



Impact of combustion-generated particles on respiratory response to viral infection

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Disclosure to Learners



- Financial relationships with relevant companies within the past 24 months:
 - None

Air Pollution and SARS-CoV-2 – Epidemiological Studies from around the World



- Large portions of air pollution throughout the world are generated via combustion (wildfires, burning of fossil fuels, biomass burning, etc.) (Lightly et al., 2000)
- During the SARS outbreak in 2002, SARS patients coming from areas of high air pollution were more than twice as likely to die from the disease (Cui et al., 2003)
- Long-term exposure to air pollution was associated with increased COVID-19 severity (Conticini et al., 2020)
- Higher historical PM_{2.5} exposures are positively associated with higher county-level COVID-19 mortality (Wu et al., 2020)
- Environmental factors (air quality index) are associated with daily number of COVID-19 cases (Ma et al., 2021)

Why could what we breathe affect viral infections?

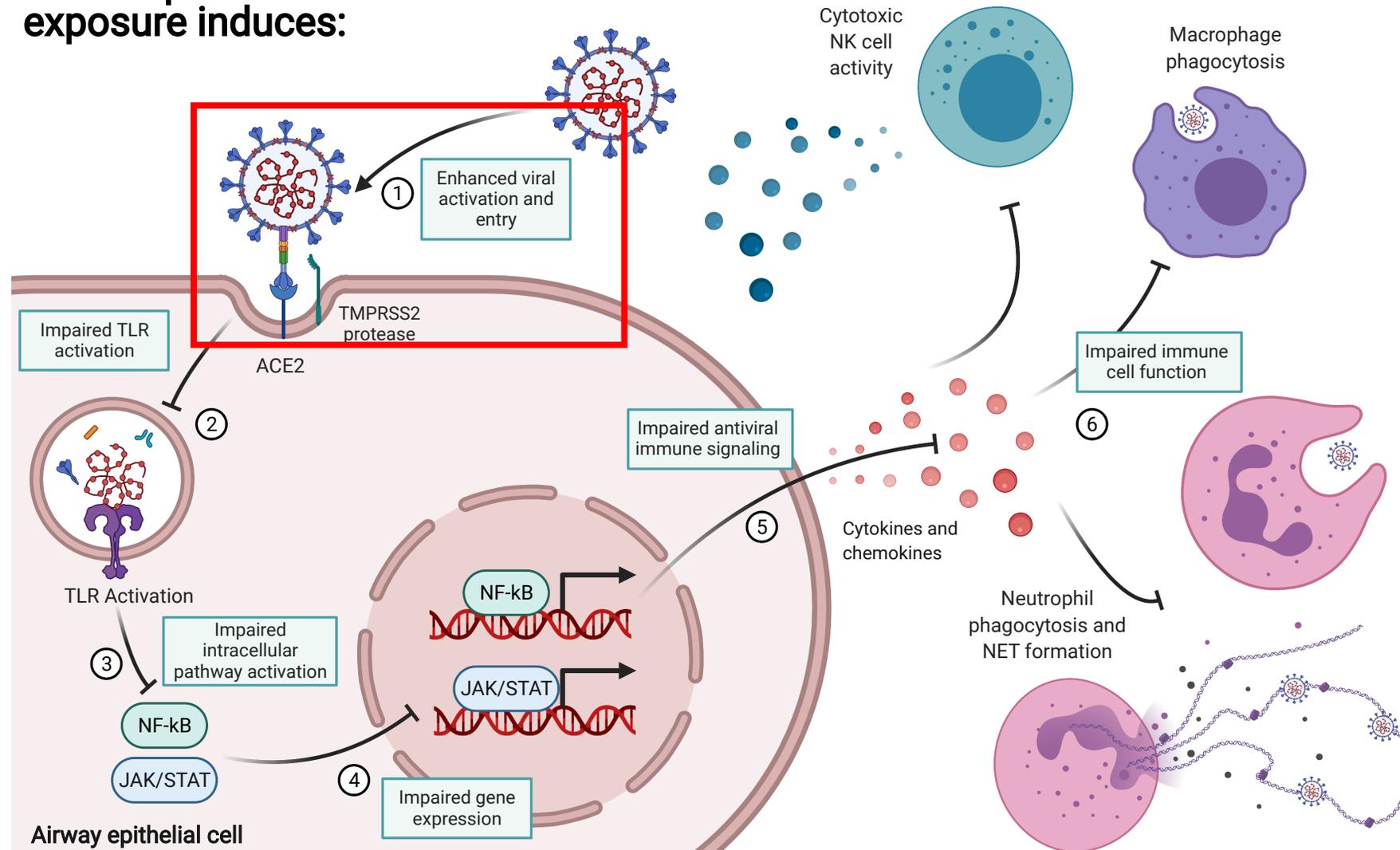


- **Concurrent exposures:**
 - We breathe over 12,000 Liters of air per day – contains both infectious pathogens and air pollutants
- **Target the same tissue**
 - Respiratory epithelium is the primary target for both viral pathogens and inhaled pollutants
- **Rely on similar host defense functions**
 - Innate immune cells are the primary and initial response for pollutants and viral infections

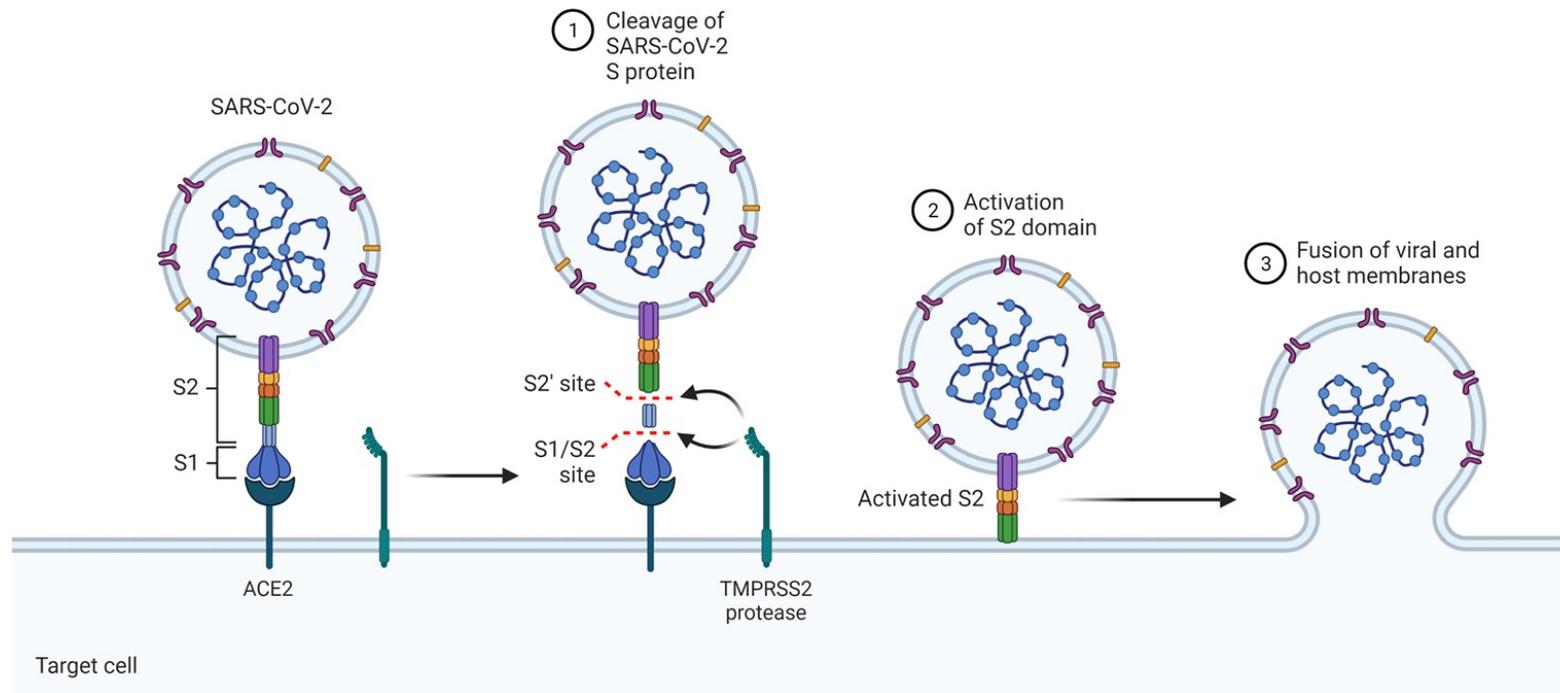
Potential Interactions Between Air Pollutants and SARS-CoV2



Inhaled pollutant exposure induces:



Viral Entry



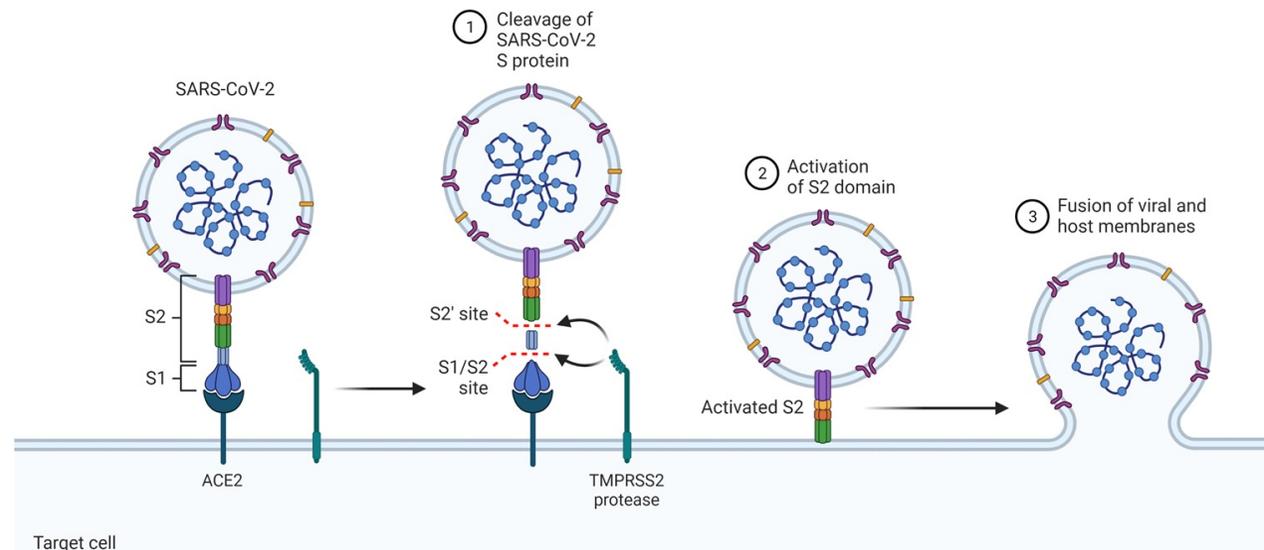
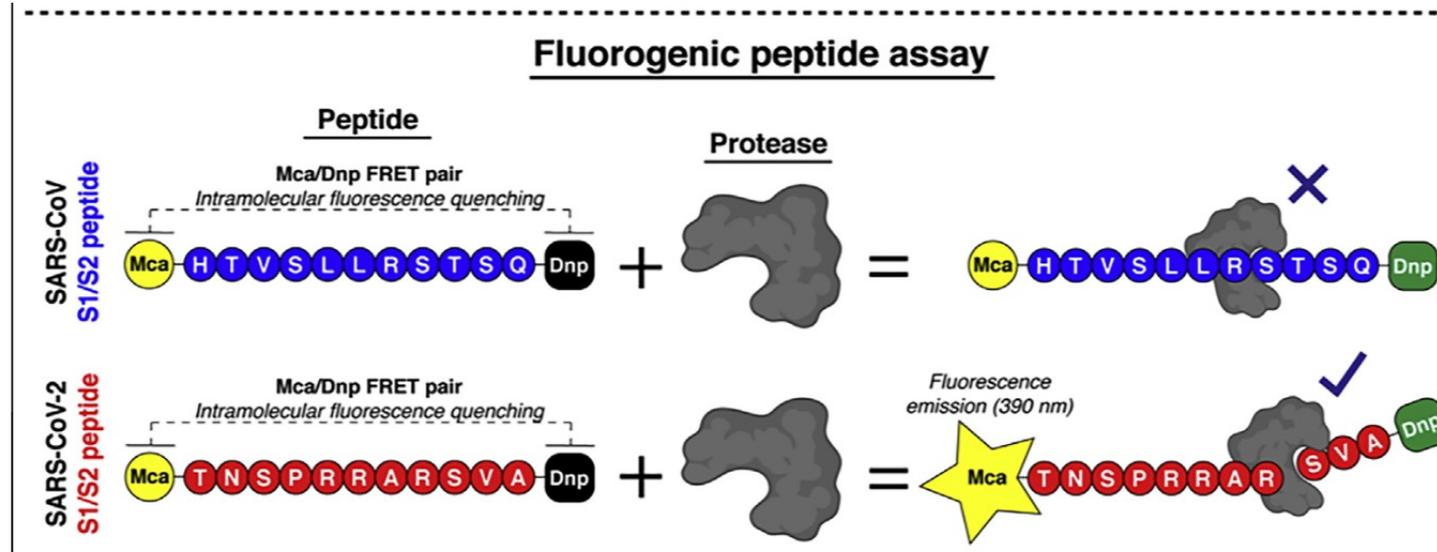
Viral activation for infection requires cleavage of the viral fusion protein

Pollutant-induced Modification of Viral Receptors/Proteolytic Activation

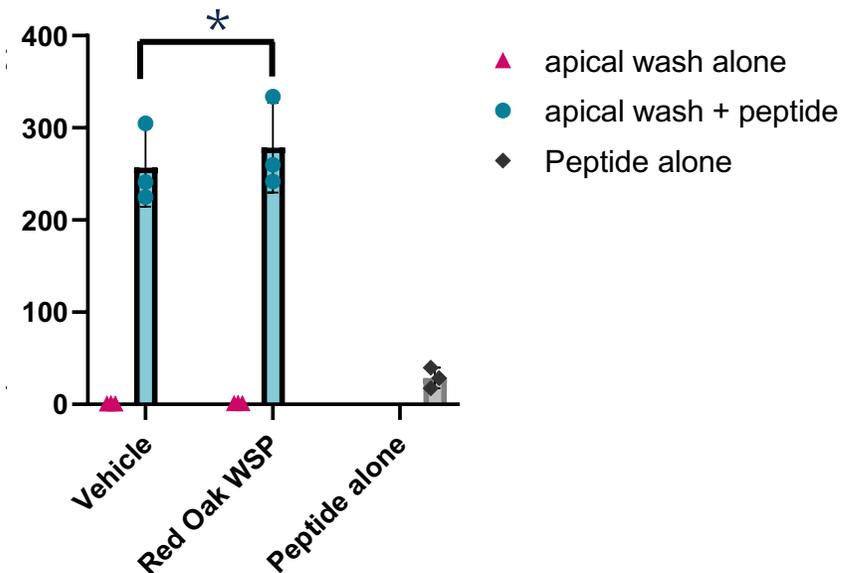
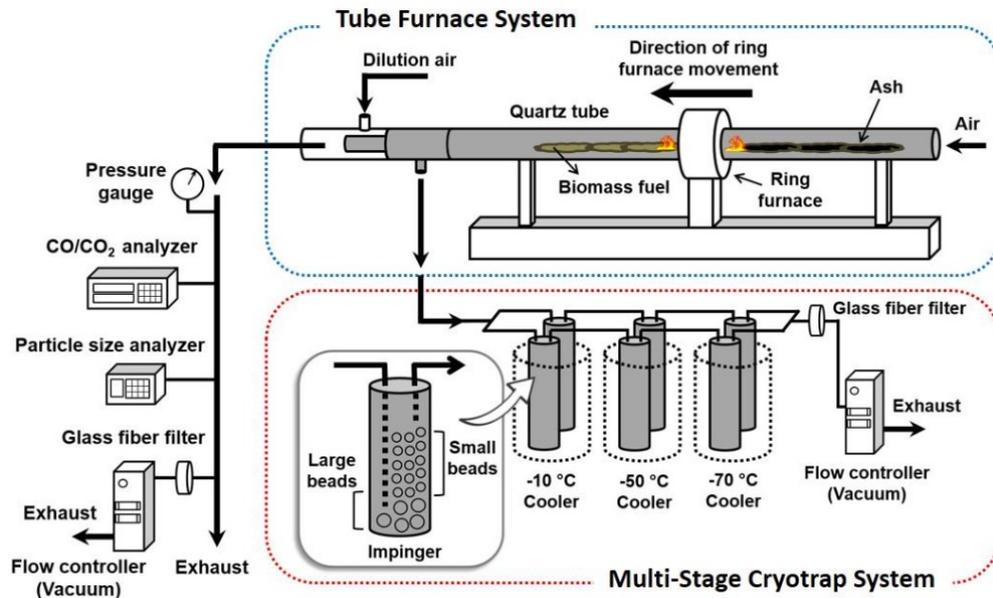
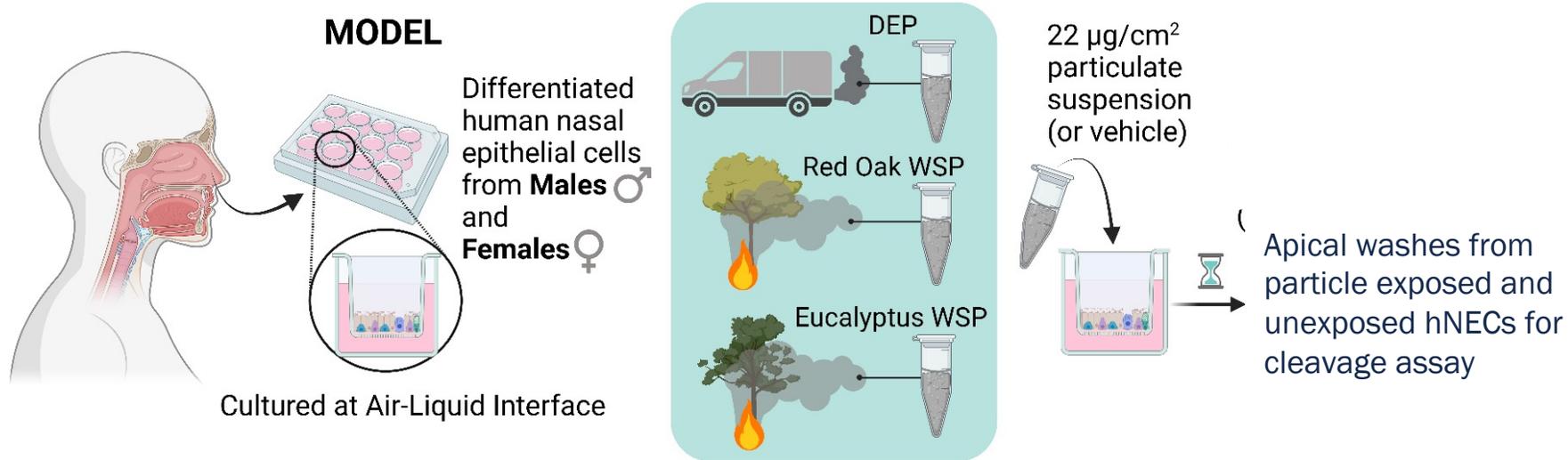


- Proteolytic Cleavage/Activation of SARS-CoV2 requires proteases, such as TMPRSS2, furin, cathepsins, etc.
- Exposure to diesel exhaust increased expression of ACE2 and TMPRSS2 in human pluripotent stem cell-derived alveolar epithelial cells and alveolar organoids (Kim et al., 2020)
- Expression of ACE2 and TMPRSS2 might be regulated by several consensus motifs for binding of the aryl hydrocarbon receptor (AhR), a common pathway activated by ambient air pollutants (Watzky et al., 2020;Lawal 20217)
- Exposure to ozone increases secreted levels of TMPRSS2 and decreases levels of SLPI (antiprotease), which was linked to increased viral entry of influenza virus (Kesic et al., 2012)

Internally Quenched Fluorescent Peptides to Assess Cleavage Activity



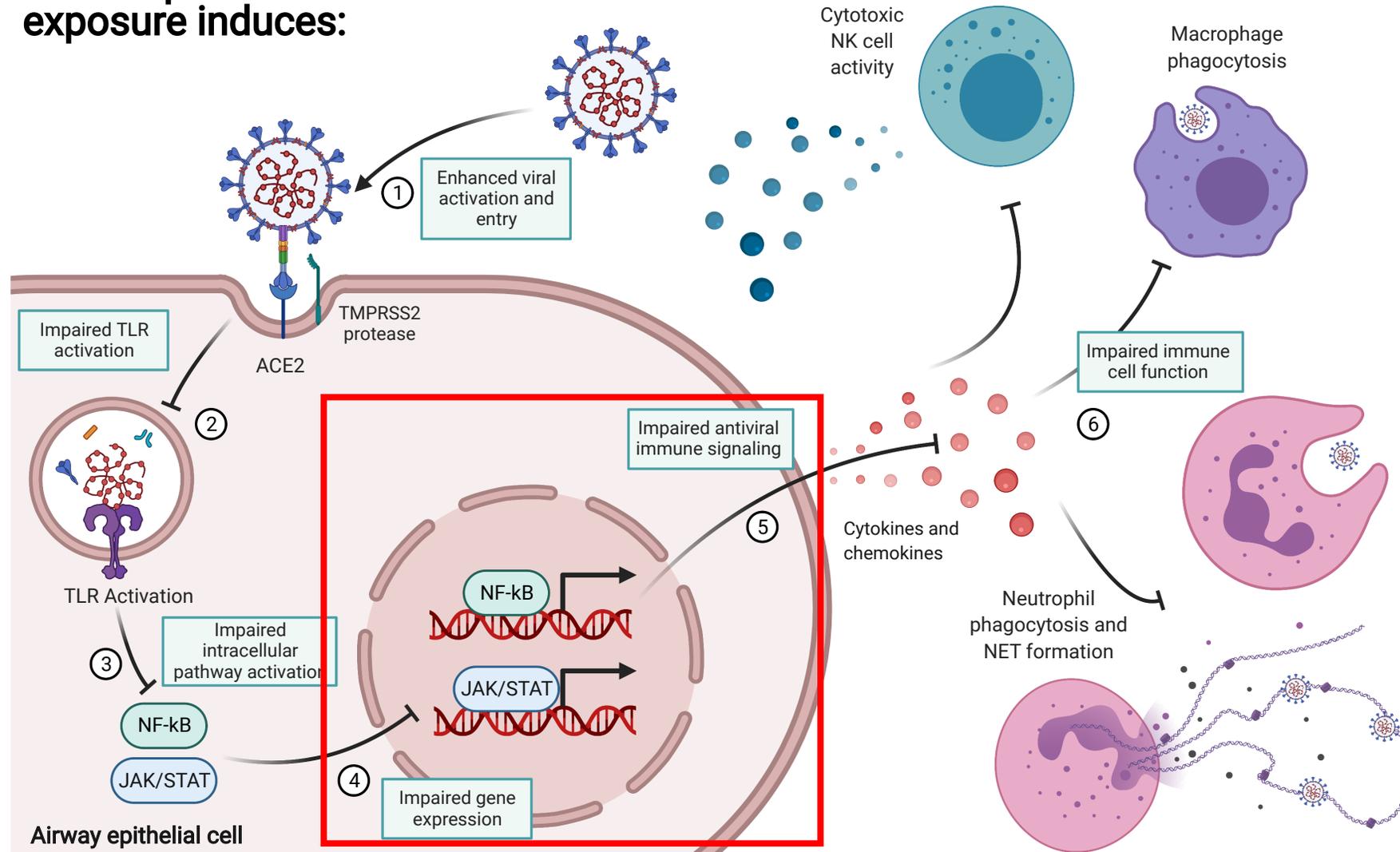
WSP Enhance Cleavage of SARS-CoV2 Peptide



Potential Interactions Between Air Pollutants and SARS-CoV2



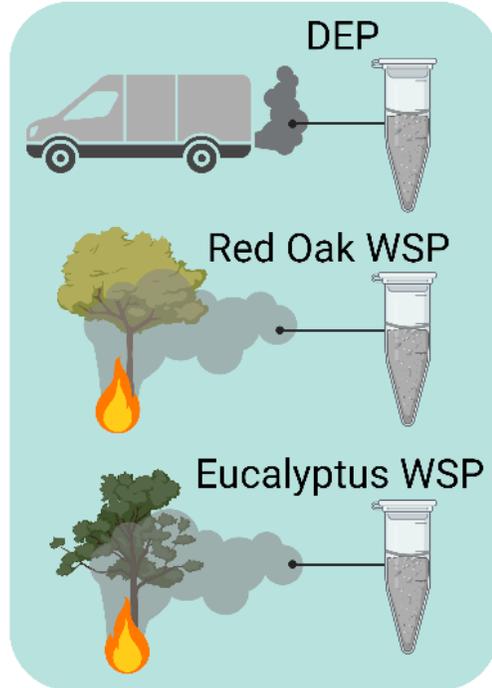
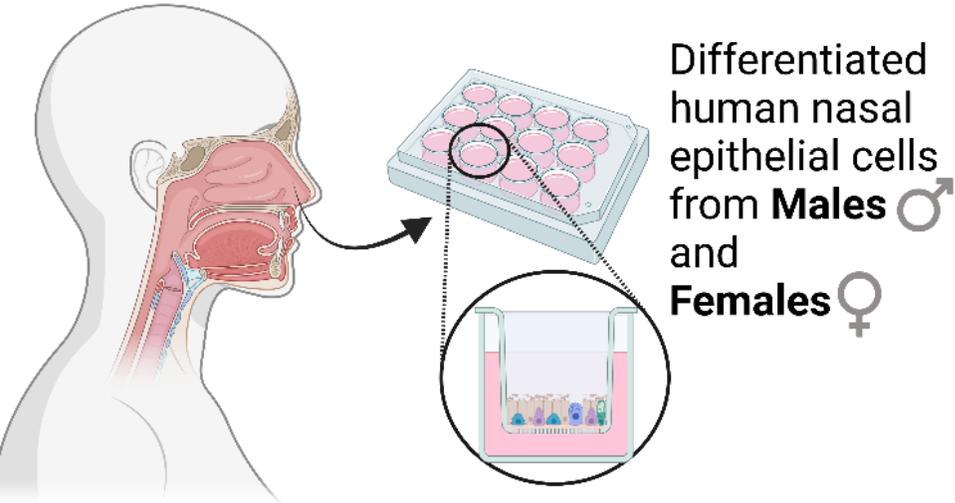
Inhaled pollutant exposure induces:



Experimental Outline

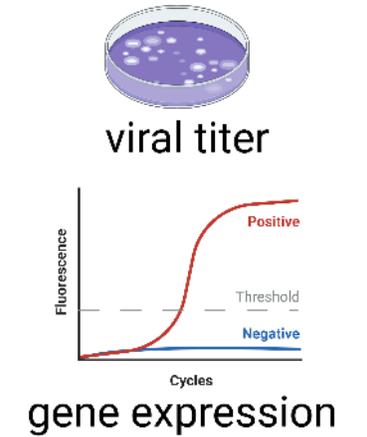
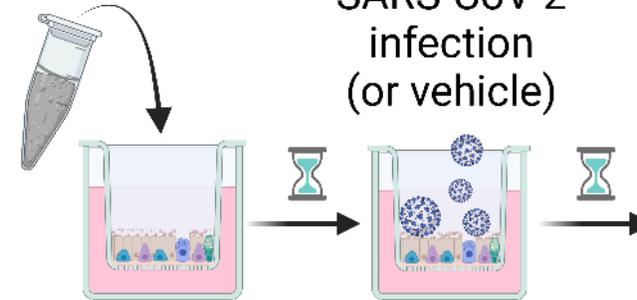


MODEL



22 $\mu\text{g}/\text{cm}^2$
particulate
suspension
(or vehicle)

METHOD



WSP Modify SARS-CoV2 Antiviral Gene Expression in a Sex-dependent Manner



RESULTS

Particulate exposure did not affect viral load

SARS-CoV-2

Red Oak WSP +

CONCLUSIONS



 | Research Article

Woodsmoke particle exposure prior to SARS-CoV-2 infection alters antiviral response gene expression in human nasal epithelial cells in a sex-dependent manner

Stephanie A Brocke, Grant T Billings, Sharon Taft-Benz, Neil E. Alexis, Mark T Heise, and Ilona Jaspers *

02 FEB 2022 // <https://doi.org/10.1152/ajplung.00362.2021>



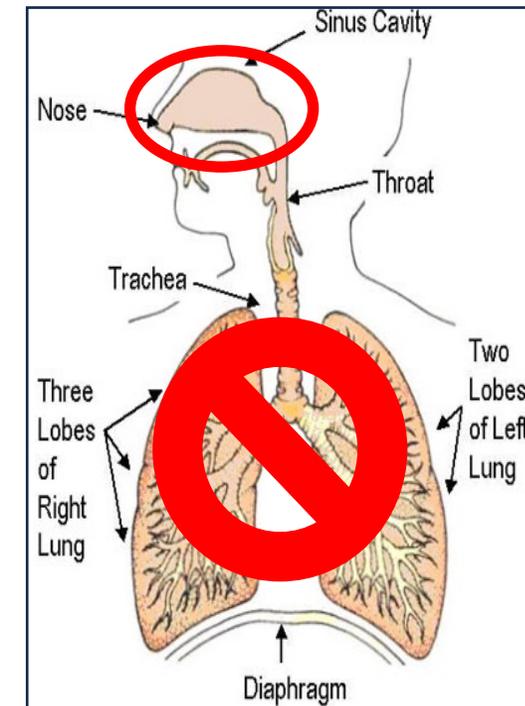


Human *in vivo* studies of Influenza Infections

fluMist
Influenza Virus Vaccine
Live, Intranasal



- **FluMist™ is a cold-adapted Live Attenuated Influenza Virus (LAIV) vaccine**
- **“cold-adapted”, thus replication limited to nasal cavity (32°C)**
- **It generates a replicative but self limited viral infection with innate and immune host defense responses**
- **Provides a safe tool to study influenza virus infections *in vivo***



Model Pollutant Exposures and LAIV - Woodsmoke



- PM derived from Woodsmoke is of increasing public health concern in the US and globally
- Healthy study participants were exposed to either 500 $\mu\text{g}/\text{m}^3$ of wood smoke particulate or air for two hours
- LAIV-induced CXCL10 (a critical IFN-inducible chemokine) levels were suppressed in the nasal mucosa of all participants
- An exposure by sex interaction was observed, with males showing greater inflammation-related gene expression, while in females' host-defense related gene expression was mildly decreased

Wood Smoke Exposure Alters Human Inflammatory Responses to Viral Infection in a Sex-Specific Manner A Randomized, Placebo-controlled Study

Meghan E. Rebuli¹, Adam M. Speen¹, Elizabeth M. Martin^{1,2}, Kezia A. Addo¹, Erica A. Pawlak³, Ellen Glista-Baker³, Carole Robinette³, Haibo Zhou⁴, Terry L. Noah^{1,3,5}, and Ilona Jaspers^{1,3,5}

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Summary



- Both acute and chronic exposure to air pollutants (particulate and gas phase) affect respiratory host defense
- Epidemiological studies suggest a link between pollutant exposure and COVID-19, but more studies are needed to determine whether other co-factors are modifying this effects
- Organotypic *in vitro* models have uncovered plausible mechanisms by which inhaled air pollutants could affect the susceptibility to SARS-CoV2
- Model virus infections combined with controlled acute pollutant exposures have identified pollutant-induced modification of respiratory virus infection; sex is a biological variable that needs to be considered

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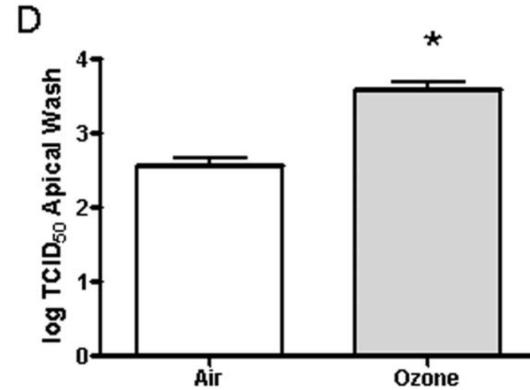
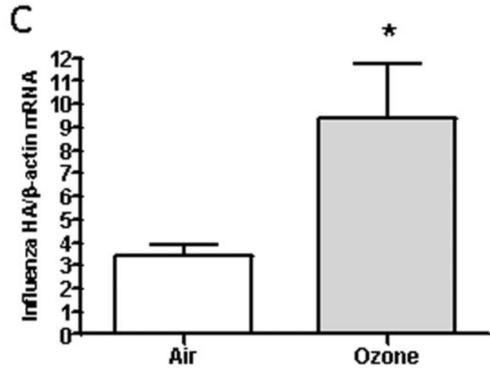
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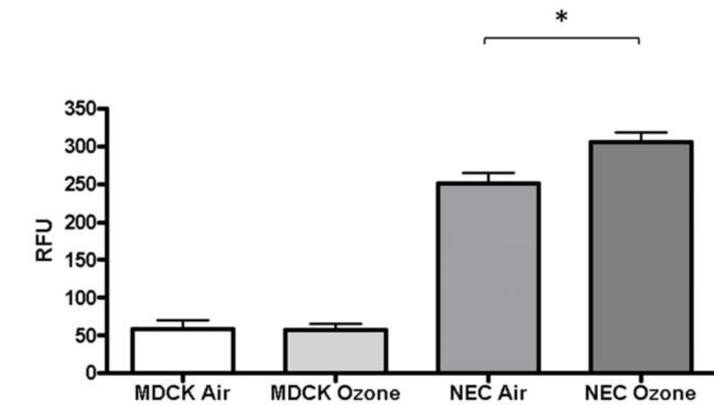
Biorender.com



Ozone Exposure enhances influenza infection by increasing protease expression



Ozone exposure increased Influenza A infection of HNECs



Ozone exposure enhanced influenza virus-like particle entry

B Secreted

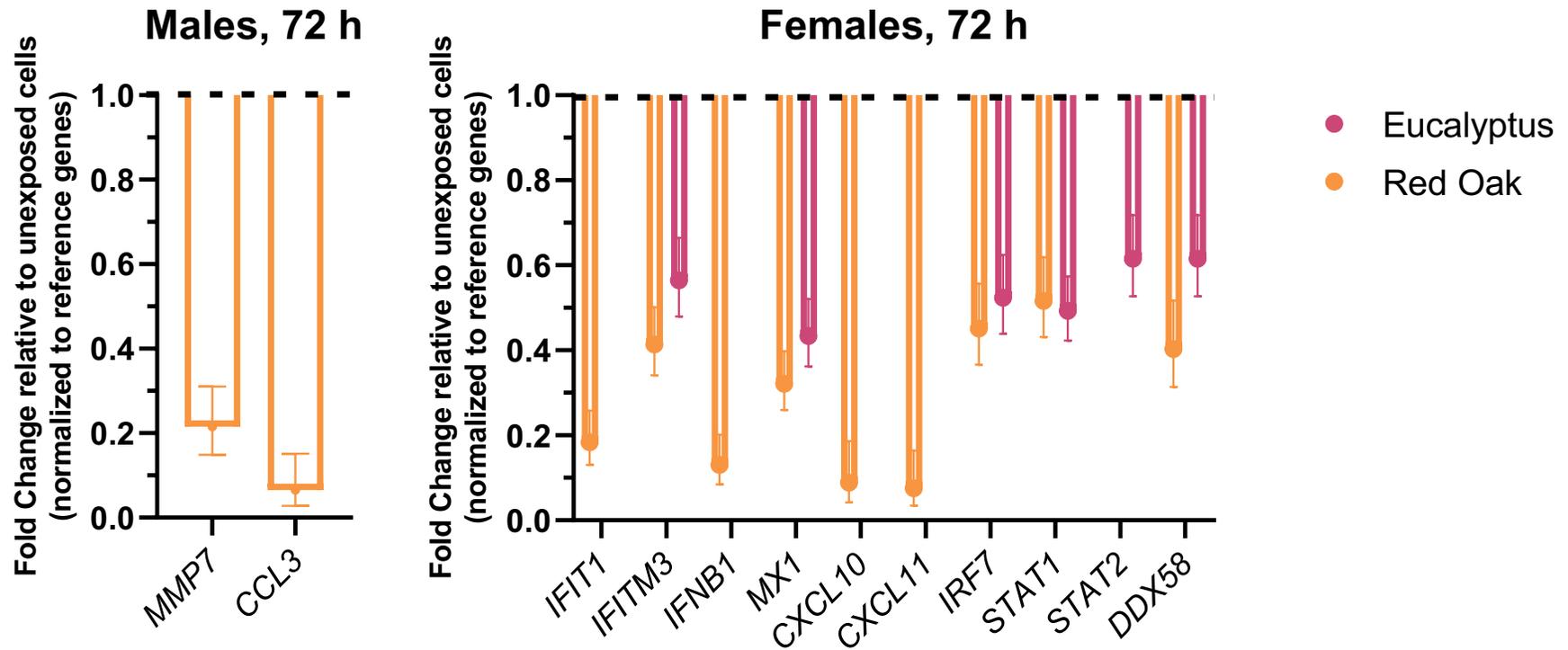


Ozone exposure increased expression of TMPRSS2 and HAT (proteases which activate influenza A) and decreased expression of their antiprotease SLPI

Woodsmoke particle exposure suppresses antiviral gene expression, 72 h p.i.



Fold change of statistically significant changes in gene expression,
Sex disaggregated





Diesel Exhaust Exposure and Nasal Response to Attenuated Influenza in Normal and Allergic Volunteers

Terry L. Noah^{1,2}, Haibo Zhou^{2,3}, Hongtao Zhang³, Katie Horvath⁴, Carole Robinette^{1,2}, Matthew Kesic², Megan Meyer⁵, David Diaz-Sanchez⁶, and Ilona Jaspers^{1,2}

¹Department of Pediatrics, ²Center for Environmental Medicine, Asthma and Lung Biology, ³Department of Biostatistics, ⁴Curriculum in Toxicology, and ⁵Department of Microbiology and Immunology, University of North Carolina at Chapel Hill, Chapel Hill, North Carolina; and ⁶United States Environmental Protection Agency, Chapel Hill, North Carolina

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