

We thank the reviewers for their time and comments. Below are detailed responses to each comment. The responses are italicized and the modified texts are in red.

Response to anonymous referee #1 comments:

- 1) Page 4 Line 73: The authors state that there are many gaps. What are the gaps? What is the specific gap this work is attempting to address?**

Thank you for your comment. There are currently too many gaps to include a comprehensive list. The current work focuses on addressing the relative toxicities of different SOA systems, which was mentioned in lines 97 – 104. We have added an additional sentence in this section to clearly state the gap the current work is attempting to address.

Line 75: “Despite these findings, there are still many gaps in knowledge regarding PM-induced health effects. **The current work will focus on the relative toxicities of different SOA systems, as** field studies have repeatedly shown that SOA often dominate over primary aerosols **(e.g., PM emitted directly from combustion engines)** even in urban environments...”

- 2) Page 4 Line 76-81: the authors state that health studies focus on primary emissions rather than SOA, but then cited more SOA studies than primary studies. Seems contradictory. In fact, there is now a lot of attention on SOA. I suggest rephrasing.**

We have rephrased this section accordingly.

Line 86: “**Furthermore, in recent years, there have been an increasing number of studies on the health effects of SOA** formed from the oxidation of emitted hydrocarbons, **demonstrating their potential contribution to PM-induced health effects...**”

- 3) Page 5 line 95: Why were IL-6 and TNF-alpha chosen as the biomarkers? There are many other markers (such as HO-1, IL-17). Are these biomarkers better indicators of oxidative stress and better linked to health endpoints than others? Given that there is a nuanced response shown in Fig. 4, perhaps the choice of IL-6 and TNF-alpha was deliberate, but as a reader I am not sure why.**

We chose to measure IL-6 and TNF- α due to their central roles in cellular responses to stimuli and high production in MH-S cells. We have included a brief justification for choosing these specific biomarkers.

Line 106: “...cytokines indicative of the inflammatory response. **TNF- α is a hallmark biomarker involved in triggering a number of cellular signaling cascades. More specifically, TNF- α is involved in the activation of NF κ B, which regulates the expression of a variety of genes involved in inflammation and cell death, and the activation of protein kinases, which**

regulate various signaling cascades (Witkamp and Monshouwer, 2000). IL-6 has both pro- and anti-inflammatory effects, and may directly inhibit TNF- α (Kamimura et al., 2004). Furthermore, both cytokines are produced at relatively high levels in MH-S cells, ensuring a high signal-to-noise ratio and thus reliable measurements (Matsunaga et al., 2001; Chen et al., 2007).”

- 4) **Page 8 Line 149: 45% relative humidity is still quite dry. I would not label it as “humid”.**

We prefer to label these experiments as “humid”, as they can be considered relatively humid compared to our “dry” experiments (45% RH vs. 5% RH).

- 5) **Page 8 lines 154-158: does an acidic seed affect the background ROS production? Or is there sufficient buffer that cells are exposed to the same pH?**

It is unlikely for the acidic seed to affect background ROS/RNS production because the mass of seed per volume of media is low. Additionally, no changes in media color were observed during the extraction process. Since the cell culture media (RPMI-1640) contains phenol red, which is an indicator of pH, any significant changes in pH would result in an observable change in color. RPMI-1640 also uses a sodium bicarbonate buffer system to maintain physiological pH, so cells should be exposed to the same pH for all samples.

- 6) **Page 8 line 161: What is zero air? Is this purified air? How is the air purified?**

We have modified all instances of “zero air” to “pure air” and included how the air is purified at the first mention of pure air.

Line 171: “Chambers were flushed with pure air (generated from AADCO, 747-14) for ~24 hrs...”

Line 183: “...passing pure air over the solution until it fully evaporated.”

Line 186: “Naphthalene was injected by passing pure air over solid naphthalene flakes...”

- 7) **Page 8 line 169: presumably this concentration of OH is yielded only upon irradiation for the specific set of chamber lights.**

Yes, this is the OH concentration yielded upon irradiation with the specific set of chamber lights ($j\text{NO}_2 = 0.28 \text{ min}^{-1}$) (Boyd et al., 2015). This value is comparable to typical values for OH concentrations obtained in previous chamber studies (e.g., Eddingsaas et al., 2012; Loza et al., 2014; Ng et al., 2007; Chan et al., 2009; Chan et al., 2011).

- 8) **Page 10 line 212: why is 24 hrs chosen? What happens if cytokine levels were measured earlier or later? Are there recovery effects of exposure?**

We chose to measure both cytokines at 24 hrs to enable comparison at the same time point as ROS/RNS measurements (optimized in Tuet et al. (2016)) and because the production levels of both cytokines are relatively high at this time point for MH-S cells. Previous studies have shown that TNF- α and IL-6 production peak around 4 and 24 hrs, respectively (Haddad, 2001). Measuring at an earlier or later time point results in a decreased response, which may indicate recovery effects. We have modified the manuscript to clarify.

Line 236: “following manufacturer’s specifications (ThermoFisher). **This time point was chosen to enable comparison with ROS/RNS levels (also measured at 24 hrs, optimized in Tuet et al. (2016)) and to ensure a high signal for both cytokines. Previous literature have shown that TNF- α and IL-6 production peak around 4 and 24 hrs, respectively (Haddad, 2001). However, while TNF- α production peaks earlier, the signal at 24 hrs is well above the detection limit of the assay, and previous studies have utilized this time point to measure both cytokines (Haddad, 2001; Matsunaga et al., 2001). Nonetheless, it should be noted that these measurements represent a single time point in the cellular response....”**

- 9) **Page 12 line 247: H₂O₂ is unlikely to be taken up by inorganic seeds particles on a Teflon filter (as shown by the authors’ results), but may be taken up if there are organics coated on the filter. Is it possible there is further heterogeneous reactions of H₂O₂ on the organics, given the H₂O₂ concentrations are 3ppm?**

Since H₂O₂ uptake by inorganic seed particles was not observed (as shown by the blank results), it is unlikely that more H₂O₂ was taken up by SOA given the hygroscopicity parameter values ($\kappa = 0.53$ for ammonium sulfate vs. $\kappa = 0.006 - 0.2$ for organic compounds) (Petters and Kreidenweis, 2007).

- 10) **Page 12-13 lines 259-268: This is a central finding of this manuscript: the carbon backbone seems to play a bigger role than formation conditions. While I do not dispute the results, this finding is hard to rationalize. Formation conditions will affect mostly the functional groups that go onto the molecule (there may be small changes in the backbone with fragmentation pathways), while precursor identity will determine the size and shape of the backbone. ROS is likely produced through electron transfer to/from the functional groups interacting (or reacting) with O₂, H₂O, antioxidants and NADPH. It is therefore difficult to imagine that the functional group matters less than the backbone structure. Also, by that logic, reactions that change the molecular structure (such as oligomerization, fragmentation) would change the cellular ROS quite significantly. Is there any evidence of that?**

Thank you for your comment. We believe there was some confusion in this section. When we discussed the “carbon backbone”, we intended for “carbon backbone” to include both the

carbon chain length and functionalities. Furthermore, we refer to the “carbon backbone” of oxidation products, rather than the precursor compound. We have modified the manuscript to clarify these points. We also note that the referenced section (lines 259 – 268) refers to findings from a previous study, where the chemical oxidative potentials of these SOA systems were measured (Tuet et al., 2017). In that study, precursor identity was found to influence oxidative potential more significantly than formation condition. We bring this up here to highlight potential differences between chemical and cellular assays. In the current study, both precursor identity and formation condition influenced the level of cellular response, and products with similar functionalities and carbon chain length may induce similar responses. Oligomerization and fragmentation reactions influence the O:C ratio (and hence \overline{OS}_c), of SOA. We did observe a correlation between \overline{OS}_c and ROS/RNS production, shown in Fig. 3.

Line 27: “...which suggests that the **chemical structure (carbon chain length and functionalities) of photooxidation products** may be important...”

Line 295: “DTT may only be sensitive to larger differences arising from different precursors, whereas cellular assays...”

Line 305: “...for SOA precursors **whose products share similar chemical structures (i.e., similar carbon chain length and functionalities)**...”

Line 529: “...SOA systems whose products share similar functionalities and carbon chain length are likely to induce...”

- 11) Page 16 line 326: this is an interesting explanation. If fatty acids are really changing cell functions that significantly, meat cooking organic aerosols, which are composed almost entirely of fatty acids, would elicit very strong responses.**

Thank you for the suggestion. We have added this as a potential implication.

Line 380: “...lesser response compared to pentadecane SOA exposure. **These observations, particularly those for pentadecane SOA, suggest that aerosols from meat cooking may have health implications, as fatty acids comprise a majority of these aerosols (Mohr et al., 2009; Rogge et al., 1991).**”

- 12) Page 16 Line 343: Naphthalene is not “completely” different. For example, IL-6 and TNF-alpha are still somewhat positively correlated at low levels. Perhaps it is just a more distinct pattern.**

Thank you for the suggestion. We have modified the manuscript accordingly.

Line 384: “Naphthalene exhibits a **different, more distinct** pattern **compared to** the rest of the SOA systems...”

- 13) Page 18 Line 395-396 and Fig. 3b: what does significant correlation mean? There is an asterisk in Fig. 3b. Does that mean the trend is statistically significant? If so, please provide statistical justification (e.g. 95% confidence interval?). Does it have to be a linear model? Does the correlation still stand if naphthalene SOA (which is the outlier) points are removed? It would seem reasonable to me to remove the naphthalene system if there is reason to believe it has a very different toxicological mechanism.**

The method for determining statistical significance was described in the methods section. We have modified the manuscript and figure caption accordingly. The correlation does not hold if naphthalene SOA is removed. However, since other SOA systems (i.e., pentadecane and β -caryophyllene) may also participate in toxicological pathways unique to those SOA systems, we did not exclude naphthalene from the correlation. Furthermore, it is interesting that there exists a correlation between oxidation state and ROS/RNS even though different toxicological mechanisms may be involved.

Line 436: “Nevertheless, a significant correlation ($p < 0.05$) was observed...”

Line 592: “...colored by SOA system. * indicates significance, $p < 0.05$.”

- 14) Page 19 lines 404-425: What is the relationship between ROS/RNS and cytokines for these SOA systems? It seems that plotting them against each other would help explain trends in each SOA system, or at least establish whether or not ROS/RNS are linked to upregulation of these cytokines.**

We show the relationship between ROS/RNS and cytokines in Fig. 4, where the ROS/RNS level is influenced by a balance between both cytokines due to pro- and anti-inflammatory effects. We did plot ROS/RNS against cytokine measurements, however, individual correlation plots did not reveal any additional information as the inflammatory markers are involved in pathways with many overlaps and crosstalk. These relationships were only apparent when all three measurements were plotted, as shown in Fig. 4.

- 15) Page 3 Line 52: “anti-oxidant” should be “antioxidant”**

We have modified the manuscript accordingly.

Line 52: “...redox reactions using an **antioxidant** species...”

Line 53: “The **antioxidant** is oxidized...”

Line 401: “...products that promote electron transfer reactions with **antioxidants**...”

16) Page 7 line 127: “form” should be “from”

We have modified the manuscript accordingly.

Line 149: “SOA formed **from** the photooxidation...”

17) Page 21 line 464: “RNS/RNS” should be “ROS/RNS”

We have modified the manuscript accordingly.

Line 505: “...produce low levels of **ROS/RNS**...”

References:

Boyd, C. M., Sanchez, J., Xu, L., Eugene, A. J., Nah, T., Tuet, W. Y., Guzman, M. I., and Ng, N. L.: Secondary organic aerosol formation from the β -pinene+NO₃ system: effect of humidity and peroxy radical fate, *Atmos. Chem. Phys.*, 15, 7497-7522, 10.5194/acp-15-7497-2015, 2015.

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Chan, M. N., Surratt, J. D., Chan, A. W. H., Schilling, K., Offenberg, J. H., Lewandowski, M., Edney, E. O., Kleindienst, T. E., Jaoui, M., Edgerton, E. S., Tanner, R. L., Shaw, S. L., Zheng, M., Knipping, E. M., and Seinfeld, J. H.: Influence of aerosol acidity on the chemical composition of secondary organic aerosol from β -caryophyllene, *Atmos. Chem. Phys.*, 11, 1735-1751, 10.5194/acp-11-1735-2011, 2011.

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Kamimura, D., Ishihara, K., and Hirano, T.: IL-6 signal transduction and its physiological roles: the signal orchestration model, in: *Reviews of Physiology, Biochemistry and Pharmacology*, Springer Berlin Heidelberg, Berlin, Heidelberg, 1-38, 2004.

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Witkamp, R., and Monshouwer, M.: Signal transduction in inflammatory processes, current and future therapeutic targets: A mini review, *Veterinary Quarterly*, 22, 11-16, 10.1080/01652176.2000.9695016, 2000.

We thank the reviewers for their time and comments. Below are detailed responses to each comment. The responses are italicized and the modified texts are in red. The main comments have been addressed by including a discussion on the limitations of this study and by clarifying our statistical analysis method. The revisions do not affect the conclusions of the manuscript.

Response to anonymous referee #2 comments:

- 1) Limitations of this study: I didn't see any discussion regarding the limitations of this study, and they mainly cited their own DTT papers throughout the discussion. This would be my most major criticism. You have to be careful to say that your acute exposures here will really translate to the in vivo condition. Specifically, why does one need to be careful in extending the results obtained from in vitro exposures to the in vivo condition? What are the potential issues with extracting filters for resuspension into cell culture? Does the chemistry change, and if so, how might that affect the toxicological response?**

Thank you for your suggestion. We are aware that there are limitations regarding all health studies and have modified the manuscript to include several examples of these limitations. We note that the main objective of this study was to provide perspective on the relative toxicities of different SOA systems. Further studies are required to establish whether results from in vitro assays represent in vivo animal exposures, and from there, whether results from animal exposure studies can be generalized to actual human exposures.

Line 552: “**Additionally**, this study confirms...”

Line 562: “...to fully interpret ROS/RNS measurements. **Finally, several limitations must be considered before generalizing results from this study to in vivo exposures. For instance, only one cell type was explored in this study, whereas an organism consists of multiple tissues comprised of multiple cell types. Interactions between different cell types and tissue systems were not considered in this study. Furthermore, the doses investigated may not fully represent real world exposures due to differences in exposure routes and potential recovery from doses due to clearance. Nevertheless, this study provides perspective on the relative toxicities of different SOA systems which future studies can build upon.**”

- 2) Rationale for using murine alveolar macrophages: I think the authors should provide the rationale for using murine alveolar macrophages for this study. Would certain phenotype of this cell line differ from human alveolar macrophages? How easily relatable is it to human cells? What are limitations of cell lines versus primary cells and would that matter?**

Thank you for your comment. We have included rationale for using this cell type in the manuscript. We chose murine alveolar macrophages as they are the first line of defense against environmental insults, and the particular cell line (MH-S) retains many properties

of primary alveolar macrophages (e.g., phagocytosis, cytokine production, ROS/RNS production) (Sankaran and Herscowitz, 1995; Mbawuiké and Herscowitz, 1989). Furthermore, we have successfully utilized this cell line to investigate the production of ROS/RNS as a result of exposure to ambient PM samples (Tuet et al., 2016). To our knowledge, immortalized human alveolar macrophages do not exist. Mice have also been widely used as a model organism for studying human responses (Rosenthal and Brown, 2007; Takao and Miyakawa, 2015). As for the choice between cell lines and primary cells, primary cells are harvested from multiple animals, which increases the response variability. Results may therefore be less reproducible compared to cell lines.

Line 137: “Exposures were conducted using immortalized murine alveolar macrophages (MH-S, ATCC®CRL-2019™) as they are the first line of defense against environmental insults (Oberdörster, 1993; Oberdörster et al., 1992). The particular cell line also retains many properties of primary alveolar macrophages, including phagocytosis as well as the production of ROS/RNS and cytokines (Sankaran and Herscowitz, 1995; Mbawuiké and Herscowitz, 1989). MH-S cells were cultured...”

- 3) I noticed that the authors’ cell culture and exposure media contain fetal bovine serum (FBS), which is known to potentially interfere with the ELISA assays. Normally people use serum-free media to avoid such interferences. Do the authors have any control experiments to show that FBS wouldn’t interfere with their ELISA measurement?**

We normalized all ELISA responses to a control (cell culture supernatant from cells exposed to stimulant-free media supplemented with FBS) to capture any interferences. For our time point (24 hrs), FBS supplemented media is necessary to prevent serum starvation, which is known to induce oxidative stress (Kuznetsov et al., 2011; Wright et al., 2012). We also disagree that serum-free media is generally used for ELISA measurements, as many previous studies have performed exposures using supplemented media (e.g., Mukherjee et al., 2009; Chen et al., 2007; Sullivan et al., 2000).

- 4) They use the cell media to extract filters. Since cell media contain a lot of supplementary materials/nutrients, would this affect the fraction of SOA materials extracted? Also, for the reactive products, would they be hydrolyzed before cell exposure?**

For oxidative potential measurements, it is known that using different extraction methods (e.g., different solvent, filtration, removing the filter) results in different components extracted and hence yields different oxidative potential measurements (Gao et al., 2017). However, there are limitations for each method. For instance, using an organic solvent requires the subsequent removal of the solvent via evaporation, which may result in loss of unstable components (e.g., semi-volatile organics). In this study, we chose to adapt an extraction method best suited for cellular exposure. While media contains species that would indeed alter the fraction of material extracted, these species are also present in the

alveolar fluid and the extract obtained is biologically relevant. We would also like to note that plain media (without FBS) was used for extraction and that FBS was supplemented after filtration of extracts. We did not investigate the hydrolysis of reactive products due to extraction, however this would be a potential issue for all extraction methods used in offline analysis. Further studies comparing offline and online analysis are required to investigate this.

- 5) **Lines 265-266: I think the redox activity is likely more sensitive to the functionality/electronic configuration of the functional groups, instead of carbon backbone. If it is carbon backbone, it looks to me that DTT is removed by other mechanisms such as absorption, but not through redox mechanisms.**

The referenced section refers to a previous study, where the chemical oxidative potentials as determined by DTT consumption were measured for these SOA systems (Tuet et al., 2017). In this study, we focus on the cellular responses and we find that the precursor identity and formation condition are both important and affect the cellular responses significantly. We note that there may have been some confusion in this section, as we intended “carbon backbone” to include both carbon chain length and functionalities. We have modified the manuscript to clarify our findings.

Line 27: “...which suggests that the **chemical structure (carbon chain length and functionalities) of photooxidation products** may be important...”

Line 295: “DTT may only be sensitive to larger differences arising from different precursors, whereas cellular assays...”

Line 305: “...for SOA precursors **whose products share similar chemical structures (i.e., similar carbon chain length and functionalities)**...”

Line 529: “...SOA systems whose products share similar functionalities and carbon chain length are likely to induce...”

- 6) **How these inflammatory responses relate to each other? Are they involved in the same biological network? They probably need to provide a more detailed biological background for the biomarkers they measured. For example, TNF-alpha induces IL-8 via NF-κB. This is well known in the toxicological literature. In some of the toxicological literature, TNF-alpha is used as positive control to stimulate IL-8 in BEAS-2B cells. I don't see a clear connection between the endpoints they measured in this paper and this needs to be more justified. Without a connection to a specific biological system, it makes it hard (especially for an atmospheric chemist I'm sure) to understand what your results really mean.**

Thank you for your suggestion. We have included justification on our cytokine measurements.

Line 106: "...cytokines indicative of the inflammatory response. **TNF- α is a hallmark biomarker involved in triggering a number of cellular signaling cascades. More specifically, TNF- α is involved in the activation of NF κ B, which regulates the expression of a variety of genes involved in inflammation and cell death, and the activation of protein kinases, which regulate various signaling cascades (Witkamp and Monshouwer, 2000). IL-6 has both pro- and anti-inflammatory effects, and may directly inhibit TNF- α (Kamimura et al., 2004). Furthermore, both cytokines are produced at relatively high levels in MH-S cells, ensuring a high signal-to-noise ratio and thus reliable measurements (Matsunaga et al., 2001; Chen et al., 2007).**"

- 7) Lines 288-290: The authors cite Lin et al. (2016, ES&T Letters), but I think this discussion is really unclear. What genes are similar? What pathways do the authors mean? They should make them clear. Note that Lin et al. (2016, ES&T Letters) only measured oxidative stress-associated genes, but not inflammatory genes in that paper. I noted that Lin et al. (2017, ES&T) just had a newly accepted paper where they found most genes are associated with the Nrf2 pathway, but not much inflammatory response from isoprene SOA exposure under non-cytotoxic conditions. Also, in Lin et al. (2017, ES&T) time course experiments, they found that IL-8 expression is time sensitive. The expression maximized at 9 hr and much lowered at 24 hr, which was also shown in Arashiro et al. (2016, ACP). Their cellular materials were collected 24 hr post-exposure, so they might have missed the peak. How do the authors justify the 24 hr post exposure time? Did they conduct a series of time course experiments to see where things might peak in terms of cellular response? The authors and readers need to realize you may only captured 1 slice in time in how the cells responded.**

We have modified the manuscript to clarify this discussion. Specifically, we include an example of a gene whose fold change was similar between the two types of SOA studies in Lin et al. (2016) and discuss how that gene is related to the inflammatory cytokines measured in this study. Oxidative stress plays a crucial role in the inflammatory process, and as such, the oxidative stress related genes measured in Lin et al. (2016) may influence cytokine production. We thank the reviewer for pointing out Lin et al. (2017) and have cited the paper accordingly. We are aware that cytokine production peaks at different time points for different cytokines. In our case, TNF- α peaks around 4 hrs, while IL-6 peaks much later at 24 hrs (Haddad, 2001). We chose to measure both cytokines at the latter time point to allow comparison. Previous studies have shown that the level of TNF- α is sufficiently high at the latter time point for accurate determination (Haddad, 2001; Matsunaga et al., 2001). The manuscript has been modified to include this justification as well.

Line 93: "**However, the cellular exposure studies involving SOA focused on SOA formed from a single precursor and included** different measures of response (e.g. ROS/RNS,

inflammatory biomarkers, gene expression, etc.) (Arashiro et al., 2016; Lund et al., 2013; McDonald et al., 2010; McDonald et al., 2012; Baltensperger et al., 2008; Lin et al., 2017).”

Line 318: “...the fold change of several genes reported in Lin et al. (2016) are actually similar (e.g., *ALOX12*, *NQO1*). Several of these genes directly affect the production of inflammatory cytokines measured in this study. For instance, studies have observed that arachidonate 12-lipoxygenase (*ALOX12*) products induce the production of both TNF- α and IL-6 in macrophages (Wen et al., 2007). As such, a similar response level regardless of SOA formation condition may be observed depending on the biological endpoints measured. Thus, it is possible that the inflammatory cytokines measured in this study are involved in pathways concerning those genes, resulting in a similar response level regardless of SOA formation condition.”

Line 231: “following manufacturer’s specifications (ThermoFisher). This time point was chosen to enable comparison with ROS/RNS levels (also measured at 24 hrs, optimized in Tuet et al. (2016)) and to ensure a high signal for both cytokines. Previous literature have shown that TNF- α and IL-6 production peak around 4 and 24 hrs, respectively (Haddad, 2001). However, while TNF- α production peaks earlier, the signal at 24 hrs is well above the detection limit of the assay, and previous studies have utilized this time point to measure both cytokines (Haddad, 2001; Matsunaga et al., 2001). Nonetheless, it should be noted that these measurements represent a single time point in the cellular response...”

8) Line 305: what kind of chemical structure do they mean here?

Thank you for the comment. We have modified the manuscript to clarify.

Line 340: “These observations further imply that the chemical structures (e.g., carbon chain lengths and functionalities) of oxidation products...”

9) Line 322-329: I am not sure about the insertion of pentadecane oxidation products to the membrane. They should at least provide some references to support such a statement. I would expect some cellular response, specifically cytotoxicity, from these products since they are detergent like, which could potentially rupture the cell membrane. Did they see cell death from MTT data for pentadecane oxidation products?

Thank you for the suggestion. We have included references to support this hypothesis. We did not observe decreases in cellular metabolic activity as measured by the MTT assay (mentioned in lines 282 – 286 in the revised manuscript).

Line 363: “...could potentially insert into the cell membrane (Loza et al., 2014), as previous studies have shown that fatty acids can feasibly insert into the cell membrane bilayer (Khmelinskaia et al., 2014; Cerezo et al., 2011).”

- 10) The mechanism of PAH-DNA adduct formation is well known through metabolic activation to diol epoxides. This is not mentioned at all in current discussion.**

We mentioned the formation of DNA adducts briefly in the section on naphthalene SOA (lines 420 – 424). The specific mechanism by which these adducts are formed is beyond the scope of this study, but would be interesting to investigate in future studies.

- 11) Statistical Analysis: One more critical comment relates to the authors statistical analysis. Where are their linear regression results and the associated p values? Also, with multiple groups, one-way ANOVA should be used instead of student's t test to get p-values (same idea as the increasing type I error with multiple testing). Lastly, when they talked about the trend, I didn't see any statistical support to differentiate between groups. Are the results really statistically significant?**

Based on the reviewer's comment, we believe the trend referenced refers to Fig. 3. The Pearson's correlation coefficient is given in the original figure. For clarity, we have modified the manuscript and figure caption to reflect that correlations were evaluated using a 95% confidence interval. Since only two variables (cellular response and bulk aerosol composition, e.g., ROS/RNS and \overline{OS}_c) were tested, the student's t-test and one-way ANOVA are actually equivalent (Park, 2009).

Line 436: "Nevertheless, a significant correlation ($p < 0.05$) was observed..."

Line 592: "...colored by SOA system. * indicates significance, $p < 0.05$."

- 12) I'm curious why the authors didn't gravimetrically weigh the filters before and after sampling to insure actual mass on filter for dose-response purposes? If you use the SMPS, you must make assumptions about density to calculate the mass. How was density accurately determined if you did use that approach? Was the SMPS sheath flow conditioned to the appropriate RH used in the chamber?**

Mass loadings were low for isoprene and pentadecane SOA. To be consistent, we choose to determine mass by integrating the SMPS volume concentrations for all SOA systems. An aerosol density of 1 g cm^{-3} was assumed to facilitate comparison between studies, since SOA density varies with precursor identity and formation condition. We have added this clarification to the manuscript. For all experiments, the SMPS was connected to the chamber for 2-3 hrs before the start of the experiment to condition the recirculating sheath flow.

Line 201: "...multiplying by the total volume of air collected. SMPS volume concentrations were converted to mass concentrations by assuming a density of 1 g cm^{-3} to facilitate comparison between studies..."

- 13) Related to #12 above, were extraction efficiencies of aerosol mass from the filters determined by spiking them with representative internal standards? Extracting filters with cell media I may not actually remove a lot of materials (such as oligomers of SOA) from the filters. Why wasn't organic solvents used, then dried, and then the dried extracts reconstituted with cell media for the exposures? Toxicologists might find your dosing completely uncertain as its hard to gauge how well you removed the SOA from the filters without this information. This is a very important point for Figures like Figure 3. The AMS sees most of the SOA mass but filter extractions may not actually remove all of it for the exposure assessment done here.**

Extraction efficiencies were not measured in this study. While different extraction methods are known to result in different constituents being extracted from the PM sample, there are limitations for each method. These are discussed in a recent publication by Gao et al. (2017). For example, using an organic solvent and drying the extract for reconstitution may result in loss of unstable constituents. For this study, we chose an extraction method best suited for cellular exposure.

- 14) I have a curiosity question. Did the authors observe brown color on some of their filters (like from naphthalene SOA or isoprene SOA)? If so, did you seen any trends with brown carbon and your toxicological endpoints?**

We only observed brown color on our naphthalene SOA filters. We did not measure brown carbon in this study.

- 15) Line 81-84: This seems to be an incomplete sentence or poorly worded sentence. Please revise.**

Thank you for your comment. We have modified the sentence.

Line 91: “**However**, the cellular exposure studies **involving** SOA focused on SOA formed from a single precursor and **included** different measures of response...”

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1 **Inflammatory responses to secondary organic aerosols (SOA) generated from biogenic and**
2 **anthropogenic precursors**

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10 organic aerosol

11 Abstract

12 Cardiopulmonary health implications resulting from exposure to secondary organic
13 aerosols (SOA), which comprise a significant fraction of ambient particulate matter (PM), have
14 received increasing interest in recent years. In this study, alveolar macrophages were exposed to
15 SOA generated from the photooxidation of biogenic and anthropogenic precursors (isoprene, α -
16 pinene, β -caryophyllene, pentadecane, *m*-xylene, and naphthalene) under different formation
17 conditions ($\text{RO}_2 + \text{HO}_2$ vs. $\text{RO}_2 + \text{NO}$ dominant, dry vs. humid). Various cellular responses were
18 measured, including reactive oxygen/nitrogen species (ROS/RNS) production and secreted levels
19 of cytokines, tumor necrosis factor- α (TNF- α) and interleukin-6 (IL-6). SOA precursor identity
20 and formation condition affected all measured responses in a hydrocarbon-specific manner. With
21 the exception of naphthalene SOA, cellular responses followed a trend where TNF- α levels
22 reached a plateau with increasing IL-6 levels. ROS/RNS levels were consistent with relative levels
23 of TNF- α and IL-6, due to their respective inflammatory and anti-inflammatory effects. Exposure
24 to naphthalene SOA, whose aromatic ring-containing products may trigger different cellular
25 pathways, induced higher levels of TNF- α and ROS/RNS than suggested by the trend. Distinct
26 cellular response patterns were identified for hydrocarbons whose photooxidation products shared
27 similar chemical functionalities and structures, which suggests that the carbon backbone chemical
28 structure (carbon chain length and functionalities) of photooxidation products may be important
29 for determining cellular effects. A positive nonlinear correlation was also detected between
30 ROS/RNS levels and previously measured DTT activities for SOA samples. In the context of
31 ambient samples collected during summer and winter in the greater Atlanta area, all laboratory-
32 generated SOA produced similar or higher levels of ROS/RNS and DTT activities. These results

33 suggest that the health effects of SOA are important considerations for understanding the health
34 implications of ambient aerosols.

35 Introduction

36 Particulate matter (PM) exposure is a leading global risk factor for human health (Lim et
37 al., 2012) with numerous studies reporting associations between elevated PM concentrations and
38 increases in cardiopulmonary morbidity and mortality (Li et al., 2008; Pope III and Dockery, 2006;
39 Brunekreef and Holgate, 2002; Dockery et al., 1993; Hoek et al., 2013; Anderson et al., 2011;
40 Pope et al., 2002). A possible mechanism for PM-induced health effects has been suggested by
41 toxicology studies, wherein PM-induced oxidant production, including reactive oxygen and
42 nitrogen species (ROS/RNS), initiates inflammatory cascades thus resulting in oxidative stress and
43 cellular damage (Li et al., 2003a; Tao et al., 2003; Castro and Freeman, 2001; Gurgueira et al.,
44 2002; Wiseman and Halliwell, 1996; Hensley et al., 2000). Furthermore, prolonged stimulation of
45 these inflammatory cascades may lead to chronic inflammation, for which there is a recognized
46 link to cancer (Philip et al., 2004). Together, these findings suggest that a possible relationship
47 exists between PM exposure and observed health effects.

48 Various assays have been developed to study PM-induced oxidant production, including
49 cell-free chemical assays that measure the oxidative potential of PM samples (Kumagai et al.,
50 2002; Cho et al., 2005; Fang et al., 2015b) as well as cellular assays that measure intracellular
51 ROS/RNS produced as a result of PM exposure (Landreman et al., 2008; Tuet et al., 2016). Cell-
52 free assays simulate biologically relevant redox reactions using an ~~anti-oxidant~~antioxidant species
53 (e.g. dithiothreitol, DTT; ascorbic acid, AA). The ~~anti-oxidant~~antioxidant is oxidized via electron
54 transfer reactions catalyzed by redox-active species in the PM sample and its rate of decay serves

55 as a measure of the concentration of redox-active species present (Fang et al., 2015b). On the other
56 hand, cellular assays utilize a fluorescent probe (e.g. carboxy-H₂DCFDA) that reacts with
57 ROS/RNS and the measured fluorescence is proportional to the concentration of ROS/RNS
58 produced as a result of PM exposure (Landreman et al., 2008; Tuet et al., 2016). Both types of
59 assays have been utilized extensively to characterize a variety of PM samples and identify sources
60 that may be detrimental to health (Verma et al., 2015a; Saffari et al., 2015; Fang et al., 2015a;
61 Bates et al., 2015; Li et al., 2003b; Tuet et al., 2016). In particular, numerous studies suggest that
62 organic carbon constituents, especially humic-like substances (HULIS) and oxygenated
63 polyaromatic hydrocarbons (PAH), may contribute significantly to PM-induced oxidant
64 production (Li et al., 2003b; Kleinman et al., 2005; Hamad et al., 2015; Verma et al., 2015b; Lin
65 and Yu, 2011). Furthermore, recent measurements of ROS/RNS production and DTT activity
66 using ambient samples collected in summer and winter around the greater Atlanta area showed
67 that there is a significant correlation between summertime organic species and intracellular
68 ROS/RNS production, suggesting a possible role for secondary organic aerosols (SOA) (Tuet et
69 al., 2016). The same study also reported a significant correlation between ROS/RNS production
70 and DTT activity for summer samples, while a relatively flat ROS/RNS response was observed
71 for winter samples spanning a similar DTT activity range (Tuet et al., 2016). These results
72 highlight a need to consider multiple endpoints as a simple correlation may not exist between
73 different endpoints, especially cellular responses that may result from complicated response
74 networks.

75 Despite these findings, there are still many gaps in knowledge regarding PM-induced
76 health effects. ~~While~~The current work will focus on the relative toxicities of different SOA
77 systems, as field studies have repeatedly ~~showed~~shown that SOA often dominate over primary

78 aerosols (e.g., PM emitted directly from combustion engines) even in urban environments (Zhang
79 et al., 2007; Jimenez et al., 2009; Ng et al., 2010), ~~many prior health studies have focused on the~~
80 ~~effects of primary emissions (e.g. PM emitted directly from combustion engines) (Kumagai et al.,~~
81 ~~2002; Bai et al., 2001; McWhinney et al., 2013a; Turner et al., 2015) rather than those of SOA~~
82 ~~formed from the oxidation of emitted hydrocarbons (McWhinney et al., 2013b; Rattanavaraha et~~
83 ~~al., 2011; Kramer et al., 2016; Lund et al., 2013; McDonald et al., 2010; McDonald et al., 2012;~~
84 ~~Baltensperger et al., 2008; Arashiro et al., 2016; Platt et al., 2014). The cellular exposure studies~~
85 ~~that do explore SOA focused on SOA formed from a single SOA precursor and include~~
86 Furthermore, in recent years, there have been an increasing number of studies on the health effects
87 of SOA formed from the oxidation of emitted hydrocarbons, demonstrating their potential
88 contribution to PM-induced health effects (McWhinney et al., 2013b; Rattanavaraha et al., 2011;
89 Kramer et al., 2016; Lund et al., 2013; McDonald et al., 2010; McDonald et al., 2012;
90 Baltensperger et al., 2008; Arashiro et al., 2016; Platt et al., 2014; Gallimore et al., 2017).
91 However, the cellular exposure studies involving SOA focused on SOA formed from a single
92 precursor and included different measures of response (e.g. ROS/RNS, inflammatory biomarkers,
93 gene expression, etc.) (Arashiro et al., 2016; Lund et al., 2013; McDonald et al., 2010; McDonald
94 et al., 2012; Baltensperger et al., 2008; Lin et al., 2017). As a result, there is a lack of understanding
95 in terms of the relative toxicity of individual SOA systems. Recently, Tuet et al. (2017)
96 systematically investigated the DTT activities of SOA formed from different biogenic and
97 anthropogenic precursors and demonstrated that intrinsic DTT activities were highly dependent on
98 SOA precursor identity, with naphthalene SOA having the highest DTT activity. As a result, a
99 systematic study on the cellular responses induced by these SOA systems may provide similar

100 insights. Furthermore, cellular responses may complement these previously measured DTT
101 activities to elucidate a more complete picture of the health effects of PM.

102 In the present study, alveolar macrophages were exposed to SOA generated under different
103 formation conditions from various SOA precursors. Cellular responses induced by SOA exposure
104 were measured, including intracellular ROS/RNS production and levels of tumor necrosis factor-
105 α (TNF- α) and interleukin-6 (IL-6). Intracellular ROS/RNS production serves as a general
106 indicator of oxidative stress, whereas TNF- α and IL-6 are pro-inflammatory cytokines indicative
107 of the inflammatory response (Henkler et al., 2010; Kishimoto, 2003; Wang et al., 2003). TNF- α
108 is a hallmark biomarker involved in triggering a number of cellular signaling cascades. More
109 specifically, TNF- α is involved in the activation of NF κ B, which regulates the expression of a
110 variety of genes involved in inflammation and cell death, and the activation of protein kinases,
111 which regulate various signaling cascades (Witkamp and Monshouwer, 2000). IL-6 has both pro-
112 and anti-inflammatory effects, and may directly inhibit TNF- α (Kamimura et al., 2004).
113 Furthermore, both cytokines are produced at relatively high levels in MH-S cells, ensuring a high
114 signal-to-noise ratio and thus reliable measurements (Matsunaga et al., 2001; Chen et al., 2007).

115 Precursors were chosen to include major classes of biogenic and anthropogenic compounds known
116 to produce SOA upon atmospheric oxidation (Table S1). The selected biogenic precursors include:
117 isoprene, the most abundant non-methane hydrocarbon (Guenther et al., 2006); α -pinene, a well-
118 studied monoterpene with emissions on the order of global anthropogenic emissions (Guenther et
119 al., 1993; Piccot et al., 1992); and β -caryophyllene, a representative sesquiterpene. Both
120 monoterpenes and sesquiterpenes have been shown to contribute significantly to ambient aerosol
121 (Eddingsaas et al., 2012; Hoffmann et al., 1997; Tasoglou and Pandis, 2015; Goldstein and
122 Galbally, 2007). Similarly, the anthropogenic precursors include: pentadecane, a long-chain

123 alkane; *m*-xylene, a single-ring aromatic; and naphthalene, a poly-aromatic. These compounds are
124 emitted as products of incomplete combustion (Robinson et al., 2007; Jia and Batterman, 2010;
125 Bruns et al., 2016) and have considerable SOA yields (Chan et al., 2009; Ng et al., 2007b; Lambe
126 et al., 2011). In addition to precursor identity, the effects of humidity (dry vs. humid) and NO_x
127 levels (different predominant peroxy radical (RO₂) fates, RO₂ + HO₂ vs. RO₂ + NO) on SOA
128 cellular inflammatory responses were investigated, as different formation conditions have been
129 shown to affect aerosol chemical composition and mass loading, which could in turn result in a
130 different cellular response (Chhabra et al., 2010; Chhabra et al., 2011; Eddingsaas et al., 2012; Ng
131 et al., 2007b; Loza et al., 2014; Ng et al., 2007a; Chan et al., 2009; Boyd et al., 2015). Finally,
132 correlations between bulk aerosol composition, specifically elemental ratios, and cellular
133 inflammatory responses were investigated to determine whether there is a link between different
134 inflammatory responses and aerosol composition.

135 Methods

136 **Alveolar macrophage cell line.** ~~Immortalized murine alveolar macrophages (MH-S,~~
137 ~~ATCC®CRL-2019™)~~ Exposures were conducted using immortalized murine alveolar macrophages
138 (MH-S, ATCC®CRL-2019™) as they are the first line of defense against environmental insults
139 (Oberdörster, 1993; Oberdörster et al., 1992). The particular cell line also retains many properties
140 of primary alveolar macrophages, including phagocytosis as well as the production of ROS/RNS
141 and cytokines (Sankaran and Herscowitz, 1995; Mbawuiké and Herscowitz, 1989). MH-S cells
142 were cultured in RPMI-1640 media supplemented with 10% fetal bovine serum (FBS, Quality
143 Biological, InC.), 1% penicillin-streptomycin, and 50 μM β-mercaptoethanol (BME) at 37°C and
144 humid air containing 5% CO₂. For exposure experiments, MH-S cells were seeded at a density of
145 2 x 10⁴ cells well⁻¹ onto 96-well plates pre-treated with 10% FBS in phosphate buffered saline

146 (PBS, Cellgro). For seeding and all assay procedures thereon, FBS-supplemented cell culture
147 media without BME addition was used as BME is a reducing agent that may interfere with
148 inflammatory measurements.

149 **Chamber experiments.** SOA formed ~~form~~from the photooxidation of biogenic and
150 anthropogenic precursors were generated in the Georgia Tech Environmental Chamber (GTEC)
151 facility. Details of the facility have been described elsewhere . Briefly, the chamber facility
152 consists of two 12 m³ Teflon chambers suspended within a 21 x 12 ft temperature-controlled
153 enclosure. Black lights and natural sunlight fluorescent lamps surround the chambers, and multiple
154 sampling ports allow for injection of reagents, as well as gas- and aerosol-phase measurements.
155 Gas-phase O₃, NO₂, and NO_x concentrations were monitored using an O₃ analyzer (Teledyne
156 T400), a cavity attenuated phase shift (CAPS) NO₂ monitor (Aerodyne), and a chemiluminescence
157 NO_x monitor (Teledyne 200EU) respectively, while hydrocarbon decay was monitored using a gas
158 chromatography-flame ionization detector (GC-FID, Agilent 7890A). Hydrocarbon decay was
159 also used to estimate hydroxyl radical (OH) concentrations. For aerosol-phase measurements, a
160 Scanning Mobility Particle Sizer (SMPS, TSI) was used to measure aerosol volume concentrations
161 and distributions, while a High Resolution Time-of-Flight Aerosol Mass Spectrometer (HR-ToF-
162 AMS, Aerodyne; henceforth referred to as the AMS) was used to determine bulk aerosol
163 composition (DeCarlo et al., 2006). AMS data was analyzed using the data analysis toolkit
164 SQUIRREL (v. 1.57) and PIKA (v. 1.16G). Elemental ratios, including O:C, H:C, and N:C, were
165 obtained using the method outlined by Canagaratna et al. (2015) and used to calculate the average
166 carbon oxidation state (\overline{OS}_C) (Kroll et al., 2011). Temperature and relative humidity (RH) were
167 also monitored using a hydro-thermometer (Vaisala HMP110).

168 Experiments were designed to probe the effects of humidity, RO₂ fate, and precursor
169 identity on cellular inflammatory responses induced by different SOA formed under these
170 conditions (Table 1). All chamber experiments were performed at ~25 °C under dry (RH < 5%) or
171 humid (RH ~ 45%) conditions. Chambers were flushed with pure air ([generated from AADCO,](#)
172 [747-14](#)) for ~24 hrs prior to each experiment. During this time, chambers were also humidified for
173 humid experiments by means of a bubbler filled with deionized (DI) water. Seed aerosol was
174 injected by atomizing a 15 mM (NH₄)₂SO₄ seed solution (Sigma Aldrich) to obtain a seed
175 concentration of ~20 μg m⁻³. It should be noted that experimental conditions deviate for experiment
176 7 (isoprene SOA under RO₂ + HO₂ dominant, “humid” conditions) due to low SOA mass yields.
177 For this experiment, an acidic seed solution (8 mM MgSO₄ and 16 mM H₂SO₄) and a dry chamber
178 were used to promote SOA formation via the isoprene epoxydiol (IEPOX) uptake pathway. This
179 pathway has been shown to contribute significantly to ambient OA and has a higher SOA mass
180 yield compared to the IEPOX + OH pathway (Surratt et al., 2010; Lin et al., 2012; Xu et al., 2015).

181 SOA precursor was then introduced by injecting a known amount of hydrocarbon solution
182 [isoprene, 99%; α-pinene, ≥ 99%; β-caryophyllene, > 98.5%; pentadecane, ≥ 99%; *m*-xylene, ≥
183 99%; naphthalene, 99% (Sigma Aldrich)] into a glass injection bulb and passing [zeropure](#) air over
184 the solution until it fully evaporated. For pentadecane and β-caryophyllene, the glass bulb was also
185 heated gently during hydrocarbon injection to ensure full evaporation (Tasoglou and Pandis,
186 2015). Naphthalene was injected by passing [zeropure](#) air over solid naphthalene flakes as described
187 in previous studies (Chan et al., 2009). OH precursor was then introduced via injection of hydrogen
188 peroxide (H₂O₂) for RO₂ + HO₂ experiments or nitrous acid (HONO) for RO₂ + NO experiments.
189 For H₂O₂, a 50% aqueous solution (Sigma Aldrich) was injected using the same method described
190 for hydrocarbon injection to achieve an H₂O₂ concentration of 3 ppm. This amount yielded OH

191 concentrations on the order of 10^6 molec cm^{-3} . For HONO injections, HONO was first prepared
192 by adding 10 mL of 1%wt aqueous NaNO_2 (VWR International) dropwise into 20 mL of 10%wt
193 H_2SO_4 (VWR International) in a glass bulb. Zero air was then passed over the solution to introduce
194 HONO into the chamber (Chan et al., 2009; Kroll et al., 2005). Photolysis of HONO yielded OH
195 concentrations on the order of 10^7 molec cm^{-3} . NO and NO_2 were also formed as byproducts of
196 HONO synthesis. Once all the H_2O_2 evaporated ($\text{RO}_2 + \text{HO}_2$ experiments) or NO_x concentrations
197 stabilized ($\text{RO}_2 + \text{NO}$ experiments), the UV lights were turned on to initiate photooxidation.

198 **Aerosol collection and extraction.** Aerosol samples were collected onto 47 mm Teflon™
199 filters (0.45 μm pore size, Pall Laboratory). The total mass collected onto each filter was
200 determined by integrating the SMPS time-dependent volume concentration over the filter
201 collection period and multiplying by the total volume of air collected. SMPS volume
202 concentrations were converted to mass concentrations by assuming a density of 1 g cm^{-3} to
203 facilitate comparison between studies. To account for potential H_2O_2 or HONO uptake,
204 background filters were also collected. These filters were collected when only seed particles and
205 OH precursor (H_2O_2 or HONO) were injected into the chamber under otherwise identical
206 experimental conditions. All collected samples were placed in sterile petri dishes, sealed with
207 Parafilm M®, and stored at $-20 \text{ }^\circ\text{C}$ until extraction and analysis (Fang et al., 2015b). Collected
208 particles were extracted following the procedure outlined in Fang et al. (2015a) with modifications
209 for cellular exposure. Briefly, filter samples were submerged in cell culture media (RPMI-1640)
210 and sonicated for two 30 min intervals (1 hr total) using an Ultrasonic Cleanser (VWR
211 International). In between sonication intervals, the water was replaced to reduce bath temperature.
212 After the final sonication interval, sample extracts were filtered using 0.45 μm PTFE syringe filters

213 (Fisherbrand™) to remove any insoluble material and supplemented with 10% FBS (Fang et al.,
214 2015b).

215 **Intracellular ROS/RNS measurement.** ROS/RNS were detected using the assay
216 optimized in Tuet et al. (2016). Briefly, the assay consists of five major steps: (1) pre-treatment of
217 96-well plates to ensure a uniform cell density, (2) seeding of cells onto pre-treated wells at $2 \times$
218 10^4 cells well⁻¹, (3) incubation with ROS/RNS probe (carboxy-H₂DCFDA, Molecular Probes C-
219 400) diluted to a final concentration of 10 μ M, (4) exposure of probe-treated cells to samples and
220 controls for 24 hrs, and (5) detection of ROS/RNS using a microplate reader (BioTek Synergy H4,
221 ex/em: 485/525 nm). Positive controls included bacterial cell wall component lipopolysaccharide
222 (LPS, 1 μ g mL⁻¹), H₂O₂ (100 μ M), and reference filter extract (10 filter punches mL⁻¹, 1 per filter
223 sample, from various ambient filters collected at the Georgia Tech site, while negative controls
224 included blank filter extract (2 punches mL⁻¹) and control cells (probe-treated cells exposed to
225 media only, no stimulants).

226 A previous study on the ROS/RNS produced induced by exposure to ambient PM samples
227 found that ROS/RNS production was highly dose-dependent and could therefore not be
228 represented by measurements taken at a single dose (Tuet et al., 2016). Here, we utilize the dose-
229 response curve approach described in Tuet et al. (2016). For each aerosol sample, ROS/RNS
230 production was measured over ten dilutions and expressed as a fold increase in fluorescence over
231 control cells. A representative dose-response curve is shown in Fig. 1. For comparisons to other
232 inflammatory endpoints and chemical composition, ROS/RNS production was represented using
233 the area under the dose-response curve (AUC), as AUC has been shown to be the most robust
234 metric for comparing PM samples (Tuet et al., 2016).

235 **Cytokine measurement.** Secreted levels of TNF- α and IL-6 were measured post-exposure
236 (24 hrs) using enzyme-linked immunosorbent assay (ELISA) kits following manufacturer's
237 specifications (ThermoFisher). This time point was chosen to enable comparison with ROS/RNS
238 levels (also measured at 24 hrs, optimized in Tuet et al. (2016)) and to ensure a high signal for
239 both cytokines. Previous literature have shown that TNF- α and IL-6 production peak around 4 and
240 24 hrs, respectively (Haddad, 2001). However, while TNF- α production peaks earlier, the signal
241 at 24 hrs is well above the detection limit of the assay, and previous studies have utilized this time
242 point to measure both cytokines (Haddad, 2001; Matsunaga et al., 2001). Nonetheless, it should
243 be noted that these measurements represent a single time point in the cellular response. All
244 measurements were carried out using undiluted cell culture supernatant. For each aerosol sample,
245 TNF- α and IL-6 were measured over seven dilutions and represented as a fold increase over
246 control. Similarly, the AUC was used to represent each endpoint for comparison purposes.

247 **Cellular metabolic activity.** The MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-
248 diphenyltetrazolium bromide) assay (Biotium) was used to assess cellular metabolic activity.
249 Briefly, supernatants containing sample extracts were removed after the exposure period and
250 replaced with media containing MTT. Cells were then returned to the incubator for 4 hrs, during
251 which the tetrazolium dye was reduced by cellular NAD(P)H-dependent oxidoreductases to
252 produce an insoluble purple salt (formazan). Dimethyl sulfoxide was then used to solubilize the
253 salt and the absorbance at 570 nm was determined using a microplate reader (BioTek Synergy H4).

254 **Statistical analysis.** Linear regressions between bulk aerosol composition and cellular
255 inflammatory responses were evaluated using Pearson's correlation coefficient, and the
256 significance of each correlation coefficient was determined using multiple imputation, which
257 calculated the total variance associated with the slope of each regression. Details of this method

258 are described in Pan and Shimizu (2009). Briefly, response parameters (i.e. AUCs for each
259 endpoint) were assumed to follow a normal distribution. Ten “estimates” were obtained for each
260 response using the average and standard deviation determined from the dose-response curve fit.
261 These estimates were then plotted against bulk aerosol composition (e.g. O:C, H:C, and N:C) to
262 obtain ten fits, and the slopes and variances generated from these fits were used to calculate the
263 between and within variance. Finally, a Student’s *t*-test was used to calculate and evaluate the
264 associated *p*-values using a 95% confidence interval.

265 Results and Discussion

266 **Effect of SOA precursor and formation condition on SOA inflammatory response.** To
267 investigate whether SOA formed from different precursors elicited different inflammatory
268 responses, levels of ROS/RNS, TNF- α , and IL-6 were measured after exposing alveolar
269 macrophages to SOA generated from six VOCs generated under three formation conditions (Table
270 1). The AUC per mass of SOA (μg) in the extract for ROS/RNS, TNF- α and IL-6 are shown in
271 Fig. 2, shaped by SOA formation condition. It should be noted that all responses were normalized
272 to probe-treated control cells to account for differences between endogenous levels of ROS/RNS
273 produced in cells (Henkler et al., 2010). Uncertainties associated with AUC were determined by
274 averaging the AUCs obtained by fitting dose-response data with each point removed
275 systematically, following the methodology described in Tuet et al. (2016). ROS/RNS production
276 was also measured for background filters and found to be within the uncertainty of control cells,
277 indicating that there was no evidence for significant H₂O₂ or HONO uptake onto seed particles
278 (Fig. S1). Furthermore, exposure to filter extract did not result in decreases in metabolic activity
279 as measured by the MTT assay for all SOA systems investigated (Fig. S2). Since results from MTT
280 may represent the number of viable cells present, changes in inflammatory endpoints did not likely

281 result from changes in the number of cells exposed (i.e. decreases in response cannot be attributed
282 to cell death).

283 For all inflammatory responses measured (levels of ROS/RNS, TNF- α , and IL-6), SOA
284 precursor identity and formation condition influenced the level of response, as demonstrated by
285 the range of values obtained from different SOA precursors and different formation conditions
286 (Fig. 2). Despite having a clear effect, no obvious trends were observed for each variable (precursor
287 or formation condition) on individual responses. This is in contrast to that observed for the
288 oxidative potential as measured by DTT (OP^{WS-DTT}) for these samples, where only precursor
289 identity influenced OP^{WS-DTT} substantially (Tuet et al., 2017). However, this may not be surprising
290 as DTT is a chemical assay, which only accounts for the potential of species to participate in redox
291 reactions (Cho et al., 2005), whereas cellular assays account for many complicated cellular events
292 involved in intricate positive and negative feedback loops. Due to the considerably different
293 classes of compounds chosen as SOA precursors, aerosol compositional changes between different
294 precursors were generally larger than those between different formation conditions of the same
295 precursor (see Fig. 3a) (Tuet et al., 2017). DTT may only be sensitive to larger differences arising
296 from different precursors (~~i.e. a different carbon backbone~~), whereas cellular assays could also be
297 sensitive to differences between different formation conditions and chemical composition of the
298 same precursor. Moreover, while Tuet et al. (2017) showed that the intrinsic OP^{WS-DTT} spanned a
299 wide range, with isoprene and naphthalene SOA generating the lowest and highest OP^{WS-DTT} , these
300 bounds were less clear for cellular responses. While isoprene and naphthalene SOA still generated
301 the lowest and highest inflammatory responses in general, a few exceptions exist (e.g. ROS/RNS
302 levels induced by pentadecane SOA formed under dry, $RO_2 + HO_2$ dominant conditions, Fig. 2).

303 Though no apparent trends in individual inflammatory responses were observed as a
304 function of SOA precursor identity or formation condition, several patterns among all three
305 inflammatory responses were observed for SOA precursors with whose products share similar
306 chemical structures (i.e., similar carbon ~~backbones~~-chain length and functionalities). Exposure to
307 isoprene SOA induced the lowest levels of TNF- α and IL-6 among the aerosol systems studied
308 (Fig. 2). Furthermore, isoprene SOA generated from different pathways (i.e. photooxidation under
309 different RO₂ fates and reactive uptake of IEPOX) (Surratt et al., 2010; Xu et al., 2014; Chan et
310 al., 2010) produced similar responses for each inflammatory endpoint. These results suggest that
311 different isoprene SOA products (Surratt et al., 2010; Xu et al., 2014; Chan et al., 2010) may induce
312 similarly low inflammatory responses and are consistent with the intrinsic OP^{WS-DTT} obtained for
313 these SOA samples, where isoprene SOA generated the lowest OP^{WS-DTT} of all SOA systems
314 studied and the OP^{WS-DTT} was similar for all SOA formation conditions explored (Tuet et al., 2017).
315 This finding is in contrast to a previous study by Lin et al. (2016), where methacrylic acid epoxide
316 (MAE)-derived SOA was found to be substantially more potent than IEPOX-derived SOA.
317 However, while exposure to MAE-derived SOA induced the upregulation of a larger number of
318 oxidative stress response genes than IEPOX-derived SOA, the fold change of several genes
319 reported in Lin et al. (2016) are actually similar: (e.g., *ALOX12*, *NQO1*). Several of these genes
320 directly affect the production of inflammatory cytokines measured in this study. For instance,
321 studies have observed that arachidonate 12-lipoxygenase (*ALOX12*) products induce the
322 production of both TNF- α and IL-6 in macrophages (Wen et al., 2007). As such, a similar response
323 level regardless of SOA formation condition may be observed depending on the biological
324 endpoints measured. Thus, it is possible that the inflammatory cytokines measured in this study

325 are involved in pathways concerning those genes, resulting in a similar response level regardless
326 of SOA formation condition.

327 Similarly, exposure to SOA generated from the photooxidation of α -pinene and *m*-xylene
328 resulted in similar inflammatory responses for all three formation conditions (Fig. 2). These
329 cellular assay results are consistent with results from the DTT assay where the OP^{WS-DTT} was not
330 significantly different between SOA formed under different formation conditions (Tuet et al.,
331 2017). Response levels induced by these two SOA systems are also similar across all three
332 inflammatory measurements investigated (Fig. 2). This suggests that products from both
333 precursors may induce similar cellular pathways resulting in the production of similar levels of
334 inflammatory markers. Indeed, there are several similarities between products formed from the
335 photooxidation of α -pinene and *m*-xylene. For instance, a large portion of α -pinene and *m*-xylene
336 oxidation products under both $RO_2 + HO_2$ and $RO_2 + NO$ pathways are ring-breaking products
337 with a similar carbon chain length (Eddingsaas et al., 2012; Vivanco and Santiago, 2010; Jenkin
338 et al., 2003). As a result of this similarity, products from both SOA systems may interact with the
339 same cellular targets and induce similar cellular pathways, resulting in a similar response
340 regardless of precursor identity and formation condition. These observations further imply that the
341 chemical structures (e.g., carbon chain lengths and functionalities) of oxidation products may be
342 important regardless of PM source/precursor.

343 A different pattern was observed for β -caryophyllene and pentadecane SOA, where the IL-
344 6 response spanned a much larger range than ROS/RNS and TNF- α (Fig. 2). This is in contrast to
345 the trends observed for the OP^{WS-DTT} for β -caryophyllene and pentadecane SOA, where OP^{WS-DTT}
346 was similar regardless of formation condition (Tuet et al., 2017). This suggests that there are
347 differences between organic peroxides and organic nitrates formed from certain precursors that

348 influence cellular responses, but are not captured by redox potential measurements. Less is known
349 about the effects of humidity on SOA formation and chemical composition for all SOA systems
350 investigated, as most laboratory chamber studies in literature have been conducted under dry
351 conditions. Specifically here, very high levels of IL-6 were observed post-exposure to pentadecane
352 SOA formed under humid conditions. Prior studies reported opposing findings with some showing
353 a significant effect of water on aerosol formation and chemical composition (Nguyen et al., 2011;
354 Wong et al., 2015; Healy et al., 2009; Stirnweis et al., 2016), while others found little influence
355 (Edney et al., 2000; Boyd et al., 2015; Cocker III et al., 2001). It is clear that humidity effects are
356 highly hydrocarbon-dependent and further studies into the specific products formed under humid
357 conditions are required to understand how these differences in chemical composition may translate
358 to different cellular endpoints. Nonetheless, the known products formed from the photooxidation
359 of these hydrocarbons may provide some insight into the inflammatory responses observed. While
360 there are no prior studies involving pentadecane oxidation products, it is expected that the
361 oxidation products will be similar to those reported in the oxidation of dodecane (i.e. same
362 functionalities with a longer carbon chain) (Loza et al., 2014). It is therefore likely that pentadecane
363 oxidation products resemble long chain fatty acids and could potentially insert into the cell
364 membrane (Loza et al., 2014); as previous studies have shown that fatty acids can feasibly insert
365 into the cell membrane bilayer (Khmelinskaia et al., 2014; Cerezo et al., 2011). This insertion
366 could potentially affect membrane fluidity, which is known to affect cell function substantially
367 although the specific effect depends strongly on the particular modification and cell type of interest
368 (Baritaki et al., 2007; Spector and Yorek, 1985). In some cases, these alterations lead to the
369 induction of apoptosis, which involves pathways leading to the production of TNF- α (Baritaki et
370 al., 2007; Wang et al., 2003). TNF- α can then induce the production of IL-6, which once produced

371 can also inhibit the production of TNF- α in a feedback loop (Kishimoto, 2003; Wang et al., 2003).
372 These cellular events are consistent with the observed inflammatory response induced by
373 pentadecane SOA exposure, where there is a high IL-6 response and a lower TNF- α response. The
374 low ROS/RNS response observed is also in line with these cellular events, as IL-6 exhibits anti-
375 inflammatory functions, which can neutralize ROS/RNS production. These responses are less
376 pronounced for β -caryophyllene aerosol, which may be due to the shorter carbon chain observed
377 in known products (Chan et al., 2011). While β -caryophyllene and pentadecane are both C15
378 precursors, β -caryophyllene is a bicyclic compound and many SOA products retain the 4-
379 membered ring, resulting in a shorter carbon backbone (Chan et al., 2011). As a result, fewer
380 products may insert into the cell membrane, leading to a lesser response compared to pentadecane
381 SOA exposure. These observations, particularly those for pentadecane SOA, suggest that aerosols
382 from meat cooking may have health implications, as fatty acids comprise a majority of these
383 aerosols (Mohr et al., 2009; Rogge et al., 1991).

384 Naphthalene exhibits a ~~completely~~ different, more distinct pattern ~~from compared to~~ the
385 rest of the SOA systems investigated, with a large range observed for both TNF- α and IL-6 under
386 different formation conditions (Fig. 2). Higher levels of ROS/RNS were also observed as a result
387 of exposure to naphthalene aerosol irrespective of SOA formation condition. Similarly, the OP^{WS-}
388 DTT of naphthalene SOA previously measured by Tuet et al. (2017) was an outlier among all SOA
389 systems investigated, as the measured OP^{WS-DTT} was at least twice that of the next highest SOA
390 system. These observations are consistent with the formation of specific SOA products such as
391 naphthoquinones, which are known to induce redox-cycling in cells and are formed under both
392 $RO_2 + HO_2$ and $RO_2 + NO$ pathways (Henkler et al., 2010; Kautzman et al., 2010). Consequently,
393 aerosol generated from naphthalene may induce higher levels of inflammatory responses than

394 other SOA due to this process (Henkler et al., 2010; Lorentzen et al., 1979). However, as shown
395 by the high levels of IL-6, exposure to naphthalene SOA may also induce anti-inflammatory
396 pathways not captured by OP^{WS-DTT} measurements. Moreover, a clear increasing trend is apparent
397 for TNF- α and IL-6 produced upon naphthalene SOA exposure, with a higher level of both
398 cytokines observed for aerosol formed under RO₂ + NO dominant and humid conditions.
399 Previously, the effect of different RO₂ fates on SOA OP^{WS-DTT} was attributed to the different
400 products known to form under both pathways (Tuet et al., 2017). The same explanation applies for
401 cellular measurements as SOA products that promote electron transfer reactions with ~~anti-~~
402 ~~oxidants~~santioxidants can result in redox imbalance as measured by OP^{WS-DTT} and the induction of
403 related cellular pathways such as ROS/RNS and cytokine production (Tuet et al., 2017). Finally,
404 naphthalene SOA induced cellular responses outside of those observed for other aerosol systems,
405 with higher levels of all inflammatory markers than other SOA systems. As shown previously for
406 OP^{WS-DTT}, naphthalene may be an outlier due to aromatic ring-containing products, which may
407 then induce different cellular pathways compared to other aerosol systems investigated, the
408 products of which do not contain aromatic rings. Additionally, many known aerosol products
409 formed from the photooxidation of naphthalene have functionalities that resemble those of
410 dinitrophenol, which is known to decouple phosphorylation from electron transfer (Terada, 1990).
411 It is therefore possible that the aromatic functionality present in the majority of naphthalene SOA
412 products results in the involvement of very different cellular pathways, leading to outlier
413 inflammatory endpoint responses. Various products of naphthalene oxidation such as
414 nitroaromatics and polyaromatics are known to have mutagenic properties and may induce the
415 formation of DNA adducts (Baird et al., 2005; Helmig et al., 1992). As such, it is possible that

416 these products may induce health effects via other pathways as well and naphthalene SOA
417 exposure may have effects beyond redox imbalance and oxidative stress.

418 Bulk aerosol elemental ratios (O:C, H:C, and N:C) were determined for each SOA system
419 investigated. Different types of organic aerosol are known to span a wide range of O:C, which may
420 be utilized as an indication of oxidation, and the van Krevelen diagram was used to visualize
421 whether changes in O:C and H:C ratios corresponded to changes in levels of inflammatory
422 response (Fig. 3a, S3) (Chhabra et al., 2011; Lambe et al., 2011; Ng et al., 2010). Changes in the
423 slope within the van Krevelen space provide information on SOA functionalization (Heald et al.,
424 2010; Van Krevelen, 1950; Ng et al., 2011). Beginning from the precursor hydrocarbon, a slope
425 of 0 indicates alcohol group additions, a slope of -1 indicates carbonyl and alcohol additions on
426 separate carbons or carboxylic acid additions, and a slope of -2 indicates ketone or aldehyde
427 additions.

428 As seen in Fig. 3a, the laboratory-generated aerosols span a large range of O:C and H:C
429 ratios. Both SOA formation condition and precursor identity influenced elemental ratios, however,
430 precursor identity generally had a larger effect as evident by the clusters observed for different
431 SOA precursors. Despite these differences in chemical composition, there were no obvious trends
432 between O:C or H:C and any inflammatory endpoint measured. This is similar to that observed for
433 chemical oxidative potential as measured by DTT, where a higher O:C did not correspond to a
434 higher oxidative potential for both laboratory-generated and ambient aerosols (Tuet et al., 2017).
435 This is likely due to the different formation conditions used to generate SOA, which may not be
436 directly comparable. Nevertheless, a significant correlation ($p < 0.05$) was observed between
437 ROS/RNS and \overline{OS}_c (Fig. 3b). This positive correlation is not surprising, as a higher average
438 oxidation state would likely correspond to a better oxidizing agent. Future studies should evaluate

439 the effect of the degree of oxidation for SOA formed from the same SOA precursor under the same
440 formation condition to investigate whether atmospheric aging of aerosol (which typically leads to
441 increases in the degree of oxidation) affects inflammatory responses. Finally, the N:C ratio was
442 also determined for SOA systems formed under conditions that favor the RO₂ + NO pathway (Fig.
443 S4) and were found to span a large range. Similarly, there was no obvious trend between N:C ratios
444 and the inflammatory endpoints measured.

445 **Relationship between inflammatory responses.** To visualize whether there exists a
446 relationship between inflammatory markers measured, levels of TNF- α and IL-6 are shown in Fig.
447 4, sized by ROS/RNS. With the exception of naphthalene SOA, the inflammatory cytokine
448 responses for all aerosol systems investigated follow an exponential curve (Fig. 4, shown in black)
449 where there appears to be a plateau for TNF- α levels. Along this curve, ROS/RNS levels also
450 appear to increase with increasing inflammatory cytokine levels to a certain point, after which
451 ROS/RNS levels decrease. These observations are in line with the interconnected effects of both
452 cytokines. While both TNF- α and IL-6 have pro-inflammatory effects that may lead to the increase
453 of ROS/RNS production, the individual pathways are also involved in many complicated
454 stimulation and inhibition loops and there is extensive cross-talk between both pathways. For
455 instance, TNF- α induces the production of glucocorticoids, which in turn inhibits both TNF- α and
456 IL-6 production (Wang et al., 2003). IL-6 also directly inhibits the production of TNF- α and other
457 cytokines induced as a result of TNF- α (e.g. IL-1) and stimulates pathways that lead to the
458 production of glucocorticoids (Kishimoto, 2003). As a result, increases in IL-6 may be
459 accompanied by decreases in TNF- α , resulting in the observed plateau. Furthermore, ROS/RNS
460 levels may represent a fine balance between anti-inflammatory and pro-inflammatory effects. Both
461 cytokines are involved in the acute phase reaction and can affect ROS/RNS levels via pro-

462 inflammatory pathways. IL-6 also exhibits some anti-inflammatory functions and may thus lower
463 ROS/RNS levels as well. These interconnected pathways could account for the observed parabolic
464 pattern for ROS/RNS production. Exposure to naphthalene SOA resulted in responses outside of
465 those observed for other aerosol systems, likely due to the formation of aromatic ring-retaining
466 products as discussed in the previous section.

467 **Comparison with ambient data.** To evaluate how the oxidative potential and ROS/RNS
468 production of the SOA systems investigated compare in the context of ambient samples, the
469 measurements obtained in this study were plotted with those obtained in our previous study
470 involving ambient samples collected around the greater Atlanta area (Fig. 5) (Tuet et al., 2016).
471 These ambient samples were analyzed using the same methods for determining oxidative potential
472 (DTT assay (Cho et al., 2005; Fang et al., 2015b)) and ROS/RNS production (cellular carboxy-
473 H₂DCFDA assay (Tuet et al., 2016)). Furthermore, the same extraction protocol (water-soluble
474 extract) was followed in both studies (Tuet et al., 2016). Results from both studies are therefore
475 directly comparable. Previously, a significant correlation between ROS/RNS production and
476 oxidative potential as measured by DTT was observed for summer ambient samples. In the same
477 study, correlations between ROS/RNS production and organic species were also observed for
478 summer ambient samples, and it was suggested that these correlations may reflect contributions
479 from photochemically produced SOA (Tuet et al., 2016).

480 Fig. 5 shows that laboratory-generated SOA oxidative potential is comparable to that
481 observed in ambient samples, with the exception of naphthalene SOA, which produced higher
482 DTT activities due to its aromatic ring retaining products (Tuet et al., 2017; Kautzman et al., 2010).
483 Laboratory-generated SOA also induced similar or higher levels of ROS/RNS compared to
484 ambient samples. There are many possible explanations for the observed higher response for some

485 SOA samples. For instance, individual, single precursor SOA systems were considered in this
486 study, whereas ambient aerosol contains SOA from multiple precursors as well as other species
487 that are not considered in this study (e.g. metals). Interactions between SOA from different
488 precursors is likely to occur and may result in different response levels. Complex interactions
489 between SOA and other species present in the ambient (e.g. metals or other organic species) are
490 also likely involved (Tuet et al., 2016). Previous studies have also suggested the possibility of
491 metal-organic complexes. For instance, Verma et al. (2012) showed that certain metals were
492 retained on a C-18 column, which is utilized to remove hydrophobic components, suggesting that
493 these metals were likely complexed and removed in the process. Further chamber studies involving
494 photochemically generated SOA and metals may elucidate these interactions. Furthermore, there
495 are likely species present in the ambient that do not contribute to ROS/RNS production. That is,
496 while certain species contribute to the mass of PM, there is little to no ROS/RNS production
497 associated with these species. Ambient samples where these species comprise a significant fraction
498 will have a low per mass ROS/RNS production level. Finally, only three SOA formation conditions
499 were investigated in this study. There are multiple other possible oxidation mechanisms that lead
500 to the formation of SOA in the ambient, which were not accounted for in this study. Nonetheless,
501 despite the low ROS/RNS levels observed post SOA exposure, there is an association between
502 ROS/RNS production and DTT activity (Fig. 5). These results suggest that our previous findings
503 based on ambient filter samples may be extended to SOA samples. That is, while the relationship
504 between ROS/RNS production and DTT activity is complex, DTT may serve as a useful screening
505 tool as samples with low DTT activities are likely to produce low levels of ~~RNS~~ROS/RNS (Tuet
506 et al., 2016).

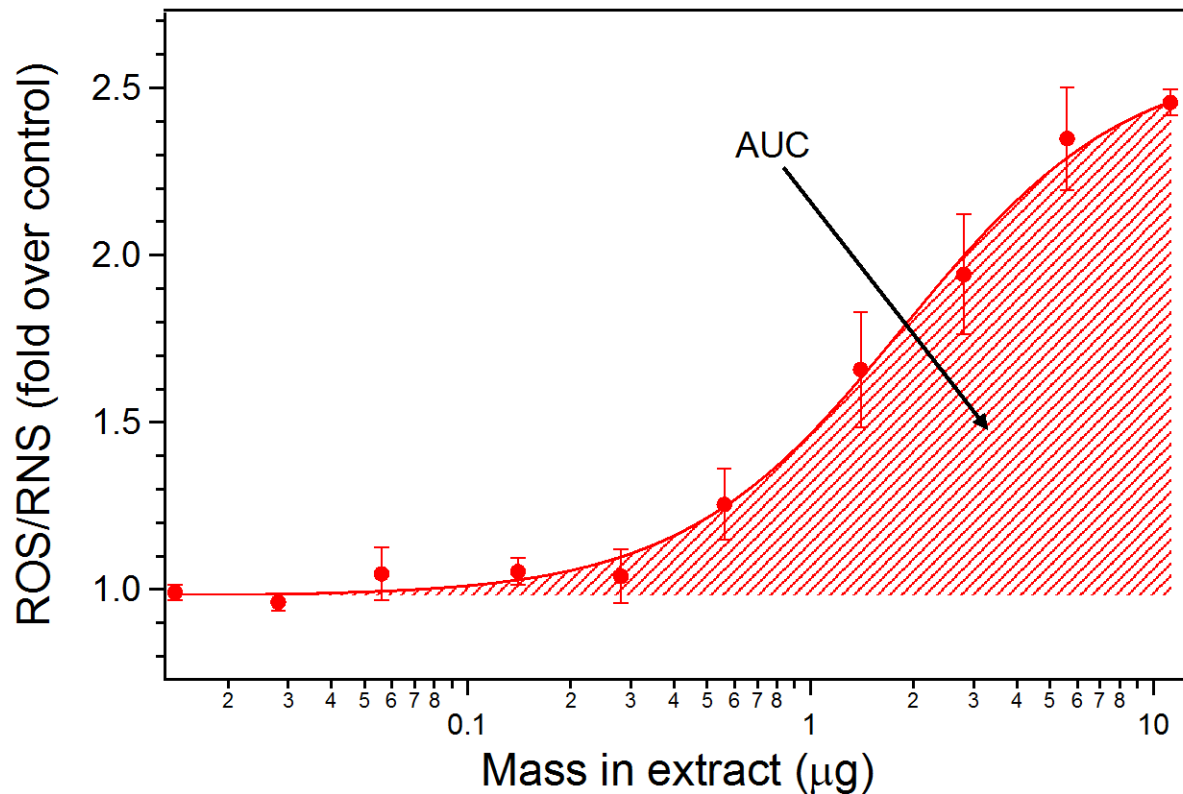
507 **Implications.** Levels of ROS/RNS, TNF- α , and IL-6 were measured after exposing cells
508 to the water-soluble extract of SOA generated from the photooxidation of six SOA precursors
509 under various formation conditions. Although previous epidemiological and ambient studies have
510 found correlations between metals and various measures of health effects (Verma et al., 2010;
511 Pardo et al., 2015; Burnett et al., 2001; Huang et al., 2003; Akhtar et al., 2010; Charrier and
512 Anastasio, 2012), the measured levels of TNF- α , IL-6, and ROS/RNS obtained in this study
513 demonstrate that organic aerosols alone can induce a cellular response. This was previously
514 observed for the oxidative potential as measured by DTT activity as well, where the same
515 laboratory-generated organic aerosol samples catalyzed redox reactions and resulted in
516 measureable DTT decay in the absence of metal species (Tuet et al., 2017).

517 Results from this study also show that SOA precursor identity and formation condition
518 influenced response levels, with naphthalene SOA producing the highest cellular responses of the
519 SOA systems investigated. As discussed previously, the aromatic functionality present in many
520 naphthalene photooxidation products may be an important consideration for health effects. It may
521 therefore be worthwhile to investigate other anthropogenic aromatic ring-containing precursors as
522 well and to closely study the cellular effects of naphthalene SOA products given its high response.
523 Several patterns were also noted for SOA systems whose products shared similar functionalities
524 and chemical structures. For instance, photooxidation productions from pentadecane and β -
525 caryophyllene share similarities with long chain fatty acids and may participate in membrane
526 insertions, whereas many known products of naphthalene photooxidation are mutagens capable of
527 inducing cellular pathways beyond those that affect cellular redox balance (Baird et al., 2005;
528 Helmig et al., 1992). Given these observations, it may be possible to roughly predict responses
529 based on known SOA products as SOA systems whose products share similar functionalities and

530 carbon chain length (~~i.e. similar carbon backbone~~) are likely to induce similar cellular pathways
531 and produce similar levels of various inflammatory endpoints. Exposure studies involving
532 individual classes of SOA products may elucidate further details as to whether these types of
533 predictions would be plausible. Moreover, such studies could be used to determine whether the
534 hypothesized cellular pathways are indeed involved and whether certain cellular functions are
535 indeed affected by specific products (e.g. membrane insertion by pentadecane photooxidation
536 products and oxidative phosphorylation decoupling by naphthalene photooxidation products).

537 Mixture effects may be another important consideration as ambient PM contains SOA
538 formed from multiple SOA precursors. As a result, precursor emissions and their corresponding
539 SOA formation potential must be considered to fully assess PM health effects. Furthermore, it may
540 be worthwhile to investigate various prediction models for multi-component mixtures to bridge
541 the gap between laboratory studies and real ambient exposures. For instance, concentration
542 addition may not apply as ambient aerosol is formed in the presence of multiple precursors and the
543 SOA produced may induce response levels completely different from those observed for single
544 precursor SOA systems that comprise the mixture. Interactions between organic components and
545 metal species have also been suggested in previous studies (Verma et al., 2012; Tuet et al., 2016)
546 and may influence responses significantly. While these interactions were not considered in the
547 current study, there may be evidence to support the plausibility of mixture effects as ambient PM
548 samples produced lower levels of ROS/RNS than that of any single SOA system investigated.
549 Laboratory chambers can serve as an ideal platform to investigate mixture effects, as experiments
550 can be conducted under well-controlled conditions where the aerosol chemical composition and
551 health endpoints can be determined.

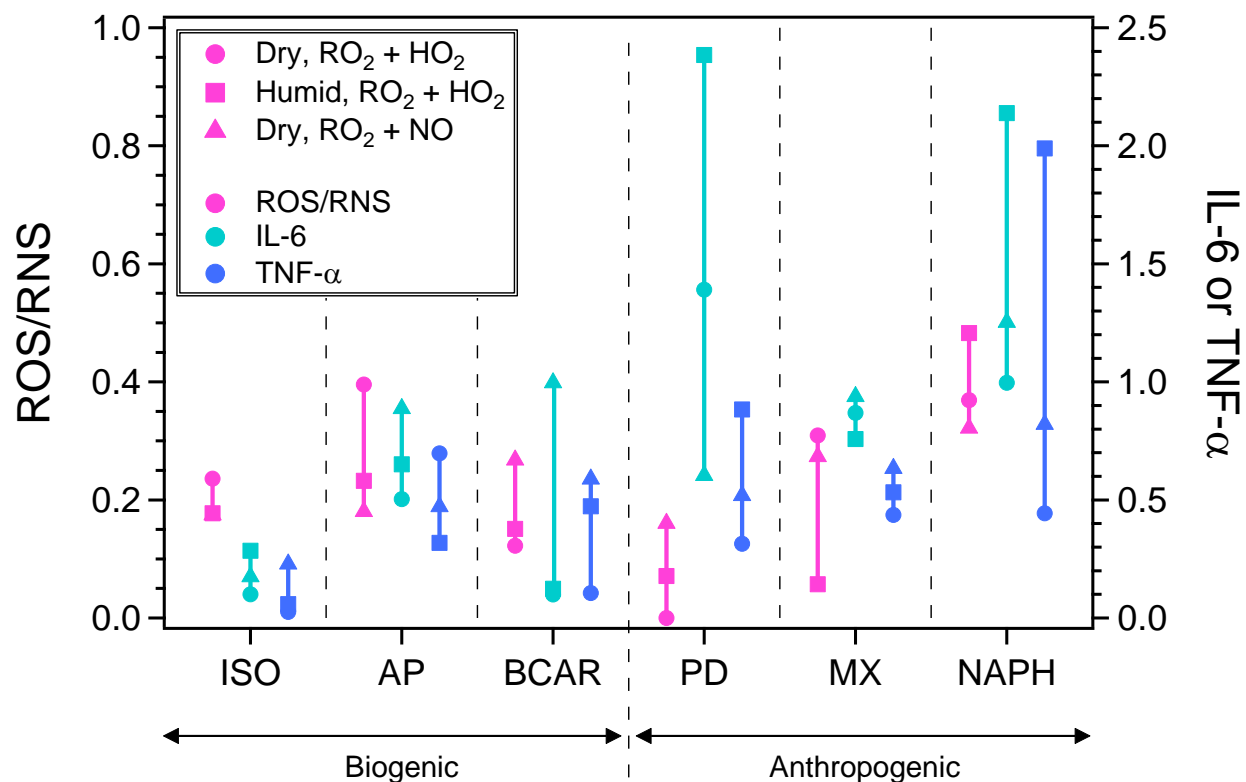
552 ~~Finally~~Additionally, this study confirms that while there is not one simple correlation
553 between oxidative potential and cellular responses for different PM samples, the DTT assay may
554 serve as a useful screening tool as a low DTT activity will likely correspond to a low cellular
555 response. Furthermore, while ROS/RNS may serve as a general indicator of oxidative stress, there
556 may be instances where a low level of ROS/RNS does not necessary indicate a lack of cellular
557 response. In the current study, ROS/RNS levels were associated with levels of inflammatory
558 cytokines for the majority of SOA systems investigated. However, aerosol formed from the
559 photooxidation of pentadecane induced low levels of ROS/RNS production and relatively high
560 levels of both cytokines (i.e. higher than expected given the ROS/RNS level measured). These
561 results suggest that at least one additional measure (e.g. inflammatory cytokines) may be required
562 to fully interpret ROS/RNS measurements. Finally, several limitations must be considered before
563 generalizing results from this study to *in vivo* exposures. For instance, only one cell type was
564 explored in this study, whereas an organism consists of multiple tissues comprised of multiple cell
565 types. Interactions between different cell types and tissue systems were not considered in this
566 study. Furthermore, the doses investigated may not fully represent real world exposures due to
567 differences in exposure routes and potential recovery from doses due to clearance. Nevertheless,
568 this study provides perspective on the relative toxicities of different SOA systems which future
569 studies can build upon.



570

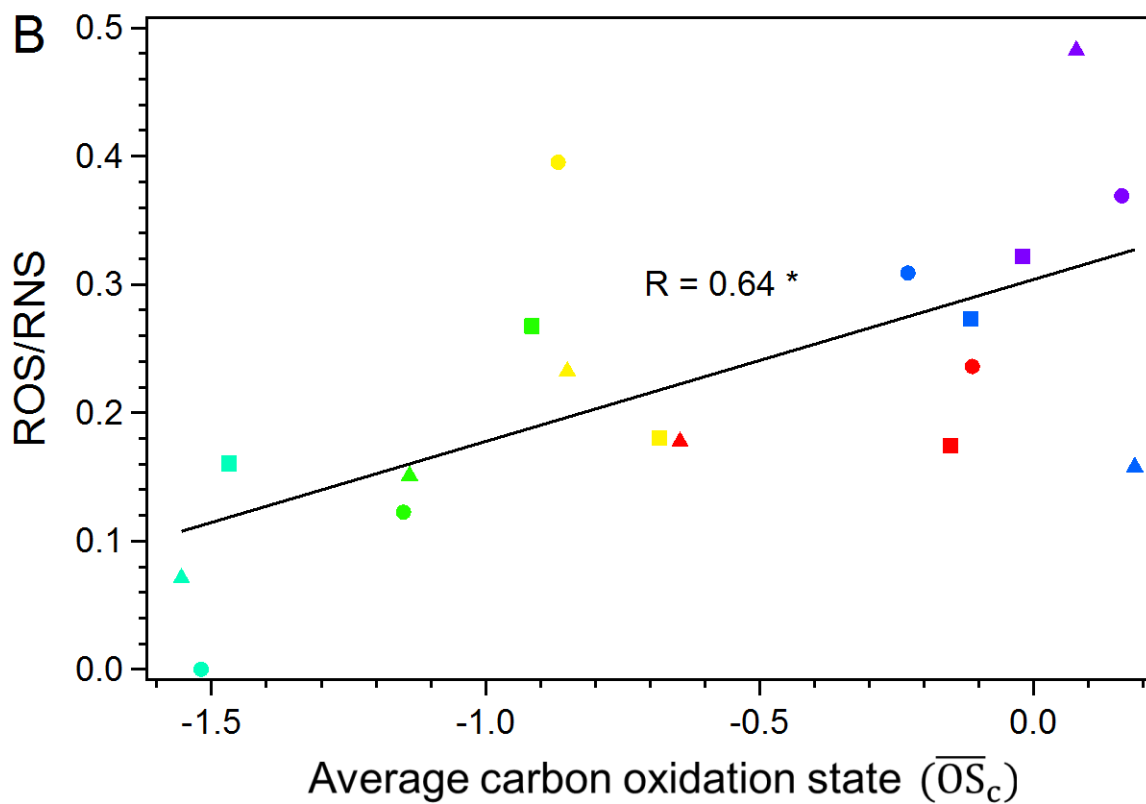
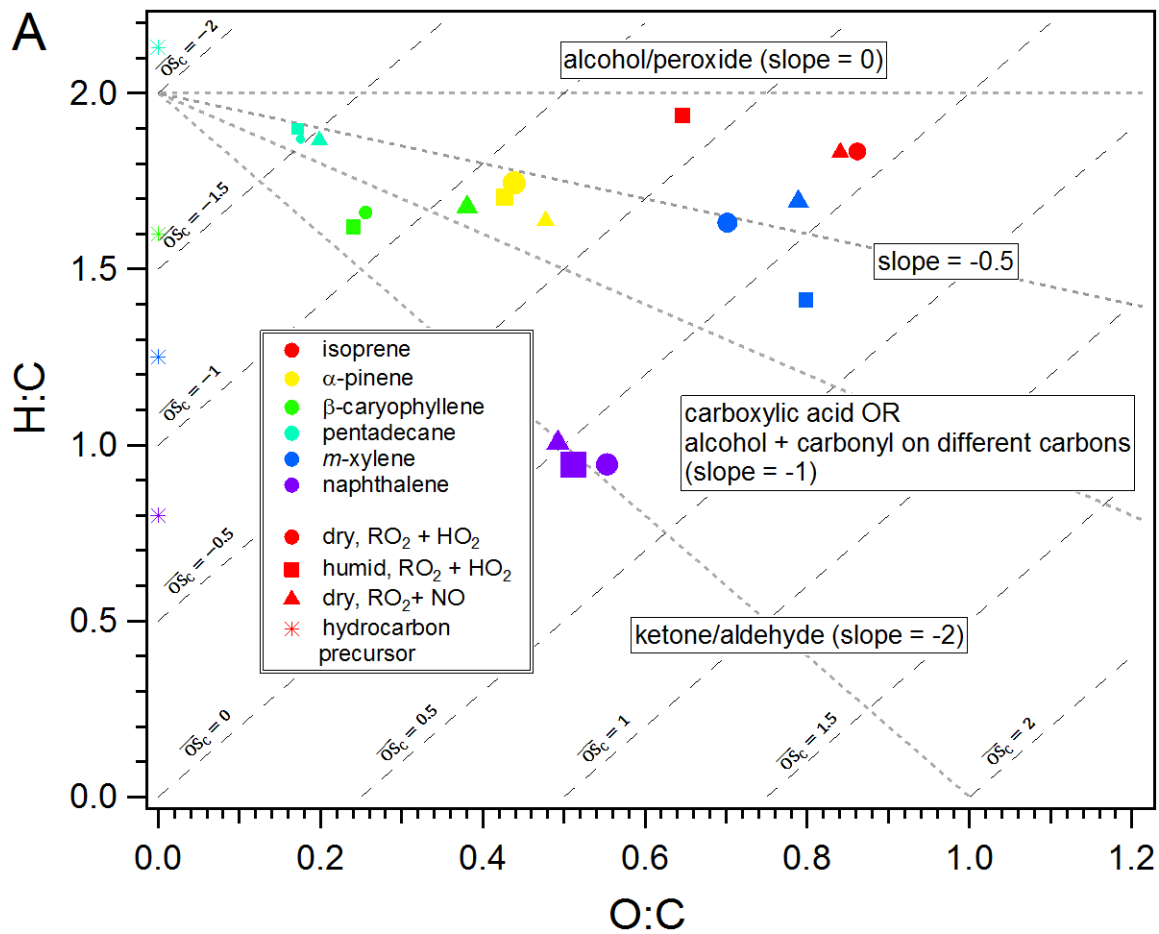
571 **Figure 1.** Representative dose-response curve of ROS/RNS produced as a result of filter
 572 exposure (naphthalane SOA formed under dry, RO₂ + NO dominant conditions). ROS/RNS is
 573 expressed as a fold increase over control cells, defined as probe-treated cells incubated with
 574 stimulant-free media. Dose is expressed as mass in extract (µg). Data shown are means ±
 575 standard error of triplicate exposure experiments. The Hill equation was used to fit the dose-
 576 response curve and the area under the dose-response curve (AUC) is shown.

577

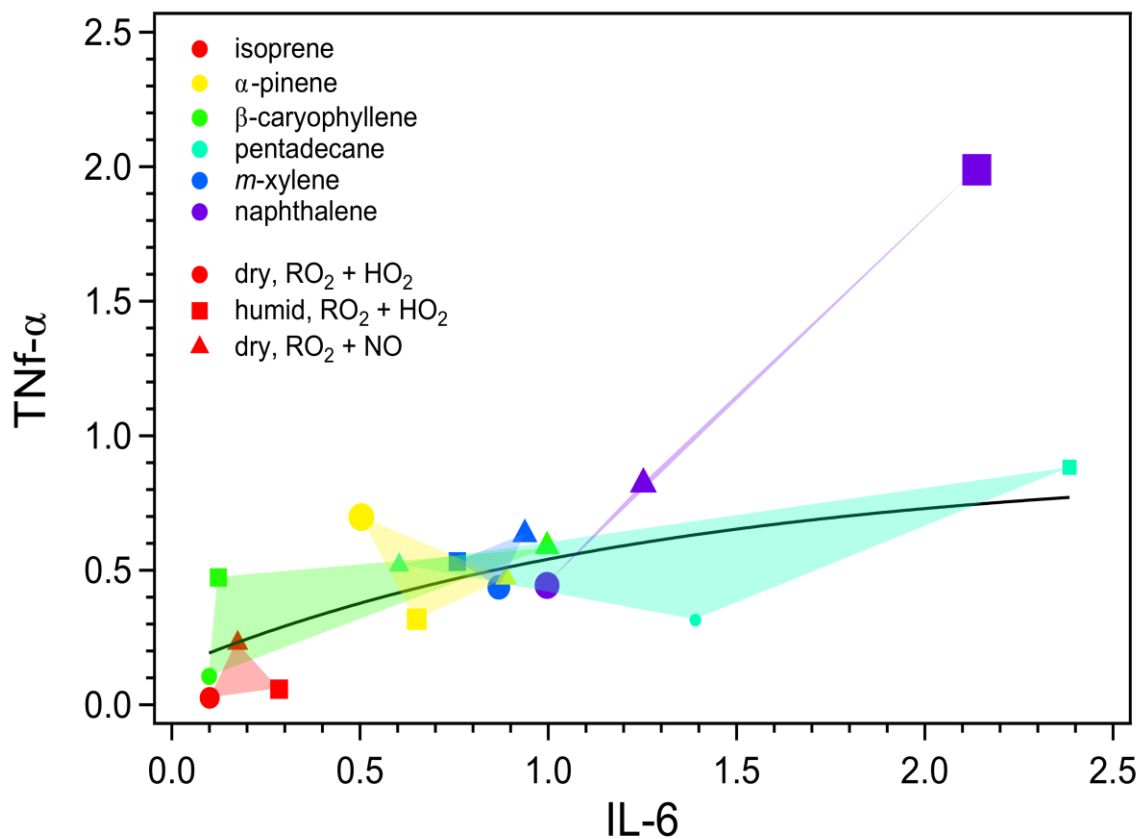


578

579 **Figure 2.** Area under the dose-response curve for various inflammatory responses induced as a
 580 result of SOA exposure: **ROS/RNS**, **IL-6**, and **TNF- α** . SOA were generated from various
 581 precursors (ISO: isoprene, AP: α -pinene, BCAR: β -caryophyllene, PD: pentadecane, MX: *m*-
 582 xylene, and NAPH: naphthalene) under various conditions (circles: dry, RO₂ + HO₂; squares:
 583 humid, RO₂ + HO₂; and triangles: dry, RO₂ + NO). Lines connecting the same inflammatory
 584 response for SOA generated from the same precursor under different formation conditions are also
 585 shown.

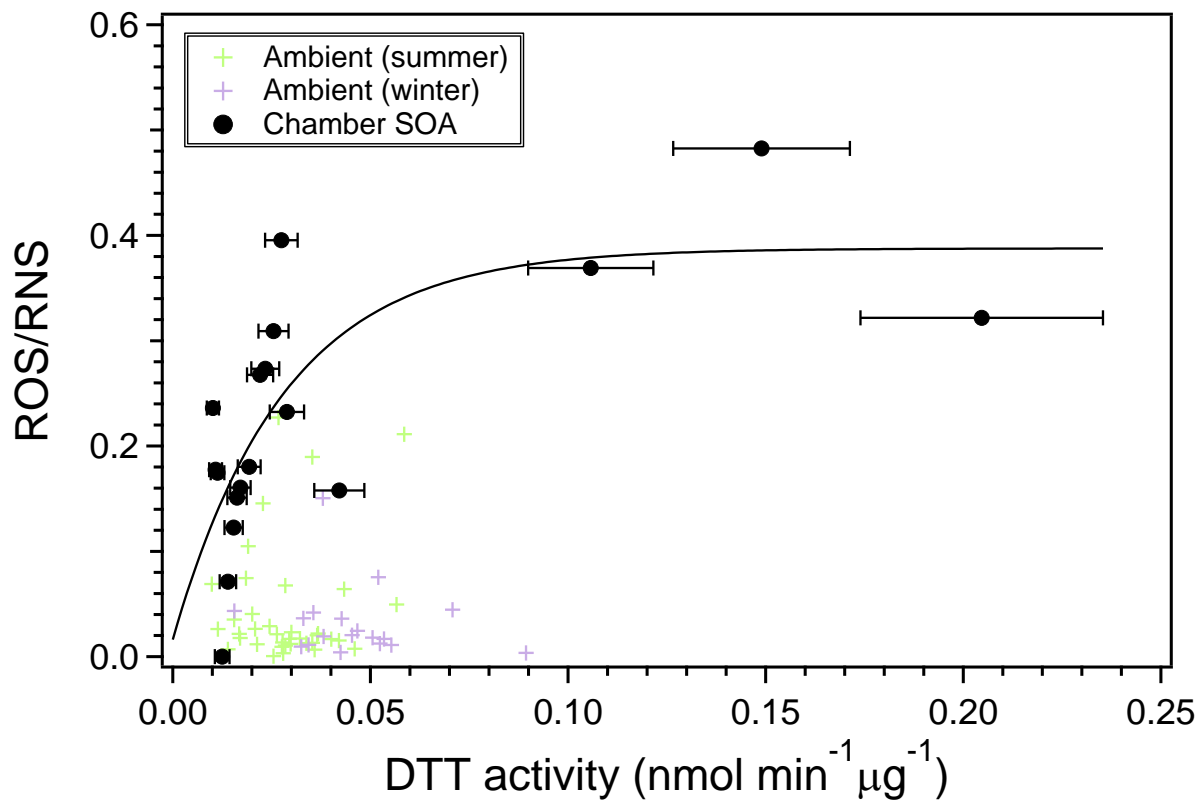


587 **Figure 3.** van Krevelen plot for various SOA systems sized by ROS/RNS levels (panel A) and
588 correlation between ROS/RNS levels and average carbon oxidation state (panel B). Data points
589 are colored by SOA system (red: isoprene, yellow: α -pinene, green: β -caryophyllene, light blue:
590 pentadecane, blue: *m*-xylene, and purple: naphthalene), shaped according to formation conditions
591 (circle: dry, RO₂ + HO₂; square: humid, RO₂ + HO₂; and triangle: dry, RO₂ + NO). SOA precursors
592 are shown as stars, colored by SOA system. * indicates significance, $p < 0.05$.



593

594 **Figure 4.** Area under the dose-response curve per mass of SOA for various inflammatory
 595 responses induced as a result of SOA exposure. Data points are sized according to ROS/RNS level.
 596 SOA were generated from various SOA precursors (red: isoprene, yellow: α -pinene, green: β -
 597 caryophyllene, light blue: pentadecane, blue: *m*-xylene, and purple: naphthalene) under various
 598 conditions (circles: dry, $\text{RO}_2 + \text{HO}_2$; squares: humid, $\text{RO}_2 + \text{HO}_2$; and triangles: dry, $\text{RO}_2 + \text{NO}$).
 599 A fitted curve excluding naphthalene data is shown as a guide. Shaded regions for each system,
 600 colored by SOA precursor, are also shown to show the extent of clustering and provide a
 601 visualization for the different patterns observed.



602
 603 **Figure 5.** ROS/RNS production and intrinsic DTT activities for chamber SOA and ambient
 604 samples collected around the greater Atlanta area. All samples were analyzed using the method
 605 outlined in Cho et al. (2005) and Tuet et al. (2016). Ambient samples are colored by season as
 606 determined by solstice and equinox dates between June 2012 and October 2013 (Tuet et al.,
 607 2016). A fitted curve for laboratory-generated samples is shown as a guide.

608 **Table 1.** Experimental conditions.

Experiment	SOA precursor	OH precursor	Relative humidity (%)	[HC] ₀ (ppb)
1	isoprene	H ₂ O ₂	<5%	97
2	α -pinene	H ₂ O ₂	<5%	191
3	β -caryophyllene	H ₂ O ₂	<5%	36
4	pentadecane	H ₂ O ₂	<5%	106
5	<i>m</i> -xylene	H ₂ O ₂	<5%	450
6	naphthalene	H ₂ O ₂	<5%	178
7	isoprene	H ₂ O ₂	<5% ^a	97
8	α -pinene	H ₂ O ₂	40%	334
9	β -caryophyllene	H ₂ O ₂	42%	63
10	pentadecane	H ₂ O ₂	45%	106
11	<i>m</i> -xylene	H ₂ O ₂	45%	450
12	naphthalene	H ₂ O ₂	44%	431
13	isoprene	HONO	<5%	970
14	α -pinene	HONO	<5%	174
15	β -caryophyllene	HONO	<5%	21
16	pentadecane	HONO	<5%	74
17	<i>m</i> -xylene	HONO	<5%	431
18	naphthalene	HONO	<5%	145

609 ^a Acidic seed (8 mM MgSO₄ and 16 mM H₂SO₄) was used instead of 8 mM (NH₄)₂SO₄

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614 ABBREVIATIONS

615 PM: particulate matter; SOA: secondary organic aerosol; ROS/RNS: reactive oxygen/nitrogen
616 species; TNF- α : tumor necrosis factor- α ; IL-6: interleukin-6

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