



Published in final edited form as:

J Pharmacogenomics Pharmacoproteomics. ; 5(4): .

Respiratory Viral Infections and Subversion of Cellular Antioxidant Defenses

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Abstract

Reactive oxygen species (ROS) formation is part of normal cellular aerobic metabolism, due to respiration and oxidation of nutrients in order to generate energy. Low levels of ROS are involved in cellular signaling and are well controlled by the cellular antioxidant defense system. Elevated levels of ROS generation due to pollutants, toxins and radiation exposure, as well as infections, are associated with oxidative stress causing cellular damage. Several respiratory viruses, including respiratory syncytial virus (RSV), human metapneumovirus (hMPV) and influenza, induce increased ROS formation, both intracellularly and as a result of increased inflammatory cell recruitment at the site of infection. They also reduce antioxidant enzyme (AOE) levels and/or activity, leading to unbalanced oxidative-antioxidant status and subsequent oxidative cell damage. Expression of several AOE is controlled by the activation of the nuclear transcription factor NF-E2-related factor 2 (Nrf2), through binding to the antioxidant responsive element (ARE) present in the AOE gene promoters. While exposure to several pro-oxidant stimuli usually leads to Nrf2 activation and upregulation of AOE expression, respiratory viral infections are associated with inhibition of AOE expression/activity, which in the case of RSV and hMPV is associated with reduced Nrf2 nuclear localization, decreased cellular levels and reduced ARE-dependent gene transcription. Therefore, administration of antioxidant mimetics or Nrf2 inducers represents potential viable therapeutic approaches to viral-induced diseases, such as respiratory infections and other infections associated with decreased cellular antioxidant capacity.

Keywords

Respiratory syncytial virus; Oxidative stress; Nrf2; ROS; Free radicals

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Introduction

Molecular oxygen is essential for supporting the life processes of all aerobic organisms. Under physiological conditions, oxygen is combusted in a highly controlled manner by the cell's metabolic machinery to obtain chemical energy in form of ATP, and this process leads to the formation of reactive oxygen species (ROS) [1,2]. ROS are unstable molecules, which in small quantities are involved in cellular signaling, but become toxic when produced in large quantities by initiating oxidation of cellular components such as proteins, lipids, and DNA [1]. ROS are broadly classified into two groups, radical and non-radicals. Members of the radical group, often called free-radicals, have at least one unpaired electron in the outer orbital and therefore are highly reactive, as they readily donate or accept an additional electron to achieve stability [3,4]. This group includes compounds such as superoxide ion radical (O_2^-), hydroxyl radical (OH^\cdot), nitric oxide radical (NO^\cdot), peroxy (ROO^\cdot) and alkoxy radicals (RO^\cdot) [1,5,6]. The non-radicals group includes compounds such as hypochlorous acid (HClO), hydrogen peroxide (H_2O_2), organic peroxides and aldehydes. In addition to endogenous ROS, exogenous compounds such as air pollutants, cigarette smoke, radiation, heavy metals etc., can generate ROS [7]. In order to protect from the continuous exposure of exogenous and endogenous ROS, organisms have developed a complex antioxidant system which include enzymatic (superoxide dismutase, catalase, glutathione peroxidase, etc.) and non-enzymatic (transferrin, ferritin, vitamin A and C, etc.) defenses. Failure to keep the equilibrium between ROS formation and antioxidant defenses leads to oxidative stress. This is characterized by an augmented generation of oxidant species and reduced antioxidant cellular capacity [2,8–11]. At molecular level, the oxidative damage to DNA cause polysaccharide ring cleavage, base modification or chain breakage, leading to mutations and altered/ failed gene transcription; damage to proteins can modify functional groups, such as addition of nitro radicals and carbonyl groups, resulting in altered activity, aggregation, fragmentation and/or cleavage; damage to lipids leads to formation of lipid aldehydes, lipid peroxides, causing changes in fluidity and permeability of membranes [6,12,13].

While ample information is available about the mechanism(s) of increased ROS generation, little is known about the regulating changes in antioxidant enzymes (AOE) expression [8]. At the gene expression level, many of the genes coding for AOE are controlled by the redox sensitive transcription factor NF-E2-related factor 2 (Nrf2), binding to promoter antioxidant responsive element (ARE) sites. These ARE elements are also present in the regulatory regions of many genes encoding phase-2 detoxification enzymes and various cytoprotective proteins, such as NAD(P)H:Quinoneoxidoreductase (NQO1) [14–16]. Nrf2 is a cap'n collar basic leucine-zipper transcription factor, which under normal physiologic conditions is sequestered in the cytoplasm by Kelch-like ECH associated protein 1 (Keap1), forming a complex bound to the cytoplasmic membrane through actin [17,18]. In the presence of elevated levels of ROS and cellular oxidative stress, Nrf2 is released from this complex by conformational change in cysteine disulfide bonds of Keap1 [19–22]. Nrf2 is then phosphorylated at serine 40 by protein kinase C and translocate to the nucleus [23], where it forms DNA-protein complexes with transcription factors belonging to the small musculoaponeurotic fibrosarcoma (Maf) and transcriptional co-activators, such as CREB

binding protein (CBP) and p300, to initiate transcription of ARE-dependent genes [17]. Once the cellular redox status returns to equilibrium, Keap1 sequesters Nrf2 and directs it to Cul3-ubiquitin mediated proteasome degradation [17,21]. In the past few years, the Nrf2-Keap1/ARE system has been the focus of intense investigation, because of its possible role in the pathogenesis of several diseases [16,24].

Generation of oxidative stress has been reported in about 200 diseases [24] and oxidative stress is thought to play an important pathogenic role in pulmonary disorders such as chronic obstructive pulmonary disease (COPD) [25] and asthma [26–28], cancer [29,30], neurological diseases, including Alzheimer's [24,31], cardiovascular [32] and metabolic disorders, such as diabetes [33], vision disorders [34] and aging [35]. This review focuses on Nrf2 and oxidative stress associated with respiratory viral infections with emphasis on respiratory syncytial virus (RSV), although other viruses, including human immunodeficiency virus (HIV), Hepatitis B and C have been shown to induce ROS *in vitro* and *in vivo* [36–39]. Free radicals generated during various respiratory viral infections are shown in Table 1.

Respiratory syncytial virus

RSV is an enveloped, negative-sense, single-stranded RNA virus belonging to the *Paramyxoviridae* family, and is the leading cause of respiratory diseases in infants and young children. Annually in the US alone, RSV infections are responsible for more than 100,000 hospitalizations among children <1 year of age and accounts for ~1.5 million outpatient visits among children <5 years of age, with an economic burden of more than \$500 million/year [40–42]. Worldwide each year, an estimated 33.8 million new episodes of RSV-associated acute lower respiratory tract infections (ALRI) occur in children <5 years of age, with about 3.4 million children requiring hospital admission, and an estimated 66,000–199,000 fatal cases, mostly in developing countries [43]. RSV infection is also a major concern in elderly people with chronic heart and lung diseases, and in immunocompromised patients [44]. Around 1500 to 7000 deaths due to RSV infection occur in the USA alone each year, especially in people >65 years of age [45]. Although RSV has been the focus of intense investigation for several decades, no effective drug or vaccine is currently available [46]. While the mechanisms of RSV-induced airway disease and its associated long-term consequences are not fully understood, lung inflammatory response and oxidative stress are important pathophysiological features of RSV lower respiratory tract infections [47]. This review focuses on the potential role of oxidative lung damage in RSV pathogenesis and possible novel therapeutic approaches targeting ROS formation and Nrf2 activation in the context of this, as well as other respiratory viral infections.

ROS generation in RSV infection

As mentioned before, ROS formation occurs as part of aerobic cellular metabolism and plays an important role in cellular signaling, leading to the expression of a variety of molecules, including proinflammatory mediators, such as cytokines and chemokines [48]. If ROS are not neutralized by cellular antioxidant systems, they can cause extensive cellular and tissue damage. Many of the features of acute and chronic lung diseases, such as bronchoconstriction, airway hyper reactivity, enhanced mucous secretion, epithelial cell

damage, and microvascular leakage, have been shown to be associated with oxidative stress due to increased generation of ROS [28]. RSV infection has been reported to enhance ROS formation in airway epithelial cells, the primary target of infection, as measured by the fluorescent probe 2',7' dichlorodihydrofluorescein diacetate [49–52]. RSV infection leads to the release of superoxide, H₂O₂ and myeloperoxidase (MPO) in the extracellular environment by inducing the recruitment and activation of neutrophils and eosinophils into the airways [53,54]. ROS generation in airway epithelial cells during RSV infection was recently summarized in a review by Garofalo et al. [47]. Several NADPH oxidase inhibitors, including diphenyleneiodonium chloride (DPI), apocynin, and 4-(2-aminoethyl) benzene sulfonyl fluoride (AEBSF), inhibit RSV-induced cellular signaling in airway epithelial cells, in particular chemokine expression, as well as activation of the transcription factors interferon regulatory factor-3 (IRF-3), signal transducer and activator of transcription (STAT)-1 and the upstream kinase inhibitor of κ B kinase epsilon (IKK ϵ) [55–58].

In regard to other respiratory viruses, rhinovirus infection has been associated with superoxide and hydrogen peroxide production through NOX1 [59] and influenza infection generates superoxide through NOX2 [60], as shown in Table 1.

Oxidative stress in RSV infection

RSV-induced ROS formation is associated with significant cellular oxidative stress *in vitro* as well as *in vivo*, as a result of disruption of the fine balance between pro-oxidant and antioxidant factors. During RSV infection of airway epithelial cells, SOD 2 expression and activity progressively increases, with a progressive decrease in the expression of all the other tested AOE such as SOD 1, SOD 3, catalase, GST expression, and GPx activity. These changes in AOE expression suggest that increased amounts of superoxide, generated by RSV through NADPH oxidase, could result in accumulation of H₂O₂ by increased SOD 2 activity and reduced activity of catalase, GST and GPx [8]. The non-detoxified H₂O₂, as well as other radicals generated from H₂O₂ auto-oxidation in presence of transition metals, such as the hydroxyl radical (\cdot OH), reacts with lipids, proteins and DNA, causing structural cellular damage. Proteomics studies have also shown that several AOE, from peroxiredoxins to catalase, SOD 1, GPx 1 and various forms of GST, are significantly decreased in the lungs of infected animals compared to uninfected (Table 2 summarizes all antioxidant proteins whose expression in broncho alveolar lavage (BAL) changes in response to RSV infection). Decreased expression/activity of antioxidant proteins was confirmed *in vivo*, both in a mouse model of RSV infection as well as in children with severe bronchiolitis [11].

Lipid peroxidation refers to the oxidative degradation of lipids. It is the process in which free radicals “steal” electrons from the lipids in cell membranes, resulting in cell damage. It most often affects polyunsaturated fatty acids. The end products of lipid peroxidation are reactive aldehydes, such as malondialdehyde (MDA) and 4-hydroxynonenal (HNE), the second one being known also as “second messenger of free radicals” and major bioactive marker of lipid peroxidation, due to its numerous biological activities resembling activities of reactive oxygen species. In addition to MDA and HNE, 8-isoprostane are also considered markers of cellular oxidative stress, as they are formed *in vivo* from the free radical-

catalyzed peroxidation of essential fatty acids (primarily arachidonic acid) without the direct action of cyclooxygenase enzymes. The unbalance between ROS formation and antioxidant defenses leads to oxidative stress during the course of RSV infection, as it has been demonstrated by the increased formation of lipid peroxidation products both *in vitro* and *in vivo* models of infection [8,61], as well as in patients with primary RSV infection, in which the levels of 8-isoprostane, as well as MDA and HNE present in respiratory secretion correlate with the severity of infection [11].

In addition to RSV, the closely related human metapneumovirus (hMPV), which is also a common cause of lower respiratory tract infections in children [62], significantly affects AOE expression *in vitro* and *in vivo*. Microarray analysis of gene expression studies from hMPV infected airway epithelial cells demonstrated progressively decreased levels of SOD 3, catalase, GST and peroxiredoxin gene expression and protein levels, with a concomitant increase in SOD 2 [63], similar to what has been observed with RSV. These changes in AOE expression was also observed in a mouse model of hMPV infection [11]. Such an increase in SOD 2 expression and decreased expression of catalase, as well as decreased GSH/GSSG ratio, has also been reported in influenza infection both *in vitro* and *in vivo* [64–66]. Taken together, this information suggests that airway oxidant-antioxidant imbalance could play a very important role in the pathogenesis of RSV-induced lung disease and possibly other respiratory viral infections.

Potential regulatory mechanism of AOE expression in RSV infection

The exact mechanism of decreased expression of AOE during RSV infection, as well in the context of other viral respiratory infections, is largely unknown. Most of the AOE gene expression is regulated in part through ARE sequences and Nrf2 activity [30,67]. Transcription factor Nrf2 is an important redox-responsive protein that protects the cells from oxidative stress and injury (Reviewed in [16]). Nrf2-dependent AOE gene expression might be reduced by: (i) competition for binding to the ARE site - Bach1/small Maf protein complex or AP-1 family transcription factors like c-Fos and FRA1 can bind to ARE acting as a transcriptional repressor [67,68]; (ii) preventing Nrf2 activation through direct physical association - Activating transcription factor (ATF)3 or retinoic acid receptor α were shown to form inhibitory complexes with Nrf2, leading to displacement from ARE elements; (iii) interfering with recruitment of co activators, such as CBP, to the ARE site - NF- κ B activation can lead to decreased availability of CBP and promote the recruitment of co repressors (histone deacetylases) at Nrf2-ARE site [67]; (iv) reduced nuclear levels, which can occur due to enhanced nuclear to cytoplasm efflux or increased Nrf2 degradation [8]. A recent study has shown that RSV infection in *Nrf2*^{-/-} mice is more severe and associated with higher viral titers, augmented inflammation, enhanced mucus production and epithelial injury compared to Nrf2 wild type mice, indicating the protective role of Nrf2-ARE pathway against RSV infection [69]. RSV infection can indeed induced a progressive decrease in ARE-dependent gene transcription in A549 cells, carcinoma-derived type II-like airway epithelial cells, investigated using luciferase reporter gene assays (Figure 1A, left panel) [47]. A similar result was obtained when cells were infected with hMPV (Figure 1A, right panel). Reduced nuclear levels of Nrf2 was observed in both RSV infection (Figure 1B, left panel) [8] and hMPV infection (Figure 1B, right panel), together with increased nuclear

levels of known ARE transcriptional repressors such as Bach1 and ATF3 (Casola A, unpublished observation) [47], suggesting a potential mechanism for viral-induced down regulation of AOE gene expression. As Nrf2 positively regulates its own gene transcription, reduced Nrf2 mRNA levels were observed in airway epithelial cells at late time point of RSV infection [8]. Our recent studies indicate that RSV infection is associated with Nrf2 deacetylation, likely due to RSV-induced upregulation of histone deacetylase (HDAC) activity, and increased degradation, which occurs through the ubiquitin-proteasome pathway. Blocking proteasome and class I HDAC activity, in particular HDAC 1 and 2, rescued Nrf2 activation and ARE-dependent gene expression during RSV infection (Casola A, unpublished observation). A summary of findings and a proposed model of RSV-induced oxidative stress in airway epithelial cells are depicted in Figure 2.

In regard to other respiratory infections, Yageta et al. showed that influenza infection in Nrf2-deficient mice is associated with increased mortality, compared to wild type mice, when animal are exposed to cigarette smoke. Nrf2-deficient mice could not control the oxidative stress caused by cigarette smoke and showed enhanced peribronchial inflammation, lung permeability damage, and increased mucus secretion [70]. Inhibition of Nrf2-dependent gene expression in differentiated human nasal epithelial leads to increased influenza virus entry and replication, due to increased oxidative stress [39]. In addition, overexpression of Nrf2 in alveolar type II cells provides protection from influenza infection by reducing oxidative stress and viral replication [71]. These data suggest that Nrf2 plays an important role in influenza infection by controlling ROS formation, viral replication and lung injury.

Therapeutic approaches

Since oxidative stress seems to play an important role in the pathogenesis of RSV, and possibly other viral-associated lung diseases, antioxidant intervention would represent a rational approach for treatment of lower respiratory tract infections. Table 3 shows antioxidant therapies tested against various respiratory viral infections. Antioxidants reduce oxidative stress by quenching free radicals and help the host to function properly. Our group has tested two complementary approaches that can affect the outcome of RSV-associated lower respiratory tract infections: (i) SOD mimetics that can scavenge free radicals and reduce oxidative stress in RSV infected cells and; (ii) induction of airway antioxidant defenses by modulating AOE gene expression/activity.

SOD mimetics—SOD 1 and 2 administration and SOD 3 overexpression have been shown to protect mice lungs from influenza-induced oxidative stress damage [72]. Both SOD 1 and 2 administration, either parenterally or intranasal in a cotton rat model of RSV infection, reduced pulmonary viral titer [73]. In the past few years, quite a few classes of synthetic SOD mimetics that are based on organo-manganese complexes have been developed and explored as possible therapeutics against oxidant-related lung damage [74]. In a recent study, the effect of airway epithelial cell treatment with various Eukarionsalen-manganese complexes (EUK) on cellular signaling and oxidative stress in response to RSV infection was assessed. EUKs are synthetic salen-manganese complexes that exhibit SOD and catalase activities [75]. Salen complexes are Schiff bases, usually prepared by the

condensation of a salicylaldehyde with an amine. Based on the salen ring substitutes, EUKs are named from EUK-8 to EUK-189 and have different rates of SOD and catalase/ peroxidase activities [76,77]. Treatment of RSV-infected airway epithelial cells with EUK-8, -134 and -189, which have SOD and catalase/peroxidase activity, results in significantly reduced ROS levels (Figure 3A) and markers of oxidative cell damage (Figure 3B). In addition, EUK treatment is associated with reduced activation of the viral-induced transcription factors NF- κ B and IRF-3 and reduced secretion of cytokines and chemokines [8,78]. Enhancement of both SOD and catalase/GPx activities is important to reduce ROS levels and pro-inflammatory gene expression during RSV infection, as EUK-163, which has no significant catalase or peroxidase activity, does not have a significant effect on RSV-induced pro-inflammatory mediator secretion [8]. In addition, EUK treatment in high concentration (500 μ M) significantly reduce viral replication [78], suggesting that EUKs could represent a novel therapeutic approach to modulate RSV-induced lung damage.

Nrf2 inducers—Several compounds that stimulate Nrf2-ARE driven transcription have been identified from natural and dietary sources, metabolites, and synthetic agents. Nrf2 inducers are broadly divided into Triterpenoids that include oleanolic acid and ursolic acid (natural); oleanane triterpenoids, 2-cyano-3,12-dioxooleana-1,9,-dien-28-oic acid (Synthetic) [79,80]; Isothiocyanates including sulforaphane, found mainly in cruciferous vegetables; polyphenols including flavonoids quercetin and EGCG and the non-flavonoids curcumin, resveratrol and butylated hydroxyanisole (BHA) [81,82].

Sulforaphane modifies a number of cysteine residues in Keap1 through formation of carbamodithioate and releases Nrf2 that leads to increased nuclear localization of Nrf2 and ARE transcription [82]. Sulforaphane pretreated nasal epithelial cells during influenza virus infection showed significantly increased levels of Nrf2 and HO-1 associated with reduced hemagglutinin gene expression and viral replication [39]. In a model of RSV infection, mice treated with sulforaphane showed significantly reduced numbers of neutrophils and eosinophils in BAL after infection [69], suggesting that this compound has the potential for modulating viral-induced oxidative stress and disease.

BHA and t-Butyl hydroquinone (tBHQ) treatment induces phase II enzymes HO-1 and NQO1 *via* Nrf2-ARE transcription in rat and human hepatocytes [83]. Hence, BHA was assessed for its ability to modulate oxidative stress in a mouse model of RSV infection. BHA treatment significantly attenuated RSV-induced lung oxidative stress, as indicated by decreased markers of oxidative damage in BAL of RSV-infected mice. In addition, lungs of BHA treated mice showed reduced cytokine and chemokine secretion [61]. The beneficial effect of BHA and tBHQ in RSV-induced lung inflammation and oxidative stress could be in part ascribed to the ability of these phenolic compounds to modulate Nrf2-dependent gene expression, in addition to directly scavenging ROS formed in response to the viral infection. Preliminary studies revealed that treatment of airway epithelial cells with tBHQ significantly increased ARE-dependent gene transcription and Nrf2 protein expression. tBHQ treatment rescued Nrf2-ARE driven activity during RSV infection and also ameliorated RSV induced oxidative damage as demonstrated by reduced lipid damage (Casola A, unpublished observation).

Conclusion

Respiratory tract infections are a leading cause of morbidity and mortality worldwide. RSV and other viruses such as influenza and hMPV are a major cause of pediatric upper and lower respiratory tract infections, associated with bronchiolitis, pneumonia and flu-like syndromes, as well as asthma exacerbations. There is still no vaccine or effective treatment available for RSV infections, as well as for many other respiratory viruses, necessitating an explicit understanding of the pathogenic mechanisms associated with these infections. As oxidative stress is likely to play an important role in initiating and sustaining lung injury and inflammation, approaches that combine scavenging ROS together with the inhibition of viral replication, may be effective in modulating severe lung disease associated with RSV and other viral respiratory infections. This could be obtained by either administration of antioxidant compounds that possess antiviral activity, in addition to ROS scavenging properties, or by combining antivirals with compounds capable of increasing lung antioxidant defenses, such as AOE mimetics or Nrf2 inducers. These treatment approaches would be effective only if compounds are available at the site of infection, therefore route of administration, bioavailability, tissue distribution are all important parameters that will need to be taken into consideration when planning future therapeutic intervention.

Acknowledgments

Funding Sources

This article was funded by NIH grant fundings: R01 AI079246, P01 AI062885 and R21 AI103565. The authors do not have anything to disclose.

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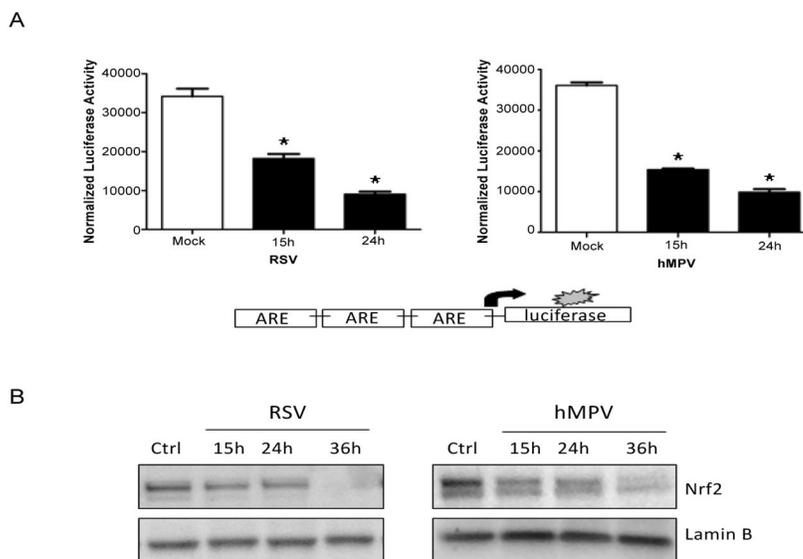


Figure 1. RSV and hMPV infection modulates ARE-dependent gene transcription

(A) A549 cells were transiently transfected with a plasmid containing multiple copies of the NQO1 ARE site linked to the luciferase gene and then infected with either RSV (Left panel) or hMPV (Right panel). Cells were harvested at different times post-infection to measure luciferase activity. Uninfected cells, transfected with reporter plasmid only and mock-infected, served as controls. For each plate luciferase was normalized to the β -galactosidase reporter activity. Data are expressed as mean \pm standard deviation of normalized luciferase activity. * $P < 0.05$ relative to RSV or hMPV infected cells. (B) Nuclear extracts prepared from A549 cells infected with RSV (left panel) or hMPV (right panel) for various periods of time post infection (p.i.) were subjected to western blot with anti Nrf2 antibody. Membranes were stripped and re-probed for lamin B as an internal control for protein integrity and loading.

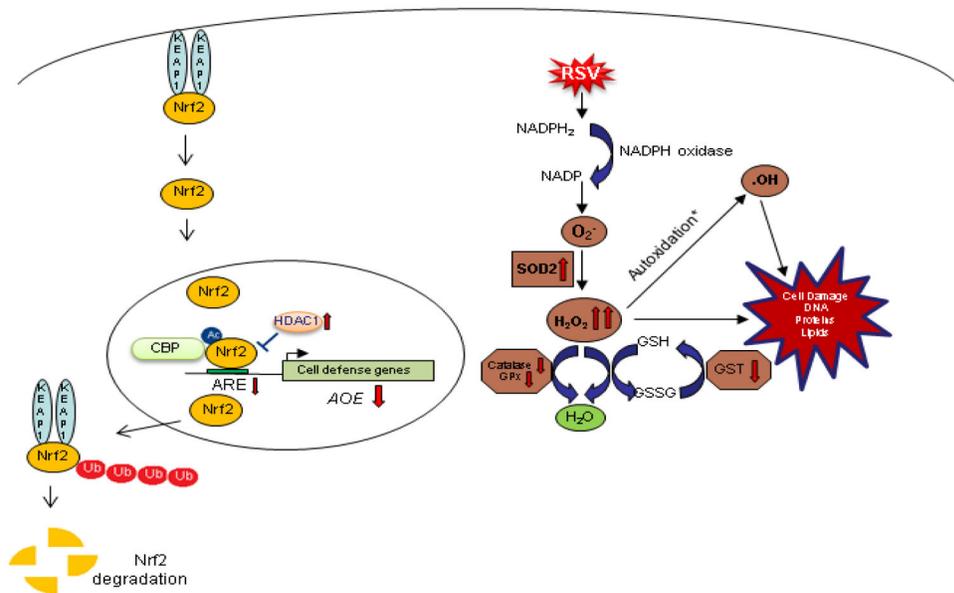


Figure 2. Schematic representation of the proposed mechanisms of oxidative cell damage during RSV infection

RSV infection of airway epithelial cells leads to increased superoxide formation and increased H₂O₂ production, due to up regulation of SOD 2 expression and activity. RSV-induced inhibition of Nrf2 activation, due to proteasome-dependent degradation, causes a progressive decrease in the expression of a variety of AOEs involved in H₂O₂ detoxification leading to accumulation of highly reactive radicals, such as hydroxyl radical, and subsequent cellular damage (* autoxidation in presence of transition metals).

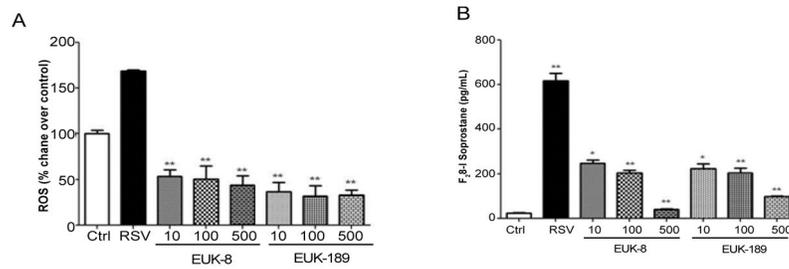


Figure 3. Effect of EUK treatment on RSV-induced ROS formation and oxidative stress
 (A) A549 cells were treated with different micromolar concentrations of EUK-8 and EUK-189, infected with RSV for 24 h, and harvested to measure DCF-DA fluorescence. Ctrl indicates uninfected cells. Mean Fluorescence Intensity is reported as percent increase over control. (B) Cell supernatants were harvested at 24 h p.i. to measure F2-isoprostanes. Results are expressed as mean \pm standard error. Results are representative of two independent experiments run in triplicate. * $p < 0.05$, ** $p < 0.01$ compared to untreated RSV-infected cells. Reprinted with permission of the American Physiological Society. Copyright © 2014 The American Physiological Society [78].

Table 1

Free radicals generated in response to respiratory viral infections.

Virus	Free radical generated	Effect	Host proteins associated with oxidative stress	Ref
Rhinovirus	superoxide (O_2^-), hydrogen peroxide (H_2O_2)	airway inflammation	activation of xanthine oxidase; reduced concentration of glutathione (GSH) and increased activity of NADPH oxidase 1 (NOX1)	[59,84–87]
Influenza Virus	O_2^- , nitric oxide (NO)	enhanced viral mutations, replication and airway inflammation	reduced concentrations of catalase, glutathione and super oxide dismutase (SOD); Increased activity of NADPH oxidase 2 (NOX2)	[60,66,88–91]
RSV	NO, O_2^- , H_2O_2	airway inflammation	virus induced nitric oxide synthase (iNOS) activity; progressive decrease of antioxidant enzymes SOD 1, SOD 3 and Catalase; reduced nuclear translocation of Nrf2 and Nrf2-ARE driven transcription	[8,11,52,92] (Casola A, unpublished observation)
hMPV	O_2^- , H_2O_2	airway inflammation	progressive decrease of antioxidant enzymes SOD 3, catalase, GST, and Prdx; reduced nuclear translocation of Nrf2 and Nrf2-ARE driven transcription	[11,63] (Casola A, unpublished observation)

Table 2

Differential expression of antioxidant proteins in bronchoalveolar lavage of respiratory syncytial virus-infected mice.

AOE	Fold Change in RSV BAL Compared to Control				
	Day 1	Day 3	Day 5	Day 9	Day 25
1-Cys peroxidase protein	-1.0	-6.1	—	-4.1	—
Catalase	—	-2.5	-2.1	—	—
Cu/Zn SOD 1	-2.3	-3.4	-2.0	-2.0	—
Glutathione peroxidase 1	-1.8	-2.3	—	1.3	—
Glutathione S-transferase	—	—	—	-6.0	—
Glutathione S-transferase omega 1	-6.8	-3.6	-2.3	-2.0	1.3
Glutathione S-transferase, alpha 4	-2.2	—	—	—	—
Glutathione S-transferase, mu 1	—	-4.0	-7.0	-1.7	1.4
Glutathione S-transferase, mu 2	—	-4.3	—	3.4	-1.3
Glutathione-disulfide reductase	—	—	—	3	—
Nonelenium glutathione peroxidase	-2.6	—	-4.2	-1.3	1.2
Peroxiredoxin 6	—	-3.1	-3.9	-4.1	1.3
Peroxiredoxin 2	2.7	2.4	-2.1	1.7	—
Thioredoxin 1	—	1.5	—	—	1.1

Shown are high probability antioxidant protein identifications and their expression (in terms of fold changes in RSV BAL compared to control mice) at different days of p.i. from peptide mass fingerprinting in MALDI-TOF/MS. BAL, bronchoalveolar lavage; —, not determined.

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Table 3

Antioxidant therapy against respiratory viruses.

Virus	Antioxidant	Effects	Ref
Rhinovirus	Glutathione	Scavenge free radicals and suppress NF- κ B induced Intercellular adhesion molecule-1 (ICAM-1) receptor for rhinovirus	[84]
	Pyrrrolidinedithiocarbonate (PDTC)	Inhibited viral replication <i>in vitro</i>	[93]
Influenza Virus	Thiol antioxidants Pyrrrolidinedithiocarbonate (PDTC)	Scavenge OH ⁻ free radicals and Inhibited synthesis negative strand RNA and viral replication <i>in vitro</i> and <i>in vivo</i>	[72,94]
	<i>N</i> -Acetyl-L-cysteine (NAC)	Inhibited induction of apoptosis and pro-inflammatory cytokines such as IL-6, IL-8 and RANTES	[95,96]
	Glutathione	Inhibited induction of apoptosis and viral replication <i>in vitro</i> and <i>in vivo</i>	[97,98]
	Hydroxyl antioxidants/Polyphenols Nordihydroguaiaretic acid (NDGA)	scavenge O ₂ ⁻ , OH ⁻ radicals and H ₂ O ₂ and Inhibited viral replication through inhibiting intracellular transport of viral glycoproteins	[99–104]
	Thujaplicin	Inhibited induction of apoptosis and viral replication <i>in vitro</i>	[100]
	Resveratrol	Blocked nuclear cytoplasmic translocation viral ribonucleoproteins and reduced expression of late viral proteins and resulted in reduced viral replication <i>in vitro</i> and <i>in vivo</i>	[101]
	Ambroxol	Suppressed proliferation of virus <i>in vivo</i>	[103]
	Ascorbic acid	inhibited the proliferation of virus <i>in vitro</i>	[104]
	Flavonoids Catechins	Scavenge O ₂ ⁻ and Inhibited viral replication by inhibiting activities of hemagglutinin, neuraminidase and suppressing viral RNA synthesis in <i>in vitro</i> and <i>in vivo</i>	[105–107]
	Quercetin 3-rhamnoside	Inhibited viral replication by inhibiting viral mRNA synthesis	[106]
	Isoquercetin	Inhibited viral replication and pro-inflammatory cytokines	[107]
	Antioxidant enzymes SOD, Catalase	scavenge O ₂ ⁻ /OH ⁻ free radicals; restores redox status <i>in vitro</i> and <i>in vivo</i> ; enhances recovery	[70,108–111]
	Nrf2 inducers Sulforaphane	antiviral activity	[39]
RSV	Thiols NAC	scavenge H ₂ O ₂ , OH ⁻ free radicals, and hypochlorous acid; suppress NF- κ B activation and viral replication	[51,112]
	Polyphenols Resveratrol	reduced IFN- γ levels associated with RSV-mediated airway inflammation and AHR; inhibit TRIF signaling pathway	[113,114]
	SOD	significantly reduced pulmonary viral titers	[73]
	SOD Mimetics	scavenge ROS and inhibit chemokine secretion <i>in vitro</i>	[8,78]
	Nrf2 inducers Sulforaphane BHA and tBHQ	antiviral activity in mouse; scavenge ROS by inducing expression of antioxidant enzymes and inhibit chemokine secretion <i>in vivo</i> and <i>in vitro</i> ; mice treated with BHA recovered faster	[61,69] Casola A. (unpublished observation)