant (p < 0.05) differences from control. Brackets indicate significant (p < 0.05) differences between PM-exposed groups.



#### Fig. 7. Lung tissue inflammation in mice exposed to particulate matter (PM) extracts.

Panels are light micrographs of hematoxylin- and eosin-stained bronchiolar (a-d) and alveolar (e-h) tissues collected on day 4 after the last of five separate 50- $\mu$ L oropharyngeal aspiration exposures to a MilliQ water control (a & e; n = 9) or a 20- $\mu$ g/50  $\mu$ L PM extract (n = 18/group) from Sacramento, CA (b & f); Jinan, Shandong (c & g); or Taiyuan, Shanxi (d & h). The total PM dose was 0 or 100  $\mu$ g/mouse over 14 days. In panels a-d, arrows indicate neutrophils, and arrowheads indicate macrophages. In panels e-h, arrows identify regions of alveolar inflammation. AW = airway, scale bar is 20  $\mu$ m.





Graphs show results from quantitative polymerase chain reaction assays performed on mRNA obtained from the lungs of mice at 1, 2, or 4 day(s) after the last 50-µL oropharyngeal aspiration (OPA) exposure to MilliQ water, or a PM extract (20 µg/50 µL) from Sacramento, CA (PM<sub>CA</sub>); Jinan, Shandong (PM<sub>SD</sub>); or Taiyuan, Shanxi (PM<sub>SX</sub>). N = 9 for controls, and 18/PM group (6/group/day). OPA occurred on five separate days over two weeks, yielding a total PM dose of 0 or 100 µg/mouse. Gene levels were analyzed for each animal and averaged for each treatment group. Gene expression is shown relative to the housekeeping gene, *EEf1a1*. Data are shown as the mean  $\pm$  SEM. One-way ANOVA and Tukey tests were performed at a significance level of p < 0.05. Asterisks (\*) indicate significant (p < 0.05) differences from vehicle control; brackets indicate significant (p < 0.05) differences between non-control groups.



# Fig. 9. Mice exhibited subacute increases in Monocyte Chemoattractant Protein (MCP)-1 levels following exposure to particulate matter (PM) extracts from California or China.

The bar graph shows the results from enzyme-linked immunosorbent assays performed on lung tissues from mice at 1, 2, or 4 day(s) after the last 50- $\mu$ L oropharyngeal aspiration (OPA) exposure to MilliQ water (Control), or a PM extract (20  $\mu$ g/50  $\mu$ L) from Sacramento, CA (PM<sub>CA</sub>); Jinan, Shandong (PM<sub>SD</sub>); or Taiyuan, Shanxi (PM<sub>SX</sub>). N = 9 for total controls, and 6/post-OPA day (1, 2, and 4) for each PM. Over two weeks, OPA occurred on five separate days, yielding a total PM dose = 0 or 100  $\mu$ g/mouse. Separate ANOVAs were performed for each time point to determine inter-group differences due to exposure. Asterisks (\*) indicate significant (*p* < 0.05) differences from control; brackets indicate a significant (*p* < 0.05) difference between PM<sub>SD</sub> and PM<sub>SX</sub>.

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Sample	PAH/Organic ratio	Organic fraction	Percentage of dissolved PM mass (%)	Enhancement factor
$\rm PM_{CA}$	0.00017	0.54	0.00918	1.00
$PM_{SD}$	0.0004	0.57	0.02280	2.48
$\mathrm{PM}_{\mathrm{SX}}$	0.0024	0.44	0.10560	11.5

The method used to determine the total PAH-to-organic PM mass ratio (Table 1) by AMS was first described by Dzepina and colleagues (2007).