Molecular Pathways: Reactive Oxygen Species Homeostasis in Cancer Cells and Implications for Cancer Therapy

Veronique Nogueira and Nissim Hay

Abstract

Reactive oxygen species (ROS) are important in regulating normal cellular processes, but deregulated ROS contribute to the development of various human diseases, including cancers. Cancer cells have increased ROS levels compared with normal cells, because of their accelerated metabolism. The high ROS levels in cancer cells, which distinguish them from normal cells, could be protumorigenic, but are also their Achilles’ heel. The high ROS content in cancer cells renders them more susceptible to oxidative stress–induced cell death, and can be exploited for selective cancer therapy. In this review, we describe several potential therapeutic strategies that take advantage of ROS imbalance in cancer cells by further increasing oxidative stress, either alone or in combination with drugs that modulate certain signaling pathways. Clin Cancer Res; 19(16); 4309–14. ©2013 AACR.

Background

Redox homeostasis

Oxidative stress is defined as an imbalance between production of oxidants or reactive oxygen species (ROS), and their elimination by protective mechanisms or antioxidants. Disturbance in this redox balance can lead to damage of important components of the cell, including proteins, lipids, and DNA, with potential impact on the whole organism, and with an increase risk in mutagenesis (1, 2). These effects of ROS are thought to contribute to the aging process (3), but could also play a role in the genesis of cancer.

ROS are byproducts of a normal cellular metabolism and play vital roles in the stimulation of signaling pathways, such as intracellular signal transduction, metabolism, proliferation, and apoptosis (4, 5). Under a sustained environmental stress, ROS are produced over a long time, and thus significant damage may occur to cell structure and functions and may induce somatic mutations and neoplastic transformation (6, 7). ROS could facilitate cancer development by direct oxidative damage to DNA, induction of lipid peroxidation, or oxidative protein damage leading to structural alterations in the DNA (1, 2). There are several free radicals ROS and nonfree radicals ROS inside cells. The major site of ROS generation inside the cells is the mitochondrial electron transport chain where electrons can escape their route and react with oxygen (8). This biologic reduction of molecular oxygen is the source of ROS such as superoxide anion (O$_2^-$), which is the major free radical and hydrogen peroxide (H$_2$O$_2$), which is the major nonfree radical ROS (8, 9). ROS can also be generated by the activation of growth factor receptors, which in turn activate NADPH oxidase. NADPH oxidase oxidizes NADPH to generate superoxide (Fig. 1).

Cellular redox balance is maintained by a powerful antioxidant system that scavenges ROS. The majority of superoxide is transferred to the mitochondrial matrix, where it is dismutated to H$_2$O$_2$ by the superoxide dismutase (MnSOD or SOD2). Some of the superoxide is transferred to the cytosol and is dismutated to H$_2$O$_2$ by the cytosolic SOD (SOD1). The antioxidant system also consists of catalase, the glutathione system [reduced glutathione (GSH), glutathione reductase (GR), glutathione peroxidase (GPX), and glutathione transferase], the thioredoxin system [thioredoxins (Trx), thioredoxin peroxidase, and peroxiredoxins (PRX)], and vitamin E and C (ref. 9; Fig. 1). GSH is the major nonenzymatic component of intracellular antioxidant defenses and plays a central role in maintaining redox balance (NADPH/NADP). The ratio of NADPH to NADP ratio is via its tight regulation of NADPH intracellular levels. The intracellular NADPH level is determined by the difference between its production and its consumption. The major sources of intracellular NADPH are the pentose phosphate pathway (PPP) and mitochondrial metabolism. The major consumers of intracellular NADPH are H$_2$O$_2$ detoxification by GPX and PRX and fatty-acid synthesis (10). Thus, the production and the consumption of NADPH inside cells should be coordinated to maintain NADPH homeostasis. This is particularly important during the formation of solid tumor and during metastasis when cancer cells undergo energetic stress.

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In the past two decades, cancer promotion and progression have been linked to oxidative stress by increasing DNA mutations or inducing DNA damage, genome instability, and cell proliferation (11–13). Indeed, one of the main features of cancer cells, when compared with the normal ones, is a persistent prooxidative state that can lead to intrinsic oxidative stress (14, 15). The enhanced oxidative stress observed in cancer cells can result not only from ROS overproduction, but also from low levels or inactivation of antioxidant mechanisms. Survival of tumor cells is highly dependent on their capacity to control expression of endogenous antioxidants to maintain a steady state level of ROS below the threshold that will induce tumor cell death. Therefore, cancer cells have evolved mechanisms to protect themselves from intrinsic oxidative stress and have developed a sophisticated adaptation system that essentially involves the rearrangement of the antioxidant functions and the upregulation of prosurvival molecules (16, 17). In addition to the increase in mutations rate, ROS could contribute to the initiation of cancer through accelerating protumorigenic signaling pathways. By oxidizing disulfide bonds of cysteine residues, ROS could change the activity of certain proteins, and particularly the tyrosine phosphatases superfamily (18). Perhaps, the best example related to cancer is the documented inactivation of the tumor suppressor PTEN by oxidation (19, 20). Another example is the inhibition of the mitogen-activated protein kinase (MAPK) phosphatase by ROS, which in turn induces activation of extracellular signal–regulated kinases (ERK). The inhibition
of PTEN by ROS hyperactivates the PI3K/Akt signaling pathway, which is perhaps the most frequently activated signaling pathway in cancer cells (reviewed in refs. 21–23). The activation of this pathway would further increase intracellular ROS, at least in part, through the inhibition of the transcription factors FOXOs, which elevate the expression of antioxidants, such as SOD2, catalase, and sestrin3 (24).

**ROS as a double-edged sword in cancer cells**

The high intracellular ROS levels in cancer cells are largely the byproducts of the highly metabolic nature of these cells. These ROS levels could be protumorigenic, but also increase the susceptibility of cancer cells to cell death. Activated signaling pathways that increase intracellular ROS level in cancer cells render these cells more vulnerable than normal cells to oxidative-stress-induced cell death. This vulnerability was documented in cells expressing oncogenic Ras (25) and in cells that display hyperactive PI3K/Akt signaling (24). The high levels of ROS in cancer cells that display hyperactive Akt is generated by increased mitochondrial metabolism and by the suppression of antioxidants gene expression, through the inhibition of FOXO transcription factor (24). In addition to the elevation of SOD2 and catalase, FOXO was shown to induce the expression of Sestrin3 (24, 26). Sestrin3 is a member of a family of proteins that includes Sestrin1 and Sestrin2, which were originally identified as antioxidants induced by the tumor suppressor p53 (27–30). The FOXO–Sestrin axis is conserved in flies, and has an additional function in both mammalian and flies cells as an activator of AMPK and the inhibitor of the mammalian target of rapamycin complex 1 (mTORC1; refs. 26, 31). Thus, the suppression of Sestrins expression in cancer cells could increase intracellular ROS and activate mTORC1. Because ROS inhibits PTEN activity, this would further activate the PI3K/Akt signaling in Pten-proficient cells and further activate mTORC1 (Fig. 2A). To maintain the intracellular level of ROS below a toxic threshold level, cancer cells adopted alternative mechanisms of antioxidation. For instance, relatively high levels of FOXM1 expression in cancer cells could compensate for the loss of FOXO activity. FOXM1 is expressed at low levels in normal cells, but its expression is markedly elevated in cancer cells (32). FOXM1 exerts multiple protumorigenic activities, but was also shown to reduce ROS levels through the transcriptional induction of SOD2, catalase, and mitochondrial Trx-dependent peroxide reductase (PRDX3; ref. 33). In addition, the expression of detoxifying enzymes such as glutathione S-transferases (GST), NAD(P)H:quinone oxidoreductase 1 (NQO1) was reported to be elevated in cancer cells (34, 35).

Energetic stress conditions could lead to oxidative stress. Cancer cells that consume high levels of glucose undergo energetic stress during the formation of solid tumors when cells detach from the matrix and translocate to the lumen (36), or during metastasis when they leave the primary tumor site to relocate to another site. The decrease in glucose uptake during these processes suppresses ATP production and activates AMPK (36, 37), but also inhibits the generation of NADPH via the PPP. The reduced level of NADPH results in increase of intracellular ROS, which could eventually cause cell death (36, 37). However, the concomitant activation of AMPK elicits alternative mechanisms that maintain intracellular NADPH levels. The activated AMPK phosphorylates and inhibits acetyl-CoA carboxylase 1 (ACC1) and ACC2. Both enzymes may have some overlapping activities, but ACC1 is largely responsible for the generation of malonyl-CoA for fatty acid synthesis (FAS), whereas ACC2 inhibits fatty acid oxidation (FAO; ref. 38). By inhibiting ACC1, AMPK inhibits FAS and therefore decreases NADPH consumption in FAS. By inhibiting ACC2, AMPK increases FAO and therefore NADPH production via the mitochondria (ref. 37; Fig. 2B). Thus, AMPK activation is critical for the adaptation of cancer cells to energetic stress, which subsequently induces oxidative stress.

**Clinical–Translational Advances**

The high levels of ROS in cancer cells in comparison with normal cells could be exploited for cancer therapy. In fact, many of the commonly used chemotherapeutic regimens induce high level of ROS that kill cancer cells (39, 40). However, most of these therapeutic regimens were not strictly developed to exploit the high level of ROS in cancer cells for cancer therapy.

The persistent prooxidative state characterizing cancer cells, as well as their multiple adaptation mechanisms, can be exploited to develop new therapeutic strategies that will specifically target cancer cells, to ensure a good therapeutic selectivity. Because the overall redox homeostasis is maintained by the balance between ROS generation and scavenging, exogenous compounds that specifically increase ROS production, or inhibit ROS elimination, can favor the accumulation of ROS in cancer cells, and hence induce cell damage or even cell death once their threshold of “tolerance” is reached (41).

**ROS and chemotherapy**

Developing cancer therapies based on escalating further the high ROS level in cancer cells to a toxic level by triggering ROS accumulation directly and/or inhibition of ROS scavenging systems represent powerful avenues for selectively killing cancer cells. Several drugs have been identified as promoting ROS generation. These reagents include: (i) mitochondrial electron transport chain modulators (e.g., arsenic trioxide, doxorubicin, topotecan, etc.); (ii) redox-cycling compounds (e.g., motexafin gadolinium); (iii) agents that disrupt the antioxidant defenses mechanism, such as GSH-depleting agents (e.g., buthionine sulfoximine, β-phenylethyl isothiocyanate (PEITC)] and inhibitors of SOD (e.g., 2-methoxyestradiol), and catalase (e.g., 3-aminoo-1,2,4-triazole) (25, 41–44). Through promoting oxidative stress or targeting the endogenous antioxidant system of cancer cells, these reagents could be used as a therapeutic weapon against cancer cells.

**Rapamycin and oxidative stress combination as cancer therapy.** The PI3K/Akt signaling pathway is thought to play a prominent role in the initiation and maintenance of human cancer, as many components of this pathway have
been found to be mutated or amplified in a broad range of human cancers (21–23), and thereby promoting resistance to therapeutic agents that induce apoptosis.

Akt is activated by extracellular signals that activate phosphoinositide 3-kinase (PI3K), and is negatively regulated by phospholipid phosphatases that negate the activity of PI3K, such as the tumor suppressor PTEN. Akt activity is also downregulated by the activation of its downstream effector, mTORC1, which in turn induces a negative feedback mechanism that inhibits Akt activity (21, 22, 45). By increasing cellular metabolism, Akt inhibits AMPK activity and increases the level of the byproducts of energy metabolism, ROS. The inhibition of FOXOs by Akt inhibits the expression of antioxidants, which in turn further increase ROS levels. There is also interplay between FOXOs and mTORC1, whereby Sestrin3 induced by FOXOs activates AMPK and inhibits mTORC1. The activation of mTORC1 by Akt and by AMPK inhibition promotes several anabolic functions, but also elicits negative feedback loops that inhibit Akt. The high levels of FOXM1 in cancer cells could compensate for the suppression of FOXOs through induced expression of antioxidants genes. 

Figure 2. Signaling pathways regulating intracellular ROS in cancer cells. A, Akt is activated by extracellular signals that activate PI3K. Akt activation inhibits the FOXO transcription factors and activates mTORC1. By increasing cellular metabolism, Akt inhibits AMPK activity and increases the level of the byproducts of energy metabolism, ROS. The inhibition of FOXOs by Akt inhibits the expression of antioxidants, which in turn further increase ROS levels. There is also interplay between FOXOs and mTORC1, whereby Sestrin3 induced by FOXOs activates AMPK and inhibits mTORC1. The activation of mTORC1 by Akt and by AMPK inhibition promotes several anabolic functions, but also elicits negative feedback loops that inhibit Akt. The high levels of FOXM1 in cancer cells could compensate for the suppression of FOXOs through induced expression of antioxidants genes. B, energy stress during certain stages of tumor development increases oxidative stress by decreasing NADPH production through the PPP. AMPK activation under energy stress conditions is required to maintain NADPH homeostasis and prevent oxidative stress–induced cell death. AMPK activation inhibits the consumption of NADPH by FAS through the phosphorylation and inhibition of ACC1, and increases NADPH production via FAO, through phosphorylation and inhibition of ACC2.

Rapamycin analogs are currently being used in clinical trials and have been already approved for certain types of cancer (46, 47). Rapamycin alone attenuates cell proliferation and rarely elicits cell death. Furthermore, it could also increase cell survival and chemoresistance via the inhibition of mTORC1, and consequently activating Akt through the inhibition of a negative feedback loop (21, 22, 45). However, by activating Akt, rapamycin further sensitizes cells to ROS-induced cell death, and thus, the combination of rapamycin and oxidative stress could be an attractive strategy to selectively eradicate cancer cells (24).

Isothiocyanates such as the PEITC are thiol modifiers that have been shown to inhibit the GSH antioxidant system by extruding GSH from the cell and by inhibiting GPX (25, 44). This leads to ROS overproduction, oxidative damage of mitochondria, and apoptosis preferentially in cancer cells, presumably due to their increased constitutive ROS levels (24, 25, 44). Clinical studies with PEITC are currently being initiated (48). A combination therapy of rapamycin and PEITC was proven efficient to selectively eradicate tumors with hyperactivated Akt in preclinical studies (24). This strategy evades the chemoresistance induced by the hyperactivation of Akt in cancer cells.

Agents that enhance proteotoxic stress, including the HSP90 inhibitor IPI-504, are also known ROS inducers. It has been shown that IPI-504 and rapamycin synergize in Ras-driven tumors by promoting irreversible endoplasmic
reticulum (ER) stress, resulting in catastrophic ER and mitochondrial damage, and tumor regression (49). The mechanism by which these agents cooperate reveals a therapeutic paradigm that can be expanded to develop additional combinations.

**FOXM1 inhibition and oxidative stress combination as cancer therapy.** The transcription factor FOXM1 is over-expressed in a majority of human tumors and is implicated in tumor angiogenesis, invasion, and metastasis (32). Increased FOXM1 expression in tumors is associated with advanced tumor stage and poor prognosis. FOXM1 could reduce intracellular ROS levels in cancer cells by inducing the expression of detoxifying enzymes, including catalase, SOD2, and PRDX3 (ref. 33; Fig. 2A).

FOXM1 transcriptional activity and expression can be inhibited by proteasome inhibitors such as bortezomib (Velcade) and MG132, or the thiazole antibiotics, Siomycin A and thioistrepton (50–52). It was shown that the suppression of FOXM1 by proteasome inhibitors sensitizes human cancer cells to cell death induced by DNA-damaging agents including doxorubicin and γ-irradiation (53). Therefore, targeting FOXM1, in combination with oxidative stress is a sound strategy to eliminate tumor cells.

**Targeting AMPK for cancer therapy.** AMPK is a master switch of metabolic adaptation (37, 54, 55), and its activation appears to be critical for cell survival during energetic stress (ref. 37; Fig. 2B). The activation of AMPK could inhibit the proliferation of cancer cells and therefore is a considered strategy for cancer therapy (55). However, the recent studies discussed earlier, which show that AMPK activation is required to combat oxidative stress and promote cancer cell survival during energetic stress, suggest that the inhibition of AMPK could be also a therapeutic strategy for cancer therapy. This could be an effective therapy at certain stages of cancer progression such as during solid tumor formation and metastasis when cancer cells undergo energetic stress. Because the absence of AMPK activation during energetic stress of cancer cells elicits ROS-mediated cell death, the combination of exogenous ROS inducers and inhibitors of AMPK could be another attractive strategy for cancer therapy.

Sestrin proteins activate AMPK and also reduce ROS. As shown recently, under energetic stress conditions, Sestrin2 expression is elevated and further potentiates AMPK activity (56, 57). Therefore, targeting Sestrins could not only elevate intracellular ROS but also attenuate AMPK activation during energetic stress, and may constitute another strategy for cancer therapy.

The use of AMPK activation for cancer therapy may stop cancer cell proliferation, but may also increase survival of cancer cells during solid tumor formation and metastasis. To circumvent this possibility and render AMPK activation, a more potent therapeutic approach, it would be useful to use AMPK activation in combination with pharmacologic activators of ACCs to counteract the prosurvival activity of AMPK activation (37).

In summary, we described elevated ROS level as one hallmark of cancer cells that could be exploited for selective cancer therapy. ROS are protumorigenic but also toxic above a certain threshold level. This threshold level could be easily achieved in cancer cells by a relatively subtle increase in ROS level that does not increase ROS level to a toxic level in normal cells. We described several potential therapeutic strategies using ROS inducers alone or in combination with other therapeutic regimens. However, the same as with many therapeutic regimens, ROS inducers could be protumorigenic for normal cells. Therefore, ROS inducers should be kept at restricted doses in order not to affect normal cells.

**Disclosure of Potential Conflicts of Interest**
No potential conflicts of interest were disclosed.

**Authors’ Contributions**
Conception and design: V. Nogueira
Development of methodology: V. Nogueira
Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): V. Nogueira
Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): V. Nogueira
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