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Mechanistic Insights into the Impact of Air Pollution on Pneumococcal Pathogenesis and

Transmission

Daan Beentjes, Rebecca K. Shears, Neil French, Daniel R. Neill, and Aras Kadioglu

Abstract

Streptococcus pneumoniae (the pneumococcus) is the leading cause of pneumonia and bacterial meningitis. A number of recent studies indicate an association between the incidence of pneumococcal disease and exposure to air pollution. Although the epidemiological evidence is substantial, the underlying mechanisms by which the various components of air pollution (particulate matter and gases such as NO₂ and SO₂) can increase susceptibility to pneumococcal infection are less well understood. In this review, we summarize the various effects air pollution components have on pneumococcal pathogenesis and transmission; exposure to air pollution can enhance host susceptibility to pneumococcal colonization by impairing the mucociliary activity of the airway mucosa, reducing the function and production of key antimicrobial peptides, and upregulating an important pneumococcal adherence factor on respiratory epithelial cells. Air pollutant exposure can also impair the phagocytic killing ability of macrophages, permitting increased replication of *S. pneumoniae*. In addition, particulate matter has been shown to activate various extra- and intracellular receptors of airway epithelial cells, which may lead to increased proinflammatory cytokine production. This increases recruitment of innate immune cells, including macrophages and neutrophils. The inflammatory response that ensues may result in significant tissue damage, thereby increasing susceptibility to invasive disease, because it allows *S. pneumoniae* access to the underlying tissues and blood. This review provides an in-depth understanding of the interaction between air pollution and the pneumococcus, which has the potential to aid the development of novel treatments or alternative strategies to prevent disease, especially in areas with high concentrations of air pollution.

Keywords: PM_{2.5}, PM₁₀, particulate matter, pneumonia, pneumococcal infections

Asymptomatic colonization of the nasopharynx is the primary lifestyle of the pneumococcus and the principal reservoir for onward transmission (1, 2). Nasopharyngeal carriage also plays a key role in the development of pneumonia and invasive pneumococcal diseases, such as septicemia and meningitis (3–7). It is estimated that 91% of people worldwide are exposed to air pollution concentrations that exceed World Health Organization guideline limits, and those living in low- to middle-income countries are exposed to the highest concentrations (8). These populations also have the greatest burden of lower respiratory infections (LRIs) (9). Exposure to air pollutants is one of the major risk factors for mortality caused by LRIs, particularly for children under 5, in whom it is second only to childhood wasting (9). A multicenter international study showed that *Streptococcus pneumoniae* was responsible for 50% of LRI fatalities globally in 2016, with more than 1.18 million deaths (9). Exposure to air pollution has been identified as a risk factor for pneumococcal carriage (10–13) and is associated with a higher incidence of pneumococcal pneumonia and invasive pneumococcal disease (14–23). These studies indicate a strong association of air pollution exposure and pneumococcal infections. The underlying mechanisms are not fully elucidated, however, and thus form the basis of this review.

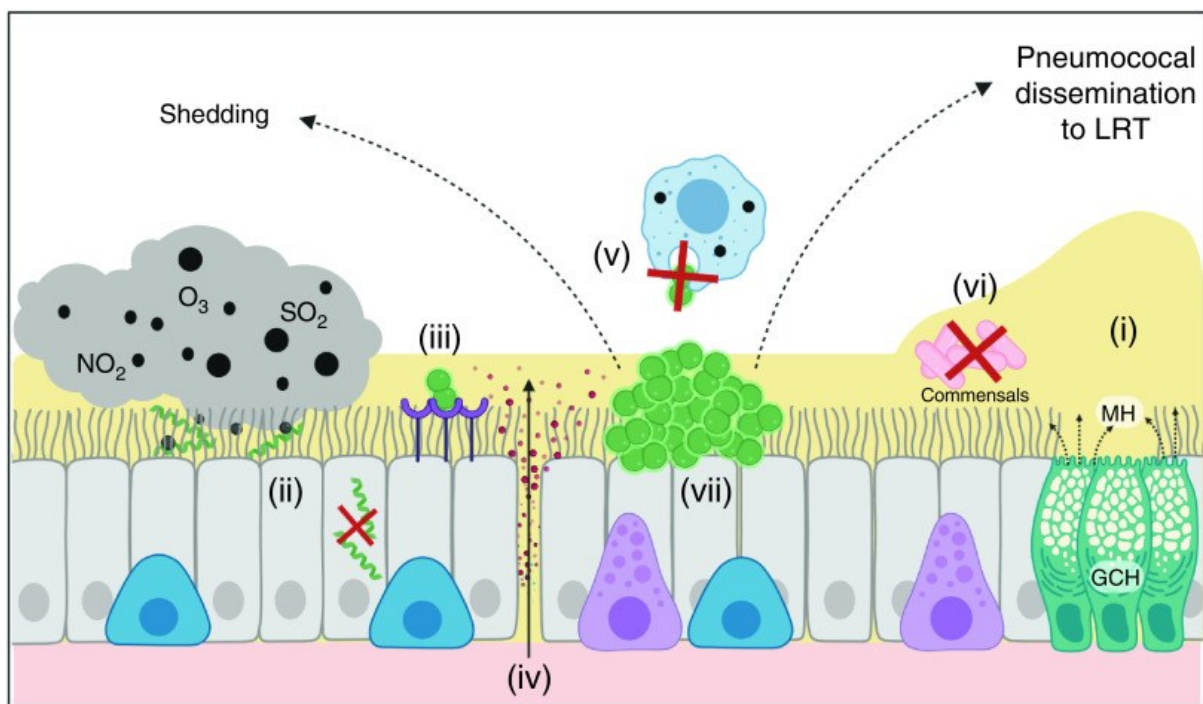
Air pollution typically contains a mixture of particulate matter (PM) and gaseous components such as ozone (O₃), volatile organic compounds, nitrogen oxides, sulfur dioxide (SO₂), and carbon monoxide (24). PM can be split into three categories based on aerodynamic diameter: coarse material (PM_{2.5–10}; ≤10 μm but >2.5 μm), fine PM (PM_{2.5}; ≤2.5 μm), and ultrafine PM (≤0.1 μm) (25).

Particles larger than PM₁₀ are typically retained in the nose and upper respiratory tract; however, both PM₁₀ and PM_{2.5} are able to filter into the lungs (24, 25). This review summarizes the effects that these gaseous and particulate components of air pollution have on the host and how this may contribute to pneumococcal pathogenesis and transmission. We begin by describing the impact of air

pollution on pneumococcal colonization and survival in the respiratory tract, followed by explaining the mechanisms by which air pollution may increase the severity of pneumococcal infections.

Increased Susceptibility to Pneumococcal Colonization

Colonization of the nasopharynx is a prerequisite for the development of pneumococcal disease and plays an important role in transmission (1, 2, 26). In addition, pneumococci that disseminate from the upper to the lower respiratory tract need to find a suitable niche for growth and survival. In this section, we discuss the most vital host responses that protect against pneumococcal colonization in the respiratory tract and how inhaled air pollution particles may aid the pneumococcus in evading these defenses (27–29). [Figure 1](#) shows a summary of the main findings.



[Figure 1.](#)

Air pollution increases susceptibility to pneumococcal colonization. Inhaled air pollution particles, including particulate matter (PM), sand, dust, and ash (black dots) and gases NO_2 , O_3 , and SO_2 (gray cloud) can enhance pneumococcal colonization through various mechanisms. (i) Reduced mucociliary

clearance as a result of impaired ciliary activity, goblet cell hyperplasia, and mucus hypersecretion.

(ii) Inactivation of antimicrobial peptides and attenuated production by epithelial cells. (iii)

Upregulation of an important pneumococcal adhesion factor, platelet-activating factor receptor on

respiratory epithelial cells. (iv) Influx of nutrients into the airway lumen by increasing the

permeability of the epithelial layer. (v) Reduced phagocytotic ability of macrophages. (vi) Microbiota

dysbiosis. (vii) Impaired host responses enabling the pneumococcus to proliferate to higher densities

in the upper respiratory tract, which is associated with increased susceptibility to the development

of disease and enhanced shedding from the host. Increased proinflammatory responses in the

respiratory tract observed after PM exposure can result in higher shedding events. GCH = goblet cell

hyperplasia; LRT = lower respiratory tract; MH = mucus hyperproduction. Created with

BioRender.com.

Decreased Mucociliary Clearance and Improved Adhesion to Airway Epithelium

The mucosal surfaces of the upper and lower respiratory tract are the first host cells that respiratory pathogens encounter. The mucus layer that covers these surfaces is key for trapping debris and

pathogens, including *S. pneumoniae*, and forms an important barrier between the outside world and the host ([30](#), [31](#)). Contained bacteria are cleared by the mucociliary activity of ciliated epithelial cells,

found on various parts of the respiratory tract ([32](#), [33](#)). The importance of a functional mucosal

surface in preventing pneumococcal diseases is highlighted by research on primary ciliary dyskinesia,

a disease that affects ciliary beating ([32](#), [34](#)). These studies show that *S. pneumoniae* is commonly

found in the respiratory tract of patients with primary ciliary dyskinesia. In addition, in chronic

obstructive pulmonary disease, mucociliary defects are common ([35](#)) and associated with a high

incidence of pneumococcal infections ([36](#), [37](#)). Various studies report that air pollution can

contribute to a dysfunctional airway mucosa and could therefore aid pneumococcal colonization. For

example, it has been observed that PM reduces ciliary beating and mucociliary transport ([38](#)).

Decreased ciliary activity was also measured in rats and healthy volunteers after exposure to carbon

monoxide, NO₂, or SO₂ (39–41). In addition, exposure to urban PM can cause mucus hypersecretion and goblet cell hyperplasia (41–43). This has been shown in both respiratory epithelial cell cultures after PM exposure and murine exposure models (41–43). The overproduction of mucus that ensues may cause obstructions in the airways and impair mucociliary clearance, hence benefiting the colonization of various bacterial species (35, 44–47). Moreover, urban PM and desert sand or dust can cause acidification and an increased viscosity of the produced mucus (43, 48, 49), resulting in a more rigid mucus layer that is more difficult to clear from the respiratory tract (47, 50–52).

Montgomery and colleagues measured gene expression in human nasal epithelial cell cultures exposed to an organic extract of PM_{2.5} and showed an upregulation of several genes involved in mucus secretion (53). The authors also observed significantly decreased expression of *FOXJ1* and *MCIDAS* (53), two transcription factors that play a vital role in cilia formation (54). This altered gene expression in epithelial cells may provide an underlying mechanism behind the impaired mucociliary function that is observed after exposure to air pollution.

In addition to causing a dysfunctional airway mucosa, exposure to urban PM or toxic gases, such as welding fumes, can enhance pneumococcal adherence to epithelial cells (27, 55). The main mechanism behind this is an upregulation of the platelet-activating factor receptor (PAFR)—a key pneumococcal adherence factor—on respiratory epithelial cells that was observed after exposure to urban PM or welding fumes (27, 55). The impact of increased PAFR expression on pneumococcal infection is highlighted in a study on electronic cigarette (E-cigarette) vapor (56). Here compared with unexposed mice, mice exposed to nicotine-containing E-cigarette vapor showed increased PAFR expression on nasopharyngeal epithelial cells and higher pneumococcal densities in the nasopharynx (56). This study indicates that oxidative stress responses that are activated after exposure to E-cigarette vapor contribute to the upregulation of PAFR. Therefore, the authors suggest that redox-promoting metals (e.g., copper) present in vapor could play an important role in the upregulation of PAFR (56). However, further research is required to determine the impact of each individual component of E-cigarette vapor.

A number of studies also suggest a role of PAFR in the development of pneumococcal disease (57, 58). For example, Rijneveld and colleagues showed that PAFR-deficient (PAFR^{-/-}) mice have lower pneumococcal densities in their lungs and a higher survival rate after induction of pneumococcal pneumonia than a wild-type control group (57). Others observed decreased dissemination of *S. pneumoniae* from the lungs to the blood and the central nervous system in PAFR^{-/-} mice compared with wild-type mice (58). These studies could provide an explanation for why welders have an increased susceptibility to invasive pneumococcal disease (59), because daily exposure to welding fumes may lead to an increased PAFR expression on epithelial cells (55).

Impaired Antimicrobial Responses

In addition to being a physical barrier, the respiratory mucosa also contains antimicrobial peptides and proteins, which contribute to the eradication of bacteria (60, 61). Antimicrobial compounds found in the respiratory tract and important in combating *S. pneumoniae* infections include (apo)lactoferrin, lysozymes, and the cationic antimicrobial peptides (CAMPs) such as LL37 and β -defensins (62–66). Several studies show that air pollution could affect the expression and bactericidal properties of these compounds, which may reduce their effectiveness against respiratory pathogens. For example, carbon black nanoparticles—an example of ultrafine PM—can induce structural changes in LL37, thereby impairing its antibacterial activity (67). Airway acidification by urban PM and desert dust, described earlier in this review, can affect the activity of CAMPs, such as β -defensins and LL37, because their activity is pH dependent (68). A different mechanism of action is observed in a study on coal fly ash. This ambient PM is negatively charged and can neutralize the electropositive antimicrobials and therefore reduce their activity (69). In addition, epithelial cells show a decreased expression of β -defensin after exposure to PM (70–72). A recent study found that the suppression of CAMPs by corticosteroids led to increased bacterial counts in the lungs of mice challenged with a pneumonia-inducing dose of *S. pneumoniae* (73). In addition, low β -defensin 2 plasma concentrations in patients with pneumonia, measured upon hospital admission, were associated

with 30-day adverse outcomes (74). These studies highlight the importance of sufficient CAMP concentrations in the context of pneumococcal infections, and air pollution could therefore play a substantial role in the progression of disease. However, further detailed investigations into the effects of air pollution and the production of CAMPs is necessary because Nam and colleagues found an enhanced β -defensin response in PM-exposed alveolar epithelial cells (75).

Dysbiosis of the Airway Microbiota

Within the airway mucosa, *S. pneumoniae* must compete for adhesion sites and nutrients with other bacterial pathogens and commensals. These residents (i.e., the airway microbiota) are able to sequester carbon sources (76) and produce a variety of antimicrobial peptides (77), which provides some protection against intruding pathogens (78). The microbiota of the respiratory tract is a highly dynamic community (79), and exposure to PM, ozone, NO₂, or SO₂ has been shown to cause a shift in its microbial composition (80–83). Although the specific impact of air pollution is understudied and discrepancies exist in the literature, exposure to air pollution generally reduces the abundance of commensals, including the genera *Flavobacterium*, *Prevotella*, *Fusobacterium*, and *Veillonella*, whereas pathogenic species, including streptococci, are more commonly found (80–84). A recent review looked into the role of air pollution in (airway) microbiota dysbiosis and proposed that an impaired immune response after air pollution exposure is one of the underlying mechanisms behind it (85). Li and colleagues also observed higher bacterial diversity (Chao 1 and phylogenetic diversity whole tree index) in the lungs of rats exposed to ambient PM (i.e., biomass fuel and motor vehicle exhaust) (81). According to an experimental human challenge study, volunteers with a more diverse nasopharyngeal microbiota (i.e., higher Shannon index) showed an increased susceptibility to pneumococcal acquisition, which could suggest that ambient PM may increase host susceptibility (86). Further investigation into the effect of air pollution on microbiota diversity is needed, however, because other studies observed no or opposite effects of air pollution (recently reviewed by Xue and colleagues [84]).

In addition, *S. pneumoniae* and other pathogenic species have been found attached to PM (87, 88). The diversity and abundance of bacterial species on PM depends on geographical location and air quality index, and they vary seasonally (88–90). Although mucociliary clearance will likely prevent the colonization of most PM-attached bacteria, the inhalation of air pollution particles may introduce pathogenic species into the respiratory tract, which could cause dysbiosis of the airway microbiota. Liu and colleagues observed that PM size has an impact on the diversity and abundance of bacterial species attached to PM (89). The authors measured a higher abundance of pathogenic bacterial species on PM_{2.5} than on PM₁₀, which may suggest that the risk of introducing pathogenic species in the airways is relatively higher after inhalation of PM_{2.5}. Given that PM_{2.5} can penetrate more deeply into the lungs than PM₁₀, PM size could also affect the location where pathogenic species are deposited within the airways (91). In addition, Shears and colleagues suggested that adherence of *S. pneumoniae* to PM causes upper airway colonized pneumococci to be dragged down from the nasopharynx to the lungs (92). Attachment of pneumococci to PM could therefore increase susceptibility to pneumonia (92).

Improved Pneumococcal Survival

A substantial problem for *S. pneumoniae* to overcome is the limited nutrient availability within the respiratory tract. Restricting nutrient availability is an important defense mechanism for the host against pathogens (93, 94), and *S. pneumoniae* must adapt to find new ways of acquiring suitable carbon sources (95). PM, such as diesel exhaust particles (DEPs), can aid nutrient acquisition for bacterial pathogens within the respiratory tract by increasing the permeability of the airway epithelial barrier (96, 97), causing an influx of essential nutrients such as glucose into the airway lumen (93). An impaired epithelial barrier is also observed in epithelial cell cultures and *in vivo* after exposure to NO₂ or ozone, respectively (98, 99). In addition, it has been observed that DEPs can serve as a carbon source for *S. pneumoniae* under nutrient-restricted conditions (92).

A recent study also described a direct effect of air pollution on pneumococcal phenotype. The authors observed improved pneumococcal competence after exposure to environmental dust (100). Competence is a tool used by *S. pneumoniae* to acquire new DNA through transformation, which is believed to be the main driver of pneumococcal evolution. New traits acquired through the addition of fresh genetic material aids adaptation to new environments or competition with other bacteria for nutrients and adhesion sites (101). Inhaled dust particles may therefore enhance pneumococcal colonization and survival in the airway mucosa.

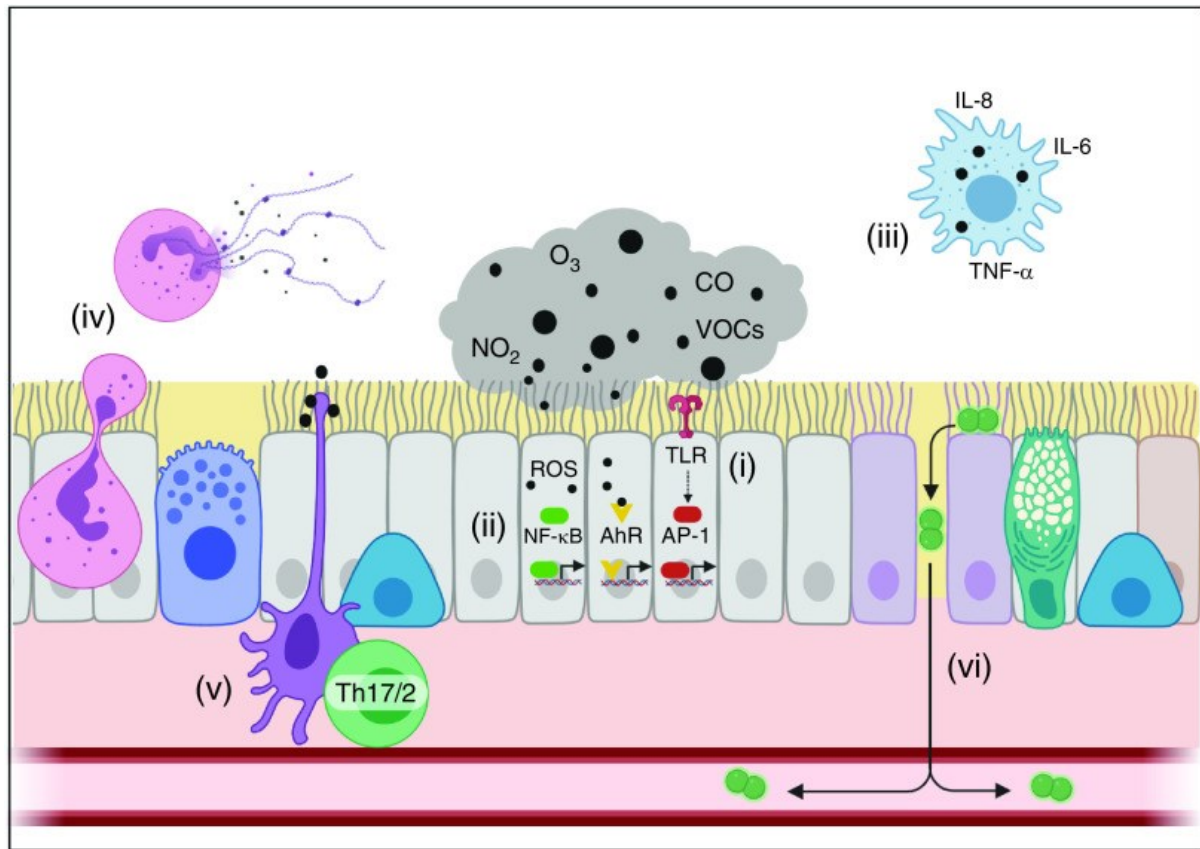
Reduction in Macrophage Phagocytic Killing

One of the key roles of alveolar macrophages is the removal of pathogens, dead host cells, and debris (including PM) from the lungs (102). Rylance and colleagues observed carbon loading of macrophages isolated by BAL from Malawian adults using wood-burning stoves (103), whereas Shears and colleagues reported carbon loading of alveolar macrophages isolated from DEP-exposed mice (92). An observational study performed by Kulkarni and colleagues found a significant correlation between the amount of PM that children were exposed to on a daily basis and PM uptake by alveolar macrophages isolated by BAL, as well as an inverse relationship between carbon loading of macrophages and lung function (as measured by FEV₁ and FVC) (104). Reduced phagocytic killing function of PM congested macrophages is also well documented (21, 92, 103–105). PM loading of macrophages is associated with reduced phagocytic killing ability in a number of different PM types, including black carbon generated from wood burning in Malawi, concentrated ambient particles collected in Boston, DEPs, and desert sand, and appears to affect all types of macrophages, including monocyte-derived macrophages (MDMs), human sputum-derived macrophages, and murine J774.2 cells (a macrophage cell line) (21, 92, 103, 105). This defect in phagocytic killing appears to be a result of reduced internalization of bacteria rather than impaired intracellular killing (104, 106), although a lower oxidative burst capacity has been reported for highly congested MDMs (103). Shears and colleagues reported that the alveolar macrophages of mice exposed for a total of 3 days

to DEPs were still loaded with PM 2 weeks later (92). These data suggest that clearance of PM by airway macrophages is a slow process and that the impaired phagocytic killing of PM-loaded macrophages may be long-lasting (92).

Exacerbation of Proinflammatory Responses

Innate and adaptive immune responses are required to control bacterial growth in the airways. These processes begin with the production of proinflammatory cytokines and the recruitment of immune cells to the site of infection. However, an overactivated immune response may lead to tissue damage, thereby enhancing the dissemination of pathogens such as *S. pneumoniae* to underlying tissues and blood, a key step in the development of invasive pneumococcal disease (21, 92). Because air pollution exposure is associated with increased incidence of pneumococcal disease (14–22), we summarize the impact that air pollution has on the immune responses that play a key role in controlling pneumococcal infections. The main findings are summarized in [Figure 2](#).



[Figure 2.](#)

Exacerbations of pneumococcal infections. The gaseous (gray cloud) and particulate matter components (black dots) of air pollution can reach the lower respiratory tract and enhance proinflammatory responses through various mechanisms. (i) Stimulation of Toll-like receptors (TLRs) on the surface of airway epithelial cells and intracellular aryl hydrocarbon receptor (AhR). (ii) Increased reactive oxygen species (ROS) production that indirectly activates proinflammatory signaling pathways. (iii) Increased proinflammatory cytokine production by macrophages. (iv) Enhanced neutrophil recruitment and NETosis. (v) Improved dendritic cell activation and T-cell priming (with T helper type 17/2 [Th17/Th2] responses favored). (vi) Epithelial damage as a result of uncontrolled immune responses could lead to increased dissemination of *S. pneumoniae* to the underlying tissues and systemic circulation. AP-1 = activation protein-1; NF-κB = nuclear factor-κB; TNF-α = tumor necrosis factor-α; VOC = volatile organic compound. Created with BioRender.com.

Activation of Proinflammatory Pathways through Toll-Like Receptors, Aryl Hydrocarbon Receptor

Signaling, and Oxidative Stress

Stimulation of Toll-like receptors (TLRs), reactive oxygen species (ROS), and polycyclic aromatic hydrocarbon (PAH)-sensing pathways by components of air pollution can activate proinflammatory signaling, including through mitogen-activated protein kinase and nuclear factor- κ B (NF- κ B), leading to downstream inflammation (24, 107–109). PM containing LPSs and/or fungal spores is likely to be more immunogenic, because these are natural TLR ligands (110–112). PAHs found within PM, such as DEPs, are able to diffuse through cell membranes, where the PAHs are sensed by the aryl hydrocarbon receptor (AhR) (24, 113). Examples of PAHs that are abundant in air pollution include phenanthrene and pyrene (114–116). Binding of PAHs to the AhR allows the AhR to translocate to the nucleus, which leads to increased CYP1a1 (a cytochrome P450 family 1 member) expression and further generation of cytotoxic compounds (117). In fact, PAHs have been described as the most genotoxic components of air pollution (118). Others report that lipophilic components of DEPs are able to stimulate the AhR in endothelial cells, leading to increased IL-1 α , IL-1 β , COX2, and matrix metalloprotein 1 gene expression (119).

Induction of oxidative stress is thought to be due to both the heavy metal (e.g., copper) and organic components of PM (24). Kim and colleagues showed that mice have increased inflammatory responses and cell injury in the lungs after inhalation of copper particles (120). These copper-exposed mice also showed decreased bacterial clearance in the lungs when infected with *Klebsiella pneumoniae*, which highlights the negative impact of heavy metal exposure in the context of bacterial infections. Other metals that can be found attached to PM, such as iron, can catalyze the production of highly reactive hydroxyl ions from H₂O₂, which can react very quickly with proteins, DNA, and other biomolecules, resulting in tissue damage (113). Metals such as cobalt, chromium, and vanadium can undergo redox cycling, whereas cadmium, lead, mercury, and nickel deplete glutathione (an antioxidant) and protein-bound thiol groups, leading to ROS generation (113). These

metals can be bound to the surface of PM, particularly PM_{2.5}, which has a larger surface area than the equivalent amount of PM₁₀ (121). NF-κB, Nrf2, and activation protein-1 (AP-1) pathways are all regulated by redox status. These pathways are involved in the transcription of a wide range of genes, including those involved in oxidative stress and inflammation (113). The nuclear redox status can affect histone acetylation, making certain regions of the genome more accessible to transcription factors, including NF-κB, a key transcription factor controlling expression of a wide range of proinflammatory cytokines (24, 113). There is also some evidence to suggest that ROS can interact with Ca²⁺ channels, which may also impact NF-κB signaling, because Ca²⁺ can act as a second messenger in this pathway (24, 122). Nrf2 controls a range of antioxidant genes and also interacts with NF-κB and mitogen-activated protein kinase pathway members (24). Expression of Nrf2 and its upstream regulators is decreased after PM exposure, which may contribute to the increased oxidative stress and altered immune responses observed (123). Induction of AP-1 signaling is required for transcription of γ-glutamyl cysteine synthetase, an enzyme involved in antioxidant glutathione synthesis (113).

In a human experimental study involving weekly 4-hour indoor exposures of 13 volunteers to wood smoke, the authors reported detection of the ROS species malondialdehyde in the breath of volunteers, demonstrating that exposure to wood smoke in a real-world setting can induce oxidative stress in the lungs (124). A recent *in vitro* study demonstrated that exposure of human small airway epithelial cells to biomass smoke activates the AP-1 and AhR sensing pathways, but not NF-κB, resulting in short-term increases in IL-8 and granulocyte-macrophage colony-stimulating factor (125).

In addition to activating the proinflammatory pathways described above, ROS can damage proteins and DNA within the cell, which activates an intracellular damage-associated molecular pattern sensing apparatus such as the NLRP3 (NLR family pyrin domain containing 3) inflammasome (112, 126). Hirota and colleagues demonstrated that PM₁₀-induced NLRP3 activation results in IL-1β and CCL20 production, dendritic cell (DC) activation, and lung neutrophilia (126). Elevated CO₂ in air

pollution may also trigger NLRP3 activation and IL-1 β production by neutrophils (127). In addition, damage to lung tissue caused by oxidative stress attracts inflammatory cells from the circulation, which may further exacerbate the damage (128). Park and colleagues generated several forms of anthropogenic PM under controlled laboratory conditions and investigated the ability of these forms of PM to cause DNA damage and oxidative stress responses using *in vitro* exposure models (129). The authors showed that the toxicity differs between PM derived from different sources and that diesel engine exhaust particles are the most toxic compared with other forms of anthropogenic PM.

Increased Cytokine Responses by Macrophages

PM exposure has been shown to increase proinflammatory chemokine and cytokine production by macrophages (92, 103). Rylance and colleagues reported an increase in IL-6 and IL-8 from MDMs exposed to black carbon, demonstrating that black carbon itself has inflammatory properties (103). This is supported by the observation of elevated serum IL-6, IL-8, and MCP-1 in firefighters after smoke exposure (130). These inflammatory properties may be due partly to microbial components that are absorbed onto the surface of PM (110–112). However, Shears and colleagues found that bone marrow–derived macrophages exposed to endotoxin-depleted DEPs produced higher concentrations of tumor necrosis factor- α and IL-6 when stimulated with pneumococcal antigens compared with sham-exposed bone marrow–derived macrophages also stimulated with pneumococci (92). These findings suggest that the PM itself (without interference from endotoxic material that may be adsorbed on the surface) promotes inflammation in the context of microbial challenge. Two previous studies found that *in vitro* coculture of PM-exposed epithelial cells and macrophages resulted in greater production of proinflammatory cytokines than either cell type in isolation, suggesting that there may be an additive effect (131, 132).

Neutrophilia and Neutrophil Extracellular Trap Formation

Increased IL-8 in the lung and circulation after exposure to air pollution is a common finding across *in vitro*, murine, and human exposure models (103, 125, 130, 133). IL-8 is a key neutrophil

chemoattractant, and although early influx of neutrophils is critical for controlling bacterial infections in the lung, it is likely that PM-induced neutrophilia, in the absence of infection, might be detrimental, because uncontrolled neutrophil influx results in further release of proinflammatory cytokines, recruitment of additional immune cells, and tissue pathology (134). However, a recent human exposure model found that short-term exposure of healthy nonsmokers to diesel exhaust leads to increased lymphocytic but not neutrophilic inflammation (135). Instead, the authors reported increased neutrophil extracellular trap (NET) formation in the lungs, which may contribute to the pathophysiology of diesel exhaust (135). Another study found that O₃ increases lung neutrophilia and NET production, demonstrating that different components of air pollution may have different effects on neutrophil recruitment and function (136). NETs are composed of DNA, histones, proteases such as elastase, and antimicrobial molecules including myeloperoxidase (134). NETs are an important part of the neutrophil's antimicrobial armor; however, they can also cause significant collateral damage to respiratory tissues (134). Neutrophil-derived NAD⁺ reduced oxidase and myeloperoxidase can generate superoxide anions and hypochlorous acid, which, coupled with transition metals absorbed onto the surface of PM, could catalyze further redox cycling, causing more oxidative stress and contributing air pollution–induced lung damage (24, 134).

Effects on Adaptive Immunity

Because exposure to air pollution drives inflammation, with influx of neutrophils, macrophages, eosinophils, and lymphocytes, and increased proinflammatory cytokine production by these cells, it is perhaps not surprising that there are alterations in the adaptive immune response, too. Matthews and colleagues reported an increase in DC maturation and T cell priming when DCs were exposed to DEPs or in response to DEP-exposed epithelial cells (137). In particular, the authors reported an increase in IFN- γ ⁺ IL-17A⁺ CD4 T cells, as well as increased IFN- γ ⁺ CD8 T cells (138, 139). In contrast, an older report found that exposure of human T cells to DEPs can affect intracellular T cell signaling, leading to reduced T-bet (the key T helper type 1 [Th1] transcription factor) and Txk (a tyrosine

kinase that controls the transcription of IFN- γ expression (140). This may cause Th2 polarization (24, 140), resulting in an increased Th2-induced cytokine response. Several studies show that increased or chronic Th2-induced inflammation in the respiratory tract is associated with an impaired immunity to pneumococcal infections (141, 142). Altered T cell signaling after DEP exposure may therefore increase the risk of pneumococcal infections.

In addition to its role in the production of proinflammatory cytokines, described above, the AhR can act as a regulator of T regulatory (Treg) and Th17 cell differentiation at epithelial sites (143). A recent report demonstrated that PAHs in urban dust promote Th17 differentiation of T cells and lead to increased IFN- γ ⁺ DCs through activating the AhR (144). A balance between Th17 and Treg responses is critical for controlling *S. pneumoniae* infections, where memory Th17 cells are required for pathogen clearance (145), whereas Treg cells are key for controlling inflammation and minimizing tissue damage (146). O'Driscoll and colleagues showed that not all PAHs drive AhR-induced Th17 responses to the same degree (147). The authors found that of the five sources of PM tested, barley smoke promoted the strongest Th17 response *in vitro*, followed by urban dust and fine and ultrafine DEPs, whereas pine wood smoke was the least potent inducer of IL-17 (147). In another report, DEPs were shown to stimulate production of the alarmins IL-33, IL-25, and thymic stromal lymphopoietin by primary bronchial epithelial cells through interactions with the AhR (148). These cytokines trigger downstream activation of Th2 immunity (149), which, as mentioned above, could increase the susceptibility to pneumococcal infections (141, 142).

Summary: Implications for Transmission and Development of Pneumococcal Diseases

Collectively, published research demonstrates that various forms of air pollution can impair the mucociliary function of the airway mucosa, disrupt innate immune responses, and cause microbiota dysbiosis. On the basis of ability to impair these primary defense mechanisms of the host (32, 34, 62, 150, 151), air pollution increases host susceptibility to pneumococcal colonization, which

would in turn increase transmission ([Figure 1](#)). In addition, enhanced colonization of the upper respiratory tract could increase the risk of LRI.

Air pollution can result in higher nutrient concentrations in the airway lumen and reduce innate immune responses by attenuating the activity of antimicrobial peptides and impairing the phagocytotic ability of macrophages. As a result, inhaled air pollution particles may enable *S. pneumoniae* to replicate to higher densities in the respiratory tract. A higher pneumococcal count in the nasopharynx is associated with a higher incidence of pneumococcal pneumonia ([152](#), [153](#)) and is a driving force behind pneumococcal shedding from a host ([154](#)). Air pollution may therefore play a role in transmission and in the development of pneumococcal disease ([Figure 1](#)). This idea is supported by the study of Jusot and colleagues ([21](#)), in which significantly higher densities of pneumococci were found in the nasopharynx of mice exposed to dust. These dust-exposed mice also showed a higher susceptibility to develop pneumonia than an unexposed control group. In addition to the mechanisms described above, Shears and colleagues described an additional role of PM in the development of pneumonia ([92](#)). The authors showed that *S. pneumoniae* can directly attach to DEPs, which suggests that inhalation of DEPs can act as a physical conduit, dragging down pneumococci from the nasopharynx to the lungs.

Various components of air pollution can cause oxidative stress responses and activate multiple inflammatory pathways in epithelial cells. This may lead to increased production of proinflammatory cytokines and recruitment of innate immune cells, such as macrophages and neutrophils. When exposed to air pollution, these macrophages and neutrophils show an increased production of inflammatory cytokines and NETs, respectively, which may further enhance inflammation of the respiratory tract. Increased inflammatory responses are likely to cause significant tissue damage, which, if severe enough, could grant *S. pneumoniae* access to the underlying tissues and the systemic circulation ([21](#), [92](#)). By exacerbating inflammatory responses during active pneumococcal infections, air pollution could therefore contribute to the development of invasive pneumococcal disease.

Adaptive immune responses, including those mediated through Th17 and Treg cells, are also stimulated by air pollution. Neill and colleagues showed that a balance between immune suppression by Treg responses and inflammatory responses are key in controlling *S. pneumoniae* growth in the airways and preventing tissue damage ([146](#)).

Inflammation of the respiratory tract is also one of the underlying mechanisms behind pneumococcal shedding ([155](#)). The increased inflammatory responses cause more mucus production, which increases nasal secretions ([156](#)). Increased mucus production, as described earlier in this review and elsewhere ([41–43](#)), might therefore have an impact on pneumococcal transmission. However, impaired mucociliary clearance that has been observed after air pollution exposure ([38–41](#)) may contain the pneumococci within the airways. This could limit pneumococcal shedding from a host and may therefore impair transmission. Air pollution is also associated with the exacerbation of rhinitis, a prevalent respiratory disease ([157](#)). Sneezing—a common rhinitis symptom—increases the expulsion of respiratory aerosols and droplets and has therefore been suggested to play a role in pneumococcal transmission ([2](#), [158](#), [159](#)). Increased sneezing induced by air pollution could therefore impact transmission. Other respiratory reflexes such as coughing may also have a role, especially given that influenza-like symptoms, which encompass coughing, are associated with higher prevalence of pneumococcal carriage ([160](#)). However, a recent meta-analysis showed no link between air pollution exposure and cough ([161](#)).

This review summarizes several host responses that are affected by air pollution and describes how these altered host responses could benefit pneumococcal colonization, pathogenesis, and transmission. There is substantial evidence that air pollution causes a dysfunctional airway mucosa, impairs macrophage function, and exacerbates proinflammatory immune responses, because this has been observed in several *in vitro* and *in vivo* (human) exposure models ([38–42](#), [67](#), [69](#), [92](#), [103–105](#), [124–126](#), [130](#), [133](#), [136](#)). However, discrepancies exist in the literature for other effects of air pollution, such as the production of antimicrobial peptides and microbiota dysbiosis. In addition, the

link between air pollution and nutrient concentrations in the respiratory tract and the impact on pneumococcal proliferation are suggestive and require further research. Only a few studies exist that directly assess the impact of air pollution exposure on the development of (invasive) pneumococcal disease, and, because pneumococcal transmission is a relatively understudied topic, the effect of air pollution, including the associated effects on the host, has not yet been investigated in pneumococcal transmission models.

Although the incidence of pneumococcal disease is higher in areas with high air pollution ([14–18](#), [20–22](#)), the association is confounded by the strong link between burden of disease and low socioeconomic status ([162](#), [163](#)). Living in a high-pollution area is also linked to low socioeconomic status, making it difficult to determine if other effects associated with low socioeconomic status, such as chronic lung disease and poor diet, may also play a role ([162](#), [163](#)). It is known that welders are at increased risk for invasive pneumococcal disease when compared with nonwelder groups with a similar socioeconomic status; however, further research is required to determine if this is a specific risk from welding fumes ([19](#)). To control for confounding variables and determine direct impacts of air pollution on pneumococcal disease, we rely on *in vitro* and *in vivo* exposure models.

Effective vaccines against *S. pneumoniae* have led to a global reduction in the number of deaths caused by this pathogen (by 7.24% since 1990) ([9](#)). Some studies suggest that vaccination might reduce the adverse effects of air pollution exposure. For example, a study of 6,740 Chinese children showed that influenza vaccination reduced the negative effects of air pollution on lung function ([164](#)). In addition, the risk of acute coronary syndrome in the elderly is associated with exposure to air pollutants, but this risk is lower in the elderly population that received yearly influenza vaccines ([165](#)). Further research is required to determine if pneumococcal vaccines can reduce the adverse effects of air pollution. Conversely, air pollution may have an impact on the efficacy of vaccines. For example, lower neutralizing antibody titers to an inactivated severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) vaccine were measured in pollutant-exposed individuals, and reduced

antibody responses to a candidate *Haemophilus influenzae* vaccine antigen were observed in mice exposed to cigarette smoke ([166](#), [167](#)). The impact of air pollution on the effectiveness of pneumococcal vaccines has yet to be investigated. The effect of air pollution is likely to be more relevant for the pneumococcal polysaccharide vaccine than for the pneumococcal conjugate vaccine (PCV), because pneumococcal conjugate vaccine is administered early in life, whereas the effects of air pollution accumulate over a lifetime.

Despite effective vaccine programs, *S. pneumoniae* was still responsible for approximately 1.18 million deaths globally in 2016 ([14](#)), highlighting the need for other strategies to reduce mortality. Given the important role of air pollution in host susceptibility and development of pneumococcal disease, reducing air pollution concentrations or preventing inhalation of air pollution would be an effective strategy to reduce respiratory disease burden ([10](#), [11](#), [21](#)). In addition, treating pollutant-induced effects (e.g., an overactive immune response) may aid the management of pneumococcal disease. For example, a systematic review by Corrales-Medina and Musher ([168](#)) showed that treatment with immunomodulatory compounds has the potential to reduce the morbidity of pneumonia. These strategies could be especially effective in regions with high air pollution concentrations. Although epidemiological data show a strong correlation between air pollution and respiratory infections, much less is known about the underlying mechanisms ([14–23](#)). We rely primarily on animal exposure models, but it is difficult to evaluate or summarize the impact of air pollution because studies use different exposure concentrations and different generation/sampling methods ([129](#)). In addition, it is often difficult to extrapolate these data to “real life.” This is partly because of the technical challenge of simulating the dose and duration of real-life exposure in a laboratory setting. To fully understand the effects of air pollution—and that of the various chemical and biological components—we are in need of better real-world data to help us improve our current exposure models.

Footnotes

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