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Abstract

Reactive oxygen species (ROS) are a family of oxygen molecules with an unpaired electron and play an important role in homeostasis and pathogenesis. The reactive molecules modify lipids, proteins and nucleic acids, and modulate a wide range of cellular functions. The importance of ROS in infection has been established through clinical and in vitro studies. Here we review the role of oxidative stress in HIV pathogenesis, the impact of ROS on immune responses in HIV patients, and ROS-mediated regulation of HIV infection. Future studies on the interplay between ROS and HIV infection may offer a new strategy for prevention and treatment.

Keywords: Reactive oxygen species; HIV; Immune response

Introduction

Reactive oxygen species (ROS), metabolic byproducts in both prokaryotic and eukaryotic cells, were thought to be detrimental to the host by damaging cellular macromolecules including DNA, proteins and lipid, leading to pathogenesis of diseases [1]. However, it has been well recognized that ROS are crucial for maintaining host physiological functions [2] and play an important role in control of infection [3-9]. The association of ROS with HIV pathogenesis and disease progression has been speculated since late 80's [10,11]. Several clinical trials on modulation of ROS in HIV-infected patients have been conducted (reviewed in [12]) although the results have not been consistent. Considering the complex role of ROS in infection and immune response, understanding the balance between redox biology and oxidative stress will offer insights into prevention and treatment for infection and diseases.

ROS and their Functions

ROS are a family of reactive free radicals. Physiologically relevant ROS include superoxide (O_2^{\bullet}) , hydrogen peroxide (H_2O_2) , hypochlorous acid (HOCl), singlet oxygen $({}^{1}O_2)$ phenoxyl radicals (PhO[•]), lipid peroxides (ROOH), hydroxyl radical (OH[•]), nitric oxide (*NO), nitrogen dioxide(*NO₂), and peroxynitrite (ONOO⁻) [13,14]. ROS originate from specific intracellular locations and are generated by different mechanisms. ROS reactions are dependent on the kinetics, thermodynamics, and location (reviewed in [1,2,9]). The largest source of ROS is from the mitochondria during oxidative phosphorylation. The respiratory chain proteins complexes I and III can "leak" electrons which quickly react with mitochondrial O_2 ; this reaction forms superoxide (O_2^{\bullet}) (reviewed in [15]). Superoxide is quickly converted into H_2O_2 by superoxide dismutase. H_2O_2 is less reactive and can act as a second messenger in mitochondrial signaling. ROS production is illustrated in (Figure 1).

Other major sources of ROS are enzymes located in specific organelles. NADPH oxidases (NOXs) are located at cellular membranes and generate superoxide. NOXs are important for both pathogen killing and cellular signaling. Xanthine oxidases are found in the peroxisomes and generate hydrogen peroxide for catabolic oxidation. Oxidants are also generated in the endoplasmic reticulum to facilitate protein folding and disulfide bond formation (reviewed in [3]).

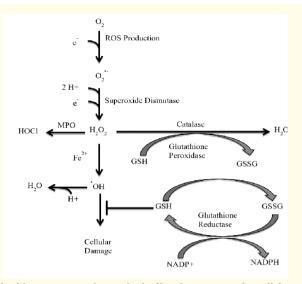


Figure 1: Catabolism of ROS. ROS are highly reactive and must be buffered to protect the cell from damage. ROS enzymatic sources, such as mitochondrial proteins and NOX, pass electrons to oxygen (O_2) ; this results in superoxide $(O_2 \bullet -)$ formation. Superoxide can be converted to hydrogen peroxide (H_2O_2) by superoxide dismutase. Hydrogen peroxide is a less reactive than superoxide and stable enough to exist as a signaling molecule. The fate of hydrogen peroxide will be determined by location and the surrounding microenvironment. For example, in the phagosome myeloperoxidase (MPO) can convert H_2O_2 into hypochlorous acid (HOCL), yet in the cytosol H_2O_2 can be converted to water by catalase. H_2O_2 exposure to ferrous iron (Fe²⁺) results in Fenton's reaction generating the very reactive hydroxyl radical •OH). Hydroxyl radical is buffered by single molecule antioxidants, such as glutathione. Antioxidants are enzymatically recycled.

ROS are highly reactive and must be regulated to prevent damage to cellular macromolecules including DNA, lipids and proteins. ROS are regulated by antioxidants and redox proteins. Single molecule antioxidants, such as glutathione or thioredoxin, are oxidized by ROS. Oxidized antioxidants are then recycled enzymatically to replenish the reduced form of the molecule; this makes the antioxidant available to reduce ROS again. An imbalanced reduction/oxidation ratio results in oxidative stress. Oxidative stress is also reduced by redox proteins such as catalase and glutathione peroxidase. These proteins catalytically convert ROS to harmless byproducts such as water (reviewed in [13-14]).

ROS are antimicrobial. Phagocytic immune cells, such as macrophages and neutrophils, contain NOX in the membrane of the phagosome. NOX pumps ROS into the phagosome where engulfed pathogens can be damaged. ROS target thiols for depletion and metal centers for OH production, a highly reactive oxygen intermediate; this increases damage of nucleotide bases, proteins, and lipids (reviewed in [16]).

ROS play an important role in cellular signaling. The pleiotropic role of ROS is illustrated by the function of the NOX proteins. NOX signaling has been linked to apoptosis, proliferation, homeostasis and gene regulation.

ROS act as a second messenger in the MAP kinase cascade to control homeostasis and gene regulation. ROS can also directly modulate redox sensitive transcription factors such as NF κ B, AP-1 and p53 (reviewed in [17]). Proteins that are also redox sensitive can be oxidized at cysteine, methionine, tyrosine, phenylalanine and tryptophan resides. These small redox reactions can have major implications for the cell. For example, latency associated peptide 1 β (LAP-1 β) is redox sensitive chaperone protein for transforming growth factor β 1 (TGF- β 1). When LAP-1 β is oxidized at its amino acid 253 methionine residue, a conformational change is triggered that releases TGF- β 1, leading to TGF- β 1 activation [18]. Activated TGF- β 1 is then free to bind tyrosine I and II receptors and initiate a signal cascade.

Recent studies on the involvement of ROS in inflammasome activation highlight the role of ROS in immune responses and immuneassociated diseases [19-21]. ROS triggered by mitophagy/autophagy blockade leads to ROS-generating mitochondria, resulting in NLRP3 inflammasome activation [22]. The crosstalk between ROS and toll-like receptor activation or cytokines such as TNF α or IL-6 modulates innate immune response and inflammation [6,7,23-25] that can alter the course of infection and disease progression.

ROS in HIV Infection

Oxidative stress was first observed in HIV pathology within a few years of the discovery of HIV. Glutathione (GSH) is depleted in plasma, lymphocytes, monocytes and lung epithelial lining in HIV-infected patients [26-29]. GSH depletion is inversely correlated to increased GSSG (glutathione disulfide) in plasma of HIV positive patients, suggesting that free radical defense was compromised [30,31]. Thioredoxin (TRX) is depleted in the dendritic cells of the lymph nodes but elevated in the plasma of HIV-infected patients [32,33]. Treitinger, *et al.* [34] have found reduced SOD levels in plasma and monocytes of HIV-infected patients although glutathione peroxidase levels are comparable compared to the control subjects. Spontaneous H_2O_2 production in monocytes of HIV-infected patients is associated with viral load. However, intracellular TRX levels are suppressed in asymptomatic patients while increased in patients with AIDS compared to healthy controls [35]. Clinical data are summarized in (Table 1).

Citation	ART	Sex	AIDS (CDC Classification)	CD4 Counts (Mean)	Oxidative Stress
van der Ven., <i>et al</i> . 1998	Y	30M; 5 W	11 AIDS; 17 asymptomatic; 7 symptomatic	232±224/mmE3 no-AIDS; 83±55/mmE3 AIDS	GSH Elevation
Allard., <i>et al.</i> 1998	Y/N	47M; 2W	24 AIDS; 25 asymptomatic	214±41 E6/L placebo; 269±46 E6/L supplement	Vitamin E & C decrease ROS levels
Masutani., <i>et al</i> . 1992	NA	6 M; 0W	5 AIDS; 1 ARC	NA	TRX depletion in lymph nodes
Staal., <i>et al</i> . 1992	NA	121M; 11W; 2T	71 AIDS; 31 asymptomatic; 32 ARC	74±120/μl AIDS; 283±243/ μl ARC; 516±318/μl asymp- tomatic	GSH and cysteine depletion
Malorni <i>., et al</i> . 1998	N	10M; 4W	1 Acute HIV; 7 AIDS; 6 asymptomatic	789/mmE3 acute HIV; 94±56/ mmE3 AIDS; 423±92/mmE3 asymptomatic	Inverse correlation of ROS levels and CD4+ cell count
Nakamura. <i>, et al</i> . 1996	Y	М	12 AIDS; 32 ARC; 40 asymptomatic	<500 cells/μl	TRX elevated
Lopez Galera., et al. 1996	Y/N	30 M; 4W	16 AIDS; 4 ARC; 14 asymptomatic	NA	GSH depletion/ GSSG eleva- tion
Aukrust., <i>et al</i> . 1995	NA	18 M; 4W	12 symptomatic; 10 asymptomatic	120 E6/L	GSH depletion/ GSSG eleva- tion
Elbim. <i>, et al</i> . 1999	Y/N	26M; 9W	9 AIDS; 10 asymptomatic; 7 asymptomatic	<500/µl symptomatic; >500/ µl asymptomatic; <500/µl asymptomatic	Enhanced ROS in monocytes
Mburu., <i>et al</i> . 2013	N	5M; 15W	20 asymptomatic	411/mmE3	Vitamin C & NAC decreased early apoptosis and CD25

Table 1: Summary of Clinical Data Assessing the Impact of ROS on HIV Patients.

•Abbreviations: Y: Yes On ART, N: No ART, Y/N: Cohorts Yes On ART and No ART, NA: Information Not Available, M: Men, W: Women, T: Transexual, ARC: AIDS-related Complex

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The changes in ROS in HIV-infected patients provided the rationale for clinical studies to assess antioxidant regimens as treatment for patients with HIV. There was a decrease in viral load after three months of Vitamin E & C supplementation [36]; however these effects were not widely reproduced. While vitamin supplementation did have beneficial effects of reduced mitochondrial dysfunction, oxidative stress and metabolic complications in HIV-infected patients receiving HAART, vitamins failed to reduce viral loads or consistently preserve CD4+ T cell counts in patients with anti-retroviral therapy [37]. A clinical trial that aims to assess the impact of micronutrient and antioxidant supplementation on disease progression, decline of immunity, and the start of ART treatment in untreated asymptomatic HIV positive adults is currently in a Phase III trial [38].

The immunological profiles in HIV-infected patients are impacted by oxidative stress and possibly modulated by antioxidant treatment. High oxidative stress is found in serum of HIV-infected patients and is inversely associated with IL-7-mediated STAT5 activation in CD8+ T cell subsets and CD4+ central memory [39]. Oxidative stress-mediated apoptosis is enhanced in lymphocytes of asymptomatic HIV patients [40]; however, LPS-induced apoptosis is reduced in CD4+ T cells of asymptomatic HIV patients with combined Vitamin C and NAC treatment [41]. Vitamin C and NAC treatment also reduced LPS-induced up-regulation of activation marker, CD25 on CD4+ T cells in these patients. Lipid peroxidation, an indicator of oxidative stress, is inversely correlated with CD4+ T cell counts in HIV positive patients; however, pretreatment with NAC prevented altered morphology of PBMC *in vitro* [42]. In chronically infected monocytic cell line, U1, glutathione and NAC inhibited reverse transcription when stimulated with PMA, TNF, or IL-6 [43]. In Ul cells, NAC also showed significant suppression of HIV mRNA accumulation with PMA or TNF stimulation. Similar findings showed PMA reactivation of HIV-LTR was blocked by NAC in stably transfected cell lines 293.27.2 and H9 T cells [44]. NAC and GSH were also found to suppress reverse transcription in primary cord blood monocyte-derived-macrophage (MDM) and adult MDM treated post-infection [45,46].

HIV proteins have been shown to modulate ROS in vitro as summarized in (Table 2). HIV accessory protein TAT (trans-activator of transcription) increases H_2O_2 production. TAT-induced H_2O_2 signal along with a Ca²⁺ influx induces CD95L thereby, mediating activation induced cell death [47]. TAT also up-regulated NOX 2, a ROS-generating enzyme, in a phosphatidylinositol-3-kinase/Akt dependent manner, leading to NFkB activation [48]. TAT increases TNF-mediated cytotoxicity and NFkB transcription in a dose dependent manner, by reducing mitochondrial SOD expression and function [49]. TAT also reduces the GSH:GSSG glutathione ratio, leading to decreased redox capacity of the cell [49]. Interestingly, TAT induces the master regulator of the antioxidant response elements (ARE), Nrf2 genes, which encode many antioxidant enzymes [50]. Viral accessory protein, Vpr, increases ROS thereby modulating IL-6 and mediating reactivation of latent virus in primary MDM [51]. HIV env glycoprotein gp41 can also increase ROS in bystander cells and is critical for HIV env-mediated apoptosis, which depends on ROS production [52].

HIV	ROS	Cellular Target	Cell type	Reference	
TAT	↑	Increase NFkB Transcription	MAGI-CCR5	Zhang, 2011	
TAT	↑	Increase TNF	Jurkat T cell & HeLa-tat	Westendorp, 1995	
TAT	↑	Decrease Mitochondrial SOD	Jurkat T cell & HeLa-tat	Westendorp, 1995	
TAT	↑	Increase Activation-Induced Cell Death (ACID)	Jurkat T cell & Primary T cells	Gulow, 2005	
gp41	Î	HIV Env-mediated apoptosis	CHO & Primary T cells	Garg, 2005	
Vpr	Î	IL-6 mediated viral reactivation	Primary MDM & Ul monocytes	Hoshino, 2010	

Table 2: Summary of Studies Showing HIV Proteins Enhance ROS Production. 1 indicates induction

ROS modulates HIV infectivity *in vitro*. ROS has been shown to enhance HIV entry into Jurkat T cells at H_2O_2 concentrations between 1-10 μ M, which is correlated with increased surface chemokine receptor CXCR4 expression [53]. However, HIV entry is decreased at H_2O_2

concentrations above 50 μM and CXCR4 expression is also decreased at higher H2O2 concentrations. Surface receptor modulation by ROS has also been shown in macrophage [54]. ROS increase viral replication by reactivating 5'LTR (long terminal repeat) in latently infected cells through canonical NFκB signaling [29, 55]. These studies are summarized in (Table 3).

ROS	Mechanism	HIV Infection	Cell Type	Reference
1	Enhance HIV Entry	1	Jurkat T cell	Lan, 2013
1	Increase LTR Induction	1	Jurkat T cell	Staal, 1990
1	Increase CXCR4 Expression	1	Primary Macrophage	Saccani, 2000
Ļ	Decrease RT & viral protein synthesis	\downarrow	U1 Monocytes	Kalebic, 1991

Table 3: Summary of Studies Illustrating ROS's Modulation of HIV Infectivity. 1 induction; 4 inhibition

The role of ROS in the inflammasome activation of HIV is an exciting area and has not yet been thoroughly investigated. ROS production is required for NLRP3 inflammasome activation in both primary macrophage and THP-1 cells, a human monocytic cell line [56]. Basal levels of NLRP3 inflammasome genes in monocyte-derived dendritic cells from HIV+ patients was increased three fold compared to healthy controls [57]. Additionally, single nucleotide polymorphisms in the NLRP3 gene are associated with increased HIV susceptibility[58]. Inflammasome caspase-1 mediated pyroptosis has been found to cause 95% of CD4+ cell death in human lymphoid aggregate culture infected with HIV ex vivo [59]. Therefore, further studies on the interplay between ROS and inflammasome will offer insights into the dynamic of immune activation in response to HIV infection [60,61].

Conclusions

ROS are clearly involved in HIV pathogenesis and disease progression. It may also play a role in HIV transmission, which has not been thoroughly investigated. Because of their diverse species, locations and pleotropic functions in immune responses, it has been challenging to alter the symptoms or course of diseases in HIV-infected patients. Further understanding the interplay between ROS and immune response in context of HIV infection at the molecular level may help develop new strategies for HIV prevention and treatment.

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