INTRODUCTION

Ozone (O$_3$) is a prevalent environmental pollutant and a major public health concern with hospital admissions and emergency room visits increasing following days of high ambient O$_3$ concentrations (Delfino, Murphy-Moulton, Burnett, Brook, & Becklake, 1997; Stieb, Burnett, Beveridge, & Brook, 1996). Ground level O$_3$ is formed by
pholysis of reactive intermediates largely created from the burning of fossil fuels (Uysal & Schapira, 2003). O₃ is responsible for acute respiratory irritation leading to difficulty in breathing (Schnell & Prather, 2017), especially in patients with pre-existing respiratory disease such as asthmatics and in the elderly population (Dauchet et al., 2018; Fry et al., 2012; Rush, Wiskar, Fruhstorfer, Celi, & Walley, 2018). Even O₃ concentrations below the recommended 8 hr standards of 0.063 ppm (Vanos, Cakmak, Kalkstein, & Yagouti, 2015) are sufficient to reduce lung function in asthmatic children. According to the latest report issued by the Canadian Census Health and Environment Cohort (CanCHEC), Saskatchewan or the Eastern Prairie region is amongst the highest risk zones for O³ exposure and particulate matter related lung cancer, ischemic heart disease and COPD (Cakmak et al., 2018). Thus, it is imperative to understand the biological effects of O₃ at concentrations feasible in the environment.

Acute lung injury (ALI) often results from excessive lung damage that leads to alveolar barrier disruption with secondary lung edema and progressive loss of lung function. Bacterial pneumonia or endotoxin (or lipopolysaccharide, LPS) exposure induce immediate (i.e., within 1 hr) release of IL-1β dependent cytokines and chemokines and lead to enzymatic cleavage of plasminogen/plasmin into angiostatin fragments (Falcone, Khan, Layne, & Fernandes, 1998; Hamacher et al., 2002). Innate immune chemokines such as tumor necrosis factor (TNFα), interleukin-1β (IL-1β), keratinocyte chemotactic peptide-1 (MIP1α) lead to increased expression of adhesion molecules like platelet endothelial cell adhesion molecule-1 (PECAM-1 or CD-31), P-selectin (CD-62P), and CD11b (integrin αM). These molecules mediate platelet, neutrophil, and mononuclear infiltration into the lungs (Berman & Muller, 1950; Grommes et al., 2012; Schenkel, Chew, Chipala, Harbord, & Muller, 2006). Animal models have also helped characterize potential mechanisms of pulmonary (Dowell, Lohrbauer, Hurst, & Lee, 1970; Michaudel, Fauconnier, Jule, & Ryffel, 2018) Bhattacharya & Westphalen, 2016) as well as multiple organ injury (Bouthillier et al., 1998; Erickson et al., 2017; Henriquez et al., 2018; Kasahara et al., 2014; Thomson, Pilon, Guenette, Williams, & Holloway, 2018) resulting from O₃ exposure. These studies show that a 3–6 hr exposure to 0.8 or 2 ppm O₃ induces recruitment and accumulation of lung monocytes and neutrophils at 24 hr (Francis et al., 2017; Tighe et al., 1950). However, there is paucity of data on the early (within 1 hr) lung cellular profiles in response to O₃ exposure. To this end, we ascertained the immediate alveolar as well as vascular (pulmonary and systemic) inflammatory cell recruitment and lung chemokine gradients in response to O₃ exposure in concentrations upto 0.05 ppm. We hypothesized that a low level of O₃ is capable of inducing the hallmarks of lung injury, namely CD11b-dependent Gr1 leukocyte recruitment, lung edema, histological evidence of lung damage, chemokine release and enhanced expression of plasminogen and angiostatin fragments and adhesions molecules (CD-31, CD-62P).

As O₃ potently induces instant cell death, we asked the question if 2 hr of continuous exposure to 0.05 ppm O₃ concentration will cause enough damage to the alveolar cytoskeleton to induce an adaptive response, which is accompanied by mitochondrial activation leading to upregulation of cell survival proteins such as ATP synthase (ATPβ) and signaling molecules such as reactive oxygen species (ROS). We, thus, performed intravital lung imaging to assess O₃-induced damage to the alveolar actin cytoskeletal arrangement, and the effect on mitochondrial ROS generation and mitochondrial potential. In addition, we stained lung sections for regional lung ATPβ expression and quantification.

2 | MATERIALS AND METHODS

2.1 | Mice

The study design was approved by the University of Saskatchewan’s Animal Research Ethics Board and adhered to the Canadian Council on Animal Care guidelines for humane animal use. Six-eight week old male C57BL/6NJ (Stock No. 005304) mice were procured from Jackson Labs.

2.2 | Reagents and chemicals

All chemicals utilized for the experiments were purchased from Sigma-Aldrich Chemicals unless otherwise indicated. Intravenous dyes such as Alexa 555 tagged phalloidin (Life Technologies Inc.) was superfused to highlight intracellular actin dynamics. Intracellular reactive oxygen species (ROS) were detected by the IMAGE-iT LIVE Green ROS Detection Kit which consists of 5- (and-6)-carboxy-2’,7’-dichlorodihydrofluorescein diacetate (carboxy-H2DCFDA, Life Technologies Inc.) was superfused to highlight intracellular reactive oxygen species (ROS). We, thus, performed intravital lung imaging to assess O₃-induced damage to the alveolar actin cytoskeletal arrangement, and the effect on mitochondrial ROS generation and mitochondrial potential. In addition, we stained lung sections for regional lung ATPβ expression and quantification.

2.3 | Ozone exposures

For O₃ exposures, mice were continuously exposed in an induction box for the desired times. These mice had free access to food and water while housed in the custom induction box. O₃ (±0.02 ppm) was generated, at 3 L/minute, from ultra-high-purity air using a silent-arc discharge O₃ calibrator cum generator (2B Technologies). Constant chamber air temperature (72 ± 3°F) and relative humidity (50 ± 15%) were
maintained. O₃ concentrations were measured using a real-time O₃ monitor (2B Technologies).

**2.4 | Experimental design**

**2.4.1 | Ozone time and dose titration**

Six control mice were exposed to room air (RA) in a custom induction chamber for 2 hr (designated as group I or 0 hr). In groups II and III, six mice were continuously exposed for a short (2h) duration at 0.005 ppm and 0.05 ppm O₃, respectively. In group IV, three mice were continuously exposed for a longer (4 hr) duration and at a higher concentration of 0.5 ppm O₃. In group V, three mice were continuously exposed for the longest (24 hr) duration but to the lowest (0.005 ppm) O₃ concentration to investigate the effects of prolonged O₃ exposure at concentrations that are considered safe for humans. Immediately after respective O₃ exposures (i.e., 0, 2, 4, or 24 hr), mice were anesthetized intraperitoneally with ketamine (200 mg/kg) and xylazine (10 mg/kg) and prepared for further maneuvers. The experimental design is summarized in Figure 1a. Peripheral blood was collected by cardiac puncture. Mice were then tracheostomized, a custom noncollapsible polyethylene cannula was placed just before the end of tracheal bifurcation. Broncho-alveolar lavage (BAL) was collected with three consecutive washes, each with 0.5 ml PBS (phosphate buffered saline). The descending thoracic aorta was snipped at the mid-thoracic region. After clearing residual blood, lung vascular perfusate (LVP) was collected by perfusion through the right ventricle with PBS (0.25 ml × 2). The right lung lobes were tied, resected, and flash frozen in liquid nitrogen and then stored at −80°C for further analysis. The left lung was perfused with 0.5 ml freshly prepared 4% paraformaldehyde from a 20 cm water column to enable in situ fixation for 10 min and later cryo-embedded. The collected samples are shown in the schematic in Figure 1a. Thus, for groups I-III (3 × 3 = 9 mice),

**FIGURE 1** Experiment design. (a) Experiment design for room air (RA) (i.e., 0 hr or control (Group I, N = 6), ozone (O₃) at 0.005 ppm for 2 hr (Group II, N = 6), 0.05 ppm for 2 hr (Group III, N = 6), 0.5 ppm for 4 hr (Group IV, N = 3), or 0.005 ppm for 24 hr (Group V, N = 3 per group) in the C57BL/6NJ sub-strains (black mice). Schematic of cellular compartments assayed from the lungs (Bronchoalveolar lavage/BAL, lung vascular perfusate/LVP) and peripheral/cardiac blood of each mouse. BAL and LVP were assayed for Total leukocyte counts/TLC, differential leukocyte counts/DLC, total proteins, chemokines and BAL CD11b and Gr1 localization. Left lungs were stained with H&E for histology. (b) Experimental design for intravital lung imaging in control or room air (RA) 0 hr (Group I) and ozone (O₃) (0.05 ppm for 2 hr) imaged at 2 (Group II) or 24 (Group III) h. N = 9 per group: From each group, 3 mice each were imaged for actin (Rhodamine phalloidin, 1:1,000 dilution), reactive oxygen species (ROS) (5-(and-6)-carboxy-2',7'-dichlorodihydrofluorescein diacetate (carboxy-H₂DCFDA), 1:1,000 dilution), and mitotracker potential (Reduced mitotracker, 1:1,000 dilution)
three additional mice were added in order to perform BAL immune-fluorescence (Gr1 and CD11b) staining, apart from three mice per group (groups I-V, 3 × 5 = 15 mice) used for bright-field cytology and protein quantifications from BAL, vascular perfusate and blood samples, and lung histology.

2.4.2 | Intravital imaging of ozone lung exposure

The objective of this experiment was to compare O₃-induced dynamic changes in lung architecture (actin), mitochondrial activity, and ROS production by intravital microscopy to those of control mice exposed to room air (RA). Nine control mice were exposed to RA in a custom induction chamber for 2 hr. Immediately after RA exposure, the mice were prepared for intravital lung microscopy. These mice are designated as Group I or 0 hr time-point. For Group II (or 2 hr time-point), nine mice were exposed to 0.05 ppm O₃ for 2 hr. Immediately after O₃ exposure (that is at 2 hr time-point), mice were prepared for intravital lung microscopy. For Group III (or 24 hr time-point), nine mice were exposed to 0.05 ppm O₃ for 2 hr and recovered for 22 hr in a normal RA cage, after which the mice were prepared for intravital lung microscopy (that is at 24 hr time-point). For each of these three groups, lungs from three mice were superfused with Alexa 555-phalloidin for actin visualization, three mice with the ROS dye (carboxy-H₂DCFDA) for ROS imaging, and three mice for reduced mitotracker orange for active cellular mitochondria visualization. The experimental design for the live imaging experiments is summarized in Figure 1b.

2.5 | End-points

We evaluated end-points “a through f” for the O₃ time and dose titration study and the endpoint “g” for the intravital lung imaging study.

2.5.1 | Total (TLC) and differential (DLC) leukocyte counts

Blood, BAL and lung vascular perfusate samples were centrifuged for 10 min at 3,000 rpm. The supernatants were flash frozen and stored at −80°C until further analysis. The cells were reconstituted in PBS. TLC was performed by counting BAL, blood, or lung perfusate cells on a hemocytometer. Trypan blue dye exclusion was utilized to quantify live and dead BAL cells under light microscopy. Blood TLC is presented as x10⁶ cells per mL. BAL and lung perfusate TLC are expressed in x10⁸ and x10⁶ cells per collection, respectively. Acetic acid (2%) was added to lyse RBCs, in a 1:2 ratio for blood TLC and 1:2 ratio for lung vascular perfusate TLC. The cells (not more than 1 x 10⁶ per slide) were then centrifuged (Shandon cytopsin, Thermoscientific) to prepare cytopsins and stained with Diffquick for DLCs. A minimum of 200 cells were counted for differential leukocyte cell counts (DLCs) or debris. Debris was identified as incomplete or damaged cellular fragments, extra-cellular nuclear material or condensed cells. A clump of such material was accounted as a count of one when quantifying debris.

2.5.2 | BAL cytopsin CD11b and Gr1 staining

BAL cytopsin samples from groups I, II, and III were stained for the CD11b and Gr1 surface markers (2 µg anti-CD11b PE, and 2 µg anti-Gr1 Alexa 488, respectively). Nuclei were stained with DAPI (4’,6-diamidino-2-phenylindole). Cytopsin preparations were also stained with rat IgG2b kappa isotype control antibody to validate the staining protocol. Images were acquired under a wide-field upright microscope equipped with a Sony camera.

2.5.3 | Lung immune-fluorescent staining

Lung cryosections from groups I-IV were stained with two different primary antibody combinations. Combination 1 was comprised of direct immune-fluorescent staining with Gr1-488 and CD11b-PE-labeled antibodies. The rest of the combinations were comprised of indirect immunofluorescent staining protocols. Combination 2 was comprised of either mouse ATPβ and rabbit plasminogen antibodies or mouse ATPβ and rabbit CD31 (to visualize ATPβ localization with respect to endothelial cells). Combinations 3 and 4 were stained from the short RA or O₃ exposure groups I and III to ascertain the immediate changes in lung plasminogen, angio-statin, CD62P, and vwf protein fluorescence. Combination 3 comprised of rabbit vwf and rat Gr1 antibodies. Combination 4 comprised of goat CD62P (P-selectin) and rabbit angiostatin antibodies. The respective secondary antibody combinations were anti-mouse Alexa 488 and anti-rabbit Alexa 568 for combination 2, anti-rabbit Alexa 488 and anti-rat Alexa 568 for combination 3, and anti-goat Alexa 488 and anti-rabbit Alexa 568 for combination 4.

Briefly, all sections were hydrated in tris buffered saline (TBS). Before overnight incubation with primary antibodies at 4°C, sections were permeabilized with 0.1% triton X-100 for 5 min, followed by washes in TBS buffer. Nonspecific binding was blocked with 5% bovine serum albumin for 30 min at room temperature. For combination 2, which comprised of ATPβ antibody raised in mouse, an additional 20 min room temperature blocking step was
introduced with Fc blocking buffer. Following overnight primary antibody incubation in a humidified chamber at 4°C and TBS washes to remove the unbound primary antibody, secondary antibody was then incubated for 30 min at room temperature. DAPI was incubated during the last 10 min before the secondary antibody wash. Sections were cover slipped with ProlongGold™ anti-fade mounting media and sealed after overnight drying in the dark at room temperature.

2.5.4 | BAL, lung vascular perfusate, and blood protein analysis

To quantify O₃-induced perturbations of the vascular barrier or the relative oncotic pressure in the various compartments, that is, the alveolar (interstitial), lung vascular perfusate (vascular), and peripheral blood (systemic) compartments, we measured total protein content in the collected fluids. Supernatant fractions were analyzed for their total protein concentration using a standard colorimetric assay (Pierce 660 nm protein assay, Thermoscientific).

2.5.5 | BAL, lung vascular perfusate, and bone marrow chemokine analysis

BAL supernantants, from groups I-V, were analyzed for chemokines using a 33-plex magnetic bead-based immunoassay (Bio-rod Laboratories Ltd.). To understand the chemokine gradients established by 24 hr of continuous O₃ exposure, in addition to avoiding prohibitive cost of 33-plex assays. The following chemokines were analyzed: CXCL13 (B-lymphocyte chemoattractant), CCL27 (IL-11 R-alpha-locus chemokine (ILC)), CXCL5 (epithelial-derived neutrophil-activating peptide 78 (ENA-78)), CCL-11 (eotaxin-1), eotaxin-2 (CCL-24), CX3CL1 (fractalkine), GM-CSF (CSF-2), CCL1, IFNγ (interferon gamma), IL-10 (interleukin-10), IL-16 (interleukin-16), IL-1β (interleukin-1 beta), IL-2 (interleukin-2), IL-4 (interleukin-4), IL-6 (interleukin-6), CXCL10 (interferon gamma-inducible protein 10 (IP-10)), CXCL11 (Interferon-gamma-inducible protein 9 (IP-9)), KC (keratinocyte chemoattractant), MCP-1 (monocyte chemoattractant protein-1), MCP-3 (monocyte chemoattractant protein-3), MCP-5 (monocyte chemoattractant protein-5), MDC (macrophage-derived chemokine (CCL22)), MIP-1α (macrophage inflammatory protein-1 alpha), MIP-1β (macrophage inflammatory protein-1 beta), MIP-2 (macrophage inflammatory protein-2), MIP-3α (macrophage inflammatory protein-3 alpha), MIP-3β (macrophage inflammatory protein-3 beta), RANTES (regulated on activation, normal T cell expressed and secreted (CCL5)), CXCL-16, CXCL-12/SDF-1alpha (stromal cell-derived factor 1), TARC (thymus and activation regulated chemokine (TARC)), TECK (Thymus-Expressed Chemokine (CCL25)), and TNFα (tumor necrosis factor alpha).

2.5.6 | Hematoxylin and eosin (H&E) histology

H&E staining was performed on cryoembedded lung sections for all groups.

2.5.7 | Lung intravital microscopy

Briefly, anesthetized and intubated mice were ventilated with positive end expiratory pressure (PEEP) settings of 2 cm H₂O pressure, a respiratory rate of 100–120 breaths per minute and a tidal volume of 200 µl. Lungs were exposed by carefully cutting the ribs from the left latero-dorsal side after which the lungs were gently suctioned under 25–30 mm Hg vacuum to a custom-made optically transparent thoracic window (Looney et al., 2011). An Olympus BX51W1 fixed-stage Gibraltar upright microscope was upgraded to laser powered sources of 488 and 561 nm excitation wavelengths (Coherent Light Source), for performing intravital microscopy. The lasers were coupled through a liquid light guide. A DAPI/GFP/TRITC triple handpass dichroic excitation emission filter, corresponding to 405/488/561 nm, was mounted on an Olympus BX2 cube. The transmitted visible light was collected by a consumer-grade Canon HV20A CMOS RGB camera. These signals were displayed and recorded on a computer monitor utilizing a video capture card. The fluorescent emission was collected using a scientific grade ICCD XR/Mega 10 Extreme camera (Stanford Photonics, Inc.). The effective pixel size of the ICCD camera was 10 µm with an image capture rate of 30 fps. To minimize laser toxicity for the two-light fluorescent signal acquisition, a camera trigger with a frame rate of 15 fps for the first laser excitation was followed by a 2-s pause and then captured with the second laser excitation at 15 fps. The signals were displayed and recorded onto a computer monitor via image acquisition and control software (PiperControl; Stanford Photonics Inc.). Videos were acquired under 10× air or 20× water objectives with numerical apertures of 0.30 and 0.95, respectively. Videos were acquired for a maximum of 30 min to avoid laser damage to tissue. This time was considered adequate without inducing any significant laser-induced tissue damage as assessed by changes in the average baseline ROS concentration and reduced mitotracker intensity over time.
2.6 | Image analysis

Collected image data (.tiff) files were processed and analyzed in Fiji ImageJ (https://imagej.net/Fiji/Downloads). For DiffQuick stained BAL, lung perfusate, and blood cytospins, the oil objective images were inverted and adjusted for enhanced local contrast to visualize damaged cells, extracellular nuclear material, platelets and platelet clumps. A total of 200 cells were counted manually to calculate the DLC.

For CD11b and Gr1 stained BAL cytospins, between 50 and 100 cells were manually thresholded and the corresponding regions of interest were selected in the CD11b and Gr1 channels. A log fold drop in fluorescent intensity was considered as the negative population of the corresponding fluorescent signal. Percentage of positive or negative cells were calculated and reported. For the positive cells, the integrated fluorescent pixel intensities and pixel frequency distributions (histograms) were objectively analyzed for CD11b and Gr1 expression by each cell. Cellular fluorescent intensity distributions (integrated fluorescence intensity, maxima, standard deviations in the fluorescence intensities, maximum fluorescence intensity) as well as pixel frequency and spatial distributions (skewness and kurtosis, respectively) were plotted for Alexa488 (Gr1) and the PE (CD11b) channels. Cell perimeters were also plotted to evaluate any change in cell size after O₃ exposure. The average intra-assay and inter-assay coefficient of variation (CV) for integrated fluorescence intensity of Gr1 in control group are 0.502 and 0.137, respectively. The average intra-assay and inter-assay CV for integrated fluorescence intensity of CD11b in control group are 0.503 and 0.116, respectively.

For lung immune-fluorescent quantification, bronchiolar, vascular (arteriolar and venular) as well as alveolar septal (alveolar) regions were manually thresholded, and perimeter normalized fluorescence intensity was analyzed.

The actin, ROS, and mitotracker image sequences were registered using the rigid registration plugin in Fiji. The 16-bit images were only corrected for brightness and contrast and not thresholded. Features such as air-spaces were selected using the feature’s pixel threshold and analyzed over multiple frames (100–200) to detect respiratory motion. For ROS and mitotracker image sequences, each sequence was compared on a common scale (to cover the entire histogram range) and visualized on a color scale. Motion analysis of alveolar area, ROS or mitotracker positive lung parenchyma in a sub-pleural field of view was executed after thresholding the air-spaces from stained area, converting the desired sequence into binary images and then computing the percentage of area covered by the desired read-out (i.e., alveolar space, ROS, or mitotracker positive) out of the full field of view.

For ROS and reduced mitotracker image analysis, the difference images (T1-T2, that is, start and end images for 10 min of acquisition) were an inverse of the difference image is displayed for visualization of the respective activities. The (average and maximum) fluorescent intensity gray values are reported from the original 16-bit images obtained from ROS, reduced mitotracker or actin stained sub-pleural alveoli. The image sequences were converted and saved as. avi (for the ICCD camera) or.mp4 (for the Canon camera) movie files.

2.7 | Statistical analysis

The results are expressed as mean ± SEM. A minimum of three mice were used per group. Two-way ANOVA was used to identify the effect of 2 factors, cell type (eosinophils, neutrophils, mononuclear cells, platelets, platelet clumps, debris, and total cell counts) and ozone exposure (0 hr, 0.005 2 hr, 0.05 2 hr, 0.5 4 hr, 0.005 24 hr), after homoscedasticity testing. Bonferroni’s multiple comparisons (which corrects for the variance from multiple factors (7 cell types × 5 exposure groups) as well as adjusts the p-values) were then applied for analyzing effects between groups for each cell-type. For chemokine data analysis, one-way ANOVA p-values were adjusted for false discovery rate according to the Benjamini and Hochberg correction. Pulmonary capillary diameters were analyzed by Mann–Whitney U test as data were not normally distributed. The alveolar perimeter and circularity were analyzed by unpaired two-tailed t test. For rest of the imaging experiments, we analyzed the results using a one-way or two-way ANOVA followed by Sidak’s multiple comparisons, depending on the number of derived parameters in analysis. p-values < .05 were considered significant. All data analysis was performed on Graphpad Prism (v8.3.0) software.

3 | RESULTS

3.1 | Low O₃ levels induce marked alveolar cell death, and platelet and neutrophil pooling in the LVP

At baseline (Group I, time 0 hr), >90% of cells in the BAL were mononuclear. Exposure to 0.005 (Group II) and 0.05 ppm O₃ (Group III) for 2 hr induced an 8.3% (p = .0427) and 34.6% (p < .01) reduction, respectively, in the recovered BAL cells, when compared to control or 0 ppm O₃ (Group I) (Figure 2a). A further increase in the O₃ dose to 0.5 ppm and exposure time to 4 hr (Group IV) reduced BAL cell recovery by 93.6% of control. Longer (24 hr) exposure to 0.005 ppm O₃ (Group V) also suppressed the BAL cell recovery (Figure 2a) by 36.1% of control. BAL cells recovered
after O₃ exposures were either mononuclear or dead cells (Figure 2a,b). Although not shown, we confirmed this finding with viable cell counts following trypan blue dye exclusion. Longer (24 hr) O₃ exposure induced accumulation of phlogistic debris in the BAL, comprised of extracellular nuclear material and dead cells (Figure 2b).

Along with the decrease in BAL cell counts, O₃-induced significant increase in lung vascular perfusate and peripheral blood TLC (p < .01, Figure 2d,e). In addition, exposure to increasing concentrations of O₃ also led to a dramatic change in the nature of the cells seen in the vascular perfusate. Short (2 hr) 0.05 ppm O₃ exposures (Group III) induced significant accumulation of the neutrophils (20%–40%, p < .01), platelets (>30%, p < .01) as well as platelet clumps (p < .01) in the lung vascular perfusate (Figure 2d) and peripheral blood (Figure 2c,e). Longer O₃ exposures (0.5 ppm 4 hr or 0.005 ppm 24 hr) led to a significant increase in the platelets (>40%–90% of the total cells) in lung vascular perfusate (Figure 2d) and peripheral blood (Figure 2c,e).

The suppression in neutrophil counts, accumulation of chromatin cellular debris and platelet clumps in the vascular compartments indicate extensive cell death, dysregulated hemostasis, and platelet activation after prolonged O₃ exposures (4 and 24 hr).

### 3.2 Low levels of O₃ affects total protein content in BAL, LVP, and peripheral blood

Exposure to 0.05 ppm O₃ for 2 hr (Group III) induced a 3.23-fold increase in the BAL protein (Figure 3a). Longer exposures of 4 (Group IV) or 24 hr (Group V) produced a net reduction in BAL protein content (Figure 3a) indicating hypo-osmotic alveolar fluid. The 0.05 ppm 2 hr O₃ exposure...
reduced lung vascular perfusate protein \( (p < .01, \text{Figure 3b}) \) again indicating hypo-osmotic lung vascular compartment. Longer exposures to O\(_3\) (Group IV and V), instead, increased the protein in lung vascular perfusate (Figure 3b). Although the longer O\(_3\) exposures reduced total proteins in BAL compared to the lung vascular perfusate, these numbers correlated with the total cell counts in both the compartments (Figure 2a,b, Figure 3a,b).

The shorter (2 hr) 0.05 ppm or long (24 hr) 0.005 ppm O\(_3\) exposures did not affect peripheral blood protein content compared to group I (0 ppm) but O\(_3\) exposures of 0.5 ppm for 4 hr (Group IV) reduced total protein in peripheral blood \( (p < .01, \text{Figure 3c}) \) indicating hypo-osmotic peripheral blood.

### 3.3 Low levels of O\(_3\) induce lung hemorrhage and bronchiolar damage

Short (2 hr) O\(_3\) exposures of 0.005 ppm (group II) or 0.05 ppm (group III) induced signs of alveolar, peri-vascular, and peri-bronchiolar hemorrhage in lung tissues compared to group I (Figure 4). Endothelial and bronchiolar cell damage was marked in groups IV and V (Figure 4a), indicating dose and time-dependent progression of lung damage. Notably, group IV showed frank hemorrhage and group V showed fibrin clusters within alveoli. O\(_3\) at 0.05 ppm for 2 hr, also induced coagulative inflammatory changes in multiple organs (data not shown).
3.4 Low levels of O₃ induce recruitment of distinct CD11b positive as well as negative mononuclear cells

0.05 ppm O₃ (Group III) exposure for 2 hr induced recruitment of larger BAL cells when compared to control (Group I) or 0.005 ppm of O₃ exposure for 2 hr (Group II) (Figure 5a). More than 95% of the BAL cells were CD11b- and Gr1-positive cells in the control Group I (Figure 5b). Both 0.005 and 0.05 ppm O₃ exposures for 2 hr, induced a significant decrease in the total number as well as proportion of BAL CD11b (up to 40.38%, \( p < .01 \)) and Gr1 (up to 21.27%, \( p < .01 \))-positive cells (Figure 5b). It must be noted that > 95% of the Gr1-negative population in Groups II and III were also negative for CD11b.

Among the CD11b-positive cells, O₃ induced recruitment of distinct (brighter) mononuclear BAL CD11b cells, especially in group III compared to control (group I, Figure S1a–c, https://figshare.com/s/83fe2fdabb32e0d954a). The brighter CD11b cells were also larger (Figure 5a, Figure S1g, https://figshare.com/s/83fe2fdabb32e0d954a, H) These CD11b-positive cells had higher maximum (Figure 5c) and total fluorescence intensity (Figure 5d), were positively skewed (i.e., a higher bright CD11b frequency distribution, as shown in Figure 5e), had higher kurtosis (i.e., a discrete CD11b spatial distribution, as shown in Group III Figure 5f), and more variation in the CD11b distribution compared to control (Figure 5g).

O₃ exposure did not induce recruitment of more Gr1 cells when compared to control (group I) (Figure S1d,e,f, https://
CD31 is an adhesion molecule expressed on endothelial cells as well as some leukocytes. The vascular CD31 expression was not significantly affected by O₃ exposures when compared to the control group I (Figure 6c). Alveolar and bronchiolar CD31 expression were reduced after O₃ exposures, when compared to control group I. However, in the bronchiolar region, longer, that is, 4 hr 0.5 ppm O₃ exposure, did not affect CD31 protein expression ($p > .05$, Figure 6c, Figure S2b, https://figshare.com/s/83fe2fdabb32e0d954a).

O₃ exposure at 0.05 ppm for 2 hr (Group III), enhanced vascular ATPβ (Figure 6d, Figure S2b, https://figshare.com/s/83fe2fdabb32e0d954a), vascular and alveolar plasminogen (Figure 6e, Figure S2c, https://figshare.com/s/83fe2fdabb32e0d954a) and bronchiolar angiotatin (Figure 6f, Figure S2d, https://figshare.com/s/83fe2fdabb32e0d954a) normalized fluorescence. ATPβ is predominantly localized in endothelial cells, and is elevated by fourfold after 0.05 ppm of O₃ exposure for 2 hr (Figure 4d, $p < .01$). An interesting pattern was observed in the alveolar region where 2 hr exposure to 0.05 ppm O₃ enhanced the alveolar ATPβ intensity while 4 hr exposure to 0.005 ppm O₃ reduced the alveolar ATPβ fluorescence intensity (Figure 4d), likely due to extensive alveolar damage. Group III CD62P (Figure 6g, Figure S2e, https://figshare.com/s/83fe2fdabb32e0d954a) and vwf (Figure 6h, Figure S2f, https://figshare.com/s/83fe2fdabb32e0d954a) fluorescent intensities, showed large standard deviation, when compared to control Group I (in vascular as well as alveolar regions).

Thus, lung CD11b and Gr1 protein quantification corroborates our BAL findings. Interestingly, we observed an increase in the bronchiolar expression of angiotatin, while plasminogen and ATP synthase β subunits were observed largely in the vascular and alveolar regions.

### 3.5 | O₃ upregulates lung CD11b, ATPβ, plasminogen, angiotatin, CD62P, and vwf expression

Shorter O₃ exposures of 2 hr reduced CD11b expression in vascular, bronchiolar as well as alveolar regions (Figure 6a), with significant suppression at 0.05 ppm (Group III) (note Groups II and III had majority of CD11b positive cells in BAL). The longer (4 hr) and higher (0.5 ppm, Group IV) O₃ exposure led to significant CD11b expression in the vascular and bronchiolar, but not the alveolar regions, when compared to control (Group I) (Figure 6a, Figure S2a https://figshare.com/s/83fe2fdabb32e0d954a). The alveolar CD11b expression was the least after short 0.05 ppm (Group III) O₃ exposures when compared to baseline (Group I), or after short 0.005 ppm (Group II) (Figure 6a). The alveolar CD11b expression was higher after 4 hr of 0.5 ppm exposure (Group IV) compared to Group II (Figure 6a). As longer O₃ exposures yielded negligible BAL cell recovery, this is also, why we did not perform IF BAL CD11b and Gr1 staining in groups IV and V. The lung CD11b quantification shows that there is a reciprocal relationship between BAL and lung CD11b-positive cells. The percentage of CD11b-positive cells that are recovered from BAL are correspondingly absent from the alveolar region, after 2 hr of 0.005 or 0.05 ppm O₃ exposure.

O₃ at 0.005 (Group II) or 0.05 ppm (Group III) for 2 hr and 0.5 ppm for 4 hr (Group IV) significantly blocked Gr1 expression in all the lung regions (Figure 6b, Figure S2a https://figshare.com/s/83fe2fdabb32e0d954a) when compared to control (Group I).

**FIGURE 6** Lung immunostaining. Lung cryosections (5 µm) were analyzed for normalized expression ratio of protein markers in vascular, bronchiolar and alveolar regions for groups I-IV for (a) CD11b, (b) Gr1, (c) CD31 (Platelet endothelial cell adhesion molecule-1, PECAM-1), (d) ATP synthase β subunit (ATPβ), and groups I and III for (e) Plasminogen (PLG), (f) Angiostatin (ANG), (g) P-selectin (CD62P), and (h) von Willebrand factor (vWF). *$p < .05$, **$p < .01$ represent pair-wise statistical significance. The data points for each labelled protein depict multiple lung regions analyzed in three mice. Data are analyzed with a two-way ANOVA after averaging the read-outs of multiple regions from a single mouse and analyzing the average fluorescent intensity from 3 mice.
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**Figure 1.**

**a**
- Normalized CD1b Expression Ratio
- Lung region \( p < 0.01 \)
- Ozone exposure \( p < 0.01 \)
- Interaction \( p < 0.01 \)
- Group I
- Group II
- Group III
- Group IV

**b**
- Normalized Gr1 Expression Ratio
- Lung region \( p < 0.01 \)
- Ozone exposure \( p < 0.01 \)
- Interaction \( p < 0.01 \)
- Group I
- Group II
- Group III
- Group IV

**c**
- Normalized CD31 Expression Ratio
- Lung region \( p = 0.39 \)
- Ozone exposure \( p < 0.01 \)
- Interaction \( p < 0.01 \)
- Group I
- Group II
- Group III
- Group IV

**d**
- Normalized ATP1a Expression Ratio
- Lung region \( p = 0.01 \)
- Ozone exposure \( p < 0.01 \)
- Interaction \( p = 0.07 \)
- Group I
- Group II
- Group III
- Group IV

**e**
- Normalized PLG Expression Ratio
- Lung region \( p = 0.26 \)
- Ozone exposure \( p < 0.01 \)
- Interaction \( p = 0.06 \)
- Group I
- Group III

**f**
- Normalized ANG Expression Ratio
- Lung region \( p = 0.08 \)
- Ozone exposure \( p < 0.01 \)
- Interaction \( p = 0.17 \)
- Group I
- Group III

**g**
- Normalized CDO2P Expression Ratio
- Lung region \( p = 0.56 \)
- Ozone exposure \( p = 0.83 \)
- Interaction \( p = 0.55 \)
- Group I
- Group III

**h**
- Normalized VILF Expression Ratio
- Lung region \( p < 0.01 \)
- Ozone exposure \( p = 0.50 \)
- Interaction \( p = 0.33 \)
- Group I
- Group III
FIGURE 7  Ozone (O₃)-induced lung (bronchoalveolar lavage (BAL), lung vascular perfusate, and bone marrow) chemokine 33-plex analysis. A Concentrations (pg/ml) of the 18 of 33 chemokines analyzed in bronchoalveolar lavage (BAL) fluid from groups I-V, lung vascular perfusate (LVP) and bone marrow (BM) from group V: (a) IL-1β, (b) IL-2, (c) IL-6, (d) IL-10, (e) IL-16, (f) TNFα, (g) IFNγ, (h) GM-CSF, (i) SDF-1a, (j) Eotaxin-1, (k) Eotaxin-2, (l) RANTES, (m) CXCL5, (n) KC, (o) MIP-1β, (p) MIP-2, (q) MIP-3β, and (r) CCL27. BAL from group I is shown in blue, group II BAL in red, group III BAL in green, group IV BAL in purple, group V BAL in orange, group V LVP in black and group V BM in brown bar graphs. Data are analyzed by one-way ANOVA and the p-value for false discovery rate of multiple variables was adjusted as per Benjamini and Hoshberg correction. * represents p < .05, ** represents p < .01
2 hr of 0.05 ppm O₃ exposure reduced IL-6 (Figure 7c), IL-10 (Figure 7d), TNFα (Figure 7f), RANTES (Figure 7l), KC (Figure 7n), CCL27 (Figure 7r), MIP-1β (Figure 7o), and MCP1 (Figure S4i, https://figshare.com/s/83fe2dabbc32e0d954a) concentrations in BAL when compared to group I. An increase in exposure time from 2 (group II) to 24 hr (group V), at 0.005 ppm O₃, released both immuno-suppressive as well as stimulant chemokines in BAL, like IL-1β (3.28 fold, Figure 7a), IL-2 (10.58 fold, Figure 7b), IL-6 (26.08 fold, Figure 7c), IL-10 (9.81 fold, Figure 7d), TNFα (23.49 fold, Figure 7f), IFNγ (5.53
fold, Figure 7g), GM-CSF (17.49 fold, Figure 7h), SDF-1a (4.45 fold, Figure 7i), RANTES (5.95 fold, Figure 7i), CXCL5 (Figure 7m), KC (18.18 fold, Figure 7m), MIP-1β (3.77 fold, Figure 7o), MIP2 (4.91 fold, Figure 7p), MIP3β (5.08 fold, Figure 7q), CCL-27 (2.69 fold, Figure 7r), eotaxin-1 (Figure 7j), CXCL10 (2.45 fold, Figure S4a, https://figshare.com/s/83fe2fdabb32e0d954a), CXCL11 (8.07 fold, Figure S4b), MCP-1 (14.79 fold, Figure S4i, https://figshare.com/s/83fe2fdabb32e0d954a), CX3CL1 (6.84 fold, Figure S4f, https://figshare.com/s/83fe2fdabb32e0d954a), TARC (3.19 fold, Figure S4g, https://figshare.com/s/83fe2fdabb32e0d954a), TECK (2.87 fold, Figure S4h, https://figshare.com/s/83fe2fdabb32e0d954a) and (Figure S3, https://figshare.com/s/83fe2fdabb32e0d954a, Figure 7a).

To assess the chemokine gradients established in group V, we also assessed the chemokines in lung vascular perfusate and the bone marrow. Concentrations of CXCL5 (3.34 fold, Figure 7m), IL-16 (53.68 fold, Figure 7e), RANTES (4.38 fold, Figure 7l), SDF-1a (1.55 fold, Figure 7i) were highest in the bone marrow compartment, when compared to the BAL and vascular compartments (Figure S3, https://figshare.com/s/83fe2fdabb32e0d954a). Remarkably, the concentrations of pan leukocyte chemokine, SDF-1a (Figure 7i), mononuclear cell chemokines, CXCL10 (Figure S4a, https://figshare.com/s/83fe2fdabb32e0d954a), CXCL13 (Figure S4c, https://figshare.com/s/83fe2fdabb32e0d954a), the neutrophil chemokine CXCL5 (Figure 7m) and the eosinophil chemokines eotaxin-1 (Figure 7j) and eotaxin-2 (Figure 7k), were highest in lung vascular perfusate after 0.005 ppm O3 exposures for 24 hr.

Thus, even 0.005 ppm of O3 exposure, for 24 hr, induce the release of BAL IL-1 dependent, chemokines, bone marrow IL-16 and RANTES and lung vascular SDF-1a, eotaxin-1, -2, and CXCL5. Short 2 hr exposures to 0.05 ppm O3 induced release of BAL IL-16 and eotaxin-2, thus highlighting neutrophil, eosinophil, and platelet cell dysregulation.

3.7 | 0.05 ppm O3 induces a heterogeneous loss of sub-pleural alveolar cytokeletal actin and dysregulated ventilation

Intravital lung microscopy from experiment 2 revealed pulmonary capillary vasodilatation immediately after 0.05 ppm O3 exposure (median diameter of 10.99 μm, Figure 8b,c,f) compared to the control group (median diameter of 6.10 μm, p < .01, Figure 8a). We superfused phalloidin to observe the alveolar actin arrangement at 0, 2, and 24 hr after 0.05 ppm O3 exposure. At 0 hr, we observed peripherally organized contractile actin filaments around alveolar spaces (Figure 8d and Figure S5a, https://figshare.com/s/83fe2fdabb32e0d954a). Moreover we observed the collective rhythmic motion of these filaments during respiration (Video 1 https://figshare.com/s/0b26c16f50f8dafef4c1, Figure 8h). O3 exposure led to heterogeneous alveolar damage (Figure 8e, Figure S5b-d, https://figshare.com/s/83fe2fdabb32e0d954a) marked by an increase in alveolar or air-space perimeter (Figure 8g) and reduction in circularity (Figure 8h). The sub-pleural alveoli showed either diffuse, discontinuous actin with lots of pores (Figure 8j, Figure S5b-d, https://figshare.com/s/83fe2fdabb32e0d954a), or extremely thin and condensed stress fibers (Figure 8e, Videos 2 https://figshare.com/s/0b26c16f50f8dafef4c1 and 3 https://figshare.com/s/0b26c16f50f8dafef4c1). Video-microscopy revealed that O3 prolongs inspiratory alveolar distention (Videos 2 https://figshare.com/s/0b26c16f50f8dafef4c1 and 3 https://figshare.com/s/0b26c16f50f8dafef4c1, Figure 8h).

3.8 | 0.05 ppm O3 induces immediate loss of sub-pleural ROS and mitochondrial potential followed by resurgence at 24 hr

In situ labeling of lungs for ROS, which is a marker of total cellular ROS generation and oxidative stress, showed several important features. Under baseline or RA exposure, specific lung parenchymal cells stain positive for ROS as indicated by significantly high activity in circular cells (Figure 9a). However, it is worth noting that these ROS-positive cells comprise 10%–20% of the parenchymal area (Figure 9b). Although there is not a significant difference in ROS-positive area before or after O3 exposure, the activity is representative of the respiratory cycles at 0 and 24h. ROS activity was also higher at 0 hr when measured as difference in ROS (green channel) signal over time (10 min, T1-T2) (Figure S6a, https://figshare.com/s/83fe2fdabb32e0d954a, b,c) and maximum intensity in the green channel (Figure 9a, Figure S7d) compared to at 2 hr O3 exposure. 35% of the lung parenchyma was ROS positive (Figure 9b). At 24 hr after 2 hr 0.05 ppm O3 exposure, lungs regain their ability to generate ROS (Figure 9a and Figure S6a-c, https://figshare.com/s/83fe2fdabb32e0d954a).

A similar pattern was observed with in situ reduced mitotracker dye staining (red channel) of lung alveolar cells. However, 20%–60% of the parenchymal cells are positive for reduced mitotracker dye at baseline (Figure 9c,d Video 4 https://figshare.com/s/0b26c16f50f8dafef4c1) and show a complete block of activity (Figure 9c, Figure S7a, https://figshare.com/s/83fe2fdabb32e0d954a) and maximum mitotracker dye staining (Figure 9c, Video 5 https://figshare.com/s/0b26c16f50f8dafef4c1, Figure S7a–d, https://figshare.com/s/83fe2fdabb32e0d954a) after 2 hr O3 exposure to 0.05 ppm O3. At 24 hr after 2 hr 0.05 ppm O3 exposure,
**FIGURE 8** Ozone (O₃)-induced alterations in in situ alveolar actin morphology. Representative stabilized lung thoracic window RGB images of sub-pleural C57BL/6NJ mice bright-field (without any dye or antibody treatment) exposed to (a) room air (i.e., 0 hr or control) or (b) 2 hr 0.05 ppm O₃. (c) Note the magnified view of the rectangular field from (b) showing marked vasodilatation (40.05 µm capillary diameter) in response to 0.05 ppm O₃. Representative 40× views of rhodamine phalloidin stained red fluorescent emission filtered light alveolar images visualized through ICCD camera (Stanford Photonics XR/MEGA 10 Extreme) in C57BL/6NJ mice exposed to (d) 0 hr (i.e., room air), showing well-organized septal alveolar actin filaments or (e) 2 hr 0.05 ppm O₃, showing heterogeneously damaged alveolar regions stretched out as a membrane. The diffuse contractile actin is shown by red double edged arrows which denote the diffuse actin morphology and numerous pores of Kohn or open circular air spaces are marked by yellow asterisks (*). (f) Distribution of pulmonary capillary diameter (µm) before (0 hr, shown as blue scatter plots) and after (shown as red scatter plots) 0.05 ppm ozone (2 hr) exposure, ** represents p < .01. (g) Distribution of alveolar perimeter (µm) before (0 hr, shown as blue scatter plots) and after (shown as red scatter plots) 0.05 ppm ozone (2 hr) exposure, (h) Alveolar space circularity before (0 hr, shown as blue scatter plots) and after (shown as red scatter plots) 0.05 ppm ozone (2 hr) exposure *p < .05, ** p < .01. N = 3 per group. (i) Representative line plots showing changes in alveolar area (µm²) of a single alveolus over 20 respiratory cycles, before (0 hr, shown in blue for original data points and green for smoothened data points) and after (shown in red for original data points and purple for smoothened data points) 0.05 ppm ozone (2 hr) exposure. ** represents p < .01. (j) Representative line plots showing percentage of area occupied by air-spaces in one sub-pleural field of view (FOV) over 20 respiratory cycles, before (0 hr, shown in blue) and after (shown in red) 0.05 ppm ozone (2 hr) exposure. ** represents p < .01
20%–30% of the parenchymal cells were distinctly positive for reduced mitotracker; this points to regeneration of redox potential (Figure 9c,d, Figure S7a–d Video 6 https://figshare.com/s/0b26c16f50f8daf1f4c1). We confirmed our live actin, ROS and mitotracker findings by staining cytospins of the BAL (Figure S8a-c, https://figshare.com/s/83fe2fdabbcc32ed954a), lung vascular perfusate as well as peripheral blood cells (data not shown).

Our intravital observations show loss of inter-septal connective tissue, compromised respiratory motion, immediately after 0.05 ppm of 2 hr O₃ exposure. This is also accompanied by a block of baseline cellular levels of ROS and mitochondrial potential. However, room air exposure of up to 22 hr after O₃ exposure regains the levels of ROS and mitochondrial potential in the alveolar sepal region.

4 | DISCUSSION

Our current study shows that low-dose O₃ exposure (i.e., below 0.063 ppm) is capable of inducing acute lung injury and systemic inflammation in C57BL/6NJ mice. Our model presents sensitive read-outs to assess sterile lung inflammation as summarized in a schematic format in Figure 10.

Systematic analysis of the BAL, lung vascular perfusate, and peripheral blood compartments revealed compartment-specific effects of 24 hr O₃ exposure. The short exposures led to decrease in BAL mononuclear cells, majority of which were degenerated and an increase in neutrophils in lung vascular perfusate and peripheral blood. Thus, neutrophils are recruited into the lungs but they die in the alveolar space (as indicated by their absence in recovered BAL cells).

BAL total protein content is an index of protein (airway surface liquid) secreted by the epithelial cells as well as lung edema as a result of vascular barrier disruption during inflammation. The reduction in BAL total protein in groups IV and V when compared to control group I is likely due to extensive epithelial cell death and thus loss of resulting protein secretion by airway surface liquid as noted in earlier studies (National Toxicology Program, 1994). The presence of debris in the BAL, and platelets in lung vascular perfusate and peripheral blood also translated into a higher total protein content in both the lung vascular perfusate and peripheral blood after prolonged sub-lethal (0.005 ppm) 24 hr O₃ exposure. Because platelets are a high source of proteins in blood (Crosley et al., 2009), our results indicate progressively increasing vascular damage and platelet dysregulation by O₃.
by cell debris in lung vascular perfusate), which translated into attenuated total protein content in peripheral blood. As well, total proteins in BAL and peripheral blood increased after exposure for 2 hr. The variable CD62P and vwf expressions in lung also indicate O₃-induced endothelial cell and platelet dysregulation. Overall, we observed acute alveolar cell death, which was supported by systematic findings in the BAL, lung vascular perfusate, peripheral blood as well as lung histology. Short (2 hr) O₃ exposures induced significant bronchiolar and endothelial cell damage, which progressed to development of frank hemorrhage and fibrin clumps at longer (4 and 24 hr) O₃ exposures, indicating endothelial dysfunction and platelet activation.

Other than acute cell death, discrete CD11b bright and large macrophages were prevalent in the BAL but absent in the lung vascular perfusate or peripheral blood (data not shown) following 0.05 ppm O₃ exposure for 2 hr. Moreover the fact that bronchiolar and vascular lung regions had more Cd11b positive cells suggests that these cells are highly adherent, after long (4 hr) O₃ exposures. As monocyte chemotactants are not appreciably increased in the BAL or vascular compartment, we suspect these bright and large CD11b-positive mononuclear cells to be derived from the visceral pleural lining macrophages (Cailhier et al., 2006). Pleural cavity cells such as macrophages and mesothelial cells are reported to follow haptokinetic migration to reach the lung parenchyma (Mubarak et al., 2012) and lymph nodes (Lehnert, 1992). Systematic studies in the future will be crucial to identify the important cell types or molecular signaling involved in the initiation of the repair processes that involve CD11b-positive (pleural cavity, alveolar, or monocyte-derived) macrophages.

Alveolar partitioning of neutrophils was a common feature after short and long O₃ exposures (Figure 10b). O₃, being a toxic chemical, instantly kills cells creating a zone of necrosis. Rapid neutrophil swarming outside the necrosis zone is typical of sterile inflammation models in tissues (McDonald et al., 2010; Wang et al., 2017). Neutrophils do eventually migrate into the necrotic zone and into the alveolar spaces but initially accumulate in the interstitial space around the alveoli. Most notably, the lung vascular perfusate comprised > 90% neutrophils after only 2 hr of 0.005 or 0.05 ppm O₃ exposure. The fact that O₃ did not increase the number and proportion of Gr1 positive cells in BAL, but these cells had lower fluorescence intensity deviation and a higher frequency of bright Gr1 distribution compared to the control group I, points toward
phagocytosis of the dead Gr1 positive cells (neutrophils, eosinophils, monocytes, iNKT cells and macrophages). High BAL IL-16 levels after 2 as well as 24 hr O3 exposures provide further evidence of secondary neutrophil necrosis. Activated macrophages, neutrophils as well as platelets release enzymes such as MMPs (matrix metallocproteinases) which degrade tissue extracellular matrix in the necrotic zone, cleave plasmogen to angiostatin fragments and clear paths for new vessels and collagen deposition (Benelli, Morini, Brigati, Noonan, & Albini, 2003; Benelli et al., 2002). In our O3 model, the high expression pattern of angiostatin near the bronchiolar regions, was juxtaposed to its precursor’s (plasminogen) or its binding partner’s (ATPβ) vascular and alveolar expression (Aulakh, 2018; Aulakh, Balachandran, Liu, & Singh, 2013). This shows that plasminogen is preferentially cleaved to angiostatin at sites of damage, like the bronchi.

Analysis of the 33-plex chemokine panel run on BAL fluid, lung vascular perfusate and bone marrow supernatants also revealed that O3 induces release of both immune stimulatory (IL-1β, IL-2, TNFα, IL-6, IFNγ) as well as suppressive (IL-10) chemokines. High alveolar concentrations of CXCL10, CX3CL1, and IL-16 (Center, Kornfeld, & Cruikshank, 1997; Ichikawa et al., 2013; Roth et al., 2015; Roth, Solbach, & Laskay, 2016), correlates with the secondary necrosis and phlogistic debris observed in the BAL and lung vascular perfusate cytospins from O3 exposed groups. The release of TARC and CXCL16 points toward probable activation of iNKT cells, which are capable of sensing damaged membrane phospholipids as well as releasing neutrophil chemoattractants like KC (Thanabalasuriar, Neupane, Wang, Krummel, & Kubes, 2016). A predominant release of the chemokines neutrophil as well as eosinophil chemokines such as, CXCL10, IL-16, CXCL5, RANTES, and SDF-1a (Garcia-Cuesta et al., 2019; Ichikawa et al., 2013; Kawaguchi, Zhang, & Nakanishi, 2019; Pan, Parkyn, Ray, & Ray, 2000) from bone marrow points to robust inflammatory response to 0.005 ppm O3 concentrations. Notably, the eosinophil chemoattracting agent, eotaxin-2, is released in BAL after 2 hr 0.05 ppm O3 exposure, and even higher levels are observed in vascular perfusate after 24 hr 0.005 ppm O3 exposure, indicating the recruitment of eosinophils. We also observed eosinophils in the H&E stained cryosections of spleen and thymus from O3 exposed mice (data not shown).

Imaging the dynamics of lung damage is required to understand the orchestration of the lung inflammatory response to O3. Inflammation is a sum of the vascular response and remodeling of the surrounding tissues as observed by altered diffuse alveolar actin morphology showing numerous pores in the alveoli, which are probably the pores of Kohn (Scarpelli, 2003). In addition, the lung video sequences showed perturbed respiratory motion as a direct evidence of O3-induced mechanical lung damage. Similar actin cytoskeletal changes have been reported secondary to cigarette smoke-induced damage to the alveolar epithelial cell monolayer (Nishida et al., 2017). Our observations of lung ROS and reduced mitotracker in live animals are also in agreement with the ex vivo ROS, mitotracker and actin staining of BAL, perfusate and peripheral blood cells. Whereas immediately after O3 exposure, the pulmonary epithelium is not mitochondrially active and incapable of producing ROS, O3 damaged tissue adapts by regaining mitochondrial sensitivity and ROS as seen in mice exposed to 0.05 ppm O3 for 2 hr and recovered for 22 hr under room air. Although live imaging showed that immediately after O3 exposure, when lung parenchymal cells are mitochondrially inactive and do not produce ROS, the lung tissue showed enhanced immune-fluorescent vascular lung ATP synthase β subunit (ATP β) expression and neutrophils concentrated within the lung vasculature, and not alveolar air sacs (as indicated by cell counts). At 24 hr after O3 exposure, when the lung tissue regains its ROS and mitochondrial potential, the lung vascular neutrophils plateau out and alveolar macrophage recruitment takes over.

There are numerous reports on the effects of much higher (1–2 ppm) concentrations of O3 exposure at 24 hr but these studies might be overexposing the mice and not capturing the primal inflammatory events, after O3 exposures (Francis et al., 2017; Tighe et al., 1950). Our results present a controlled model to study the acute cellular and tissue response to increasing concentration and time of O3 exposures. Radiation induced lung injury is also characterized by reduction in ROS, mitochondrial shrinkage due to excessive lipid peroxidation and ferroptotic cell death (Li, Zhuang, & Qiao, 2019). Our experiments have captured an important, but often ignored, role of ROS and mitochondrial potential in normal cellular signaling and tissue hemodynamics. One limitation of our study is that we did not examine sex-specific effects of O3. Although not ideal, the male sex was chosen to reduce variation in experimental data due to fewer animals per group. Therefore, future work needs to examine if the female mice are also as sensitive to deleterious effects of O3 as males were in our work.

The current experiments were performed on the NJ mouse sub-strain because many commercially available knock-out mice are on a mixed background with varying genetic contributions from both C57BL/6J and C57BL/6N sub-strains. We were unable to image the lung vasculature, immediately after O3 exposure, as the NJ sub-strain mice used in our current study, possessed frail hemodynamics. Currently, we are imaging the J sub-strain mice.

O3 injury of lung epithelial cells, and the ensuing inflammatory response, which leads to the production of numerous cytokines and chemokines, recruitment of neutrophils and activation of platelets likely contributes to the capacity of O3 to trigger asthma, chronic obstructive pulmonary disease (COPD) (Broeckaert et al., 1999; Broeckaert, Clippie, Wattiez, Falmagne, & Bernard, 2003). It is a common component of
urban smog and levels of O₃ in smog can easily reach up to 0.05 ppm; hence the reason why we chose this concentration range in our experiments. Coping with these effects on a daily basis can pose significant burden to additional particulate and infectious exposures like LPS, bacteria, and viruses which can lead to activation of the baseline immune response. Further understanding of how repair mechanisms take effect on a chronic basis can lead to better public health policies and therapeutic management of O₃-induced adverse health effects.

In summary, this research describes a sensitive model of sterile murine lung inflammation induced by exposing mice to short and low levels of O₃ that are below the current recommendations for safe ambient ozone concentration for humans. O₃ led to concentration and time-dependent phlogistic cell death in the broncho-alveolar lavage, lung epithelial damage and hemorrhage. We recorded CD11b positive cells in the broncho-alveolar lavage, upregulation of plasminogen and ATP synthase expression in the lung vascular and alveolar compartments while angiotatin was observed in the bronchial region in mice after 2 hr of O₃ exposure at 0.05 ppb. We also observed platelet and neutrophil accumulation in the lung vasculature and an eotaxin-2, IL-16, CXCL5, CXCL12, and CXCL13 dominant inflammatory response leading to lung injury. O₃ induced actin filament disorganization, perturbed respiratory mechanics, acute suppression of the alveolar reactive oxygen species (ROS) production and mitochondrial potential in ventilated lungs. We present evidence of systemic, as well as pulmonary toxicity, at 40-fold lower O₃ concentrations than previously reported in mice. Our findings may have important health implications of ozone exposure even at levels previously thought safe for humans.

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CONFLICT OF INTEREST
We have no conflict of interest to declare.

AUTHOR CONTRIBUTIONS
GKA, ES, and JS conceived the study, designed the experiments and wrote the manuscript. GKA, JABD, and CMGS executed the experiments. GKA and JS analyzed the data. All authors have edited and approved the manuscript.

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