



Oxidative Stress Mechanisms in the Pathogenesis of Environmental Lung Diseases

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Abstract

Globally, respiratory diseases are major cause of disability and mortality, and more alarmingly, it disproportionately affects developing countries, which is largely attributed to poor quality of air. Tobacco smoke and emissions from combustion of fossil fuel and biomass fuel are the major airborne pollutants affecting human lung health. Oxidative stress is the dominant driving force by which the airborne pollutants exert their toxicity in lungs and cause respiratory diseases. Most airborne pollutants are associated with intrinsic oxidative potential and, additionally, stimulate endogenous production of reactive oxygen species (ROS) and reactive nitrogen species (RNS). Elevated ROS and RNS in lungs modulate redox signals and cause irreversible damage to critical biomolecules (lipids, proteins and DNA) and initiate various pathogenic cellular process. This chapter provides an insight into oxidative stress-linked pathogenic cellular process such as lipid peroxidation, inflammation, cell death, mitochondrial dysfunction, endoplasmic reticulum stress, epigenetic changes, profibrotic signals and mucus hypersecretion, which drive the development and progression of lung diseases. Lungs are associated with robust enzymatic and non-enzymatic (GSH, ascorbic acid, uric acid, vitamin E) antioxidant defences. However, sustained production of free radicals due to continuous exposures to airborne pollutants overwhelms lung antioxidant defences and causes oxidative injury. Preclinical studies have

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demonstrated the critical roles and therapeutic potential of upregulating lung antioxidants for intervention of respiratory diseases; however, so far clinical benefits in antioxidant supplementation trials have been minimal and conflicting. Antioxidants alone may not be effective in treatment of respiratory diseases; however it could be a promising adjunctive therapy.

Keywords

Respiratory diseases · Oxidative stress · Lipid peroxidation · Inflammation · Cell death · Mitochondrial dysfunction · Endoplasmic reticulum stress · Epigenetic changes · Profibrotic signals · Mucus hypersecretion · Lung antioxidants

5.1 Introduction

With growing industrialization and rapid urbanization, the global burden of respiratory diseases is rising at an alarming rate and is a major source of disability and death following cardiovascular diseases [1, 2]. Globally, around 10% of all disability-adjusted life years (DALYs) lost is attributed to respiratory illnesses mainly chronic obstructive pulmonary disease (COPD), asthma, tuberculosis, lower respiratory tract infection and lung cancer [3]. Respiratory diseases disproportionately affect developing countries [1, 2], which is largely attributed to poor quality of air, although socioeconomic and genetic factors contribute significantly. Despite being so widespread, environmental respiratory illnesses are highly preventable.

Lungs are at the direct interface between body and the environment, and therefore, the major threat to lung health is the air we breathe. An adult individual inhales on average 10,000 to 15,000 litres of air every day. Inhaled oxygen along with airborne pollutants dissolves in respiratory epithelial lining fluid. The common pollutants that contaminate air include tobacco smoke, particulate matter, biomass fuel smoke, industrial emissions, ozone, oxides of nitrogen, oxides of sulphur and carbon monoxide. Most of these airborne pollutants are associated with intrinsic oxidative potential, and hence, oxidative stress is thought to be a dominant mechanism by which the air pollutants exert their toxicity in lungs. Several studies have reported elevated oxidatively damaged biomolecules in the exhaled breath condensate, nasal lavage, bronchoalveolar lavage and blood of human subjects following acute exposures to airborne pollutants such as cigarette smoke, ozone or particulate matter. Prolonged exposures to these airborne toxicants also stimulate endogenous cellular production of free radicals by diverse mechanisms including mitochondrial, NADPH oxidase, xanthine oxidase and myeloperoxidase activity. To defend against oxidative insult, lungs are endowed with powerful extracellular antioxidants in respiratory tract lining fluid as well as intracellular antioxidant defences. Measures to improve the air quality that we breathe and understanding the underlying mechanisms of disease development caused by exposures to airborne toxicants are pivotal to reduce the global burden

of respiratory disease. This chapter chronicles the key mechanisms by which oxidative stress mediates development and progression of respiratory diseases. The chapter begins with an overview and sources of free radicals in the lungs and a brief description of key environmental respiratory diseases mainly COPD, asthma, acute respiratory distress syndrome and idiopathic pulmonary fibrosis. Finally, the chapter reviews the antioxidants associated with lungs and clinical studies testing antioxidant therapy for treatment of respiratory diseases.

5.2 Sources of Free Radicals in Lungs

5.2.1 Free Radicals: ROS and RNS

Free radical is a highly reactive chemical species, which interacts and damages cellular biomolecules (proteins, lipids, DNA and carbohydrates). Free radicals are molecules associated with an unpaired electron in their valency shell, which renders them intrinsically unstable and reactive. In biological systems, free radicals are oxygen-centred radicals and nitrogen-containing radicals collectively called as reactive oxygen species (ROS) and reactive nitrogen species (RNS), respectively. ROS comprises of radical species (such as superoxide, hydroxyl radicals, lipid-derived hydroperoxides) and non-radical species (singlet oxygen, hydrogen peroxide and hypochlorous acid and ozone) (Table 5.1) [4, 5]. RNS includes nitric oxide, nitrogen dioxide, nitroxyl anion and peroxyxynitrite [6]. Elevated cellular levels of ROS and RNS with reduced capacity to detoxify or neutralize them or its derivatives result in a state of oxidative and nitrative stress, respectively.

Low levels of ROS and RNS function as secondary messengers and play a pivotal role in redox cell signalling and regulate diverse beneficial normal physiological processes such as bacterial killing during phagocytosis, vasodilation, tissue repair and regeneration [4, 7, 8]. In contrast, higher levels of ROS/RNS inflict irreversible damage to biomolecules resulting in cell and tissue injury, and if unchecked, it results in onset of inflammatory diseases including respiratory disease [9–11].

Table 5.1 List of ROS and RNS

ROS	RNS
Superoxide ($O_2^{\cdot-}$)	Nitric oxide ($\cdot NO$)
Hydrogen peroxide (H_2O_2)	Nitrogen dioxide ($\cdot NO_2$)
Hydroxyl radical ($HO\cdot$)	Nitrous acid (HNO_2)
Peroxyl radical ($RO_2\cdot$)	Dinitrogen tetroxide (N_2O_4)
Alkoxy radical ($RO\cdot$)	Dinitrogen trioxide (N_2O_3)
Hydroperoxyl radical ($HO_2\cdot$)	Peroxyxynitrite ($ONOO\cdot$)
Singlet oxygen (1O_2)	Peroxyxynitrous acid ($ONOOH$)
Ozone (O_3)	Alkyl peroxyxynitrites ($ROONO$)

5.2.2 Sources of ROS and RNS in Lungs

5.2.2.1 Endogenous Sources of ROS/RNS in Lungs

The endogenous source refers to free radicals produced by cells. Adult lung comprises around 40 different cell types [12], which can be broadly classified as epithelial, endothelial and neuroendocrine cells, smooth muscles, fibroblasts and immune cells (macrophages, neutrophils and T cells). All these cells have varied intrinsic capacity to generate ROS upon stimulation. In general, ROS/RNS are produced as intermediates or byproducts of cellular metabolism catalysed by enzymes localized in different organelles primarily plasma membrane, cytosol, mitochondria, peroxisomes and endoplasmic reticulum. The key enzymes that produce ROS/RNS in lungs include cytochrome c oxidase, NADPH oxidase, myeloperoxidase, xanthine oxidase and nitric oxidase synthase. In most of the lung disorders, the mitochondria are primary source of excess ROS [4, 9, 13]. Superoxide produced by mitochondria is a result of incomplete reduction of oxygen to water due to leakage of electrons by mitochondrial respiratory chain [4, 13]. NADPH oxidase (NOX) is a multicomponent transmembrane enzyme complex that generates superoxide as end product via one electron reduction of oxygen [8–11]. In humans, there are seven isoforms of *NOX* – *NOX1*, *NOX2*, *NOX3*, *NOX4*, *NOX5*, *DUOX1* and *DUOX2*. Nox2 is localized in phagosome of phagocytes (macrophages and neutrophils) and produces massive ROS to kill the phagocytized bacteria. *NOX1*, *NOX2*, *NOX4*, *DUOX1* and *DOUX2* are distributed in various cell types in the human lung [10, 14]. Myeloperoxidase (MPO), a heme-containing enzyme, is localized in lysosomes of neutrophils and macrophages [11] and catalyses chlorination of H_2O_2 to HOCl, a highly reactive radical [11]. Additionally, MPO catalyses oxidation of thiocyanate to generate hypothyocyanite. Xanthine oxidase, localized in the plasma membrane and cytosol, catalyses oxidation of hypoxanthine to xanthine and to uric acid, during which superoxide is generated [6, 15]. The enzyme nitric oxide synthase catalyses oxidation of L-arginine to citrulline and nitric oxide [7, 16]. Nitric oxide produced reacts with superoxide to form very potent RNS intermediate, peroxynitrite ($OONO^-$). There are three isoforms of NOS, namely, *NOS1* (neuronal NOS, nNOS), *NOS2* (endothelial NOS, eNOS) and *NOS3* (inducible NOS, iNOS). Pulmonary cells constitutively express NOS1 and NOS2, while NOS3 is elevated in the lung during inflammation [7, 16, 17].

5.2.3 ROS and RNS in Lungs: Exogenous Sources

Inhalation of airborne toxicants such as tobacco smoke, biomass fuel smoke, particulate matter and gaseous emissions (nitrogen oxides and sulphur oxides), ozone, chemical toxins, pesticides and pollen grains largely constitutes the exogenous sources of oxidants in lungs [18]. Among the airborne toxicants, ambient air pollution particulate matter and cigarette smoke are the main contributors of free radicals

in human lungs [1, 3, 19, 20]. Depending on the size, the ambient particulate matter (PM) exhibits different physicochemical properties [21, 22]. Coarse particles larger than 10 microns are composed of natural materials (mineral and silicates) and are trapped and cleared by the nose and upper respiratory tract [23, 24]. Fine PM of size 2.5 microns ($PM_{2.5}$) and lesser and ultrafine particles (<0.1 microns) are produced by the combustion process from anthropogenic activities and are composed of polycyclic aromatic hydrocarbons (PAH), metals, minerals, sulphates and nitrates [23, 25]. $PM_{2.5}$ and ultrafine particles could reach deep into distal lungs and cause injury to the alveoli [24]. A single puff of cigarette smoke comprises of 10^{15} free radicals [20, 26] and over 4000 chemicals including epoxides, peroxides, semiquinones, quinones and PAH. The cigarette smoke free radicals are classified into two phases: tar (particle) and gas. Tar phase has 10^{17} relatively long-lived radical molecules per gram such as quinone/hydroquinone (Q/QH₂) radicals, which produce $O_2^{\cdot-}$ leading to the generation of H_2O_2 and $\cdot OH$ [27]. Gas phase of cigarette smoke comprises of 10^{15} organic and inorganic radicals per puff and includes $NO\cdot$, NO_x and $ONOO-$ [28]. Ozone is another powerful exogenous oxidant in the lungs [29]. Plant pollens are associated with NADPH oxidase enzyme and are shown to generate superoxide in the airways following inhalation. Besides possessing intrinsic oxidative potential, the airborne toxicants also stimulate production of ROS by activating NADPH oxidase, damaging mitochondria and by recruitment of inflammatory cells (macrophages and neutrophils).

5.3 Oxidative Stress-Associated Pulmonary Disorders

5.3.1 Chronic Obstructive Pulmonary Disease (COPD)

As per the recent estimates by a global burden of disease study [1], COPD kills over three million people and is the third leading cause of death. COPD is characterized by progressive, irreversible limitation in expiratory airflow and abnormal lung inflammation. The disease process involves airway inflammation and remodeling, mucus hypersecretion, loss of the terminal bronchioles and destruction of the lung parenchyma [30, 31]. Tobacco smoking is the primary causal agent of COPD; however, recent epidemiological evidence suggests that exposures to indoor and outdoor air pollutants are also a major etiological factor, especially in developing countries. Genetic polymorphisms in genes coding for alpha-1 antitrypsin, metalloproteinase 33, superoxide dismutase-3 (SOD3) and tumour necrosis factor- α (TNF α) are additional risk factors in 10%–20% of the smokers for developing COPD [32–37]. Although the underlying pathophysiological mechanisms are complex, the lungs of patient with COPD show persistent oxidative stress, increased levels of proinflammatory cytokines, increased CD4 and CD8 cells, elevated levels of proteases and increased apoptosis and senescence [38–44]. Oxidative stress-driven mechanisms are strongly implicated in the initiation and progression of COPD [45].

5.3.2 Asthma

Asthma is a very common airway disorder affecting both children and adults worldwide. Clinically, asthmatics show airway hyperresponsiveness, reversible airflow obstruction and abnormal airway inflammation [46]. Genetic factors in the combination of environmental exposures increase the risk of asthma. The common environmental agents that trigger asthma include aeroallergens (indoor and outdoor), tobacco smoke, dust, air pollutants, cold air and viruses. The hallmark pathological features of asthmatic airways include airway remodeling, epithelial desquamation, goblet cell hyperplasia and inflammation and are associated with eosinophils, mast cells, neutrophils, macrophages and T helper type 2 cells (Th2 cells) [47]. In asthmatics lungs, allergen exposure triggers immunoglobulin E (IgE) production from B cells, degranulation of mast cells and infiltration of eosinophil into airways. These events are mediated by Th2 cytokines, namely, IL-13, IL-4, IL-5 and IL-9. Th1 cytokines (IFN- γ , IL-12) and Th17 cytokines (IL-23) are implicated in severe or steroid resistance asthma, which is characterized by high neutrophilic inflammation. ROS through direct injury to airway epithelial cells and via redox signalling mechanism are shown to enhance sensitivity to allergens and augment Th2/Th1 cytokine secretion and, therefore, intricately involved in the pathogenesis of asthma [48].

5.3.3 Acute Respiratory Distress Syndrome (ARDS)

Acute lung injury and its severe form ARDS are common complications in patients admitted to intensive care unit. ARDS results from direct or indirect injury to lungs. Direct injury may be caused by gastric aspiration, pneumonia, inhalation of injuries gases and pulmonary contusion. Indirect injury includes sepsis, pancreatitis and trauma. ARDS is characterized by alveolar flooding with protein-rich oedema followed by a progressive fibrotic phase [49]. Death among the patients with ARDS is mainly due to respiratory failure and/or multiorgan failure. Pathogenesis of ARDS involves an early injury to alveolar epithelium and capillary endothelium, which results in leakage and flooding of alveolar and interstitial spaces with protein-rich oedema. This is also accompanied by a massive influx of neutrophils into alveolar and interstitial spaces. Neutrophils secrete proteolytic enzymes (elastase and metalloprotease), ROS, proinflammatory mediators and further lung injury [50]. The early inflammatory exudative phase is followed by a fibroproliferative phase in which fibroblast and myofibroblast infiltrate and proliferate within the alveolar and interstitial spaces leading to lung fibrosis [51]. Oxidative stress is shown to mediate epithelial-endothelial barrier dysfunction and perpetuate inflammation in ARDS patients [52, 53].

5.3.4 Idiopathic Pulmonary Fibrosis (IPF)

IPF is a progressive interstitial pulmonary fibrosis disorder with no known causal etiological factor. The lungs of patients with IPF show excessive deposition of matrix proteins such as fibronectin and collagen in the alveoli and lung parenchyma, which destroys the gas exchange surface leading to respiratory failure [54]. It is more prevalent in the USA and Europe than South America and South Asia. IPF disproportionately affects individuals above age 65 years, and often it is referred to as age-related disorder. Pathogenesis of IPF involves chronic insult to alveolar epithelial cells (AEC), senescence of AEC and fibroblast, increased differentiation of fibroblast to myofibroblast [54] and increased accumulation of myofibroblast, which is mediated by oxidative injury, mitochondrial dysfunction, proteotoxicity and endoplasmic reticulum stress [14, 55]. Emerging evidences implicate ROS by Nox4 as a key player in the pathogenesis of IPF [55].

5.4 Oxidative Stress-Driven Mechanisms in Lung Diseases

5.4.1 Oxidative Stress and Lipid Peroxidation

Lipid peroxidation (LPO) in biological systems refers to the oxidation of cellular membrane lipids; and uncontrolled LPO is the most significant early biological process induced by oxidative stress state. Excess LPO results in defective or dead cell, inactivation of critical proteins and activation of proinflammatory responses. Together, these events not only initiate but also ensue disease progression. Numerous studies have overwhelmingly showed that LPO is a universal pathogenic event in all the respiratory diseases including COPD, IPF, ARDS and asthma [56].

Membrane lipids mainly glycerophospholipids (PL) esterified with polyunsaturated fatty acid (PUFA) and to a lesser extent free PUFA are the targets for oxidation. Free PUFAs released by action of phospholipases inside the cells are substrates for enzymes such as cyclooxygenase, lipoxygenase and cytochrome P450s, and hence, free PUFA undergoes enzyme-dependent peroxidation. PL-PUFAs are predominantly oxidized by non-enzymatic process and highly depend on the radical species. Both radical species ($\cdot\text{OH}$ and $\text{O}_2\cdot^-$) and non-radical species (H_2O_2 , HOCl , ozone and singlet oxygen) may oxidize PL-PUFA in selective or non-selective manner. The chemical reactions mediating the oxidation of PL-PUFA or free PUFA are similar; however, the products generated may vary. LPO process involves three phases – initiation, propagation and termination. During the initiation phase, non-radical lipid molecule becomes a lipid radical. The radical species abstract hydrogen from bisallylic methylene and produce a carbon-centred radical ($\text{L}\cdot$) within PUFA, which subsequently reacts with molecular oxygen and forms lipid peroxy ($\text{LOO}\cdot$) radical. During the propagation phase, the $\text{LOO}\cdot$ radical abstracts hydrogen from bisallylic methylene of another PUFA molecule and transforms itself into a lipid hydroperoxide and, concomitantly, generates a new $\text{L}\cdot$ radical, and in this manner, the peroxidation chain reaction sustains. During termination process,

antioxidant molecule, such as vitamin E, donates hydrogen, reduces lipid radicals without transforming itself into radical and thus terminates the LPO chain reaction. Lipid hydroperoxides may further participate in additional oxidative reactions such as Fenton reaction catalysed by Fe or Cu, intra- and intermolecular oxidative modification and oxidative fragmentation. Finally, LPO produces diverse reactive aldehyde byproducts including widely studied malondialdehyde (MDA) and 4-hydroxy-2-nonenal (4HNE). The oxidation of PL-PUFA also yields diverse species of oxidized phospholipids (Ox-PLs) (aldehyde, alkene, peroxy and alkane derivatives), which exhibit varying carbon chain length, hydrophobicity, reactivity, physical stability and biological activity [57]. The oxidized PUFA chain of glycerophospholipid may be released by the action of enzymes such as phospholipase A2 and PAF-acetyl hydrolases [57]. Since the cell membranes are composed of 40–50% of phosphocholine (PC) phospholipids, PC-derived Ox-PLs species are the most abundant Ox-PLs detected in injured lungs.

LPO products such as MDA, 4HNE and Ox-PLs produced in the lungs are not bystanders; rather they actively take part in the pathogenesis of lung disease by inducing cell death, epithelial-endothelial barrier dysfunction, inflammation and immune responses [58]. Ox-PLs and 4HNE are shown to mediate cytotoxicity through disruption of membrane integrity and activating cell death signalling programs such as apoptosis [59, 60] and ferroptosis [61, 62]. Ox-PLs generated following particulate matter exposure caused disruption of the endothelial barrier [62]. Ox-PLs are demonstrated to be dominant mediators of acute lung injury following gastric aspiration and viral infection [63].

A large body of evidence has reported elevated levels of MDA, 4HNE or Ox-PLs or their corresponding protein adducts in the bronchoalveolar lavage fluid, lung tissue and/or serum of patients with various lung diseases such as asthma, COPD, ARDS and IPF [43, 64–66]. Immunohistochemical analysis revealed greater accumulation of 4HNE in the airways, alveolar epithelium and inflammatory cells of the lungs of COPD patient when compared to smoker non-COPD patient with similar smoking history [67]. The bronchoalveolar lavage fluid from COPD patients shows higher levels of Ox-PLs when compared to healthy subjects [43, 64, 65]. The lung parenchyma of IPF patients showed greater accumulation of LPO products [54, 55, 68]. In most respiratory diseases, the levels of LPO byproducts increased with the severity of the pulmonary diseases, which suggest that the LPO is the central pathological event.

5.4.2 Oxidative Stress in Activating Inflammatory Response

It is proven beyond doubt that oxidative stress is involved in the initiation, promotion and augmentation of inflammation by affecting multiple redox-sensitive signal transduction pathways, including Toll-like receptor (TLR) signalling, MAPK kinase signalling and inflammasome which ultimately leads to activation of proinflammatory transcription factors particularly nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) and AP-1 [32, 69].

TLR signalling is central in activating pulmonary inflammatory responses following infectious stress and oxidative stress [70]. During infection, TLRs recognize highly conserved microbial motifs referred to as pathogen-associated molecular pattern (PAMP) and activate downstream inflammatory signals [71]. In mammals, there are 13 TLRs, which are present either on the plasma membrane (TLR1, TLR2, TLR4, TLR5, TLR6 and TLR10) or in endosome compartment (TLR3, TLR7, TLR8, TLR9) inside the cell. Upon activation by PAMPs, TLRs undergo hetero- or homodimer and trigger a downstream signal transduction by recruiting adaptor molecules, myeloid differentiation factor 88 (MYD88) or Toll-receptor-associated activator of interferon (TRIF). Except TLR3, a majority of TLRs recruit MYD88 that interacts with IRAK4 and phosphorylate IRAK1. Phosphorylated IRAK1 activates TNFR-associated factor 6 (TRAF6) through phosphorylation, which subsequently stimulates protein kinase C and transforming growth factor (TGF)- β -activated kinase 1 (TAK1). Activated TAK1 activates I κ K complex and MAPK kinase family members (ERK1/2, p38 MAPK and c-Jun terminal kinase (JNK)) by phosphorylation mechanisms. TLR3 recruits adaptor molecule TRIF and activate TRAF3, which then activates TANK-binding kinase 1 (TBK1). Activated TBK1 initiates interferon regulatory factor 3 to transcribe IFN- β cytokine which by autocrine or paracrine mechanism elicits interferon signalling pathway activation.

Transcription factor NF- κ B is a central node in regulating inflammation leading to the pathogenesis of COPD, asthma, ARDS and IPF. NF- κ B family is constituted of five members – NF- κ B1 (P50) (its precursor p105), NF- κ B2 (P52) (its precursor p100), p65/RelA, RelB and C-Rel – and exists as homo- or heterodimer. Only p65, Rel-B and C-Rel members have a transactivating domain. In an unstimulated cell, the NF- κ B dimer is sequestered in the cytoplasm by one of the three members of I κ B protein complex consisting of I κ B α , I κ B β and I κ B ϵ . Activation of NF- κ B may occur through canonical or noncanonical pathways. In canonical pathway, signals elicited by TLR(s) ligands, TNF α or IL-1 β converge at I κ K complex constituted of I κ K α , I κ K β and I κ K γ (NEMO). Upon activation I κ K complex phosphorylates I κ B on serine 32 and serine 36, which results in its proteasomal degradation and subsequently allows NF- κ B to translocate into the nucleus. NF- κ B binds to κ B element in the promoter/enhancer regions and activates transcriptional expression of cytokines, chemokines and adhesion molecules, which are involved in coordinating innate and adaptive immunity [70–72].

Oxidative stress regulates activation of TLR signalling by multiple mechanisms [70]. NADPH oxidase-dependent ROS production is shown to enhance surface trafficking of TLR4 to lipid rafts, thereby augmenting downstream signals leading to hyperactivation of NF- κ B [72–75]. Suppression of ROS generation by pharmacological NADPH oxidase inhibitor, genetic ablation of NADPH oxidase or exogenous antioxidants mitigated lipopolysaccharide (LPS)-induced TLR4 trafficking to lipid rafts and diminished downstream inflammatory responses [74, 75]. Genetic ablation of NADPH oxidase dampened lung inflammation and injury in mice exposed to gram-negative bacteria, LPS, TNF α or bleomycin, suggesting a crucial impact of NADPH oxidase-elicited ROS in producing inflammation and associated tissue injury [55, 74, 76–78]. On the other hand, ROS derived from NADPH oxidase

have also been involved in the resolution of lung inflammation [79]. Mitochondria-derived ROS also play a crucial role in enhancing TLR1, TLR2 and TLR4 signalling [80] and production of proinflammatory cytokines [81]. ROS is shown to augment TLR3 signalling partly by increasing the expression of TLR3 [82]. Oxidatively damaged biomolecules also act as danger-associated molecular patterns (DAMPs) and engage TLR4 to activate inflammatory responses. For instance, oxidized phospholipids generated in lungs following exposures to chemicals, bacteria or virus act as DAMPs and activate TLR4 directed inflammatory responses [83], which is shown to play an essential role in the initiation of acute lung injury. In contrast, excess oxidized phospholipids may also dampen TLR4 signalling by directly interacting with LPS-binding protein, CD14 and/or MD-2 and inhibit inflammation and protect from sepsis [84]. Often oxidative or nitrosative modification of proteins gives rise to modified proteins, which act as DAMPs and perpetuate inflammation. For example, protein adducts of 4HNE or MDA are shown to elicit inflammatory and immune responses [85, 86] in the lungs. 4HNE has also been shown to activate NF- κ B and p38 MAPK and promote inflammation [56]. S-nitrosylation of surfactant-D protein switches pulmonary surfactant protein-D from antioxidant to a proinflammatory mediator [87].

ROS may regulate NF- κ B activation by modulating upstream signal kinases (I κ K complex) through oxidative modification of signal transducers and/or its binding partners [88]. In early phases of oxidative stress, ROS may enhance NF- κ B activation following exogenous stimuli; however sustained oxidative stress may repress NF- κ B activation [88]. I κ K complex, mainly I κ K β , is highly susceptible for redox modification. Exposures to H₂O₂ induced oxidative inactivation of I κ K β , which prevented phosphorylation of I κ B protein and thus blocked TNF α -induced NF- κ B activation [89]. In another study, H₂O₂ posttreatment augmented I κ K kinase activity in response to TNF α and leads to higher NF- κ B activation [90]. H₂O₂ treatment was also shown to enhance NF- κ B activation in response to IL-1 cytokine by increasing NF- κ B-inducing kinase activity [91]. I κ K β is also susceptible for S-nitrosylation, which inactivates I κ K β resulting in inhibition of NF- κ B activation [92]. ROS and RNS are shown to directly modify NF- κ B or its associated proteins and alter its transcriptional activity. ROS-dependent phosphorylation of serine-276 on REL-A enhanced transcriptional activity of NF- κ B [93]. S-nitrosylation of p65 subunit inhibited NF- κ B activity. Kelleher et al. [94] demonstrated that p65 subunit is S-nitrosylated in unstimulated lung cells and LPS challenge stimulates denitrosylation of p65 resulting in activation of NF- κ B. Excess ROS may also inactivate proteasome, which impedes I κ B degradation and thus inhibits NF- κ B activation [95]. Certain cellular redox proteins play an important role in modulating upstream pathways leading to NF- κ B activation. In the nucleus, thioredoxin binds and protects oxidation of p65 subunit and enhances its DNA-binding activity. It is also shown that thioredoxin mediates denitrosylation of p65 following LPS exposure and facilitates NF- κ B activation [94] in the lungs of mice. Pretreatment with antioxidants such as N-acetylcysteine [96], GSH [97, 98] or increased expression of

antioxidant [72, 97, 98] attenuated lipopolysaccharide (LPS)-promoted NF- κ B activation emphasizing the role of oxidative stress in regulating NF- κ B activity.

Inflammasome is an intracellular multiprotein complex assembled in the cytoplasm, which recognizes microbial or environmental toxins and DAMPs (e.g. ATP) and activates inflammatory responses. Several airborne environmental pollutants such as silica, ozone, particulate matter and tobacco smoke are shown to activate inflammasome, and therefore, inflammasome signalling is implicated in the pathogenesis of several lung disorders such as acute lung injury, pulmonary fibrosis, COPD and asthma [99, 100]. Activation of inflammasome produces active caspase-1 via autoproteolytic cleavage, which then mediates proteolytic cleavage of precursor pro-IL1 β and pro-IL18 into biologically active cytokines. Among the inflammasome members, NLRP3 is redox sensitive, and therefore, intracellular ROS generated by NADPH oxidase or mitochondria have been shown to alter the activation of NLRP3 inflammasome [101]. Ablation of NADPH oxidase or depletion of mitochondrial ROS inhibited ATP-induced caspase-1 activation and IL-1 β secretion in macrophages [102, 103]. ROS may also alter NLRP3 inflammasome activation by oxidizing redox-sensitive binding partners such as thioredoxin interaction protein (TXNIP) and mitochondrial antiviral sensing (MAVS) protein to NLRP3. TXNIP is a negative regulator of thioredoxin. It is shown that ROS generated in response to a wide range of environmental stimuli oxidizes thioredoxin that liberates TXNIP. The liberated TXNIP interacts with NLRP3 and promotes NLRP3 activation [104]. MAVS regulates type 1 interferon and NF- κ B signalling following virus infection. It has been shown that ROS may induce MAVS aggregation [105] on the outer membrane of mitochondria, which enables interaction with NLRP3 and promotes activation.

MAPK kinases, namely, ERK1/2, JNK and p38 MAPK, represent key effectors of signal transduction to activate inflammatory responses in the lungs following exposure to environmental toxicants [106]. In macrophages and bronchial epithelial cells, LPS stimulation induces phosphorylation of p38 MAPK and mediates the generation of numerous proinflammatory cytokines such as TNF α , IL-6, IL-1 β [107] and also T cell (Th1 and Th17)-polarizing cytokines such as IFN γ , IL-12 and IL-23 [108]. In cigarette smoke-exposed mouse models, specific activation of p38 MAPK is shown to be a determinant of susceptibility to emphysema [106]. Several investigations suggest that ROS plays a pivotal function in activating and/or perpetuating MAPK kinase signalling. Exposure of cells to H₂O₂ induces phosphorylation of p38, ERK and JNK [109]. Although precisely how ROS activates MAPK kinase is less understood, it is postulated that ROS mediates oxidative inactivation of protein tyrosine phosphatases and MAPK phosphatases, which inactivate MAPK kinase by dephosphorylating [109, 110]. Because MAPK kinase plays a pivotal role in regulating inflammation, kinase inhibitors, particularly p38 MAPK inhibitors, are shown to be promising drug for treatment of airway disorders such as COPD and asthma [111].

5.4.3 Oxidative Stress in Programmed Cell Death

Apoptosis (a programmed cell death) is involved in removing damaged, infected and potentially neoplastic cells, and increased apoptotic cell death is involved in the pathogenesis of several lung disorders. Apoptosis can be activated by several factors including receptor-mediated signals and DNA damage; however, in most cases, ROS functions as an upstream activator of apoptosis. Apoptosis is mediated by extrinsic and intrinsic pathways [40, 112, 113]. The extrinsic pathway is mediated through interaction of death ligands of the tumour necrosis factor (TNF) family (FasL/FasR and TNF α /TNFR1) with their appropriate cell surface death receptors, while non-receptor-mediated stimuli are involved in intrinsic signalling pathways that initiate apoptosis. Oxidative stress may induce these processes by activating several signalling pathways, including MAPK (ERK, JNK and p38), cell-cycle regulators, protein kinase B and caspases [114]. For instance, 4HNE is reported to enhance the mRNA and protein expression of pro-apoptotic regulators/adaptors such as BAX and caspases [59, 115]. Further, 4HNE can directly interact with death ligand (Fas) on the cell membrane and activate apoptotic process [114]. Finally, 4HNE alters cytosolic calcium homeostasis and mitochondrial calcium uptake, resulting in apoptosis [116]. Similarly, cigarette smoke exposure is shown to activate apoptosis via ROS by activating MAPK/STAT1 pathway [117]. Several studies have reported oxidative stress-dependent apoptosis in pulmonary fibrosis, obstructive airway diseases and ARDS [12, 40, 41, 44, 113, 118].

5.4.4 Oxidative Stress in Mitochondrial Dysfunction

Besides the ‘powerhouse’ (ATP production by oxidative phosphorylation) of the cell, mitochondria physically interact and communicate with other organelles to maintain the metabolic homeostasis and many synthetic processes for normal function and survival of cell [13]. Mitochondria may also sense external stressors and alter its function to mount a protective adaptive stress response program [119]. However, prolonged exposures to environmental toxicants induce mitochondrial dysfunction mainly via oxidative stress mechanisms [120]. Mitochondrial dysfunction may present in the form of increased mitochondrial ROS, diminished oxidative phosphorylation, increased mitochondrial mass, secretion of mitochondrial DAMPs, mitochondrial DNA damage, decreased mitochondrial biogenesis and increased accumulation of defective mitochondria [13, 119, 121]. Several studies suggest that mitochondrial dysfunction is a predominant pathological feature in all lung diseases [13, 120, 121].

In lungs, owing to their dynamic function, alveolar type II epithelial cells, bronchial ciliated epithelial cells, vascular smooth muscle cells and macrophages are richer in mitochondria than other lung cell types. In normal conditions, lung cells preferentially use glucose end product, pyruvate, for oxidative phosphorylation. However, during stressful physiological or pathological conditions (such as increased surfactant production), alveolar type II epithelial cells rely on fatty acids

for energy demand. Under chronic stress conditions, mitochondrial bioenergetic metabolic function may get altered in lung cells. For example, cigarette smoke exposure is shown to damage mitochondrial structure and affect oxidative phosphorylation in lung cells [122]. Likewise, primary bronchial epithelial cells from severe COPD patients showed accumulation of abnormal mitochondria [122]. Airway smooth muscles and diaphragmatic and external intercostal muscle of patients with COPD are associated with altered mitochondrial oxidative phosphorylation [123]. Bronchial epithelium in asthmatics is associated with reduced mitochondrial oxidative phosphorylation and decreased expression and activity of cytochrome *c* oxidase [120, 124, 125]. To meet the energy demand and mount stress response, chronic stress may also induce mitochondrial biogenesis in lung cells. Alveolar type II epithelial cells showed increased mitochondrial biogenesis during acute lung injury, pneumonia and hyperoxia-induced lung injury [126]. Bronchial smooth muscles of asthmatic airways are associated with increased mitochondrial biogenesis, and this was linked to higher expression of nuclear respiratory factor 1, peroxisome proliferator-activated receptor γ coactivator (PGC)-1 α and mitochondrial transcription factor A [127]. Abnormal or defective mitochondria in the cells are constantly removed by a process called mitophagy, which is regulated by PTEN-induced kinase 1 (PINK1). Expression of PINK1 is negligible in healthy mitochondria; however its levels increase on the outer mitochondrial membrane of defective mitochondria, which recruits parkin and autophagy proteins and facilitates mitophagy. Impaired mitophagy leads to accumulation of damaged mitochondria in the cells, which promotes cellular senescence [128, 129]. Increased cellular senescence has been observed in the lungs of COPD and IPF patients [128–130]. Exposures to cigarette smoke in lung cells are shown to inhibit mitophagy, increase accumulation of damaged mitochondria [128, 129, 131] and induce cellular senescence. Alveolar type II cells of IPF patients are associated with abnormal mitochondria due to diminished PINK1 expression [132]. In mouse models, PINK1 knockdown impaired mitophagy and increased accumulation of defective mitochondria and promoted fibrosis in aging lungs [132]. On contrary, increased mitophagy may also contribute to pathogenesis of lung diseases. For example, *Staphylococcus aureus* infection increased mitochondrial expression of PINK1 and mediated acute lung injury, which was ablated in PINK1 knockout mice [133]. Mitochondrial dysfunction may also lead to leakage of cytochrome *c*, which triggers programmed cell death [116, 134]. Mitochondria have been shown to regulate various forms of cell death such as extrinsic apoptosis, intrinsic apoptosis, necroptosis and pyroptosis [134], and all these forms of cell death have been reported in various lung diseases including COPD, asthma and IPF.

At physiological concentrations, many mitochondrial-derived molecules including ROS help in normal cellular signalling. However, when secreted in excess, mitochondrial-derived molecules act as mitochondrial DAMPs (mtDAMPs) and contribute to lung injury. Mitochondrial DNA (mtDNA), a well-studied mtDAMP, is released by damaged mitochondria, which is shown to engage TLR9 and inflammasome to initiate inflammatory responses in lung cells [135, 136]. Circulatory levels of mtDNA correlate well with severity and mortality in sepsis and ARDS

patients [137]. Excess ATP released in the lungs by dead or damaged cells also acts as mtDAMPs and activates inflammatory response via NLRP3 inflammasome [119, 138]. Elevated levels of ATP are reported in bronchoalveolar lavage fluid of patients with COPD [139] and asthma as well as in mouse models of asthma [140] and pulmonary fibrosis [141]. Other mtDAMPs such as TFAM and N-formyl peptide are also implicated in driving inflammatory responses in lungs [119]. Cardiolipin, a predominant lipid located in mitochondrial inner membrane, is released by damaged mitochondria and acts as mtDAMP. Levels of cardiolipin increase during lung injury and are shown to mediate cell death and activate inflammasome signals [133, 142]. Additionally, cardiolipin in lung fluid was shown to inhibit surfactant activity and worsen lung function in mouse models of pneumonia [143]. Finally, mitochondrial-derived ROS plays diverse roles in the pathogenesis of lung diseases including perpetuating oxidative stress, augmenting TLR-NF- κ B signalling and cell death [13, 81, 119, 125, 134, 135, 144].

5.4.5 Oxidative Stress in Promoting Endoplasmic Reticulum Stress

The endoplasmic reticulum (ER) is involved in protein biosynthesis and post-translational modifications and perturbation of ER homeostasis results in ER stress which affects both these processes. To overcome the ER stress, cells initiate an evolutionarily conserved mechanism called unfolded protein response (UPR). Activation of UPR leads to decrease in protein synthesis by selectively inhibiting translation, increases protein folding machinery and removes misfolded proteins through endoplasmic reticulum-associated degradation (ERAD) pathway [145]. If UPR fails to alleviate ER stress, it activates apoptotic signalling mechanism [146] and, thus, helps in removal of damaged or stressed cells. Chronic ER stress is pathological and is associated in the pathogenesis of many lung disorders. Markers of ER stress are elevated in neutrophil-associated steroid-resistant asthma [147]. In COPD model, cigarette smoke exposure elicited ER stress and apoptosis [148, 149]. ER stress was found to be elevated in lungs of human IPF and murine models of pulmonary fibrosis [146]. ER stress is also reported to be involved in the hyperoxia-induced acute lung injury in neonates [150]. Sustained oxidative stress milieu may promote ER stress by increasing cellular stress and decreasing the efficiency of protein folding pathways [151]. A relationship has been established between ROS generation and activation of ER stress response [152]. NADPH oxidase(s) and mitochondria are reported as a probable ROS source during ER stress.

5.4.6 Oxidative Stress in Epigenetic Alterations

Chronic oxidative stress state may disturb the epigenetic state of the cell by multiple mechanisms. For example, superoxide radicals can directly mediate transfer of a methyl group from SAM to a cytosine residue without the need of DNMT by

deprotonating C5 [153]. ROS may increase DNMT expression and indirectly affect DNA methylation [154]. ROS may also directly or indirectly modify acetylation of histones by modifying activity or the expression of histone methyltransferases (HMTs) and/or histone acetyltransferases (HATs) [155]. On the other side, ROS and RNS are reported to modulate the activity of HDACs (histone deacetylase) that may influence the expression of target genes by removing acetyl groups [156–159] on histones. Recently, various ncRNAs, in particular microRNAs (miRNAs), are regulated by ROS. Interestingly, some miRNAs such as miR-9, miR-21, miR-200 and miR-210 are shown to control cellular ROS levels and are termed as redoximiRs [160]. ROS are also shown to interfere with miRNA biogenesis process as well as miRNA maturation by modulating Dicer and argonaute RISC catalytic component [161]. In the context of the lung, several lines of evidence support that ROS-dependent changes in the epigenetic background play an important role in the pathogenesis of respiratory diseases. For instance, cigarette smoke exposures inhibit HDAC2 enzyme activity through oxidative and nitrosative modification, which leads to enhanced inflammatory responses, senescence and steroid resistance in COPD [158, 159]. Another study shows that Sirtuin 1 promotes lung epithelial cell death following hyperoxia by selectively deacetylating the transcription factor nuclear factor (erythroid-derived 2)-like 2 (NRF2), accompanied by reduced levels of antioxidant enzymes [162]. In case of asthma, one study reported that exposure to environmental particulate matter could lead to demethylation of iNOS gene; subsequently this may lead to increased expression of proinflammatory iNOS, leading to lung inflammation [163]. The involvement of many factors including ROS in epigenetics of IPF has been reviewed [164].

5.4.7 Oxidative Stress in Profibrotic Mechanisms

Fibrotic lungs are associated with increased oxidative stress, as indicated by the elevated levels of biomarkers of lipid, protein and DNA damage, and several reports have implicated ROS in profibrotic processes. Activation and proliferation of fibroblasts/myofibroblasts are thought to be responsible for the excessive synthesis and accumulation of extracellular matrix (ECM) proteins, resulting in fibrosis. During the inflammatory phase of fibrosis, ROS along with growth factors (TGF- β , PDGF and CTGF) and cytokines (IL-6 and IL-13) stimulate fibroblast to produce ECM. Among these, TGF- β is the most dynamic pro-fibrogenic cytokine, which regulates important biological processes such as EMT, fibroblast activation and differentiation and ECM production [165]. ROS may influence the transformation of latent TGF- β complex into its active form, which then binds to its receptors and activates signalling pathways such as SMAD-dependent or SMAD-independent (e.g. MAPK and PI3K) pathways and enhances the transcriptional activity of various profibrotic genes such as α -SMA and COL1 [166]. On the other hand, elevated TGF- β itself reciprocally induces the production of NOX4-dependent ROS [167]. NOX4 is selectively upregulated in the lungs of IPF patients and is associated with the endothelial cell dysfunction and hypoxia [14, 55]. Elevated NOX4-generated

ROS triggers DNA oxidation and activates other ROS-dependent signalling pathways such as JNK and NF- κ B [168]. Silencing of NOX4 by siRNA inhibited TGF- β -mediated profibrotic responses in the lungs of mice [55]. Further, NOX4 knockout mice and use of NOX4 inhibitor in mice protected against bleomycin-induced acute lung injury and the onset of fibrosis [76, 169]. In fibroblasts, mitochondrial ROS has been shown to induce the expression of profibrotic genes during fibroblast differentiation [144]. ROS may also modulate integrins, transmembrane receptors that activate FAK, which in turn activate rac1 protein and initiate production of collagen and other profibrotic actors (CTGF and α -SMA) [170]. ROS and RNS may modulate activity of matrix metalloproteinases (MMPs) through the inhibition of cysteine switch and thus influence ECM degradation. ROS are also shown to induce epithelial cell senescence that may result in a diminished capacity for regeneration of epithelium [171]. In IPF, epithelial damage and epithelial cell senescence in the lung are interconnected with increased mitochondrial ROS production. Similarly, in IPF fibroblasts, ROS generation is reported to require for the maintenance and differentiation [172]. Furthermore, oxidative stress may cause ER stress, which facilitates fibrogenesis through activation of EMT, pro-apoptotic pathways and inflammatory responses [146]. In summary, oxidative stress can alter different cellular processes that amplify fibrotic responses.

5.4.8 Oxidative Stress in Airway Mucus Hypersecretion

A thin layer of gelatinous mucus covers the apical epithelial surfaces of mammalian respiratory tract, which forms a protective barrier against airborne microbes and toxins, but conversely, excessive mucus production becomes pathologic in muco-obstructive airway diseases [173]. Mucus is secreted by goblet cells in the airway epithelium and is mainly composed of mucin, which is a large filamentous glycoprotein [174]. Mucus is also rich in antioxidant scavengers such as glutathione, uric acid and ascorbic acid. Chronic airway inflammatory diseases such as chronic bronchitis and asthma are characterized by mucus hypersecretion [175], and ROS (hydroxyl radicals, superoxide anions and hydrogen peroxides) are key regulators of mucus production in goblet cells via transcriptional regulation of mucin genes [176]. Of 12 mucin genes, MUC5AC is a major inducible mucin gene in airways and reported to be highly expressed in muco-obstructive airway disorders [177, 178]. Increased intracellular ROS and exposure to hydrogen peroxide stimulate EGFR tyrosine phosphorylation and subsequent activation of ERK1/2, resulting in increased expression of MUC5AC in lung epithelial cells [176, 179]. Similarly in nasal epithelium, exogenous hydrogen peroxides exposure induces MUC5AC expression through activation of EGFR-ERK1/2 signalling [178]. *Yu et al.* [180] reported that ROS depolymerizes hyaluronan into fragments and these hyaluronan fragments interact with CD44 receptor to activate tissue kallikrein, which cleaves precursors of EGF into mature EGF. Subsequently, mature EGF binds and activates EGFR leading to activation of ERK1/2. In addition, ROS is reported to contribute to goblet cell metaplasia, a major player in mucin overproduction through JAK/STAT

pathway [181]. Likewise, activation of other pathways, such as NF- κ B, is also linked to ROS-mediated MUC5AC production in airways. However, the majority of the studies suggest that EGFR is involved in ROS-mediated mucus hypersecretion [173, 177, 178].

5.5 Antioxidant Responses in Lungs

The lungs are exceptionally exposed to greater oxidative environment than other organs. The inhaled toxicants are by themselves oxidants or may induce oxidative stress inside lung cells. To protect from the inhaled environmental oxidants, lungs are endowed with efficient antioxidant defences that includes both non-enzymatic and enzymatic antioxidant defences.

5.5.1 Antioxidant System in Respiratory Tract Lining Fluid

The airways are covered with respiratory tract epithelial lining fluid (RTLFL) which forms a physical barrier between the external environment and underlying respiratory tract epithelial cell layer. The respiratory tract lining fluid traps most of the inhaled toxicants, and by the help of mucociliary action, these trapped toxicants are cleared from the lungs. The respiratory tract lining fluid is rich in many non-enzymatic low-molecular-weight antioxidant scavengers, which directly interact and detoxify the inhaled oxidants and thereby prevent the direct contact of inhaled toxicants with the underlying epithelium. The major antioxidant molecules in the RTLFL are GSH, ascorbic acid, uric acid and vitamin E (Fig. 5.1). Additionally, airway epithelial cells secrete certain antioxidant proteins into RTLFL, which also function as antioxidant scavengers.

5.5.1.1 Glutathione (GSH)

Glutathione, a thiol-tripeptide comprised of glutamate, cysteine and glycine, is the most important antioxidant in RTLFL, and its levels in RTLFL are close to 100 times more than in the plasma [182]. GSH scavenges a number of ROS products including hydroxyl, H₂O₂, hypochlorous acid and lipid peroxy radical generated during exposures to inhaled oxidants such as cigarette smoke, ozone and allergens, and therefore, reduced bronchoalveolar lavage GSH levels has been a hallmark feature of many pulmonary diseases including COPD, asthma, ARDS and IPF [183]. Besides scavenging ROS, GSH is a co-substrate for the enzyme glutathione peroxidase and glutathione S-transferase which mediate detoxification of lipid hydroperoxides and xenobiotics, respectively. GSH in RTLFL also protects secretory antiproteases such as alpha-1-antitrypsin, alpha-2-macroglobulin and secretory leukoprotease inhibitor from oxidative inactivation [184]. Therefore, a lower level of tissue GSH intensifies oxidant-induced lung inflammatory injury. GSH also maintains thiol status of extra- and intracellular proteins and facilitates post-translational modification of proteins such as S-glutathionylation. Protein S-glutathionylation may alter the function of

many intracellular proteins. For example, S-glutathionylation inhibits DNA-binding activity of p65 or p50 subunits [185], and S-glutathionylation inactivates I κ B β [186] resulting in diminished NF- κ B activity.

5.5.1.2 Ascorbic Acid

Ascorbic acid is another major antioxidant in RTLFL. It directly reduces the oxidative potential of oxidants present in tobacco smoke [187] or particulate matter [188] as well as inhibits ROS generation by NADPH oxidase activity [189]. Besides reducing the inhaled oxidants, ascorbic acid also reduces oxidized antioxidants in RTLFL such as vitamin E, thereby maintaining the total antioxidant capacity of the lungs during oxidative insult. Ascorbic acid is rapidly used in RTLFL fluid upon exposures to environmental oxidants including ozone, nitrogen dioxide, particulate matter and tobacco smoke [190–192] as indicated by the depletion of ascorbic acid levels. Supplementation of ascorbic acid protected from cigarette smoke-induced emphysema by inhibiting protein oxidation in mouse models [192] highlighting the antioxidant potential of ascorbic acid in the lungs. Asthmatics are associated with lower levels of ascorbic acid [193, 194], and the beneficial effect of supplementation of ascorbic acid in asthmatics has been mixed and inconclusive [195]. Ascorbic acid has been shown to attenuate acute lung injury caused by inhalation of oxidant chlorine gas [196]. Ascorbic acid may also take part in pro-oxidant activity in the presence of free iron by taking part in Fenton reaction.

5.5.1.3 Uric Acid

Uric acid formed due to purine metabolism is one of the major water-soluble scavengers of singlet oxygen, ozone and peroxylnitrite (ONOO) in RTLFL [190]. Uric acid has been shown to be a major antioxidant in nasal secretion [197] and RTLFL and helps in the removal of inhaled ozone and neutralizes the oxidative potential of inhaled particulate matter in humans [198]. Uric acid also reacts and neutralizes gaseous free radical nitrogen dioxide [190, 199]. The antioxidant scavenging activity of uric acid greatly depends on ascorbic acid and hydrophilic environment. Uric acid reacts with radical species and forms urate free radical which is then quenched by ascorbic acid. In lipophilic environment, uric acid fails to stop the self-propagating lipid peroxidation reaction. Subnormal levels of serum uric acid were associated with greater risk for COPD and greater morbidity, including reduced 6-minute walk test and greater burden of exacerbations [200, 201].

5.5.1.4 Vitamin E

Vitamin E (tocopherol) is a lipophilic antioxidant scavenger in RTLFL which neutralizes ROS and attenuates self-propagating lipid peroxidation reactions in the airways. Patients with asthma and COPD are associated with lower serum levels of vitamin E [193] as compared to healthy subjects, and this formed the basis for vitamin E supplementation trials to prevent respiratory diseases. Vitamin E trials reduced levels of markers of oxidative damage in smokers [202]. Dietary intake of vitamin E improved lung function in healthy aging population [203]. Vitamin E supplements reduced endotoxin-induced sputum eosinophilia in asthma patients

[204]. In experimental mouse models, administration of vitamin E isoform γ -tocotrienol protected from cigarette smoke-induced emphysema [205] and dust mite-induced asthma. However, supplementation of vitamin E showed no benefits in the management or treatment of asthmatics [206].

5.5.2 Lung-Specific Secretory Proteins in RTLF as Antioxidants

Pulmonary surfactant which lines the alveoli surface is composed of a mixture of 90% phospholipids and 10% surfactant proteins. The surfactant proteins included high-molecular-weight hydrophilic surfactant proteins A and D and low-molecular-weight hydrophobic surfactant proteins B and C. Unsaturated phospholipids and surfactant protein are prone for oxidative inactivation following exposures to environmental oxidants such as ozone. Both surfactant proteins A and D exhibited direct antioxidant activity and protected phospholipids and LDL from copper or ferric chloride-induced oxidation [207]. Surfactant proteins A and D also protected macrophages from hydroperoxide-induced cell death [207]. Clara cell-16 (CC16) protein secreted by clara cells also exhibits antioxidant and anti-inflammatory activity [208]. Mice with genetic disruption of CC16 showed elevated oxidative damage and structural injury following exposure to cigarette smoke [209]. Low circulating levels of CC16 are shown to be associated with poor lung function growth in children [210] and smoking-dependent lung function decline in adults [211] as well as patients with COPD and asthma [208].

5.5.3 Enzymatic Antioxidant System in Lungs

Lungs are endowed with robust antioxidant protein defences to minimize oxidative stress caused by airborne environmental toxicants. Major pulmonary antioxidant enzymes include superoxide dismutase, catalase, glutathione peroxidase, glutathione reductase, hemeoxygenase-1, peroxiredoxin-1, thioredoxin and thioredoxin reductase [40, 41, 212, 213]. Exposures to environmental oxidants such as cigarette smoke lead to the coordinated activation of all these antioxidant proteins [39–41] (as illustrated in Fig. 5.1) and help in efficient detoxification of ROS and lipid peroxides generated in the lungs. The importance of individual antioxidant enzyme has been exemplified using knockout and transgenic mouse models.

All the three isoforms of superoxide dismutase, extracellular-SOD (EC-SOD), copper/zinc-SOD (Cu/Zn-SOD) and manganese-SOD (Mn-SOD), are present in lungs and provide first line of defence against superoxide radicals. The importance of each of the SOD isoforms in protecting lungs from oxidants has been well studied. EC-SOD prevented fibrosis in lungs by inhibiting oxidative degradation of the matrix proteins, type I and type IV collagen [214]. Cigarette smoke exposure and elastase instillation caused greater emphysema in EC-SOD-deficient mice; however transgenic EC-SOD mice were protected from emphysema [215]. Similarly, over-expression of human Cu/Zn-SOD in lungs protected from cigarette smoke-induced

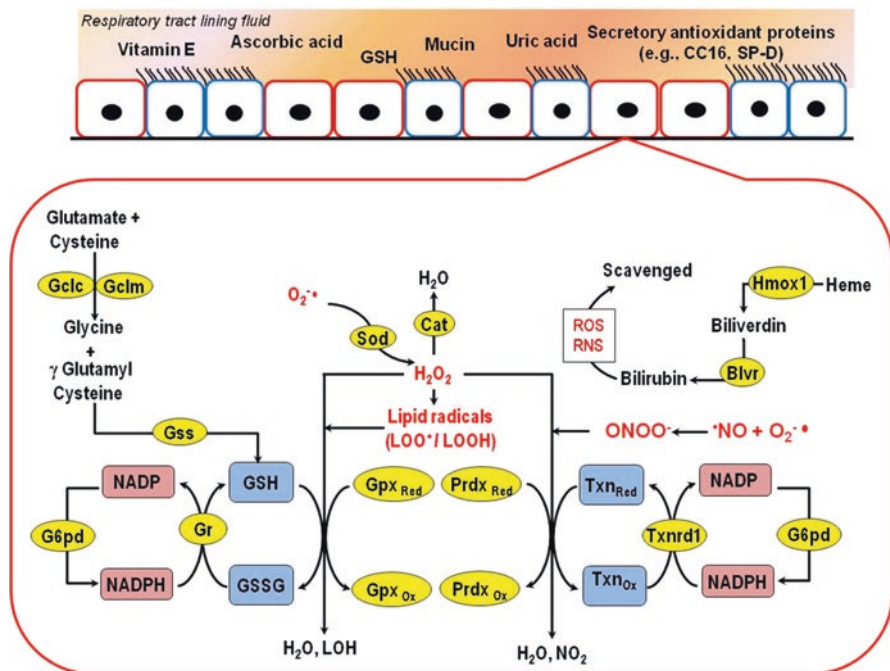


Fig. 5.1 Antioxidant defenses in lungs

emphysema in mouse model [216]. Overexpression of Mn-SOD has also been reported in alveolar macrophages during sarcoidosis and in the lung tissue of IPF patients [213]. SOD is highly sensitive for oxidative inactivation, and therefore, SOD levels are depleted in disease lungs. In IPF, there was no expression of EC-SOD in fibrotic areas where there is an enhancement of oxidative burst [214]. Enzyme activity of SOD was significantly low in asthmatics as compared to healthy subjects, which is further depleted during asthma attack [217, 218]. Oxidative and nitrosative modification of Mn-SOD was observed in the airways of asthmatic, which correlated with asthma severity [217]. Therefore, increasing SOD levels in the lungs by pharmacological approaches including SOD mimetics are thought to be a promising approach for mitigating pathogenesis of pulmonary disorders.

Glutathione peroxidase (GPx) detoxifies hydrogen peroxide and reactive lipid hydroperoxides using GSH as an electron donor. In mammalian lungs, four major selenium-containing GPx isoforms are expressed – *GPX1* (classical GPx), *GPX2* (gastrointestinal GPx), *GPX3* (extracellular GPx) and *GPX4* (phospholipid GPx). Bronchial epithelial cells and alveolar macrophages produce Gpx3 in epithelial lining fluid, which detoxify lipid hydroperoxides generated in RTLF. GPx2 is a predominant glutathione peroxidase expressed in the lungs following cigarette smoke exposure and silencing GPx2 by RNA interference enhanced cytotoxicity in bronchial epithelial cells following treatment with cigarette smoke extract [42]. In comparison with wild type, GPx2-deficient mice showed greater levels of oxidative

damage, airway inflammation and airway hyperresponsiveness in ovalbumin-induced asthma mouse model [219]. Basal levels of GPx in lungs were shown to be a key determinant of severity of pulmonary fibrosis in mouse models [220]. Peroxiredoxins are the family of peroxidase enzymes, which play a dominant role in detoxification of hydrogen peroxide within the cells. Human lung expresses all the six members of a peroxiredoxin family [221]. Thioredoxin reductase is a selenium-containing flavoprotein oxidoreductase enzyme expressed in mammalian lungs, which primarily catalyses NADPH-dependent reduction of thioredoxin, an important redox protein involved in transcriptional regulation of NF- κ B [94]. Heme oxygenase-1 (HO-1) is a highly inducible protein in the lungs which exhibits anti-apoptotic, anti-inflammatory and antioxidant activities. HO-1 catalyses heme to carbon monoxide and biliverdin and the latter is converted to bilirubin. Although the mechanism by which HO-1 mediates antioxidant and anti-inflammatory activity is less understood, the end byproducts of HO-1 enzyme activity, CO, bilirubin and Fe are shown to mediate the beneficial effects [222]. HO-1 knockout mice display greater inflammation, apoptosis and tissue injury following an ischemic reperfusion injury [223], while lung-specific expression mitigated LPS- and hyperoxia-induced lung inflammation [224, 225].

5.5.4 Regulation of Antioxidant Enzymes in Lungs

Many lines of evidence show that transcription factor Nrf2 is a central regulator of nearly all cellular antioxidant proteins in the lungs and other organs [226]. In a normal cell, Nrf2 is held in the cytoplasm by a cysteine-rich, redox sensor Keap1 protein, which functions as an adaptor molecule and bridges Nrf2 with Cul3-based E3 ubiquitin ligase [227, 228]. Under normal condition, Keap1-Cul3-based E3 ubiquitin ligase ubiquitinates Nrf2 and directs it to proteasomal degradation. However, upon exposure to ROS and electrophiles, Keap1 protein undergoes conformational change due to oxidative modification of its cysteine residues, which disrupts the interaction of Nrf2 with Cul3-dependent E3 ligase and prevents Nrf2 ubiquitination. Stabilized Nrf2 moves into the nucleus and mediates transcriptional activation of its target genes by binding to cis-element called 'antioxidant response element' in the promoter region. Genetic disruption of Nrf2 ablates transcriptional induction of antioxidant genes in the lungs and sensitizes the mice to several environmental lung diseases such as cigarette smoke-induced emphysema [41], allergen-induced asthma [229], LPS-induced acute lung injury [97] and bleomycin-induced pulmonary fibrosis [230] and sepsis [72, 74, 231]. In contrast, activation of Nrf2 by pharmacological activators and genetic disruption of Keap1 protected mice from development of these pulmonary diseases [226]. Nrf2-regulated antioxidant has been shown to be downregulated in lungs of patients with COPD [232] and IPF [233], underscoring the importance of Nrf2 pathway in protecting the lungs from oxidative stress. Besides Nrf2, NF- κ B and AP-1 also regulate transcriptional expression of certain antioxidant genes in the lungs [88, 234].

5.5.5 Antioxidant Therapy for Lung Diseases

Despite the compelling evidence from preclinical and clinical studies that pulmonary antioxidants play a pivotal role in protecting from environmental pulmonary diseases, clinical trials testing antioxidant therapy have shown modest to no significant beneficial effects. Clinical trials with N-acetyl-L-cysteine (NAC) supplementation have shown mixed results. Meta-analysis of all clinical trials using oral NAC concluded that long-term intake of NAC may reduce the acute exacerbations of chronic bronchitis [235]. However, more recent randomized double-blinded multi-centre clinical trial of oral NAC reported no beneficial effect in the prevention of lung function decline and COPD exacerbation [236]. Vitamin C and E clinical trials showed no improvement on lung function decline in COPD patients [237]. Supplementation of Nrf2 activator, sulforaphane, in the form of broccoli sprout homogenates reduced bronchoconstrictor hyperresponsiveness in asthmatics [238]. In a randomized clinical trial, supplementation of sulforaphane showed no significant upregulation of Nrf2-regulated antioxidants in lungs of COPD patients [239]. In another study, consumption of broccoli sprout showed no effect on eosinophilic inflammation as well as markers of oxidative stress in atopic asthmatic patients [240]. In smokers, consumption of broccoli sprout homogenates reduced influenza virus-induced inflammation [241]. Consumption of broccoli tea has been reported to promote rapid and sustain detoxification of air pollutants in a randomized clinical trial in China [242]. The reasons for lack of consistent benefits of antioxidant trials for pulmonary disease are still puzzling. Perhaps it could be combination of poor efficacy of a single antioxidant agent as well as limited bioavailability.

5.6 Conclusions

Oxidative stress is a central hallmark pathological feature of all the respiratory disease. Oxidative stress elicits both reversible and irreversible macromolecule damage (oxidative modification of lipids, proteins and DNA). As illustrated in Fig. 5.2, besides inducing macromolecular damage, oxidative stress propagates the disease by augmenting other pathological processes such as inflammatory responses, mitochondrial dysfunction, ER stress, profibrotic signalling, cell death and epigenetic changes. Experimental evidences suggest that antioxidant therapy may prevent or mitigate oxidative stress-mediated macromolecular damage and abnormal signal transductions and, thereby, protect from development and progression of disease. However so far, most antioxidant clinical trials have shown poor efficacy to mitigate disease progression, which may be attributed to insufficient bioavailability of antioxidant agent in the lungs and also inability to reverse pathogenic events such as epigenetic changes, macromolecule damage and mitochondrial dysfunction. Antioxidant alone may not be effective in treatment of diseases; however it could be a promising adjunctive therapy.

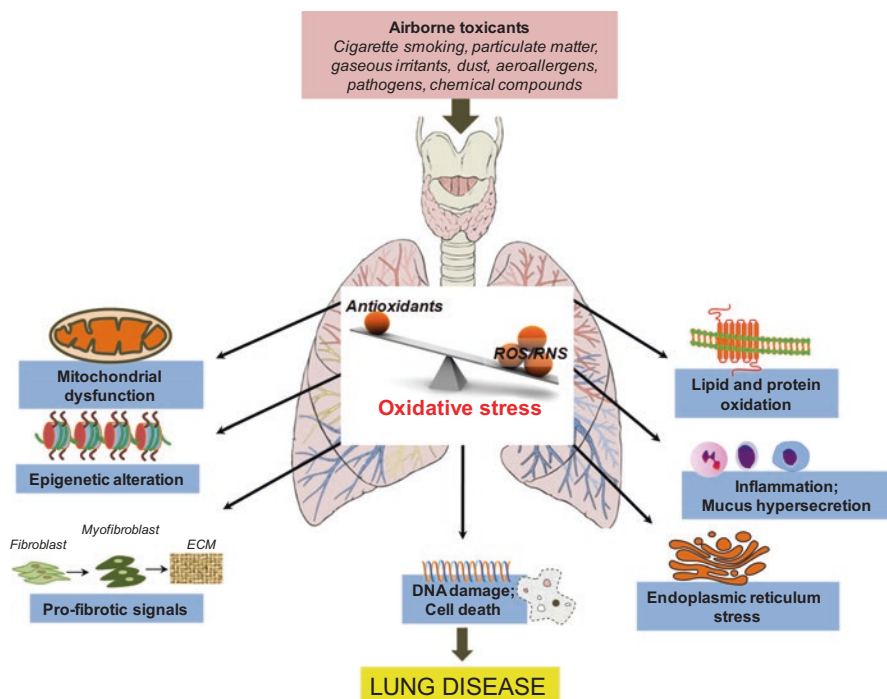


Fig. 5.2 Oxidative stress-mediated pathological events in driving lung diseases

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