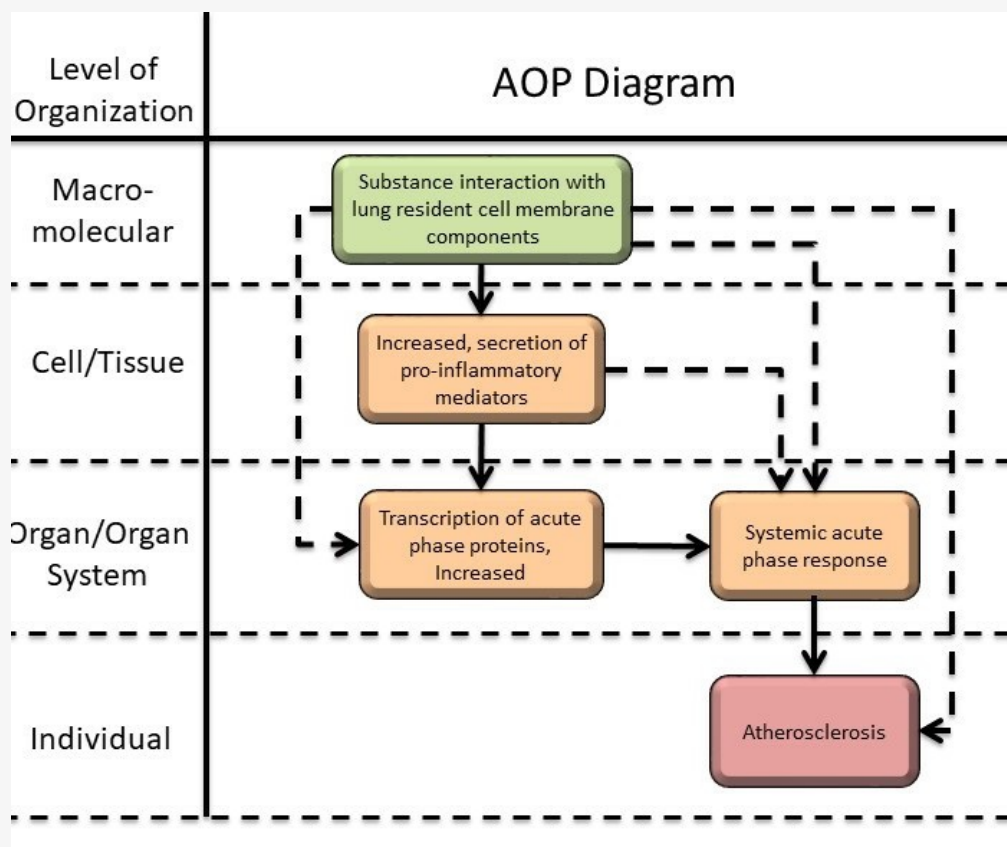


**AOP ID and Title:**

AOP 237: Substance interaction with lung resident cell membrane components leading to atherosclerosis

**Short Title: Interaction with lung cells leading to atherosclerosis****Graphical Representation****Authors**

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**Abstract**

The present adverse outcome pathways (AOP) presents the link between the interaction of stressors of the pulmonary system and atherosclerosis. After interaction with the lung cell membrane, stressors can induce the release of pro-inflammatory factors, which in turn triggers the expression of acute phase proteins genes in the lungs and other tissues. Serum amyloid A (SAA) and C-reactive protein (CRP) are the major acute phase proteins in humans, and are considered risk factors for cardiovascular disease (Table 1 presents selected differences between acute phase response in humans and mice). In particular, serum amyloid A restricts the transport of cholesterol to the liver, allowing the accumulation of cholesterol in arteries and the formation of foam cells, an early

marker of atherosclerosis.

**Table 1. Selected differences in acute phase response between humans and mice.**

Characteristic	Humans	Mice
Number of identified genes involved in acute phase response	62	62
Major acute phase proteins	CRP, SAA	Haptoglobin, SAA, serum amyloid P
Moderate and minor acute phase proteins	Haptoglobin, fibrinogen, $\alpha_1$ acid glycoprotein	CRP, fibrinogen
SAA isoforms	<i>Saa1</i> , <i>Saa2</i> and <i>Saa4</i>	<i>Saa1</i> , <i>Saa2</i> , <i>Saa3</i> and <i>Saa4</i>

References: <sup>1-4</sup>.

This AOP mainly focus on particles or particulate matter as stressors, however other inflammatory conditions that induce acute phase response, can be consider stressors and lead to atherosclerosis. In addition, most of the evidence is based on animal studies (mice) as a model for the human system, however the adverse outcome of the present AOP, atherosclerosis, is only applicable for humans. The AOP can be used for regulatory purposes and to risk assess inhalable materials having acute phase response as the critical effect.

## Background

Cardiovascular disease (CVD) is the leading cause of death worldwide, being responsible for 32% of all deaths in 2019 (WHO: <http://www.who.int>). The term CVD covers all diseases of the cardiovascular system, including atherosclerosis, which is manifested as increased plaque deposition or build-up in the arteries. Although, atherosclerosis is not a cause of death, it can lead to fatal conditions as stroke and myocardial infarction. Atherosclerosis is normally asymptotic disease and is initiated by a biological, chemical or physical insult to the artery walls. This leads to the expression of cell adhesion molecules (selectins, VCAM-1 and ICAM-1) on the endothelial lining of the arteries, which facilitates the activation, recruitment, and migration of monocytes through the endothelial monolayer <sup>5,6</sup>. Inside the intima layer, the monocytes differentiate into macrophages and internalize fatty deposits (mainly oxidized low-density lipoprotein). This results in them transforming into foam cells, which is a major component of the atherosclerotic fatty streaks. The fatty streaks reduce the elasticity of the artery walls and the foam cells promote a pro-inflammatory environment by secretion of cytokines and ROS. In addition, foam cells also induce the recruitment of smooth muscle cells to the intima. Added together, these changes lead to the formation of plaques on the artery walls. A fibrous cap of collagen and vascular smooth muscle cells protects the necrotic core and stabilizes the plaque <sup>7,8</sup>. However, blood clots can be formed if the plaque ruptures. These may travel with the bloodstream and obstruct the blood flow of smaller vessels, eg. the coronary arteries, which ultimately can lead to myocardial infarction.

Inhalation of particulate matter, chemicals and pathogens have been related to increased pulmonary inflammation. Whereas a normal immune reaction is crucial for effective elimination of threats to the body, chronic and unresolved inflammation has been linked to both adverse pulmonary and adverse systemic effects in humans. In concordance with this, various retrospective and prospective epidemiological studies have linked pulmonary exposure to respirable air particulates with increased the risk of developing CVD <sup>9-12</sup>. Inhalation of particles has been proposed to affect the cardiovascular system in several different ways, including through disruption of vasomotor function and through acceleration of plaque progression in atherosclerosis <sup>13,14</sup>.

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## Summary of the AOP

### Events

#### Molecular Initiating Events (MIE), Key Events (KE), Adverse Outcomes (AO)

Sequence	Type	Event ID	Title	Short name
	MIE	1495	<a href="#">Substance interaction with the lung resident cell membrane components</a>	Interaction with the lung cell membrane
	KE	1496	<a href="#">Increased, secretion of proinflammatory mediators</a>	Increased proinflammatory mediators
2	KE	1438	<a href="#">Transcription of acute phase proteins, Increased</a>	Increased transcription of APP

3	KE	1439	<a href="#">Systemic acute phase response</a>	Systemic APR
<b>Sequence</b>	<b>Type</b>	<b>Event ID</b>	<b>Title</b>	<b>Short name</b>
7	AO	1443	<a href="#">Atherosclerosis</a>	Atherosclerosis

## Key Event Relationships

Upstream Event	Relationship Type	Downstream Event	Evidence	Quantitative Understanding
<a href="#">Substance interaction with the lung resident cell membrane components</a>	adjacent	Increased, secretion of proinflammatory mediators	High	Low
<a href="#">Increased, secretion of proinflammatory mediators</a>	adjacent	Transcription of acute phase proteins, Increased	High	Moderate
<a href="#">Transcription of acute phase proteins, Increased</a>	adjacent	Systemic acute phase response	High	Moderate
<a href="#">Systemic acute phase response</a>	adjacent	Atherosclerosis	High	High
<a href="#">Substance interaction with the lung resident cell membrane components</a>	non-adjacent	Transcription of acute phase proteins, Increased	High	Moderate
<a href="#">Substance interaction with the lung resident cell membrane components</a>	non-adjacent	Systemic acute phase response	High	Moderate
<a href="#">Increased, secretion of proinflammatory mediators</a>	non-adjacent	Systemic acute phase response	High	Moderate
<a href="#">Substance interaction with the lung resident cell membrane components</a>	non-adjacent	Atherosclerosis	High	Moderate

## Stressors

Name	Evidence
Lipopolysaccharide	Not Specified
Graphene oxide nanoparticles	Not Specified
Carbon nanotubes	Not Specified
Insoluble nano-sized particles	Not Specified
Virus	Not Specified

## Overall Assessment of the AOP

### Domain of Applicability

#### Life Stage Applicability

##### Life Stage Evidence

Adult High

#### Taxonomic Applicability

##### Term Scientific Term Evidence Links

human Homo sapiens High [NCBI](#)

mouse Mus musculus High [NCBI](#)

#### Sex Applicability

##### Sex Evidence

Male High

Female High

This AOP is applicable to adult humans of both sexes. Although atherosclerosis is a condition that begins during childhood and progresses through life, its clinical manifestation is mostly observed in older individuals <sup>15</sup>.

The AOP is applicable to all stressors that can be inhaled and, therefore, interact with the pulmonary system, and induce pulmonary inflammation if the dose is high enough.

### Essentiality of the Key Events

	Defining question	High	Moderate	Low
Support for essentiality of KEs	What is the impact on downstream KEs and/or the AO if an upstream KE is modified or prevented?	Direct evidence from specifically designed experimental studies illustrating prevention or impact on downstream KEs and/or the AO if upstream KEs are blocked or modified	Indirect evidence that modification of one or more upstream KEs is associated with a corresponding (increase or decrease) in the magnitude or frequency of downstream KEs	No or contradictory experimental evidence of the essentiality of any of the KEs.
MIE: Substance interaction with the lung resident cell membrane components	<p><b>Moderate.</b></p> <p>It has been observed that there is a dose-response relationship between the dose of the stressor (i.e. substance interaction with lung cells), and acute phase response outcomes (KE2 and KE3).</p> <p>In addition, Danielsen et al. showed that Toll-like receptor 4 (<i>Tlr4</i>) knockout mice exposed to LPS, known to be a agonist for TLR4, did not induce an increase in cytokine/chemokines mRNA levels in lung and liver tissues (KE1) and did not produce a systemic acute phase response (KE3) <sup>16</sup>. Toll-like receptor 2 (<i>Tlr2</i>) knockout mice exposed to multiwalled carbon nanotubes did not induce increased <i>Saa1</i> mRNA levels in liver tissue (KE2) and did not induce increased SAA1 levels in plasma (KE3) <sup>16</sup>.</p>			
KE1: Increased, secretion of proinflammatory mediators	<p><b>Strong.</b></p> <p>Mice presenting IL-6 gene disruption (<i>IL-6<sup>-/-</sup></i>) shown a reduced response in liver mRNA levels (KE2) and serum levels (KE3) of the acute phase proteins haptoglobin, <math>\alpha</math>1-acid glycoprotein and serum amyloid a, after challenged by turpentine, lipopolysaccharide and bacterial infection <sup>17</sup>.</p> <p>In an <i>in vitro</i> study, blocking IL-6 receptors in hepatic cell lines resulted in a reduction of SAA1 mRNA (KE2), while blocking IL-1<math>\beta</math> and TNF-<math>\alpha</math> receptors partially reduced the expression of SAA1 mRNA <sup>18</sup>.</p> <p>In a clinical trial study, patients with a history of myocardial infarction where administered with a monoclonal antibody for IL-1<math>\beta</math> (canakinumab). The results showed that the treatment significantly reduced blood CPR levels in a dose-dependent manner (KE2)</p>			

	and KE3) after 48 months, and there was a decrease in incidence rate of recurrent cardiovascular events (AO) <sup>19</sup> .
KE2: Acute phase proteins transcription, Increased	<b>Strong.</b> Gene transcription is necessary for the synthesis of proteins (KE3). Thompson et al. showed that suppression of SAA3 in SAA1/SAA2 double knockout mice produced a significant reduction of atherosclerotic plaque area (AO) <sup>20</sup> .
KE3: Systemic acute phase response	<b>Strong.</b> Studies using animal model of atherosclerosis have shown that elevated levels of SAA induces plaque progression (AO) <sup>20,21</sup> . In prospective epidemiological studies, CRP and SAA levels are predictive of risk of cardiovascular disease <sup>22,23</sup> .

### Uncertainties or Inconsistencies

Atherosclerosis is a disease influenced by multiple factors including high levels of lipoproteins in blood, elevated blood pressure, smoking, obesity, type 2 diabetes, diet, physical activity, among others <sup>15,24,25</sup>. As described by Libby, inflammation is also involved in atherosclerosis, providing mechanisms for the risk factors to induce atherosclerotic plaque formation and progression <sup>26,27</sup>. Therefore, although inflammation and acute phase response are not the only causes of atherosclerosis, the early key events (KE1, KE2 and KE3) can be used to evaluate the particle-induced risk to developing atherosclerosis.

CRP and SAA are risk factors for cardiovascular disease <sup>23</sup>. However, Mendelian randomization studies have shown that CRP genotypes are not associated with risk of coronary heart disease and that genetically elevated levels of CRP are not associated with coronary heart disease risk <sup>28,29</sup>. In humans, measuring gene expression of acute phase proteins is not very common, as a tissue sample is needed, while measuring acute phase protein in blood is more common.

In mice studies, it is possible to measure both SAA gene expression and protein levels, however the dynamic range for Saa gene expression is larger. Although it is suggested that acute phase proteins are mainly produced in the liver <sup>2</sup>, it has been shown that in mice the liver has little upregulation of *Saa* genes after exposure to ultrafine carbon particles or diesel exhaust particle. On the other hand, the lung shows a marked expression of *Saa3* mRNA <sup>30,31</sup>.

There is an inconsistency with the results from human studies. It has been observed that in most controlled human studies, an increase in CRP and/or SAA was observed after exposure to particulate matter <sup>32-37</sup>. However, in other human studies the exposure did not induce acute phase response <sup>38,39</sup>, maybe due to low levels of exposure <sup>40</sup> or limited statistical power.

In the case of nanomaterials, it has been shown that physicochemical characteristics as size, surface area, surface functionalization, shape, composition, among others, affect the magnitude and duration of acute phase response in mice <sup>41-43</sup>. In animal models, both inflammatory and acute phase response are predicted by the total surface area of the retained, insoluble particles <sup>42,44</sup>.

## Weight of Evidence Summary

### Biological plausibility of each KER

	Defining question	High	Moderate	Low
<b>Support for Biological Plausibility of KERs</b>	Is there a mechanistic (i.e., structural or functional) relationship between KEup and KEdown consistent	Extensive understanding based on extensive previous documentation and broad acceptance - Established mechanistic	The KER is plausible based on analogy to accepted biological relationships - but scientific understanding is not	There is empirical support for a statistical association between KEs (See 3.), but the structural or functional

	with established biological knowledge?	basis	completely established.	relationship between them is not understood.
MIE => KE1: Interaction with the lung cell membrane leads to Increased proinflammatory mediators	Biological Plausibility of the MIE => KE1 is <b>High</b> .  Rationale: There is extensive evidence showing that interaction of stressors with the respiratory system induces the release of proinflammatory markers.			
KE1 => KE2: Increased proinflammatory mediators leads to Increased transcription of acute phase proteins	Biological Plausibility of the KE1 => KE2 is <b>High</b> .  Rationale: Acute phase proteins are induced by pro-inflammatory cytokines. These cytokines are produced at sites of inflammation mainly by monocytes and macrophages.			
KE2 => KE3: Increased transcription of acute phase proteins leads to Systemic acute phase response	Biological Plausibility of the KE2 => KE3 is <b>High</b> .  Rationale: After gene expression of acute phase proteins in tissues during inflammatory conditions, mRNA is translated and folded into proteins. These proteins are then release to the systemic circulation.			
KE3 => AO: Systemic acute phase response leads to Atherosclerosis	Biological Plausibility of the KE3 => KE2 is <b>High</b> .  Rationale: During acute phase response, serum amyloid A replaces apolipoprotein A-1 from high-density lipoprotein. This replacement obstructs the reverse transport of cholesterol to the liver, allowing the formation of foam cells, an early marker of atherosclerotic lesions.			
Non-adjacent MIE => KE2:  Interaction with the lung cell membrane leads to Increased transcription of acute phase proteins	Biological Plausibility of the MIE => KE2 is <b>High</b> .  Rationale: Acute phase response occurs during inflammatory condition, including the interaction of stressor with the airways. There is extensive evidence that nanomaterials induce the expression of acute phase response genes in mice.			
Non-adjacent MIE => KE3:  Interaction with the lung cell membrane leads to Systemic acute phase response	Biological Plausibility of the MIE => KE3 is <b>High</b> .  Rationale: There is plenty of evidence showing that inhalation or instillation of stressors induces systemic acute phase response in humans and mice.			
Non-adjacent KE1 => KE3: Increased proinflammatory mediators leads to Systemic APR	Biological Plausibility of the KE1 => KE3 is <b>High</b> .  Rationale: Pro-inflammatory cytokines induce the release of acute phase proteins. These proteins are releases from inflammatory sites to the systemic			

Non-adjacent MIE => AO:	circulation Biological Plausibility of the MIE => AO is <b>Moderate</b> .
Interaction with the lung cell membrane leads to Atherosclerosis	Rationale: There is evidence that the interaction of the lungs with stressor induces atherosclerotic plaque progression; however, the mechanistic relationship has not been clarified.

Please also refer to AOP173: Substance interaction with the pulmonary resident cell membrane components leading to pulmonary fibrosis, which shares MIE and KE1 with the present AOP.

#### Empirical support for each KER

	Defining question	High	Moderate	Low
<b>Empirical Support</b>	Does KEup occur at lower doses and earlier time points than KEdown and at the same dose of prototypical stressor, is the incidence of specific KEup > than that for KEdown?  Are there inconsistencies in empirical support across taxa, species and prototypical stressor that don't align with expected pattern for hypothesised AOP?	Multiple studies showing dose-dependent change in both events following exposure to a wide range of specific prototypical stressors. (Extensive evidence for temporal, dose-response and incidence concordance) and no or few critical data gaps or conflicting data	Demonstrated dependent change in both events following exposure to a small number of specific prototypical stressors and some evidence inconsistent with expected pattern that can be explained by factors such as experimental design, technical considerations, differences among laboratories, etc.	Limited or no studies reporting dose-dependent change in both events following exposure to a specific prototypical stressor (i.e., endpoints never measured in the same study or not at all); and/or significant inconsistencies in empirical support across taxa and species that don't align with expected pattern for hypothesised AOP
MIE => KE1: Interaction with the lung cell membrane leads to Increased proinflammatory mediators	Empirical Support of the MIE => KE1 is <b>Moderate</b> .  Rationale: There are limited <i>in vitro</i> studies which show a temporal and dose-dependent relationship between these two events.			
KE1 => KE2: Increased proinflammatory mediators leads to Increased transcription of acute phase proteins	Empirical Support of the KE1 => KE2 is <b>High</b> .  Rationale: There are a large number of studies showing a dose concordance and temporal concordance.			
KE2 => KE3: Increased	Empirical Support of the KE2 => KE3 is <b>High</b> .			

transcription of acute phase proteins leads to Systemic acute phase response	Rationale: There are a large number of studies showing a dose concordance and temporal concordance. However, there are inconsistencies between gene expression and translation of acute phase proteins.
KE3 => AO: Systemic acute phase response leads to Atherosclerosis	Empirical Support of the KE3 => AO is <b>Moderate</b> .  Rationale: There is a limited number of animal studies showing the relationship between the key events, in addition of epidemiological studies showing association between the key events.
Non-adjacent  MIE => KE2:  Interaction with the lung cell membrane leads to Increased transcription of acute phase proteins	Empirical Support of the MIE => KE2 is <b>Moderate</b> .  Rationale: There are a large number of studies showing a dose concordance and temporal concordance in animal studies. However, in the case of nanomaterials it has been shown that physicochemical characteristics affect the magnitude and duration of the expression of acute phase proteins in mice.
Non-adjacent  MIE => KE3:  Interaction with the lung cell membrane leads to Systemic acute phase response	Empirical Support of the MIE => KE3 is <b>Moderate</b> .  Rationale: There are plenty of studies showing a dose concordance and temporal concordance in animal and controlled human studies. It has been observed that systemic acute phase response is not always observed after exposure.
Non-adjacent KE1 => KE3: Increased proinflammatory mediators leads to Systemic APR	Empirical Support of the KE1 => KE3 is <b>Moderate</b> .  Rationale: There is plenty of studies showing a dose concordance and temporal concordance. However, there are inconsistencies between changes in blood levels of pro-inflammatory mediators and systemic APR.
Non-adjacent  MIE => AO:  Interaction with the lung cell membrane leads to Atherosclerosis	Empirical Support of the MIE => AO is <b>Moderate</b> .  Rationale: There is a number of studies showing the relationship between the key events.

## Quantitative Consideration

The table below presents a characterization of every KER.

It is important to clarify that when assessing stressors in mice studies, it is possible to measure the gene expression of acute phase proteins (KE2) in different tissues, however in humans this is not likely as a tissue sample would be required. On the other hand, in humans it is much more common and easier to measure systemic acute phase response (KE3) through a blood sample. In mice, it has been shown that *Saa3* mRNA in lung tissue and blood levels of SAA3 are correlated<sup>42</sup>. In addition, SAA levels in mice and humans seem to be in level in magnitude after exposure to zinc oxide nanoparticles<sup>42</sup>. This suggest, that systemic acute phase response in humans can be estimated from mice studies.

In the case of nanomaterials and mice studies, *Saa3* mRNA in lung tissue is also correlated to pulmonary inflammation measured as neutrophil numbers, and both of these endpoints can be estimated by calculating the dosed surface area (specific surface area multiplied by dose level)<sup>42</sup>.



Finally, the relative risk of people developing a cardiovascular disease can be calculated from blood level of acute phase proteins in epidemiological studies.

KER	Quantitative understanding
MIE => KE1: Interaction with the lung cell membrane leads to Increased proinflammatory mediators	<p>The quantitative understanding of MIE =&gt; KE1 is <b>Low</b>.</p> <p>Rationale: The quantitative prediction of the release of proinflammatory factors can be made from the interaction of the stressors with the pulmonary system.</p> <p>In the case of some stressors (nanomaterials) it is possible to make a prediction using the dosed surface area of the materials and neutrophil numbers as an indirect marker of the release of pro-inflammatory factors.</p>
KE1 => KE2: Increased proinflammatory mediators leads to Increased transcription of acute phase proteins	<p>The quantitative understanding is of KE1 =&gt; KE2 is <b>Moderate</b>.</p> <p>Rationale: In mice, the gene expression of the acute phase protein SAA after exposure to metal oxide nanomaterials can be estimated using an indirect marker of the release of pro-inflammatory factors (neutrophil numbers).</p>
KE2 => KE3: Increased transcription of acute phase proteins leads to Systemic acute phase response	<p>The quantitative understanding of KE2 =&gt; KE3 is <b>Moderate</b>.</p> <p>Rationale: In mice, the systemic levels of the acute phase protein SAA after exposure to metal oxide nanomaterials can be estimated from the gene expression in lung tissue.</p>
KE3 => AO: Systemic acute phase response leads to Atherosclerosis	<p>The quantitative understanding is of KE3 =&gt; AO is <b>High</b>.</p> <p>Rationale: The risk of developing a cardiovascular disease at population level can be calculated from blood levels of acute phase proteins.</p>
Non-adjacent MIE => KE2:  Interaction with the lung cell membrane leads to Increased transcription of acute phase proteins	<p>The quantitative understanding of MIE =&gt; KE2 is <b>Moderate</b>.</p> <p>Rationale: In mice, the gene expression of the acute phase protein SAA after exposure to metal oxide nanomaterials can be estimated from the dosed surface area.</p>
Non-adjacent MIE => KE3:  Interaction with the lung cell membrane leads to Systemic acute phase response	<p>The quantitative understanding of MIE =&gt; KE3 is <b>Moderate</b>.</p> <p>Rationale: In mice, the blood levels of the acute phase protein SAA after exposure to metal oxide nanomaterials can be estimated from the dosed surface area.</p>

Non-adjacent KE1 => KE3: Increased proinflammatory mediators leads to Systemic APR	The quantitative understanding of KE1 => KE3 is <b>Moderate</b> .  Rationale: In mice, the blood levels of the acute phase protein SAA after exposure to metal oxide nanomaterials and multiwalled carbon nanotubes can be estimated from neutrophil numbers in bronchoalveolar lavage fluid.
Non-adjacent MIE => AO: Interaction with the lung cell membrane leads to Atherosclerosis	The quantitative understanding of MIE => AO is <b>Moderate</b> .  Rationale: Epidemiological studies have shown the risk ratios of having a cardiovascular event per increase or decrease of exposure to particulate matter.

## Considerations for Potential Applications of the AOP (optional)

Particle-induced acute phase response can be regarded as a critical effect linking particle-exposure to cardiovascular disease. Dose-response relationships can be used to establish no-observed-adverse-effect levels (NOAEL) for regulatory purposes and occupational exposure limits for inhalable materials can be determined through health-based risk assessments. This approach was taken by the Danish National Research Centre for the Working Environment at request of the Danish Working Environment Authority and an occupational exposure limit for zinc oxide was proposed based on the induction of acute phase response as the critical effect (the report can be found in: [Dokumentation for helbredsbaseerede grænseværdier for kemiske stoffer i arbejdsmiljøet \(nfa.dk\)](https://nfa.dk)).

As mentioned previously, not all KE can easily be measured in humans, therefore animal studies can be used to measure early KE and perform a risk assessment of different stressors. Additionally, physicochemical properties, such as specific surface area and dissolution, are important predictors of particle-induced acute phase response that can be used for hazard assessment <sup>42</sup>

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## Appendix 1

### List of MIEs in this AOP

[Event: 1495: Substance interaction with the lung resident cell membrane components](#)

Short Name: Interaction with the lung cell membrane

#### Key Event Component

Process	Object	Action
pattern recognition receptor signaling pathway		increased
toll-like receptor signaling pathway	Toll-like receptor	increased
toll-like receptor 4 signaling pathway	Toll-like receptor 4	increased

#### AOPs Including This Key Event

AOP ID and Name	Event Type
<a href="#">Aop:173 - Substance interaction with the pulmonary resident cell membrane components leading to pulmonary fibrosis</a>	MolecularInitiatingEvent
<a href="#">Aop:451 - Interaction with lung resident cell membrane components leads to lung cancer</a>	MolecularInitiatingEvent
<a href="#">Aop:237 - Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	MolecularInitiatingEvent

#### Biological Context

##### Level of Biological Organization

Molecular

## Cell term

### Cell term

eukaryotic cell

## Evidence for Perturbation by Stressor

### Overview for Molecular Initiating Event

As stated earlier, there are many different ways by which pro-fibrotic stressors can interact with the components of cell membrane and often involve multiple interactions at the same time. Few studies investigate the exact interaction between the stressor and the cellular membrane components. Asbestos and silica crystals engage scavenger receptors present on the macrophages (Murthy et al., 2015). Bleomycin binds high affinity bleomycin binding sites present on rat alveolar macrophage surfaces, leading to macrophage activation (Denholm and Phan, 1990). However, the consequences of such interactions such as, the release of PRR agonists DAMPs (alarmins) from dying or injured cells, increased gene or protein synthesis downstream of receptor binding or in the case of NMs, their cellular uptake, are measured routinely as indicative of occurrence of such interactions (Nel et al., 2009; Cheng et al., 2013). Because of the phys-chem properties such as surface charge, NMs and asbestos like materials can bind to cellular macromolecules and cell surface/membrane components, which in turn, facilitate their uptake and intracellular sequestration by the cells (NIOSH, 2011a; Pascolo et al., 2013). Several DAMPs that can be effectively measured in biological samples and cultured cells include High Mobility Group Binding 1 (HMGB1) protein, Heat Shock proteins (HSPs), uric acid, annexins, and S100 proteins (Bianchi, 2007). Of all DAMPs, interleukin (IL)-1 $\alpha$  is the most commonly measured alarmin. IL-1 $\alpha$  is the principal pro-inflammatory moiety and is a designated 'alarmin' in the cell that alerts the host to injury or damage (Di Paolo and Shayakhmetov, 2016). It is shown that administration of necrotic cells to mice results in neutrophilic inflammation that was entirely mediated by IL-1 $\alpha$  released from the dying or necrosed cells and consequent activation of IL-1 Receptor 1 (IL-1R1) signalling (Suwara et al., 2014). IL-1 $\alpha$  is released following exposure to MWCNTs (Nikota et al., 2017) and silica (Rabolli et al., 2014). Although IL1-b is not a designated alarmin, its secretion following exposure to stressors is routinely assessed and is linked to initiation of cell or tissue injury.

Other high aspect ratio fibres such as asbestos and CNTs induce frustrated phagocytosis and acute cell injury (Boyles et al., 2015; Dörger et al., 2001; Brown et al., 2007; Kim et al., 2010; Poland et al., 2008), leading to DAMP release (Nikota et al, 2017), inflammation and immune responses.

### Domain of Applicability

#### Taxonomic Applicability

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
rat	Rattus norvegicus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

#### Life Stage Applicability

##### Life Stage Evidence

Adults High

#### Sex Applicability

##### Sex Evidence

Male High

Human, mouse, rat.

Although the expression of DAMPs following exposure to pro-fibrotic substances is not assessed across species, it is known that alarmins are released after trauma or injury, and their release is important for initiating the inflammatory response in all species including humans. The immediate acute inflammatory response involving DAMP signalling is also observed in human idiopathic pulmonary fibrosis (IPF); however, anti-inflammatory drugs have proven ineffective for treating IPF. Danger signalling axis including uric acid, adenosine triphosphate and IL-33/ST2 has been proven to promote lung fibrosis in animals.

### Key Event Description

The human lung consists of approximately 40 different resident cell types that play different roles during homeostasis, injury, repair and disease states (Franks et al., 2008; Luettich et al., 2021). Of these, resident airway epithelial cells, alveolar/interstitial macrophages and dendritic cells are well characterised for their ability to sense the danger upon interaction with harmful substances and relay the message to mount the necessary immune/inflammatory response. The resident macrophages are

present in all tissues, and in a steady state, macrophages contribute to epithelial integrity, survey the tissue for invading pathogens or chemicals and maintain an immunosuppressive environment. Their main function is to clear the incoming irritants and microbes. They are named differently based on the tissue type and their specific functions (Kierdorf et al., 2015).

### ***Substance interactions:***

The chemicals or pathogens interact with cellular membrane to gain access to the organisms' interior. A predominant interaction mechanism involves the recognition of innate immune response agonists by pattern recognition receptors (PRRs) present on resident cells such as epithelial and alveolar macrophages. PRRs are also present on other immune and parenchymal cells. PRRs can be activated by two classes of ligands. Pathogen associated molecular patterns (PAMPs) are microbial molecules derived from invading pathogens. PAMPs will not be discussed further as pathogens are not the focus for the AOP presented here. The other class of ligands are called danger associated molecular patterns (DAMPs) that include cellular fragments, nucleic acids, small molecules, proteins and even cytokines released from injured or dying cells (Bianchi, 2007). Most fibrogenic stressors discussed in this AOP act via DAMPs-driven PRR activation. High aspect ratio (HAR) materials such as asbestos or carbon nanotubes (CNTs) pierce the cellular membrane of epithelial cells or resident macrophages resulting in cell injury or non-programmed cellular death. Alveolar macrophages trying to engulf HAR fibres that are long and stiff undergo frustrated phagocytosis because of their inability to engulf the piercing fibres and subsequently lead to cell injury (Boyles et al., 2015; Brown et al., 2007; Donaldson K et al., 2010; Dörger et al., 2001; Mossman and Churg, 1998). The cellular debris from injured or dying cell then serves as ligands for PRRs (Nakayama, 2018), leading to cell activation. In case of pro-fibrotic insoluble particles such as silica, coal dust and nanomaterials (NMs), the particle adsorbed opsonins such as immunoglobulins, complement proteins, or serum proteins act as ligands to the receptors on the macrophage cell surface (Behzadi et al., 2017). The tissue response to these materials resembles that observed following foreign body invasion in lungs.

Toll-like receptors (TLRs) are highly conserved PRRs that are associated with fibrogenic stressors (Desai et al., 2018). Inhibition of TLR-4 is protective against bleomycin-induced fibrosis (Li et al., 2015). However, the exact role and mechanisms by which TLRs mediate lung fibrosis are yet to be uncovered and some studies have shown TLRs to be protective against lung fibrosis (Desai et al., 2018). Asbestos and silica crystals are suggested to engage scavenger receptors present on the macrophages. Mice deficient in class A scavenger macrophage receptor with collagenous structure (MARCO) are shown to induce reduced fibrogenic response following chrysotile asbestos exposure; although, the direct binding of MARCO by asbestos is not investigated in the study (Murthy et al., 2015). In case of soluble substances such as bleomycin, paraquat (Dinis-Oliveira et al., 2008) (N,N'-dimethyl-4, 4'-bipyridinium dichloride) and other soluble fibrogenic chemicals, direct damage of lung epithelial cells and resulting cellular debris or secreted cytokines (DAMPs) serve as triggers for downstream cascading pro-inflammatory events, tissue injury and fibrosis. Engagement of PRRs and consequent cell activation is observed in various organisms including flies and mammals (Denholm and Phan, 1990; Matzinger, 2002).

### **How it is Measured or Detected**

#### ***Detection of DAMPs or homeostasis-altering molecular processes:***

Cellular interaction with substances or particles can be measured by assessing the release of DAMPs from stressed, injured or dying cells - indicative of binding of PRRs on the cell surface. Release of DAMPs is reflective of substance interaction with resident cells and their activation, a key step in the process of inflammation.

The release of DAMPs can be measured by the techniques listed in the published literature (Nikota et al., 2017; Rabolli et al., 2014; Suwara et al., 2014).

#### ***Targeted enzyme-linked immunosorbent assays (ELISA) (routinely used and recommended):***

ELISA – permits quantitative measurement of antigens in biological samples. For example, in a cytokine ELISA (sandwich ELISA), an antibody (capture antibody) specific to a cytokine is immobilised on microtitre wells (96-well, 386-well, etc.). Experimental samples or samples containing a known amount of the specific recombinant cytokine are then reacted with the immobilised antibody. Following removal of unbound antibody by thorough washing, plates are reacted with the secondary antibody (detection antibody) that is conjugated to an enzyme such as horseradish peroxidase, which when bound, will form a sandwich with the capture antibody and the cytokine (Amsen and De Visser, 2009). The secondary antibody can be conjugated to biotin, which is then detected by addition of streptavidin linked to horseradish peroxidase. A chromogenic substrate can also be added, which is the most commonly used method. Chromogenic substrate is chemically converted by the enzyme coupled to the detection antibody, resulting in colour change. The amount of colour detected is directly proportional to the amount of cytokine in the sample that is bound to the capture antibody. The results are read using a spectrophotometer and compared to the levels of cytokine in control samples where cytokine is not expected to be secreted or to the samples containing known recombinant cytokine levels.

Interleukin (IL)-1 $\alpha$  and -1 $\beta$  is activated or secreted into the cytosol following stimulus (Di Paolo and Shayakhmetov, 2016). Targeted ELISA can be used to quantify IL-1 $\alpha$  or IL-1 $\beta$  that is released in the culture supernatant of the cells exposed to toxicants, in bronchoalveolar lavage fluid and serum of exposed animals. The assay is also applicable to human serum, cerebrospinal fluid, and peritoneal fluids.

Similarly, other alarmins can also be quantified by ELISA. Western blot is another method that can be used to quantify the release of various alarmins using specific antibodies. ELISA or real-time reverse transcription-polymerase chain reaction (qRT-PCR) assays can also be used to quantify the expression of genes or proteins that are regulated by the receptor binding – e.g. downstream of

TLR binding.

### ***Frustrated phagocytosis and cellular uptake of NMs:***

*In vitro*, interaction of NMs with the cellular membrane is investigated by assessing their uptake by lysosomes (Chen et al., 2013; Nel et al., 2009; Varela et al., 2012). Immunohistochemistry methods targeting lysosome specific proteins are regularly employed for this purpose. In co-localisation experiments, lysosomal marker Lysosomal-associated membrane protein 1 (LAMP1) antibody is used to detect particle co-localisation with lysosomes. A combination of Cytoviva hyperspectral microscope and immunolocalisation (Decan et al., 2016) or confocal microscopy to visualise co-localisation of fluorescence labelled nanoparticles with lysosomal markers have been used.

Frustrated phagocytosis is assessed using microscopic techniques such as time-lapse microscopy, backscatter electron microscopy and others (Donaldson et al., 2010; Murphy et al., 2012; Padmore et al., 2017; Pascolo et al., 2013; Schinwald et al., 2012). In addition, MIE 1668 of AOP303 notes other indirect methods for measuring frustrated phagocytosis.

### ***Cellular co-culture models of the pulmonary epithelium:***

Complex co-culture systems, such as those containing epithelial cells and immune cells, better model the environment of the lung epithelium and can be used to study the interaction of potentially pro-fibrotic fibres and particles with resident lung cells. This type of model has been used, alongside electron microscopy, to study lung cell interactions with CNTs following 24 h *in vitro* exposure (Clift et al., 2014). More recently, the EpiAlveolar model, which contains primary human alveolar epithelial cells, endothelial cells, as well as fibroblasts was assessed for its ability to predict fibrosis induced by CNTs (Barasova et al., 2020). Using laser scanning, fluorescence, and enhanced darkfield microscopy, CNT interaction with the resident cells of the model was shown, and this interaction induced the formation of holes in the epithelial model (Barasova et al., 2020). While new co-culture models are a better recapitulation of the native lung environment as compared to traditional mono-cultures, the increased complexity necessitates enhanced expertise in tissue culture techniques, and can make them less practical as compared to submerged mono culture methods.

### ***Ex vivo model of the lung – Precision cut lung slices (PCLS):***

Even closer to the *in vivo* condition than co-culture models, PCLS techniques capture the native lung architecture, cell-cell communication and cellularity of the lung. Advancement in culturing and cryopreservation techniques has increased accessibility and use of PCLS for longer term studies (Bai et al., 2016; Neuhaus et al., 2017). These slices can be cultured *ex vivo* for up to a week with minimal reduction in viability, and the technique has recently been assessed for its applicability to assess nanomaterial induced fibrosis *ex vivo* (Rahman et al., 2020). Using multi-walled carbon nanotubes (MWCNTs) and darkfield microscopy, interaction between the nanofibers and the lung epithelium could be determined. The main downside of this technique is the animal requirement, which precludes their use in a first-pass screening context for the MIE.

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## List of Key Events in the AOP

### [Event: 1496: Increased, secretion of proinflammatory mediators](#)

#### Short Name: Increased proinflammatory mediators

#### Key Event Component

Process	Object	Action
cytokine production involved in inflammatory response	Cytokine	increased
chemokine secretion	Chemokine	increased
complement activation		increased
	Interleukin	increased

#### AOPs Including This Key Event

AOP ID and Name	Event Type
<a href="#">Aop:173 - Substance interaction with the pulmonary resident cell membrane components leading to pulmonary fibrosis</a>	KeyEvent
<a href="#">Aop:320 - Binding of SARS-CoV-2 to ACE2 receptor leading to acute respiratory distress associated mortality</a>	KeyEvent
<a href="#">Aop:382 - Angiotensin II type 1 receptor (AT1R) agonism leading to lung fibrosis</a>	KeyEvent
<a href="#">Aop:392 - Decreased fibrinolysis and activated bradykinin system leading to hyperinflammation</a>	KeyEvent
<a href="#">Aop:409 - Frustrated phagocytosis leads to malignant mesothelioma</a>	KeyEvent
<a href="#">Aop:377 - Dysregulated prolonged Toll Like Receptor 9 (TLR9) activation leading to Multi Organ Failure involving Acute Respiratory Distress Syndrome (ARDS)</a>	KeyEvent

AOP ID and Name	Key Event Type
<a href="#">Aop:39 - Covalent Binding, Protein, leading to Increase, Allergic Respiratory Hypersensitivity Response</a>	KeyEvent
<a href="#">Aop:319 - Binding to ACE2 leading to lung fibrosis</a>	KeyEvent
<a href="#">Aop:451 - Interaction with lung resident cell membrane components leads to lung cancer</a>	KeyEvent
<a href="#">Aop:468 - Binding of SARS-CoV-2 to ACE2 leads to hyperinflammation (via cell death)</a>	KeyEvent
<a href="#">Aop:237 - Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	KeyEvent

## Biological Context

### Level of Biological Organization

Cellular

### Cell term

#### Cell term

eukaryotic cell

## Domain of Applicability

### Taxonomic Applicability

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
rats	Rattus norvegicus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

### Life Stage Applicability

#### Life Stage Evidence

Adults High

### Sex Applicability

#### Sex Evidence

Male High

Female High

Human, mouse, rat

Cytokines are the common pro-inflammatory mediators secreted following inflammogenic stimuli. Cytokines can be defined as a diverse group of signaling protein molecules. They are secreted by different cell types in different tissues and in all mammalian species, irrespective of gender, age or sex. A lot of literature is available to support cross species, gender and developmental stage application for this KE. The challenge is the specificity; most cytokines exhibit redundant functions and many are pleiotropic.

## Key Event Description

Pro-inflammatory mediators are the chemical and biological molecules that initiate and regulate inflammatory reactions. Pro-inflammatory mediators are secreted following exposure to an inflammogen in a gender/sex or developmental stage independent manner. They are secreted during inflammation in all species. Different types of pro-inflammatory mediators are secreted during innate or adaptive immune responses across various species (Mestas and Hughes, 2004). Cell-derived pro-inflammatory mediators include cytokines, chemokines, and growth factors. Blood derived pro-inflammatory mediators include vasoactive amines, complement activation products and others. These modulators can be grouped based on the cell type that secrete them, their cellular localisation and also based on the type of immune response they trigger. For example, members of the interleukin (IL) family including [IL-2](#), [IL-4](#), [IL-7](#), [IL-9](#), [IL-15](#), [IL-21](#), [IL-3](#), [IL-5](#) and Granulocyte-macrophage colony stimulating factor ([GM-CSF](#)) are involved in the adaptive immune responses. The pro-inflammatory cytokines include IL-1 family ([IL-1 \$\alpha\$](#) , [IL-1 \$\beta\$](#) , [IL-1 \$\gamma\$](#) , [IL-18](#), [IL-36 \$\alpha\$](#) , [IL-36 \$\beta\$](#) , [IL-36 \$\gamma\$](#) , [IL-36 \$\beta\$](#) , [IL-36 \$\alpha\$](#) , [IL-37](#)), [IL-6](#) family, Tumor necrosis factor ([TNF](#)) family, [IL-17](#), and Interferon gamma ([IFN- \$\gamma\$](#) ) (Turner et al., 2014). While [IL-4](#) and [IL-5](#) are considered T helper (Th) cell type 2 response, [IFN- \$\gamma\$](#)  is suggested to be Th1 type response.

Different types of pro-inflammatory mediators are secreted during innate or adaptive immune responses across various species (Mestas and Hughes, 2004). However, [IL-1](#) family cytokines, [IL-4](#), [IL-5](#), [IL-6](#), [TNF- \$\alpha\$](#) , [IFN- \$\gamma\$](#)  are the commonly measured mediators in experimental animals and in humans. Similar gene expression patterns involving inflammation and matrix remodelling are

observed in human patients of pulmonary fibrosis and mouse lungs exposed to bleomycin (Kaminski, 2002).

### **Literature evidence for its perturbation:**

Several studies show increased proinflammatory mediators in rodent lungs and bronchoalveolar lavage fluid, and in cell culture supernatants following exposure to a variety of carbon nanotube (CNT) types and other materials. Poland et al., 2008 showed that long and thin CNTs (>5 µm) can elicit asbestos-like pathogenicity through the continual release of pro-inflammatory cytokines and reactive oxygen species. Exposure to crystalline silica induces release of inflammatory cytokines (TNF-α, IL-1, IL-6), transcription factors (Nuclear factor kappa B [NF-κB], Activator protein-1 [AP-1]) and kinase signalling pathways in mice that contain NF-κB luciferase reporter (Hubbard et al., 2002). Boyles et al., 2015 found that lung responses to long multi-walled carbon nanotubes (MWCNTs) included high expression levels of pro-inflammatory mediators Monocyte chemoattractant protein 1 (MCP-1), Transforming growth factor beta 1 (TGF-β1), and TNF-α (Boyles et al., 2015). Bleomycin administration in rodents induces lung inflammation and increased expression of pro-inflammatory mediators (Park et al., 2019). Inflammation induced by bleomycin, paraquat and CNTs is characterised by the altered expression of pro-inflammatory mediators. A large number of nanomaterials induce expression of cytokines and chemokines in lungs of rodents exposed via inhalation (Halappanavar et al., 2011; Husain et al., 2015a). Similarities are observed in gene programs involving pro-inflammatory event is observed in both humans and experimental mice (Zuo et al., 2002).

### **How it is Measured or Detected**

The selection of pro-inflammatory mediators for investigation varies based on the expertise of the lab, cell types studied and the availability of the specific antibodies.

Real-time reverse transcription-polymerase chain reaction (qRT-PCR) – will measure the abundance of cytokine mRNA in a given sample. The method involves three steps: conversion of RNA into cDNA by reverse transcription method, amplification of cDNA using the PCR, and the real-time detection and quantification of amplified products (amplicons) (Nolan et al., 2006). Amplicons are detected using fluorescence, increase in which is directly proportional to the amplified PCR product. The number of cycles required per sample to reach a certain threshold of fluorescence (set by the user – usually set in the linear phase of the amplification, and the observed difference in samples to cross the set threshold reflects the initial amount available for amplification) is used to quantify the relative amount in the samples. The amplified products are detected by the DNA intercalating minor groove-binding fluorophore SYBR green, which produces a signal when incorporated into double-stranded amplicons. Since the cDNA is single stranded, the dye does not bind enhancing the specificity of the results. There are other methods such as nested fluorescent probes for detection, but SYBR green is widely used. RT-PCR primers specific to several pro-inflammatory mediators in several species including mouse, rat and humans, are readily available commercially.

Enzyme-linked immunosorbent assays (ELISA) – permit quantitative measurement of antigens in biological samples. The method is the same as described for the MIE. Both ELISA and qRT-PCR assays are used *in vivo* and are readily applicable to *in vitro* cell culture models, where cell culture supernatants or whole cell homogenates are used for ELISA or mRNA assays. Both assays are straight forward, quantitative and require relatively a small amount of input sample.

Apart from assaying single protein or gene at a time, cytokine bead arrays or cytokine PCR arrays can also be used to detect a whole panel of inflammatory mediators in a multiplex method (Husain et al., 2015b). This method is quantitative and especially advantageous when the sample amount available for testing is scarce. Lastly, immunohistochemistry can also be used to detect specific immune cell types producing the pro-inflammatory mediators and its downstream effectors in any given tissue (Costa et al., 2017). Immunohistochemistry results can be used as weight of evidence; however, the technique is not quantitative and depending on the specific antibodies used, the assay sensitivity may also become an issue (Amsen and De Visser, 2009).

Cell models - of varying complexity have been used to assess the expression of pro-inflammatory mediators. Two dimensional submerged monocultures of the main fibrotic effector cells – lung epithelial cells, macrophages, and fibroblasts – have routinely been used *in vitro* due to the large literature base, and ease of use, but do not adequately mimic the *in vivo* condition (Sharma et al., 2016; Sundararishnan et al., 2018). Recently, the EpiAlveolar *in vitro* lung model (containing epithelial cells, endothelial cells, and fibroblasts) was used to predict the fibrotic potential of MWCNTs, and researchers noted increases in the pro-inflammatory molecules TNF-α, IL-1β, and the pro-fibrotic TGF-β using ELISA (Barasova et al., 2020). A similar, but less complicated co-culture model of immortalized human alveolar epithelial cells and idiopathic pulmonary fibrosis patient derived fibroblasts was used to assess pro-fibrotic signalling, and noted enhanced secretion of Platelet derived growth factor (PDGF) and Basic fibroblast growth factor (bFGF), as well as evidence for epithelial to mesenchymal transition of epithelial cells in this system (Prasad et al., 2014). Models such as these better capitulate the *in vivo* pulmonary alveolar capillary, but have lower reproducibility as compared to traditional submerged mono-culture experiments.

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### **Event: 1438: Transcription of acute phase proteins, Increased**

**Short Name: Increased transcription of APP**

#### **Key Event Component**

Process	Object	Action
acute-phase response	Acute phase proteins	increased

#### **AOPs Including This Key Event**

**AOP ID and Name**

**Event Type**

**Biological Context****Level of Biological Organization**

Organ

**Organ term****Organ term**

lung

**Domain of Applicability****Taxonomic Applicability**

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

**Life Stage Applicability****Life Stage Evidence**

All life stages High

**Sex Applicability****Sex Evidence**

Male High

Female High

- Taxonomic applicability: APR is part of the immune response and is observed in vertebrate species<sup>3</sup>.
- Life stage applicability: This key event is applicable to all life stages.
- Sex applicability: This key event is applicable to male and female sexes.

**Key Event Description**

Acute phase response is characterized by the change in plasma concentration of acute phase proteins (APP), along with other physiological changes during inflammatory conditions <sup>1,2</sup>. In humans, the major APPs are C-reactive protein and serum amyloid A, while in mice the major APPs are serum amyloid A, haptoglobin and serum amyloid P <sup>1,3</sup>.

It is widely accepted that APPs are mainly produced by the liver, however several other tissues have been shown to express APPs; these include lungs, intestines, kidneys, skin and adipose tissue in humans <sup>4-7</sup>, and kidney, spleen, brain, lung and testis in mice <sup>8,9</sup>. According to National Center for Biotechnology Information (NCBI) serum amyloid A isoforms and C-reactive protein genes in mice have been shown to be expressed in tissue from adrenal gland, bladder, central nervous system, colon, duodenum, genital fat pad, heart, kidney, large intestine, limbs, liver, lung, mammary gland, ovary, placenta, small intestine, subcutaneous fat pad, testis and thymus<sup>10</sup>. In the case of humans, these genes have been shown to be expressed in tissue from adrenal gland, appendix, fat, gall bladder, heart, kidney, liver, lung, placenta, prostate, salivary gland, small intestine, stomach, thymus, thyroid, trachea and uterus<sup>10</sup>.

**How it is Measured or Detected**

Gene expression of APPs can be measured from tissue samples using quantitative Polymerase Chain Reaction (PCR). Humans and mice express four SAA isoforms (*Saa1*, *Saa2*, *Saa3* and *Saa4*), however *Saa3* is a pseudogene in humans<sup>11</sup>. CRP is expressed in humans and mice, although only moderately expressed in mice<sup>12</sup>.

It has been shown that in mice the *Saa3* isoform is the most differentially expressed APP gene in lung tissue and it is not highly expressed in the liver, while *Saa1* gene is the most differentially expressed in liver tissue after exposure to particles <sup>13-15</sup>. In humans, it has been shown that *crp*, *saa1*, *saa2* and *saa4* gene expression can be measured in lung samples taken during surgery <sup>16</sup>. In addition, microarray analysis can be used to evaluate the gene expression of several APPs at the same time <sup>13</sup>.

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**Event: 1439: Systemic acute phase response****Short Name: Systemic APR****Key Event Component**

Process	Object	Action
acute-phase response	Acute phase proteins	increased

**AOPs Including This Key Event****AOP ID and Name****Event Type**

[Aop:237 - Substance interaction with lung resident cell membrane components leading to atherosclerosis](#) KeyEvent

**Biological Context**

**Level of Biological Organization**

Organ

**Organ term****Organ term**

blood

**Domain of Applicability****Taxonomic Applicability**

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

**Life Stage Applicability**

Life Stage	Evidence
All life stages	High

**Sex Applicability**

Sex	Evidence
Male	High
Female	High

- Taxonomic applicability: APR is part of the immune response and is observed in vertebrates<sup>6</sup>.
- Life stages applicability: This key event is applicable to all life stages.
- Sex applicability: This key event is applicable to male and females sexes.

**Key Event Description**

During acute phase response, the plasma concentration of acute phase proteins (APP) changes in more than 25%. APPs that increase their concentration during APR are called positive APP, while negative APP are decreased during APR<sup>1</sup>. In humans, positive APPs include C-reactive protein, serum amyloid A, C3 and C4 complement system, mannose-binding lectin, fibrinogen, fibronectin, ferritin, haptoglobin, hemopexin, among others<sup>1,2</sup>. In humans the two major APPs are C-reactive protein (CRP) and serum amyloid A (SAA), whose concentration can increase in more than 1000-fold during acute phase response<sup>1</sup>. SAA and CRP have been shown to be correlated in humans<sup>3-5</sup>. In mice, the major APP are serum amyloid A, haptoglobin and serum amyloid P<sup>6</sup>.

**How it is Measured or Detected**

Systemic acute phase response is assessed by measuring APPs concentrations in blood plasma or serum, most often CRP and SAA. In humans, these proteins are measured by immunoassays detecting single or multiple proteins<sup>4,7-12</sup>. In addition, CRP is measured by turbidimetric<sup>13-15</sup> and nephelometric assays<sup>16</sup>.

In mice, CRP is not a major APP<sup>6</sup>, therefore SAA isoforms are measured using ELISA assays or western blot<sup>17-21</sup>.

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## List of Adverse Outcomes in this AOP

### [Event: 1443: Atherosclerosis](#)

Short Name: Atherosclerosis

### Key Event Component

Process	Object	Action
Atherosclerosis		increased

### AOPs Including This Key Event



## AOP ID and Name

## Event Type

[Aop:237 - Substance interaction with lung resident cell membrane components leading to atherosclerosis](#) AdverseOutcome

## Biological Context

## Level of Biological Organization

Individual

## Domain of Applicability

## Taxonomic Applicability

Term	Scientific Term	Evidence	Links
human	Homo sapiens	High	<a href="#">NCBI</a>

## Life Stage Applicability

Life Stage	Evidence
All life stages	High

## Sex Applicability

Sex	Evidence
Male	High
Female	High

- Taxonomic applicability: Human.
- Life stages applicability: All life stages. Although atherosclerosis is mostly observed in adult humans, this condition begins early in life, and progresses through adulthood <sup>7,8</sup>. Children with chronic inflammation diseases have shown to develop atherosclerosis in early childhood <sup>9,10</sup>.
- Sex applicability: Unspecific, atherosclerosis is manifested in males and females <sup>11</sup>.

## Key Event Description

Atherosclerosis is defined as the accumulation of fatty and fibrous material in the intima layer of arteries <sup>1</sup>. Atherosclerosis initiates with the alteration of the endothelium homeostasis and accumulation of modified low density lipoproteins (LDL) in the intima layer. The activation of endothelial cells leads to the recruitment and translocation of monocytes to the intima, where monocytes differentiate into macrophages. Following this, macrophages internalize oxidized LDL becoming foam cells. Several factors enhance plaque progression including continuous accumulation of foam cells and lipoproteins, and migration and proliferation of smooth muscle cells in the intima layer. Extracellular matrix macromolecules produced by smooth muscle cells results in the thickening of the intima. A necrotic core rich in lipids inside the atherosclerotic plaque is formed when macrophages and smooth muscle cells go through apoptosis. During progression, atherosclerotic plaques can develop calcification regions and expand toward the lumen of the artery. Advanced atherosclerosis can diminish the arteries lumen and/or form a thrombus (i.e. blood clot), reducing the blood flow, and leading to ischemia. The rupture of atherosclerotic plaques can also lead to embolism <sup>1,2</sup>.

## How it is Measured or Detected

Atherosclerosis can be detected through direct and indirect methods. Techniques that allow direct visualization of atherosclerotic plaques include ultrasonography, computed tomography angiography, magnetic resonance imaging, and optical coherence tomography <sup>1</sup>. These techniques can measure the intima thickness of arteries, along with detection of calcified components <sup>3,4</sup>. Techniques that allows the evaluation of atherosclerosis without direct visualization of plaques include angiography, aortic pulse wave velocity and the ankle-arm systolic blood pressure index <sup>1,3,5</sup>. Finally, although nonspecific, inflammatory markers are also use to evaluate atherosclerosis including blood levels of IL-6, CRP and TNF- $\alpha$  <sup>5</sup>.

The induction and/or progression of atherosclerosis after exposure to a stressor can be studied through animal models. Examples of these are the *ApoE*<sup>-/-</sup> and *LdLr*<sup>-/-</sup> mouse models and Watanabe rabbit model, where the development of atherosclerotic can be assessed <sup>6</sup>.

## Regulatory Significance of the AO

Atherosclerosis is the principal cause of cardiovascular diseases including myocardial infarction, stroke and angina pectoris <sup>1,2,12</sup>.

In turn, cardiovascular diseases are the principal cause of deaths worldwide and measures have been made by many countries to control risk factors and prevent this disease.<sup>13</sup>.

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## Appendix 2

### List of Key Event Relationships in the AOP

#### List of Adjacent Key Event Relationships

[Relationship: 1702: Interaction with the lung cell membrane leads to Increased proinflammatory mediators](#)

#### AOPs Referencing Relationship

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with the pulmonary resident cell membrane components leading to pulmonary fibrosis</a>	adjacent	Moderate	Moderate
<a href="#">Interaction with lung resident cell membrane components leads to lung cancer</a>	adjacent	Moderate	Moderate
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	adjacent	High	Low

#### Key Event Relationship Description

Innate immune response is the first line of defence in any organism against invading infectious pathogens and toxic substances. It involves tissue triggered startle response to cellular stress and is described by a complex set of interactions between the toxic

stimuli, soluble macromolecules and cells (reviewed in Nathan, 2002). The process culminates in a functional change defined as inflammation, purpose of which is to resolve infection and promote healing. In lungs, the interaction of toxic substances with resident cells results in cellular stress, death or necrosis (Pouwels et al., 2016) leading to release of intracellular components such as alarmins (Damage associated molecular patterns [DAMPs], Interleukin (IL)-1 $\alpha$ , High mobility group box 1 [HMGB1]). Released alarmins (danger sensors) bind cell surface receptors such as Interleukin 1 Receptor 1 (IL-1R1), Toll Like Receptors (TLRs) or others leading to activation of innate immune response signalling.

For example, binding of IL-1 $\alpha$  to IL-1R1 can release Nuclear factor kappa B (NF- $\kappa$ B) resulting in its translocation to nucleus and transactivation of pro-inflammatory genes including cytokines, growth factors and acute phase genes. The signalling also stimulates secretion of a variety of pro-inflammatory mediators. Overexpression of IL-1 $\alpha$  in cells induces increased secretion of pro-inflammatory mediators. Products of necrotic cells are shown to stimulate the immune system in an IL-1R1-dependent manner (Chen et al., 2007).

The secreted alarmins activate resident cells pre-stationed in the tissues such as mast cells or macrophages leading to propagation of the already initiated immune response by releasing more eicosanoids, cytokines, chemokines and other pro-inflammatory mediators. Thus, secreted mediators signal the recruitment of neutrophils, which are the first cell types to be recruited in acute inflammatory conditions. Neutrophil influx in sterile inflammation is driven mainly by IL-1 $\alpha$  (Rider P, 2011). IL-1 mediated signalling regulates neutrophil influx in silica-induced acute lung inflammation (Hornung et al., 2008). IL-1 signalling also mediates neutrophil influx in other tissues and organs including liver and peritoneum. Other types of cells including macrophages, eosinophils, and lymphocytes are also recruited in a signal-specific manner. Recruitment of leukocytes induces critical cytokines associated with the T helper type 2 immune response, including Tumor necrosis factor alpha (TNF- $\alpha$ ), IL-1 $\beta$ , and IL-13.

## Evidence Supporting this KER

### Biological Plausibility

The biological plausibility of this relationship is high. There is a mechanistic relationship between the MIE (Event 1495) and KE1 (Event 1496) which has been evidenced in a number of both *in vitro* and *in vivo* model systems in response to stressors such as, asbestos, silica, bleomycin, carbon nanotubes, and metal oxide nanoparticles (NPs) (Behzadi *et al.*, 2017; Denholm & Phan 1990; Dostert *et al.*, 2008; Mossman & Churg 1998).

Increased expression of IL-1 $\alpha$  or IL-1 $\beta$  following lung exposure to multi-walled carbon nanotubes (MWCNTs), bleomycin, micro silica particles, silica crystals, and polyhexamethylene guanidine phosphate has been shown to be associated with neutrophil influx in rodents (Gasse *et al.*, 2007; Girtsman *et al.*, 2014; Hornung *et al.*, 2008; Nikota *et al.*, 2017; Rabolli *et al.*, 2014; Suwara *et al.*, 2014). Inhibition of IL-1 function by knocking out the expression of IL-1R1 using IL-1R1 knockout mice or via treatment with IL-1 $\alpha$  or IL-1 $\beta$  neutralising antibodies results in complete abrogation of lung neutrophilic influx following exposure to MWCNTs (Nikota *et al.*, 2017), cigarette smoke (CS) (Halappanavar *et al.*, 2013), silica crystals (Rabolli *et al.*, 2014) and bleomycin (Gasse *et al.*, 2007). IL-1R1, Myeloid differentiation primary response protein (Myd88) or the IL-33/ST2 signaling are involved in pulmonary fibrosis induced by bleomycin (Gasse *et al.*, 2007; Xu *et al.*, 2016).

### Empirical Evidence

Empirical support for this KER is moderate. There are limited *in vitro* studies, which show a temporal and dose-dependent relationship between these two events, using the upregulation of specific surface receptors as a proxy for direct membrane interaction (Chan *et al.*, 2018; Denholm & Phan, 1990; Roy *et al.*, 2014). There are also studies that provide general support for the idea that an interaction with the lung resident cell membrane components leads to increased, secretion of pro-inflammatory and pro-fibrotic mediators ([Table 1](#)).

### Dose-Response Evidence:

There are a few studies which provide evidence for a dose-response relationship in this KER. An *in vitro* study demonstrated a concentration-response relationship, in which silica exposure induced increases in pro-inflammatory cytokines through scavenger receptors in cultured bone marrow-derived murine mast cells. Cells were exposed to 6.25, 12.5, 25 or 50  $\mu\text{g}/\text{cm}^2$  silica dioxide (SiO<sub>2</sub>) for 24 h. Macrophage scavenger receptor (MSR2) expression increased over time at 50  $\mu\text{g}/\text{cm}^2$  and in a concentration-dependent relationship. Moreover, Tumor necrosis factor alpha (TNF- $\alpha$ ), IL-13 and Monocyte chemoattractant protein-1 (MCP-1) increased in a concentration-dependent manner (Brown *et al.*, 2007). This provides indications that at higher concentrations of the stressor, the interaction with the lung resident cell membrane components (Event 1495) leads to an increased secretion of pro-inflammatory mediators (Event 1496).

### Temporal Evidence:

*In vitro* and *in vivo* studies have demonstrated temporal concordance of the KEs.

TLR4 signal pathway was evaluated in differentiated macrophages exposed to silica at 2.5  $\mu\text{g}/\text{cm}^2$ . After 16 and 24 h, the mRNA expression level of TLR4 increased. Moreover, the protein expression level of TLR-4 and related MyD88/Toll-interleukin-1 receptor domain containing adaptor protein (TIRAP) pathway increased at 24 h. Release of IL-1 $\beta$ , IL-6, IL-10, and TNF- $\alpha$  was induced by silica exposure at 24 h. Pre-treatment with resatorvid (TAK-242), an inhibitor of TLR4 signaling, suppressed the release of the cytokines (Chan *et al.*, 2018).

Macrophages exposed to zinc oxide (ZnO) NPs at 2.5 µg/mL for 24 h increased the expression level of TLR6 and MyD88, TNF receptor-associated factor (TRAF), and IL-1 receptor-associated kinase (IRAK). At 24 h, they also observed an increase in the mRNA and protein levels of the pro-inflammatory cytokines IL-1β, IL-6, and TNF-α. These results demonstrated that ZnO NPs induced pro-inflammatory mediators by TLR stimulation and Mitogen-activated protein kinases (MAPKs) activation (Roy et al., 2014).

The pro-inflammatory IL-1β induced granulocyte migration and can be produced as a result of cellular detection of pathogen associated molecular patterns (PAMPs). Mice exposed to 2.5 mg/mouse of silica by instillation showed an increase of mRNA expression of pro-IL-1β in bronchoalveolar lavage fluid (BALF) at 6, 12, and 24 h post-exposure in a time-dependent manner. At early time points (1 h, 3 h, 6 h), there was an increase in the release of an alarmin (IL-1α) which indicates that the alarmin was released due to cell damage leading to cytokine production and an inflammatory reaction. Moreover, at 24 h, the levels of mature IL-1β and neutrophil accumulation in BALF increased. Neutralization or deletion of IL-1α reduced the observed responses (Rabolli et al., 2014).

Epithelial damage can lead to the release of alarmins. In this stead, conditioned media from primary human bronchial epithelial cells (PBECS) exposed to thapsigargin was able to induce a pro-inflammatory response in primary human lung fibroblasts. PBECS were exposed to thapsigargin (a tumor promoter in mammalian cells) 20 µM for 2 h. After that, the cell culture medium was replaced, and cells were incubated for 24 h. At this time, the medium was recovered and used to culture lung fibroblast for 5 h. This conditioned media from epithelial cell damage contains the alarmin IL-1α, which induced increased gene expression of IL-6, IL-8, MCP-1, and Granulocyte-macrophage colony-stimulating factor (GM-CSF) in fibroblasts. These responses were reduced with anti-IL-1α treatment (Suwara et al., 2014).

Heijink et al. 2015 conducted a similar strategy to identify the relationship between DAMPs and pro-inflammatory mediator release after exposure to CS. Neutrophils treated with CS bubbled for 1 min, released high levels of HMGB1 as a consequence of necrotic cell death. The cell-free supernatant, which contains HMGB1, was used to culture human bronchial epithelial cells, and after 24 h it promoted the production of the C-X-C motif chemokine ligand (CXCL)8 or IL-8 by lung epithelial cells. Pharmacological inhibitors, such as 1-palmitoyl-2-arachidonyl-sn-glycero-3-phosphorylcholine (OxPAPC) and Receptor for advanced glycation endproducts (RAGE) antagonist peptide (RAP), reduced the effect of CXCL8 release.

HMGB1 and Heat shock protein 70 (HSP-70) can be released by damaged hepatocytes. In a study, mice were treated with acetaminophen 350 mg/Kg for 3 and 6 h. At these time points, the liver perfusate was obtained and an increase in HSP-70 and HMGB1 protein levels was observed. RAW 264.7 cells (a macrophage cell line) treated with the liver perfusate exhibited increased mRNA expression levels of MCP-1 and IL-1β (Martin-Murphy et al. 2010).

Female mice were intratracheally administered with bleomycin at 5 mg/kg to represent idiopathic pulmonary fibrosis. IL-33, a molecule that can act as a DAMP, increased in lungs after 3 and 7 days of treatment. In serum, at 7-, 14- and 28-days post-exposure, IL-4 and IL-13 increased. It was concluded that IL-33/ST2 signaling pathway is involved in pulmonary fibrosis by bleomycin (Xu et al., 2016).

### Uncertainties and Inconsistencies

Attenuation or complete abrogation of KE1 (Event 1496) and KE2 (Event 1497) following inflammogenic stimuli is observed in rodents lacking functional IL-1R1 or other cell surface receptors that engage innate immune response upon stimulation. However, following exposure to MWCNTs, it has been shown that absence of IL-1R1 signalling is compensated for eventually and neutrophil influx is observed at a later post-exposure time point (Nikola et al., 2017). In another study, acute neutrophilic inflammation induced by MWCNTs was suppressed at 24 h in mice deficient in IL-1R1 signalling; however, these mice showed exacerbated neutrophilic influx and fibrotic response at 28 days post-exposure (Girtsman et al., 2014). The early defence mechanisms involving DAMPs is fundamental for survival, which may necessitate activation of compensatory signaling pathways. As a result, inhibition of a single biological pathway mediated by an individual cell surface receptor may not be sufficient to completely abrogate the lung inflammatory response. Forced suppression of pro-inflammatory and immune responses early after exposure to substances that cannot be effectively cleared from lungs, may enhance the injury and initiate other pathways leading to exacerbated response.

### Quantitative Understanding of the Linkage

A majority of the *in vivo* studies are conducted with only one dose and thus, it is difficult to derive quantitative dose-response relationships based on the existing data. However, it is clear from the studies referenced above that greater concentrations or doses of pro-fibrotic substances result in higher release of alarmins, and consequently, higher pro-inflammatory signalling. The above studies also demonstrate strong temporal relationships between the individual KEs.

### Response-response relationship

One study has demonstrated a response-response relationship for this KER.

Human intervertebral disc cells were treated with 0, 0.5, 1, or 2 mg/ml of recombinant HMGB1 for 24 h. Protein levels were determined in cell medium supernatant by enzyme-linked immunosorbent assay (ELISA). HMGB1 stimulates the expression of IL-6 and Matrix metalloproteinase 1 (MMP-1) in a response-response relationship. A strong correlation was observed by Spearman's rank correlation coefficient between HMGB1 treatment and IL-6 or MMP-1 levels (Shah et al., 2019).

Other reports have studied both KEs, but they do not indicate if the response-response relationship was linear or not (coefficient or

correlation is not shown) (Chakraborty et al., 2017; Fukuda et al. 2017; Kim et al., 2020; Piazza et al., 2013; Yang et al., 2012;).

### Time-scale

Some studies have described how long after a change in the MIE (Event 1495; interaction substance and components), KE1 (Event 1496; pro-inflammatory mediators are secreted) is impacted (Table 2).

Table 2. Time-scale related studies relevant to the MIE (Event 1495) - KE1 (Event 1496) relationship.

Reference	<i>In vitro/in vivo</i> /population study	Design	MIE (Event 1495)	KE1 (Event 1496)
			Timepoint	Timepoint
Xu et al., 2016	<i>In vivo</i>	40 Female Kunming strain mice  Bleomycin was intratracheally administered 5 mg/Kg.  Days post-exposure	IL-33  3, 7 days	IL-4, IL-13  7, 14, and 28 days
Roy et al., 2014	<i>In vitro</i>	Primary mice macrophages exposed to 2.5 mg/ml ZnO for 24 hrs.	Increased TLR6 expression  0.5, 3, 6, 12, and 24 h	Increased IL-6, TNF- $\alpha$  24 h
Rabolli et al., 2014	<i>In vivo</i>	Female C57BL/6 mice  Exposed to silica 2.5 mg/mouse by instillation	Increased the release of IL-1 $\alpha$  1, 3, and 6 h	Increased mRNA expression of pro-IL-1 $\beta$  6, 12, and 24 h

### Known Feedforward/Feedback loops influencing this KER

Pancreatic cancer cells stimulated with S100 calcium-binding protein A8 (S100A8) and S100 calcium-binding protein A9 (S100A9) released pro-inflammatory cytokines IL-8, TNF- $\alpha$ , and Fibroblast growth factor (FGF). Cancer cell-derived conditioned media and the individual cytokines (TNF- $\alpha$  and Transforming growth factor beta [TGF- $\beta$ ]) induced the protein expression of S100A8 and S100A9 in HL-60 monocytic cell line and primary human monocytes (Nedjadi et al. 2018).

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### Relationship: 2053: Increased proinflammatory mediators leads to Increased transcription of APP

#### AOPs Referencing Relationship

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	adjacent	High	Moderate

#### Evidence Supporting Applicability of this Relationship

##### Taxonomic Applicability

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

##### Life Stage Applicability

Life Stage	Evidence
All life stages	High

##### Sex Applicability

Sex	Evidence
Male	High
Female	High

Acute phase response is present in vertebrate species <sup>14</sup>. In addition, serum amyloid A, one of the major acute phase proteins, has been conserved in mammals throughout evolution and has been described in humans, mice, dogs, horses, among others <sup>4</sup>.

#### Key Event Relationship Description

This KER presents the association between the secretion of pro-inflammatory mediators and transcription of acute phase protein in different tissues, mainly lungs and liver. The evidence of the KER presented is based on animal studies (mice), human studies and *in vitro* studies.

#### Evidence Supporting this KER

##### Biological Plausibility

The biological plausibility is high. It is known that acute phase proteins are induced by pro-inflammatory cytokines, primary interleukin (IL)-6, IL-1 $\beta$ , and tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ). These cytokines are produced at sites of inflammation mainly by monocytes and macrophages <sup>1-4</sup>. Following cytokine release, signaling cascades and transcription factors are activated, regulating the expression of acute phase reaction genes <sup>3</sup>.

Immune cells are recruited to inflammatory sites by inflammatory mediators (i.e. cytokines and chemokines)<sup>5</sup>. Pulmonary inflammation in mice is commonly assessed as the number or fraction of neutrophils in the bronchoalveolar lavage fluid (BALF) <sup>6</sup> and can be used as indirect marker of the release of pro-inflammatory factors.

##### Empirical Evidence

- IL-1 (IL-1 $\alpha$  and IL-1 $\beta$ , 10 ng/mL each) and IL-6 (500 units/mL), both in presence of 1 $\mu$ M dexamethasone, increased the relative levels of serum amyloid a (SAA) mRNA in cultured human adult aortic smooth muscle cells <sup>7</sup>.
- Human hepatoma cells exposed to IL-6, IL-1 $\beta$  and TNF- $\alpha$  for 20 h showed a reduced synthesis of albumin and increased synthesis of the acute phase proteins C3 and ceruloplasmin. In addition, mice exposed to IL-1 $\beta$  and TNF- $\alpha$  showed an increase of Saa mRNA in liver tissue <sup>8</sup>.
- After pulmonary exposure to lipopolysaccharide (LPS) (300  $\mu$ g/mL), lung tissue from female C57BL/6 mice showed upregulation of several cytokines and chemokines genes and upregulation of the acute phase proteins genes serum amyloid A and  $\alpha_1$ -protease inhibitor <sup>9</sup>.

- Mice presenting IL-6 gene disruption (IL-6<sup>-/-</sup>) shown a reduced response in liver mRNA levels of acute phase proteins haptoglobin,  $\alpha$ 1-acid glycoprotein and SAA, after challenged by turpentine, lipopolysaccharide and bacterial infection<sup>10</sup>.
- After repeated instillation of carbon black nanoparticles, female C57BL/6BomTac mice showed increased expression of chemokine genes along with increased *Saa3* gene expression in lung tissue. In addition, there were dose-response relationships with several cytokine proteins in lung tissue<sup>11</sup>.
- Intratracheal instillation of titanium dioxide in female C57BL/6 showed that 28 days after exposure, several genes of cytokines, chemokines and acute phase proteins were upregulated. Additionally, there were significant increases in inflammatory mediators in lung tissue<sup>12</sup>.

The table in the following link presents evidence of the relationship using neutrophil numbers in BALF as indirect measurement of the release of pro-inflammatory mediators: [Empirical evidence KER2](#).

## Quantitative Understanding of the Linkage

### Response-response relationship

A Pearson's correlation coefficient of 0.82 ( $p < 0.001$ ) has been calculated between log-transformed neutrophil numbers in bronchoalveolar lavage fluid and log-transformed *Saa3* mRNA levels in lung tissue, in female C57BL/6J mice 1 and 28 days after intratracheal instillation of metal oxide nanomaterials<sup>13</sup> (Figure 1).

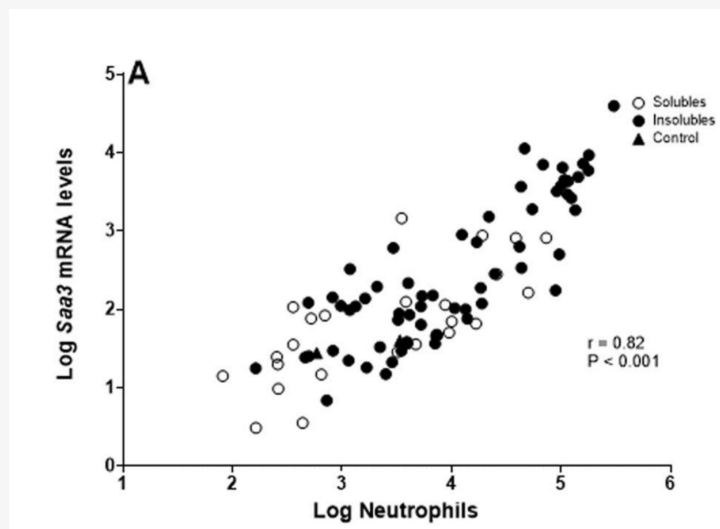


Figure 1. Correlations between neutrophil numbers and *Saa3* mRNA levels in lung tissue, including data from 1 and 28 days after exposure to nanomaterials. Reproduced from Gutierrez et al. (2023)<sup>13</sup>.

### Time-scale

It has been shown that pro-inflammatory mediators concentrations increase before acute phase proteins:

- Upregulation of cytokine genes (IL-1 $\alpha$ , IL-1 $\beta$ , IL-6 and TNF- $\alpha$ ) was shown to peak around 2h after pulmonary exposure to LPS in female C57BL/6J mice, while upregulation SAA genes showed their highest upregulation at 8-12h after exposure<sup>9</sup>.

### Known Feedforward/Feedback loops influencing this KER

Some acute phase proteins (f. ex. C-reactive protein, serum amyloid A and complement components) have pro-inflammatory functions, including induction of inflammatory cytokines, chemotaxis and activation of immune cells. On the other hand, other acute phase proteins present anti-inflammatory functions (f. ex. Haptoglobin and fibrinogen) as antioxidative and tissue repair inducer<sup>1</sup>.

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### **Relationship: 1589: Increased transcription of APP leads to Systemic APR**

#### **AOPs Referencing Relationship**

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	adjacent	High	Moderate

#### **Evidence Supporting Applicability of this Relationship**

##### **Taxonomic Applicability**

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

##### **Life Stage Applicability**

Life Stage	Evidence
All life stages	High

##### **Sex Applicability**

Sex	Evidence
Male	High
Female	High

Acute phase response is present in vertebrate species <sup>17</sup>. In addition, serum amyloid A, one of the major acute phase proteins, has been conserved in mammals throughout evolution and has been described in humans, mice, dogs, horses, among others <sup>18</sup>.

## Key Event Relationship Description

This KER presents the association between the transcription of acute phase protein genes in different tissues and induction of systemic acute phase response. The evidence of the KER presented is based on animal studies (mice).

## Evidence Supporting this KER

### Biological Plausibility

The biological plausibility is high. After gene expression of acute phase proteins in tissues during inflammatory conditions, mRNA is translated and folded into proteins <sup>1</sup>. These proteins are then release to the systemic circulation <sup>2</sup>.

### Empirical Evidence

Species	Stressor	Acute phase protein expression	Systemic acute phase response	Reference
Mouse	Carbon black nanoparticles	Yes, significant <i>Saa1</i> , <i>Saa2</i> and <i>Saa3</i> gene expression increase in lung tissue, at days 1, 3 and 28 after exposure. <i>S a a 3</i> gene expression increase in liver tissue at day 1 after exposure.	Yes, significant increase of plasma SAA at 1 and 28 days after exposure.	<sup>3</sup>
Mouse	Multiwalled carbon nanotubes (referred as CNT <sub>small</sub> )	Yes, increased differential expression of acute phase response genes in liver tissue 1 and 3 days after exposure to 162 µg. Increased differential expression of acute phase response genes in lung tissue 3 days after exposure to 18 and 162 µg, and 1 and 3 days after exposure to 54 µg.	Yes, increased plasma SAA3 1, 3 and 28 days after exposure to 162 µg, and 3 days after exposure to 18 and 54 µg.	<sup>4,5</sup>
Mouse	Multiwalled carbon nanotubes (referred as CNT <sub>large</sub> )	Yes, increased differential expression of acute phase response genes in liver tissue 1 and 3 days after exposure to 162 µg. Increased differential expression of acute phase response genes in lung tissue 1 and 3 days after exposure to 54 and 162 µg.	Yes, increased plasma SAA3 1 and 3 days after exposure to 162 µg, and 3 days after exposure to 54 µg.	<sup>4,5</sup>
Mouse	Graphene oxide	Yes, increased mRNA expression of <i>Saa3</i> in lung tissue, at all dose levels 1 and 3 days after exposure. Increased gene expression of <i>Saa1</i> in liver tissue 1 day after exposure to 18 µg, and 3 days after exposure to 162 µg.	Yes, increased SAA3 plasma levels 3 days after exposure to 54 and 162 µg.	<sup>6</sup>
Mouse	Reduced graphene oxide	Yes, increased mRNA expression of <i>Saa3</i> in lung tissue, 3 days after exposure to 162 µg. No changes in gene expression of <i>Saa1</i> in liver tissue.	No, no change in SAA3 plasma concentration 3 days after exposure.	<sup>6</sup>
Mouse	Multiwalled carbon nanotubes (NM-400 to NM-403)	Yes, increased <i>Saa1</i> mRNA expression in liver tissue with all MWCNTs, 1 day after exposure to 54 µg, and after exposure to 18 µg in the case of NM-401 and NM-403. After 28 days, only NM-400 (54 µg) produced an increase in <i>Saa1</i> mRNA levels in liver tissue. Increased <i>Saa3</i> mRNA expression in lung tissue with all MWCNTs, 1 day after exposure to 54 µg, and after exposure to 6 and 18 µg in the case of NM-402 and NM-403.	Yes, increased SAA1/2 plasma levels 1 day after exposure to NM-400, NM-401 and NM-403. No change in SAA1/2 28 and 92 days after exposure. Increased SAA3 plasma levels 1 days after exposure to all MWCNT. Increased SAA3 plasma levels 28 and 92 days after exposure to NM-401.	<sup>7</sup>

Species	Stressor	Acute phase (18-28 days) NM-400 (18 µg) and NM-403 (54 µg) produced an increase in <i>Saa3</i> mRNA levels in lung tissue.	Systemic response	acute phase	Reference
Mouse	Particulate matter from non-commercial airfield	Yes, increased expression of <i>Saa3</i> mRNA in lung tissue and <i>Saa1</i> mRNA in liver tissue after 1 day of exposure to 54 µg.  No effect after 28 and 90 days.	Yes, increased plasma SAA3 levels after exposure to 54 µg after 3 days.	8	
Mouse	Particulate matter from commercial airport	Yes, increased expression of <i>Saa3</i> mRNA in lung tissue after 1 day of exposure to 18 and 54 µg.  No effect after 28 and 90 days.	No change in plasma SAA3.	8	
Mouse	Diesel exhaust particles	Yes, increased expression of <i>Saa3</i> mRNA in lung tissue after 1 day of exposure to 54 and 162 µg, and increased expression of <i>Saa1</i> mRNA in liver tissue 1 day after exposure to 162 µg. No effect after 28 days.	Yes, increased plasma SAA3 levels after exposure to 54 µg, at 3 days.	8	
Mouse	Carbon black	Yes, increased expression of <i>Saa3</i> mRNA in lung tissue at day 1 and day 90.	No change in plasma SAA3.	8	
Mouse	Uncoated zinc oxide nanoparticles	Yes, increase on <i>Saa3</i> mRNA in lung tissue 1 day after exposure to 2 µg. No effect 3 and 28 days after exposure.	No effect on plasma SAA3.	9	
Mouse	Coated zinc oxide nanoparticles	Yes, increase on <i>Saa3</i> mRNA in lung tissue 1 day after exposure to 0.7 and 2 µg. No effect 3 and 28 days after exposure.	No effect on plasma SAA3.	9	
Mouse	Zinc oxide	Yes, increased <i>Saa1</i> mRNA expression in liver tissue 1 day after exposure to 0.7 µg. No change in <i>Saa3</i> mRNA expression in lung tissue.	No change in plasma SAA3 or SAA1/2 levels.	10	
Mouse	Copper oxide	Yes, increased <i>Saa3</i> mRNA expression in lung tissue 1 day after exposure to 2 and 6 µg. Increased <i>Saa1</i> mRNA expression in liver tissue 1 day after exposure to 6 µg.	Yes, increased plasma SAA1/2 level after exposure to 6 µg, 1 day after exposure.	10	
Mouse	Tin dioxide	Yes, increased <i>Saa3</i> mRNA expression in lung tissue and <i>Saa1</i> mRNA expression in liver tissue, 1 day after exposure to 162 µg.	Yes, increased plasma SAA3 after exposure to 162 µg, 1 day after exposure.	10	
Mouse	Titanium dioxide	Yes, increased <i>Saa3</i> mRNA expression in lung tissue and <i>Saa1</i> mRNA expression in liver tissue 1 day after exposure.	Yes, increased plasma SAA3 and SAA1/2 after exposure to 162 µg, 1 day after exposure.	10	
Mouse	Carbon black	Yes, increased <i>Saa3</i> mRNA expression in lung tissue 1 and 28 days after exposure. Increased <i>Saa1</i> mRNA expression in liver tissue 1 day after exposure	Yes, increased plasma SAA3 and SAA1/2 after exposure to 162 µg, 1 day after exposure.	10	
Mouse	Singlewalled carbon nanotubes	Yes, increased SAA1, SAP and haptoglobin gene expression in liver tissue, 1 day after exposure.	Yes, increase serum CRP, haptoglobin and SAP 1 day after exposure.	11	
Mouse	Multiwalled carbon nanotubes	Yes, increased SAA1, SAP and haptoglobin gene expression in liver tissue, 1 day after exposure.	Yes, increase serum CRP, haptoglobin and SAP 1 day after exposure. No changes after 28 days.	11	
Mouse	Serum amyloid A	Yes, significantly increase of <i>Saa3</i>	Yes, increased levels of	12	

Species	Stressor	Acute phase protein	Systemic SAA	Acute phase	Reference
Uncertainties and Inconsistencies		Acute phase protein levels in lung tissue and SAA3 protein levels in liver tissue.	Systemic SAA response		

Although it is suggested that acute phase proteins are mainly produced in the liver<sup>13</sup>, it has been shown that in mice, the liver has little upregulation of *Saa* genes after exposure to ultrafine carbon particles or diesel exhaust particle, while it is in the lung where there is a marked expression of *Saa3* mRNA<sup>14,15</sup>.

It has been observed in some studies that the increase of *Saa* genes in lung or liver tissue does not translate into an increase in plasma SAA concentration<sup>6,8,9</sup>. This might be due to a protein concentration below the methods detection levels<sup>9</sup>, while measuring gene expression provides a larger dynamic range.

## Quantitative Understanding of the Linkage

### Response-response relationship

A Pearson's correlation coefficient of 0.89 ( $p < 0.001$ ) has been calculated between log-transformed *Saa3* mRNA levels in lung tissue and log-transformed SAA3 plasma protein levels, in female C57BL/6J mice 1 day after intratracheal instillation of metal oxide nanomaterials<sup>10</sup> (Figure 1).

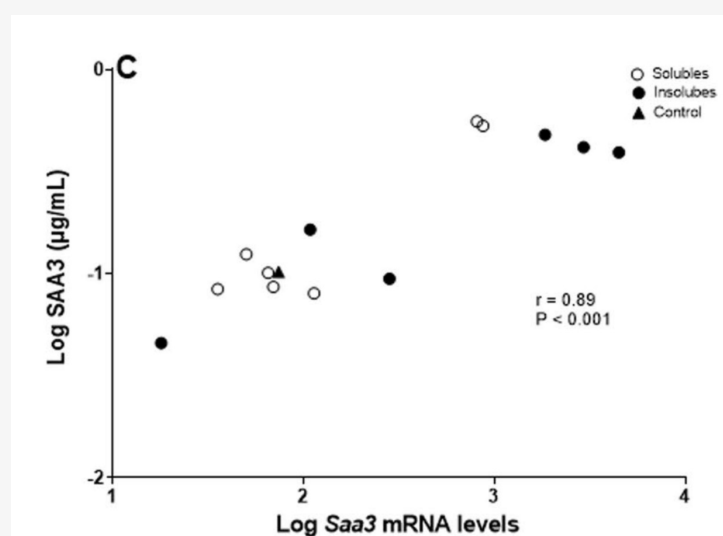


Figure 1. Correlations between *Saa3* mRNA levels in lung tissue and SAA3 plasma protein levels, including data from 1 day after exposure to nanomaterials. Reproduced from Gutierrez et al. (2023)<sup>10</sup>.

### Time-scale

After exposure to titanium dioxide nanoparticles in mice, expression of *Saa1* mRNA in the liver is short lasting, while expression of *Saa3* mRNA in lung tissue is longer lasting, as it has been observed 28 day after exposure<sup>16</sup>.

After exposure to multiwalled carbon nanotubes, it has been observed that expression of *Saa1* and *Saa3* in liver and lung tissue can be elevated 28 days after exposure, however in most cases there is no increase in plasma SAA1/2 nor SAA3 levels past day 1 after exposure<sup>7</sup>.

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### [Relationship: 2860: Systemic APR leads to Atherosclerosis](#)

#### AOPs Referencing Relationship

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	adjacent	High	High

#### Evidence Supporting Applicability of this Relationship

##### Taxonomic Applicability

Term	Scientific Term	Evidence	Links
human	Homo sapiens	High	<a href="#">NCBI</a>

##### Life Stage Applicability

Life Stage	Evidence
All life stages	High

##### Sex Applicability

Sex	Evidence
Male	High
Female	High

Although atherosclerosis is mostly observed in adult humans, this condition begins early in life, and progresses through adulthood<sup>23,24</sup>. Children with chronic inflammation diseases have shown to develop atherosclerosis in early childhood.<sup>25,26</sup> In addition, atherosclerosis is manifested in males and females<sup>27</sup>.

### Key Event Relationship Description

This KER presents the association between systemic acute phase response and atherosclerosis as adverse outcome. The evidence of the KER presented is based on animal studies (mice), epidemiological studies and *in vitro* studies.

### Evidence Supporting this KER

#### Biological Plausibility

The biological plausibility is high. During acute phase response, serum amyloid A (SAA), one of the major acute phase proteins, replaces Apolipoprotein A-1 from high density lipoprotein (HDL). This replacement obstructs the reverse transport of cholesterol to the liver, allowing the accumulation of cholesterol in cells, denominated foam cells<sup>1-3</sup>. Foam cells are early markers of atherosclerotic lesions<sup>4</sup>, and it has been shown that macrophages have a higher uptake of HDL containing SAA than HDL alone<sup>2</sup>.

The two major human acute phase response, SAA and C-reactive protein (CRP), have been shown to be correlated in humans<sup>5-7</sup>, and both are predictors of future cardiovascular event risks<sup>7</sup>.

#### Empirical Evidence

- SAA was moderately associated with angiographic coronary artery disease in women (21-86 years old) suspected on having myocardial ischemia<sup>8</sup>.
- High levels of CRP were associated with an increased risk of coronary heart disease in men and women<sup>9</sup>.
- In mouse model of periodontal disease, ApoE<sup>-/-</sup> mice were infected with a polymicrobial consortium (*Porphyromonas gingivalis*, *Treponema denticola*, and *Tannerella forsythia*). 16-weeks after infection, the infected mice presented elevated levels of SAA in comparison to control mice, in addition of increased plaque progression<sup>10</sup>.
- Male ApoE<sup>-/-</sup> mice overexpressing SAA1 presented higher levels of plasma SAA and an increase in atherosclerotic lesions (plaques) than non-SAA1 overexpressing ApoE<sup>-/-</sup> mice<sup>11</sup>.
- After one injection of adenoviral vector encoding human SAA1, ApoE<sup>-/-</sup> mice presented elevated and transient levels of human SAA along with an increase in atherosclerotic lesions<sup>12</sup>.
- Overexpression of SAA3 led to increased levels of SAA3 and atherosclerosis lesions in ApoE<sup>-/-</sup> mice in comparison to control mice. In addition, when SAA3 was suppressed in ApoE<sup>-/-</sup> × SAA1.1/2.1-DKO (ApoE<sup>-/-</sup> mice deficient in SAA1 and SAA2), there was a significant decreases in atherosclerotic lesions<sup>13</sup>.
- In an *in vitro* study, Increasing concentrations of SAA (0 – 2 µM) produced a dose-response relationship of foam cells in RAW264.7 cells<sup>14</sup>.
- Intratracheal instillation of human serum amyloid A once a week for 10 weeks in ApoE<sup>-/-</sup> mice (on Western-type diet) induced an increase in plasma SAA3 and atherosclerotic plaque progression<sup>15</sup>.

#### Uncertainties and Inconsistencies

Mendelian randomization studies have shown that CRP genotypes are not associated with risk of coronary heart disease and that genetically elevated levels of CRP are not associated with coronary heart disease risk<sup>16,17</sup>.

### Quantitative Understanding of the Linkage

#### Response-response relationship

The association between CRP and SAA levels and risk of nonfatal myocardial infarction or fatal coronary heart disease (i.e. acute events due to the progression of atherosclerosis) can be calculated from prospective, epidemiological studies<sup>7,9</sup>. This approach was used by the Dutch Expert Committee on Occupational Safety (DECOS) when establishing a health-based occupational exposure limit for diesel engine exhaust based on risk of lung cancer (<https://www.healthcouncil.nl/documents/advisory-reports/2019/03/13/diesel-engine-exhaust>).

The Nurses' Health Study (NHS) and the Health Professionals Follow-up Study (HPFS) are prospective cohort investigations respectively involving 121,700 female U.S. registered nurses who were 30 to 55 years old at baseline in 1976 and 51,529 U.S. male health professionals who were 40 to 75 years old at baseline in 1986<sup>9</sup>. In the NHS, among women without cardiovascular disease or cancer before 1990, 249 women had a nonfatal myocardial infarction or fatal coronary heart disease between the date of blood drawing and follow-up in June 1998. In the HPFS, 266 men had a nonfatal myocardial infarction or fatal coronary heart disease between the date of blood drawing and the return of a follow-up questionnaire in year 2000.

In the NHS and HPFS studies, the associations between CRP in blood and risk of nonfatal myocardial infarction or fatal coronary

heart disease for women and men were reported in Pai et al. (2004)<sup>9</sup>, whereas the association for both SAA and CRP in NHS was reported in Ridker et al. (2000)<sup>7</sup>.

The dose-response relationships are shown in Figure 1. Here, plasma levels of CRP and SAA were closely associated with future risk of coronary heart disease (CHD).

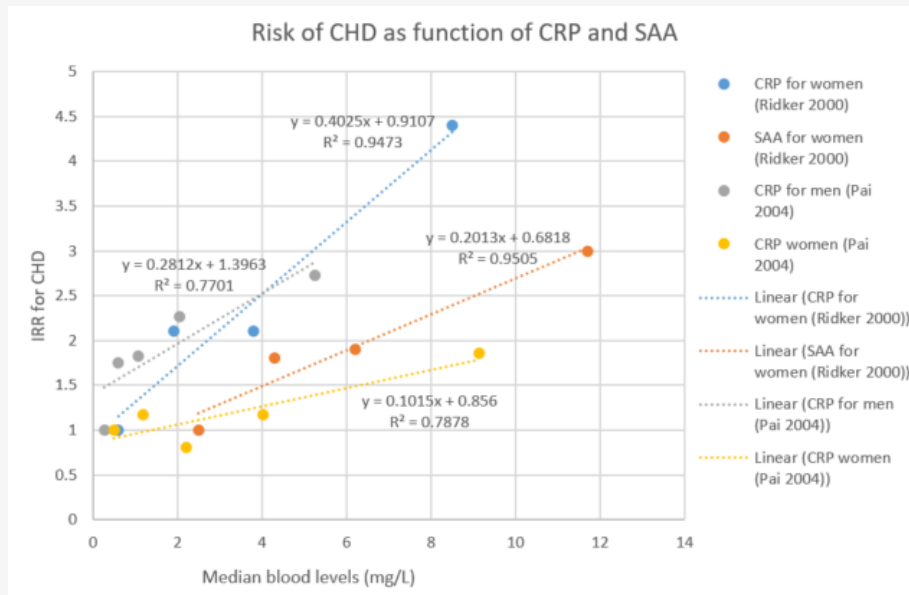


Figure 1. Association between the relative risk (RR) of CHD in NHS as function of quartiles of serum levels of CRP and SAA from Ridker et al.<sup>7</sup> and quintiles of CRP from the NHS and the HPFS studies from Pai et al.<sup>9</sup>. The trend lines are linear associations, as these gave the highest  $R^2$  values.

According to the Danish Heart Foundation (<https://hjertereforeningen.dk/alt-om-dit-hjerte/noegletal/>), when a person reached the age of 55 years, the lifetime risk of a cardiovascular event is 67% in men and 66% in women. 56,379 Danes are diagnosed with a cardiovascular disease each year. Of these, 15,087 were diagnosed with are apoplexy and 16,050 with ischemic heart disease. These diagnoses are here regarded as manifestations of plaque progression. Thus, 55% of the cardiovascular diagnoses are relate to plaque progression. The lifetime risk of these diseases is thus  $0.66 \times 0.55 = 0.363 = 36\%$ .

The relative risk of 1:100 excess cardiovascular disease was calculated as

$$RR = (1 + 36)/36 = 1.02778$$

The relative risk of 1:1000 excess cardiovascular disease was calculated as

$$RR = (1 + 360)/360 = 1.00278$$

If the relative risk of 1.02778 excess is used in the equations obtained in Figure 1 and presented in the next table, it is observed that in the studies by Ridker et. al and Pai et al., 6-54% increases in blood levels of CRP or SAA were associated with 1% increased risk of cardiovascular disease.

Biomarker	Equation of increased IRR	Increase of biomarker associated with 1% increased risk <sup>(1)</sup>	Baseline levels	Increase of biomarker in % of baseline level associated 1% increased risk
CRP women <sup>7</sup>	$\Delta IRR = 0.4025 \text{ CRP (mg/L)}$	0.07 mg/L	0.6 mg/L	$0.07/0.6 = 12\%$
SAA women <sup>7</sup>	$\Delta IRR = 0.2013 \text{ SAA (mg/L)}$	0.138 mg/L	2.5 mg/L	$0.138/2.5 = 6\%$
CRP women <sup>9</sup>	$\Delta IRR = 0.1015 \text{ CRP (mg/L)}$	0.27 mg/L	0.5 mg/L	$0.27/0.5 = 54\%$
CRP men <sup>9</sup>	$\Delta IRR = 0.2812 \text{ CRP}$	0.099 mg/L	0.27	$0.099/0.27 = 37\%$

Concentration	(mg/L)	Effect	mg/L	Effect
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(1) The biomarker level is calculated as  $0.02778/\text{slope}$ . For example, for CRP level in women  $\text{CRP} = 0.02778/0.4025 = 0.07 \text{ mg/L}$ .

#### Known modulating factors

Modulating factor	Specification	Effects on the KER	References
Life style	High body mass index	Increased level of SAA and CRP, therefore increased risk of atherosclerosis.	<sup>8</sup>
Life style	Smoking	Increased level of CRP, therefore increased risk of atherosclerosis.	<sup>8,18</sup>
Medication	Intake of non-steroidal anti-inflammatory drugs	Reduction of CRP and other pro-inflammatory markers, decrease risk of atherosclerosis.	<sup>4</sup>
Medical conditions	Chronic inflammatory diseases	Increased level of acute phase proteins, therefore increased risk of atherosclerosis.	<sup>19</sup>
Medical conditions	Infectious diseases	Increased levels of CRP, therefore increased risk of atherosclerosis.	<sup>18</sup>

#### Known Feedforward/Feedback loops influencing this KER

Atherosclerosis is an inflammatory condition<sup>20,21</sup>, therefore there are increased levels of pro-inflammatory factors, including acute phase proteins, than can sustain the progression of atherosclerosis<sup>22</sup>.

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## List of Non Adjacent Key Event Relationships

### [Relationship: 2958: Interaction with the lung cell membrane leads to Increased transcription of APP](#)

#### AOPs Referencing Relationship

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	non-adjacent	High	Moderate

## Evidence Supporting Applicability of this Relationship

### Taxonomic Applicability

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

### Life Stage Applicability

Life Stage	Evidence
All life stages	High

### Sex Applicability

Sex	Evidence
Male	High
Female	High

The expression of *Saa* mRNA in lung and liver tissue has been shown in mice after pulmonary exposure to a variety of nanomaterials (see Empirical evidence), and in humans in different tissues as lung, liver and arteries <sup>13,14</sup>.

## Key Event Relationship Description

This KER presents the association between the interaction of stressors with the lungs cells and transcription of acute phase proteins in different tissues, mainly lungs and liver. The evidence of the KER presented is based on animal studies (mice).

## Evidence Supporting this KER

### Biological Plausibility

The biological plausibility is high. Production of acute phase proteins is triggered by cellular pattern-recognition molecules after sensing pathogens, tissue damage or dysmetabolism, through a cytokine cascade <sup>1</sup>. In the lungs, this cytokine cascade is produced by epithelial cells and resident macrophages <sup>2</sup>. In the table below it is shown that a variety of stressors produced an increase in Serum amyloid A (SAA) gene isoforms in mice tissue.

### Empirical Evidence

For this KER, exposure through the respiratory system (inhalation or intratracheal instillation) of stressors is considered as interaction with lung resident cell membrane components. The table in the following link presents evidence of KER: [EMPIRICAL EVIDENCE KER5](#).

### Uncertainties and Inconsistencies

Although it is suggested that acute phase proteins are mainly produced in the liver <sup>3</sup>, it has been shown that in mice, the liver has little upregulation of *Saa* genes after exposure to ultrafine carbon particles or diesel exhaust particle, while it is in the lung where there is a marked expression of *Saa3* mRNA <sup>4,5</sup>.

In the case of nanomaterials, it has been shown that physicochemical characteristics as size, surface area, surface functionalization, shape, composition, among others, affect the magnitude and duration of the expression of acute phase proteins in mice <sup>6-12</sup>.

In humans, measuring gene expression of acute phase proteins is not very common as a tissue sample is needed, while measuring acute phase protein in blood is more common. However, *Saa* mRNA has been shown expressed in different tissues including lung, liver and arteries <sup>13,14</sup>.

## Quantitative Understanding of the Linkage

### Response-response relationship

In the case of some insoluble nanomaterials, it has been observed that log-transformed dosed surface area (dosed mass multiply by specific surface area) and log-transformed *Saa3* mRNA levels in mice lung tissue presented a Pearson's correlation coefficient of 0.70 ( $p < 0.001$ ) 1 day post-exposure. The linear regression formula obtained was  $\text{Log } Saa3\text{mRNA} = 1.080 * \text{Log Dosed surface area} + 0.9415$  ( $p < 0.001$ )<sup>7</sup> (Figure 1).

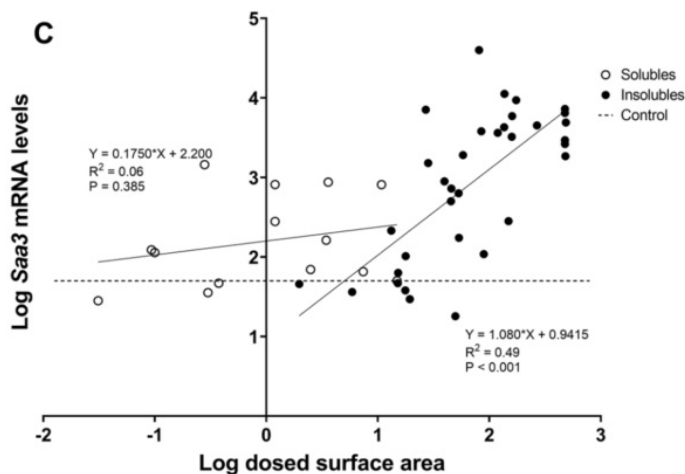


Figure 1. Correlations between dosed surface area and *Saa3* mRNA levels in lung tissue, 1 day after exposure to nanomaterials. Reproduced from Gutierrez et al. (2023)<sup>7</sup>.

### Time-scale

After exposure to titanium dioxide nanoparticles in mice, expression of *Saa1* mRNA in the liver is short lasting, while expression of *Saa3* mRNA in lung tissue is longer lasting, as it has been observed 28 day after exposure<sup>10</sup>.

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### [Relationship: 2959: Interaction with the lung cell membrane leads to Systemic APR](#)

#### AOPs Referencing Relationship

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	non-adjacent	High	Moderate

#### Evidence Supporting Applicability of this Relationship

##### Taxonomic Applicability

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>

##### Life Stage Applicability

Life Stage	Evidence
All life stages	High

##### Sex Applicability

Sex	Evidence
Male	High
Female	High

Systemic acute phase response measured as elevation of CRP and SAA in humans, and SAA in mice has been shown after exposure to several stressors (see Empirical evidence).

#### Key Event Relationship Description

This KER presents the association between the interaction of stressors with the lungs and the induction of systematic acute phase response. The evidence of the KER presented is based on animal studies (mice), controlled human studies and epidemiological studies.

#### Evidence Supporting this KER

##### Biological Plausibility

The biological plausibility is high. Pulmonary inflammation occurs when stressor interact with the airways<sup>1</sup> and acute phase response is induced during inflammatory conditions<sup>2</sup>. It has been shown (see table below) that exposure to different stressors produces an increase of acute phase proteins in blood [i.e. C-reactive protein (CRP) and serum amyloid A (SAA)] in humans and mice.

##### Empirical Evidence

For this KER, exposure through the respiratory system (inhalation or intratracheal instillation) of stressors is considered as interaction with lung resident cell membrane components. The table in the following link presents evidence of KER: [EMPIRICAL EVIDENCE KER6](#).

##### Uncertainties and Inconsistencies

In the case of nanomaterials, it has been shown that physicochemical characteristics as size, surface area, surface functionalization, shape, composition, among others, affect the magnitude and duration of acute phase response in mice<sup>3-5</sup>.

It has been observed that in most controlled human studies, an increase in CRP and/or SAA was observed after exposure to particulate matter<sup>6-10</sup>. However, in other human studies the exposure did not induce acute phase response<sup>11,12</sup>, maybe due to a low level of exposure<sup>13</sup>.

## Quantitative Understanding of the Linkage

### Response-response relationship

In the case of some insoluble nanomaterials, it has been observed that log-transformed dosed surface area (dosed mass multiply by specific surface area) and log-transformed SAA3 plasma levels in mice presented a Pearson's correlation coefficient of 0.92 ( $p < 0.001$ ) 1 day post-exposure<sup>4</sup> (Figure 1). The linear regression formula obtained was  $\text{Log SAA3} = 0.9459 \cdot \text{Log Dosed surface area} - 2.854$  ( $p = 0.01$ ). The correlation coefficient between log-transformed dosed surface area and log-transformed SAA1/2 plasma levels was 0.83 ( $p < 0.05$ ) and the linear regression formula was  $\text{Log SAA1/2} = 0.6368 \cdot \text{Log Dosed surface area} + 0.09524$  ( $p = 0.01$ )<sup>4</sup> (Figure 2).

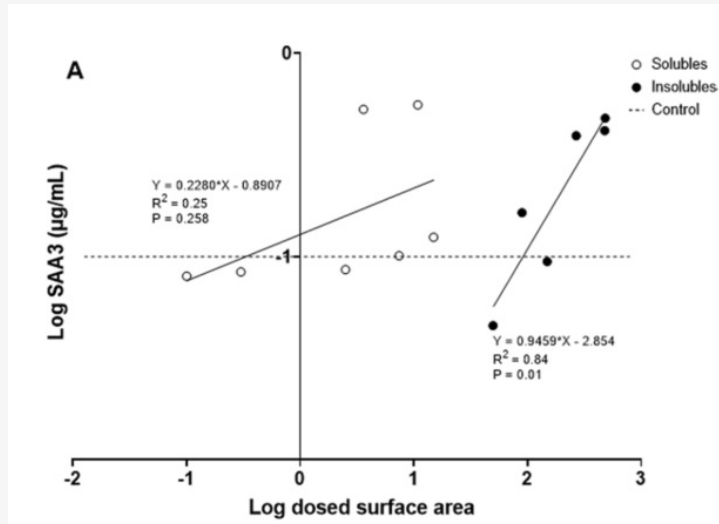


Figure 1. Correlations between pulmonary dosed surface area and SAA3 protein in plasma, 1 day after exposure to nanomaterials. Reproduced from Gutierrez et al. (2023)<sup>4</sup>.

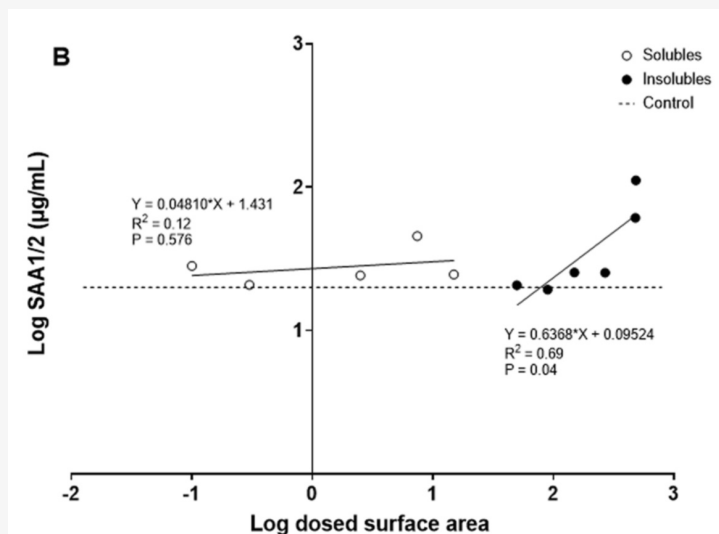


Figure 2. Correlations between pulmonary dosed surface area and SAA1/2 protein in plasma, 1 day after exposure to nanomaterials. Reproduced from Gutierrez et al. (2023)<sup>4</sup>.

### Time-scale

In mice, increased SAA levels are observed 1 and 3 days after most exposures, however increased SAA levels are not frequently observed 28 or 90 days after exposure<sup>3,14-17</sup>.

In humans, increased SAA and CRP has been observed 22h and 2 days after exposure to zinc oxide, but not 3 days after exposure<sup>7</sup>. After exposure to zinc oxide, copper oxide or a mix both, SAA levels were elevated 24h after exposure in humans, but not 6h after exposure<sup>10</sup>.

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### **Relationship: 3052: Increased proinflammatory mediators leads to Systemic APR**

#### **AOPs Referencing Relationship**

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	non-adjacent	High	Moderate

#### **Evidence Supporting Applicability of this Relationship**

#### **Taxonomic Applicability**

Term	Scientific Term	Evidence	Links
mouse	Mus musculus	High	<a href="#">NCBI</a>
human	Homo sapiens	High	<a href="#">NCBI</a>
<b>Life Stage Applicability</b>			
<b>Life Stage</b>	<b>Evidence</b>		
All life stages	High		
<b>Sex Applicability</b>			
<b>Sex</b>	<b>Evidence</b>		
Male	High		
Female	High		

Acute phase response is conserved in vertebrate species <sup>13</sup>.

### Key Event Relationship Description

This KER presents the association between the secretion of pro-inflammatory mediators and induction of systemic acute phase response. The evidence of the KER presented is based on animal studies (mice), controlled human studies and epidemiological studies.

### Evidence Supporting this KER

#### Biological Plausibility

The biological plausibility is high. The production of acute phase proteins during acute phase response is induced by the release of pro-inflammatory markers as interleukin (IL)-6, IL-1 $\beta$ , and tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) at inflammatory sites <sup>1,2</sup>. The release of inflammatory markers also induces the recruitment of immune cells to inflammation sites <sup>3</sup>.

Neutrophils in the bronchoalveolar lavage fluid (BALF) are frequently used to measure pulmonary inflammation in mice <sup>4</sup> and can be utilized as an indirect indicator of the release of pro-inflammatory factors.

#### Empirical Evidence

Evidence of the secretion of pro-inflammatory mediators is presented as change in concentration of pro-inflammatory markers in blood, or increase neutrophil numbers in blood or BALF. The table in the following link presents evidence of KER: [EMPIRICAL EVIDENCE KER8](#).

#### Uncertainties and Inconsistencies

Wyatt et al. observed a decrease in blood neutrophil numbers in humans after exposure to ambient particulate matter although an increase in SAA and CRP was observed. It was mentioned this might be due to the translocation of neutrophil from major vessels to smaller arteries <sup>5</sup>.

In the study by Meier et al., the authors obtained a negative association between PM<sub>2.5</sub> exposure and blood levels of TNF- $\alpha$  and IL-6, while SAA and CRP were positive associated with the exposure. The authors mentioned these results might be due the time point where the samples were taken <sup>6</sup>.

Barregard et al. also observed that IL-6 levels were lower after exposure to wood smoke than after exposure to clean air. The discussed this response as a possible sequestering of cytokines in the pulmonary capillary bed <sup>7</sup>.

### Quantitative Understanding of the Linkage

#### Response-response relationship

A Pearson's correlation coefficient of 0.79 (p<0.001) has been calculated between log-transformed neutrophil number in BALF and log-transformed SAA3 plasma protein levels, in female C57BL/6J mice 1 day after intratracheal instillation of metal oxide nanomaterials <sup>8</sup> (Figure 1).

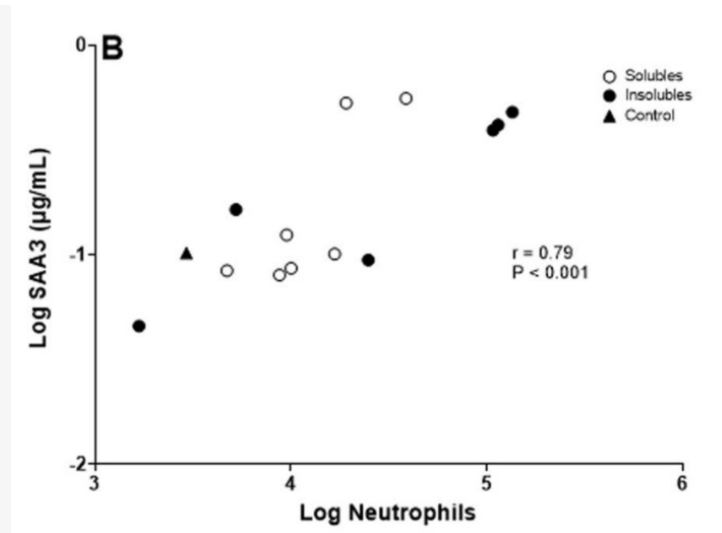


Figure 1. Correlations between neutrophil numbers and SAA3 plasma protein levels, including data from 1 day after exposure to nanomaterials. Reproduced from Gutierrez et al. (2023)<sup>8</sup>.

A linear dose-response has also been found between log<sub>10</sub>-transformed neutrophil numbers and log<sub>2</sub>-transformed SAA3 plasma protein levels in mice, 1 day after exposure to multiwalled carbon nanotubes (Figure 2)<sup>9</sup>.

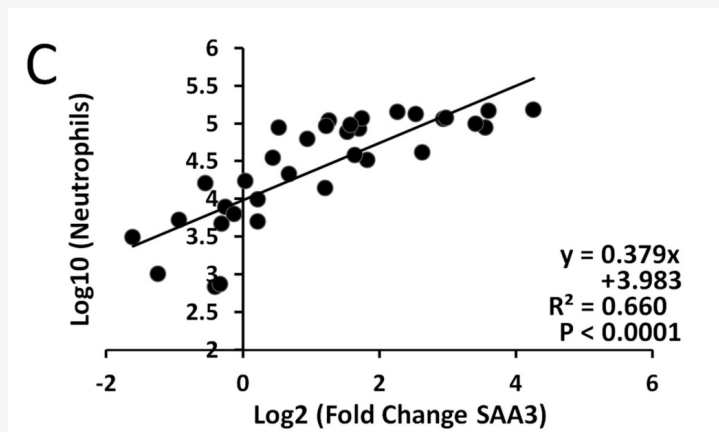


Figure 2. Transformed SAA3 protein vs. transformed neutrophil influx. Reproduced from Poulsen et al. (2017)<sup>9</sup>.

#### Time-scale

It has been shown that pro-inflammatory mediators concentrations increase before acute phase proteins:

- In humans patients with atherosclerotic renal stenosis, blood IL-6 increased in the first hour after renal artery stenting and reach its highest concentration at 6h, while C-reactive protein (CRP) increased 6h after the treatment, peaking at 24h after treatment<sup>10</sup>.
- In human infants undergoing cardiopulmonary bypass, it has been observed that blood concentrations of IL-6 significantly increased after cessation of the procedure and remained elevated 24h later, while CRP started increased 6h after bypass and kept increasing at 12h and 24h after bypass<sup>11</sup>.

#### Known Feedforward/Feedback loops influencing this KER

IL-1, IL-6 and TNF- $\alpha$  can decrease acute phase response by decreasing their own production through the induction of corticosteroids<sup>12</sup>.

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### [Relationship: 2960: Interaction with the lung cell membrane leads to Atherosclerosis](#)

#### AOPs Referencing Relationship

AOP Name	Adjacency	Weight of Evidence	Quantitative Understanding
<a href="#">Substance interaction with lung resident cell membrane components leading to atherosclerosis</a>	non-adjacent	High	Moderate

#### Evidence Supporting Applicability of this Relationship

##### Taxonomic Applicability

Term	Scientific Term	Evidence	Links
human	Homo sapiens	High	<a href="#">NCBI</a>

##### Life Stage Applicability

Life Stage	Evidence
Adults	High

##### Sex Applicability

Sex	Evidence
Male	High
Female	High

Mouse models of human atherosclerosis has been shown to present atherosclerotic lesion progression after exposure to concentrated ambient particles, welding fumes and diesel exhaust particles <sup>3,5,7</sup>.

In humans, epidemiological studies have shown that air pollution, as a stressor that interacts with the lungs, is a risk factor for cardiovascular diseases <sup>14</sup>.

#### Key Event Relationship Description

This KER presents the association between the interaction of stressors with the lungs and atherosclerosis as the outcome. The evidence of the KER presented is based on mouse models of human atherosclerosis.

## Evidence Supporting this KER

### Biological Plausibility

The biological plausibility is moderate. Exposure to different stressors have been shown to induce the progression of atherosclerotic in mouse models of human atherosclerosis (see below). In humans, it has been hypothesized that air pollution, an example of stressor that interacts with the lungs, and cardiovascular diseases are linked by three pathways: i) translocation of inflammatory mediators from the lungs to the systemic circulation, ii) activation of alveolar receptors that results in the alteration of autonomic response and changes in cardiovascular function, and iii) translocation of particles (stressors) from the lungs to the systemic circulation <sup>1,2</sup>.

### Empirical Evidence

For this KER, exposure through the respiratory system (inhalation or intratracheal instillation) of stressors is considered as interaction with lung resident cell membrane components.

- ApoE<sup>-/-</sup> and double knockout ApoE<sup>-/-</sup>LDLR<sup>-/-</sup> mice exposed to concentrated ambient particles (110 µg/m<sup>3</sup>) for 6h/d, 5d/week for 5 months develop severe atherosclerosis. ApoE<sup>-/-</sup> mice exposed to concentrated ambient particles presented a 57% increase in aortic intima surface coverage by atherosclerotic lesion than mice exposed to air <sup>3</sup>.
- Intrapharyngeal aspiration of singlewalled carbon nanotubes into ApoE<sup>-/-</sup> mice (20 µg/mouse every 2 weeks for 8 weeks) induced a significant increase in plaque progression <sup>4</sup>.
- ApoE<sup>-/-</sup> mice on a Western diet showed an increase in atherosclerotic lesion area after exposure to fumes from gas metal arc-stainless steel welding (40 mg/m<sup>3</sup>) for 3h/day for 10 days <sup>5</sup>.
- A modest increase in atherosclerotic plaque area was observed in ApoE<sup>-/-</sup> mice after intratracheal instillation of titanium dioxide nanoparticles (0.5 mg/kg) once a week for four weeks <sup>6</sup>.
- ApoE<sup>-/-</sup> mice, fed a Western diet and exposed to diesel exhaust particles through oropharyngeal aspiration (35 µg) twice a week for four weeks, presented an increased atherosclerotic lesions area <sup>7</sup>.
- Intratracheal instillation of human serum amyloid A once a week for 10 weeks in ApoE<sup>-/-</sup> mice (on Western-type diet) induced atherosclerotic plaque progression <sup>8</sup>.

In addition, several epidemiological studies have shown that exposure to particulate matter from air pollution is associated to cardiovascular diseases:

- A prospective study in six cities from USA showed that air pollution was associated with death from cardiopulmonary diseases <sup>9</sup>.
- In Dublin, there was a decrease in black smoke concentration in air, along with a significant decrease in the number of cardiovascular deaths, after the 1990 ban on coal sales <sup>10</sup>.

### Uncertainties and Inconsistencies

ApoE<sup>-/-</sup> mice seem to have a moderate plaque progression when feed a normal diet, instead of high-fat diet, and exposed to the stressor for a short period <sup>6</sup>.

## Quantitative Understanding of the Linkage

### Response-response relationship

- Following the ban of coal in Dublin, a decrease of 70% of black smoke (35 µg/m<sup>3</sup>) was observed along with a 10.3% decrease (p<0.0001) in cardiovascular deaths <sup>10</sup>.
- A prospective study following postmenopausal women from USA for 6 years observed that an increase of 10 µg of PM<sub>2.5</sub> (particulate matter with a diameter of less than 2.5 µm) was associated with 24% increased risk of cardiovascular event and a 76% increased risk of death from a cardiovascular disease <sup>11</sup>.
- Beelen et al. analyzed data from 22 European cohort studies on long-term exposure to air pollution and associations with cardiovascular diseases mortality. It was obtained that a PM<sub>2.5</sub> increase of 5 µg/m<sup>3</sup> was associated with 21% increased risk of death from cerebrovascular disease, while an increase of 10 µg/m<sup>3</sup> of PM<sub>10</sub> (particulate matter with a diameter of less than 10 µm) was associated with an 22% increased risk of death from cerebrovascular disease <sup>12</sup>.
- Results from 11 cohort studies on long-term exposure to air pollution and incidence of acute coronary events showed a 13% increased risk of coronary events associated to 5 µg/m<sup>3</sup> increase of PM<sub>2.5</sub>, and a 12% increased risk of coronary events associated to 10 µg/m<sup>3</sup> increase of PM<sub>10</sub> <sup>13</sup>.
- A cohort study of population living in Denmark between 2005 and 2017, and aged more than 50 years old, showed that a 5 µg/m<sup>3</sup> increase of PM<sub>2.5</sub> was associated to a 22% increased risk of stroke, while an increase of 1.85 µg/m<sup>3</sup> increase of PM<sub>2.5</sub> was associated to a 5.3% increased risk of myocardial infarction <sup>14,15</sup>.

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