

Review

The Pro-Metastatic Roles of ROS

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Abstract

Metastasis is a complex, multistep process in which cancer spreads from its original tumor to other sites in the body. During metastasis, tumor cells move away from the primary tumor and intravasate into the lymphatics or circulation. Surviving tumor cells can then extravasate into and remain in distant tissues until they once again begin to proliferate, forming secondary tumors. An excess of reactive oxygen species (ROS) can promote metastasis, dependent on the ROS molecule, its level of excess, and the examined step within the metastatic cascade. Here, we highlight recent studies where ROS promote epithelial-to-mesenchymal transition, cell migration and invasion, circulating tumor cell survival and disseminated tumor cell dormancy. Additionally discussed are novel in vivo ROS detection methods, FDA-approved therapies and clinical trials that manipulate ROS to improve cancer patient survival. Since metastasis is the major cause of cancer-related death, a better understanding of this process and ROS as a contributing factor will help to identify novel targets for inhibition or prevention.

Keywords: reactive oxygen species (ROS); oxidative stress; hypoxia; hydrogen peroxide; cancer; metastasis; epithelial-to-mesenchymal transition (EMT); circulating tumor cells (CTCs); anoikis; ROS detection



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1. Introduction

Cancer is broadly defined as a group of diseases characterized by the uncontrolled growth and division of abnormal cells with the capacity to invade surrounding tissues and metastasize to distant parts of the body. Before a more advanced understanding of cancer, early successes in cancer therapy involved agents that reliably inhibited or killed actively dividing cells with the major goal of shrinking the cancerous tumor. While cancer detection methods and therapies are consistently improving, the focus of cancer treatment continues to be inhibiting proliferation and killing tumor cells. Cell proliferation is more straightforward and better understood, easier to drug, and produces rapid, measurable effects, whereas metastasis is still a major challenge due to its complex biology, long timeframe and incomplete mechanistic knowledge. However, with over 80–90% of cancer deaths being due to metastatic disease rather than from complications caused directly by

the primary tumor [1,2], elucidating the multifaceted, dynamic mechanisms underlying metastasis is a critical need for the development of novel anti-metastatic strategies.

2. Oxidation–Reduction Regulation

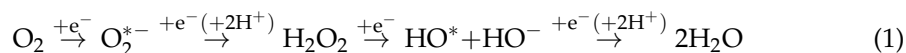
Cellular oxidation–reduction (redox) reactions occur in tandem, in which electrons are transferred between chemical species. There are many different types of redox reactions involving different reactive species, each with unique chemical characteristics and signaling roles and all fundamental to life processes. The type of reactive species depends upon its components and molecular characteristics. There are reactive oxygen (ROS), nitrogen (RNS), sulfur (RSS), and electrophile (RES) species, all of which are oxidants, or chemical species which can accept electrons. Cells in “redox homeostasis” operate within a flexible homeodynamic space, a bounded but adjustable range of redox potentials that shifts with differentiation, stress, and disease yet is actively kept within viability-supporting limits by antioxidant systems and feedback controls [3–6]. Within this homeodynamic space, transient, localized oxidizing shifts encode information via reversible cysteine modifications on proteins [6]. In addition to metabolic regulation, other major cellular processes, including epigenetics, transcription, protein homeostasis, proliferation and differentiation, and cell death, are all under redox regulation. In biology, “oxidative stress” typically refers to an excess of oxidants over antioxidant capacity. Excessive or prolonged oxidation pushes the system outside of its bounded range, leading to oxidative stress, which can damage macromolecules and alter cell signaling, disease progression or cell death [7].

2.1. Intracellular ROS Production

ROS comprise a range of molecules with oxidizing properties that are formed primarily through successive one-electron reduction steps of molecular oxygen during normal physiological and pathological processes [8]. Thus, ROS participate in important cellular signaling and homeostatic functions but also cause oxidative damage if uncontrolled. Free-radical ROS include superoxide anions ($O_2^{\bullet-}$), hydroxyl radicals ($OH^{\bullet}$), nitric oxide ($^{\bullet}NO$), and lipid radicals. Other ROS molecules with oxidizing properties that are not free radicals include hydrogen peroxide (H_2O_2), singlet oxygen (1O_2), peroxynitrite ($ONOO^{\bullet}$), and hypochlorous acid ($HOCl$).

The three primary ROS species are $O_2^{\bullet-}$, H_2O_2 , and $OH^{\bullet}$, all produced from stepwise reduction of molecular O_2 (Reaction (1)) [8]. Mitochondria are the principal organelles that generate intracellular ROS. During oxidative phosphorylation and ATP production, molecular oxygen (O_2) is reduced to water by the electron transport chain. Complexes I and III within the electron transport chain form and release superoxide anions into the mitochondrial matrix and inner membrane as well as the cytosol [9]. Complex II can also directly or indirectly be a significant source of ROS. Superoxide anions and hydrogen peroxide are produced during the forward reaction, in which succinate is oxidized to fumarate, but superoxide anions can also be generated indirectly by reverse electron transfer (RET) from Complex II to I [10]. While not a direct component of the electron transport chain but located in the mitochondrial matrix, the multisubunit enzyme, α -ketoglutarate dehydrogenase complex (KGDHC), predominately produces superoxide, which is then converted to hydrogen peroxide dependent on the redox state and chemistry of FAD or NADH and O_2 accessibility [11]. Other organelles, including the peroxisomes and the endoplasmic reticulum (ER), also minorly contribute to superoxide generation via the P450 system. In addition to organelles, the family of transmembrane NADPH oxidases

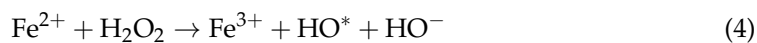
(NOXs) transfer electrons through biological membranes, resulting in oxygen reduction and generating superoxide (NOX1-3 and NOX5) [12].



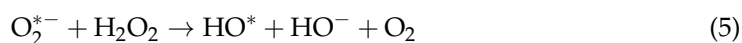
Small amounts of hydrogen peroxide can act as a second messenger, regulating redox signaling to maintain homeostatic levels. Unlike superoxide, hydrogen peroxide can passively diffuse through the plasma membrane or make its way into the cell via channel proteins called aquaporins. Hydrogen peroxide can also be generated intracellularly by the dismutation of superoxide, which may occur spontaneously or via superoxide dismutase (SOD) (Reaction (2)) [13]. SODs (SOD1-3) are enzymatic antioxidants that can diffuse between cellular compartments and appear in the cytosol, mitochondria, nucleus or extracellular space to catalyze superoxide radicals into hydrogen peroxide and molecular oxygen. Several other oxidases, including peroxisomal oxidases and monoamine oxidases on the outer mitochondrial membrane, can also convert superoxide into hydrogen peroxide. The organelles responsible for hydrogen peroxide generation are peroxisomes and ER. Peroxisomes generate large amounts of hydrogen peroxide via oxidases involved in fatty-acid β -oxidation and other catabolic pathways, while the ER produces hydrogen peroxide during oxidative protein folding and through the ER-localized NOX4 isoform. Although hydrogen peroxide is comparatively less reactive than superoxide, its accumulation is damaging. Enzymes such as catalases (CAT) and glutathione peroxidases (GPx) can break down excess hydrogen peroxide to water and oxygen [14].



The hydroxyl radical is by far the most reactive of the three primary ROS species. It can react indiscriminately with nearly any nearby molecule and has a strong affinity for aromatic or sulfur-containing molecules rich in electrons such as proteins and DNA [15]. The hydroxyl radical is formed within the cell as an unintended side reaction while trying to eliminate superoxide and hydrogen peroxide following Fenton/Haber–Weiss chemistry when in the presence of redox-active metals, particularly iron. This occurs following two steps, where the Haber–Weiss reaction (Reaction (3)) is followed by a second step, termed the Fenton reaction (Reaction (4)), yielding a net reaction (Reaction (5)).



Net reaction:



2.2. Pathological ROS Levels/Activity

Balanced redox reactions and controlled ROS production are central to energy generation and metabolism, biosynthesis and anabolic pathways, antioxidant defense and detoxification, redox signaling and regulation, and immunity [8,16–23]. Moderate levels of ROS within a homeodynamic range act as important signals for cell growth and differentiation; however, deviations outside the range are detrimental to the cell. Since the direction of redox reactions in aerobic cell metabolism is towards oxidation, high levels of reductants or antioxidants (reductive stress) are considered subphysiological, and high levels of oxidants or low antioxidant levels (oxidative stress) are considered supraphysiological [4]. Both reductive and oxidative stress are forms of redox stress, and, although

via differing mechanisms, they can damage all major biomolecules, trigger multiple forms of cell death, and contribute to many chronic diseases. Table 1 lists the main enzymatic sources, steady-state levels, approximate lifetimes, and dominant signaling roles of ROS and damaging effects from oxidative stress on different biomolecules from the three primary ROS species [8,24–26]. Characteristics for the remaining five ROS species can be found in the Supplementary Table S1 [8,26–30].

Table 1. Characteristics of the 3 primary ROS species.

ROS Species	Main Enzymatic/ Chemical Sources	Steady-State Levels and Lifetime (Physiological Conditions)	Dominant Signaling Roles	Dominant Damaging Roles
Superoxide (O ₂ ^{•−})	NADPH oxidases ETC Xanthine oxidase Uncoupled NOS ER Peroxisomal oxidases	Very low, <1–10 nM ~10 ^{−6} –10 ^{−5} s and compartment-restricted	Cytosol: very local redox modulation of enzymes Mitochondria: initiator of mtROS cascades; precursor to mitochondrial H ₂ O ₂ that tunes hypoxia signaling, metabolism, and apoptosis Extracellular: NOX-derived superoxide in immune and vascular cells regulates receptor signaling and inflammatory responses	Cytosol/mitochondria: precursor of H ₂ O ₂ and ONOO [−] driving oxidative/nitrosative damage to proteins, lipids, and DNA Extracellular: participates in LDL oxidation and matrix injury via conversion to secondary oxidants
Hydrogen peroxide (H ₂ O ₂)	Dismutation of superoxide by SOD1/2/3 Oxidase and dehydrogenase side reactions Peroxisomal oxidases	Low nanomolar to sub-micromolar, often 10–100 nM ~10 ^{−5} –10 ^{−3} s locally	Cytosol: diffusible second messenger oxidizing Cys in phosphatases and kinases Mitochondria: mtH ₂ O ₂ tunes metabolic flux, stress responses, and mitophagy; acts as exported signal to cytosol and nucleus Extracellular: modulates growth factor and cytokine receptor signaling, leukocyte recruitment, and wound responses	Cytosol/nucleus: sustained H ₂ O ₂ causes DNA base oxidation and strand breaks, redox enzyme inactivation, and protein carbonylation Mitochondria: promotes mitochondrial protein and lipid oxidation, loss of membrane potential, and release of apoptogenic factors Extracellular: contributes to oxidation of matrix proteins and lipids, promoting vascular and tissue remodeling
Hydroxyl radical (OH [•])	Fenton and Haber–Weiss reactions from H ₂ O ₂	No measurable steady-state pool, better thought of as local flux rather than measurable uniform value. ~10 ^{−10} –10 ^{−9} s (essentially instantaneous and nondiffusible)	No meaningful physiological signaling role; damage- dominated	DNA: base modifications, abasic sites, single- and double-strand breaks Proteins: side-chain oxidation, fragmentation, crosslinking, loss of enzymatic activity Lipids: lipid peroxidation in membranes, generating reactive aldehydes

ROS act as a central hub for cell signaling to multiple cell death programs, depending on the context, intensity, and subcellular site of ROS generation. Cells unable to counter excessive ROS to rebalance their redox reactions may trigger any one of the following modes

of program cell death, collectively known as oxidative cell death: apoptosis, ferroptosis, pyroptosis, paraptosis, parthanatos, oxeiptosis, necroptosis, or autophagy-dependent cell death [31,32]. Survival and/or continued proliferation of ROS-damaged cells are associated with inflammatory and age-related conditions, cardiovascular, metabolic, neurodegenerative diseases and cancer [32–39]. Focusing on cancer, ROS are involved in all stages of cancer progression; however, their role and impact on metastasis depends on tumor type, ROS species, site of generation and levels, and counteracting antioxidant or prooxidant response (Figure 1).

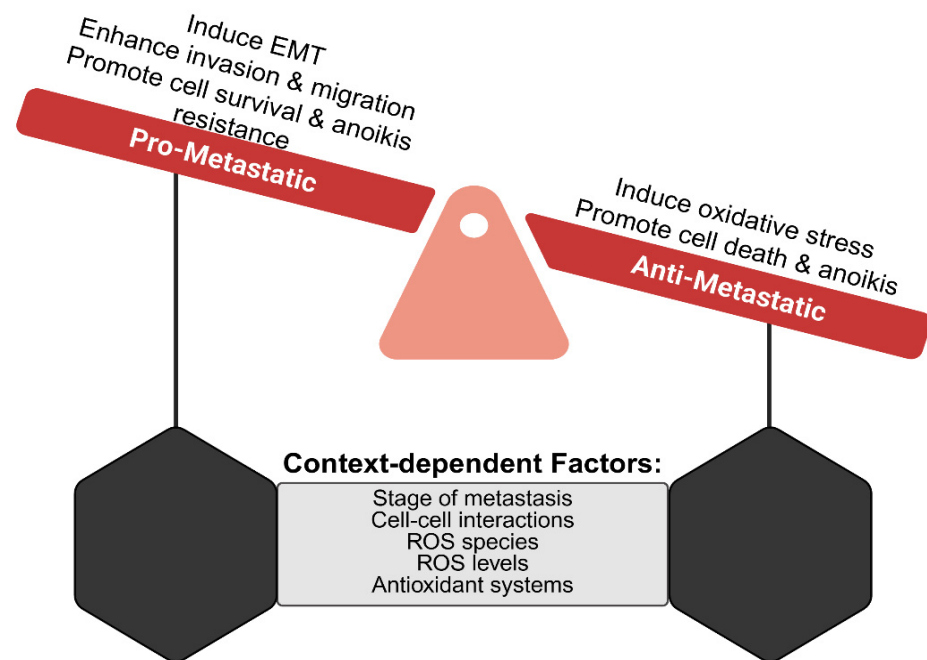


Figure 1. A shift in the balance between antioxidant or prooxidant response has the potential to promote or inhibit metastasis. ROS production and oxidative stress can promote or inhibit metastasis. The effect of ROS is dependent on multiple factors such as stage of metastasis (i.e., cell migration, CTC survival, DTC dormancy, etc.), cell–cell interaction (homotypic clustering, cell association with immune cells, etc.), the ROS species, ROS levels and the antioxidant system. Isolating the specific effects of different species of ROS during metastasis is very difficult due to multiple contributing factors. Created in BioRender. Gilchrist, D.E. (2026) <https://BioRender.com/c2448457> (accessed on 25 March 2026).

3. The Metastatic Cascade

Cancer metastasis is a complex, multistep process in which cells disseminate from the initial tumor and relocate to other tissues within the body. Disseminating tumor cells (DTCs) invade the neighboring tissue, intravasate into the bloodstream and/or lymphatics, reattach to vessel walls and then extravasate into distant tissues. After extravasation, the DTCs may either remain dormant or continue to proliferate into secondary tumors (Figure 2). This metastatic process is mediated by cancer cell intrinsic properties, such as genetic mutations, epigenetic modifications, altered signaling pathways, metabolic reprogramming, and/or mechanisms for immune evasion fundamentally driving tumor cell behavior [40,41]. In this section, we will briefly discuss the process of epithelial-to-mesenchymal transition (EMT), which promotes tumor cell dissemination and disseminating versus circulating tumor cells.

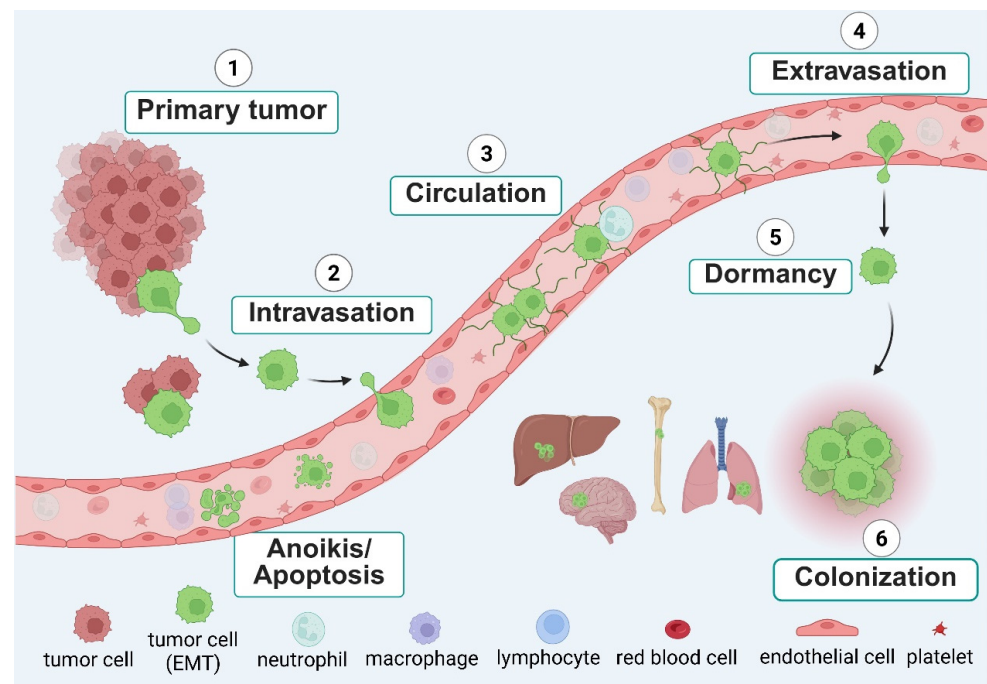


Figure 2. The metastatic cascade. Primary tumor cells that acquire more migratory and invasive characteristics via EMT can move away from neighboring cells and intravasate into the bloodstream. Most tumor cells that enter the bloodstream die due to detachment-induced death (anoikis) or fragmentation from external forces. Surviving circulating tumor cells (CTCs) may extravasate through the endothelial cell wall of blood vessels and remain dormant in distant tissues until they proliferate to form overt secondary tumors. CTC clusters made up of 2 or more tumor cells (green) are predicted to have initially moved away from the primary tumor together rather than associating in the bloodstream. Heterotypic clusters (green and pink) made up of tumor and immune cells may either migrate together from the initial tumor or associate with each other while in the bloodstream depending on the associating immune cell(s). Created in BioRender. Ju, J.A. (2026) <https://BioRender.com/c2448457> (accessed on 25 March 2026).

3.1. Epithelial-to-Mesenchymal Transition

Approximately 85% of all malignant tumors in adults are carcinomas, arising from epithelial cells which have lost their specialized tissue barrier function and instead regain cellular plasticity by undergoing an epithelial-to-mesenchymal transition (EMT) [42]. EMT is an essential process in embryonic development and tissue repair, but the process is hijacked by cancer cells during tumor progression and metastasis [43,44]. While an EMT may not be essential for every epithelial-derived cancer to metastasize, those undergoing EMT gradually lose their epithelial characteristics such as polarity and adhesion and acquire more migratory and invasive capacity similar to that of mesenchymal cells [45]. Some canonical markers for EMT changes are the loss of E-cadherin and/or the gain of vimentin, fibronectin, and N-cadherin [46]. However, simply the absence of epithelial or presence of mesenchymal markers does not represent the spectrum of EMT-associated cell behaviors. Cells with a mixture of epithelial and mesenchymal markers (partial EMT) or those with reduced epithelial and gains in mesenchymal markers (intermediate EMT) continue to maintain plasticity and are able to switch between both cell states, fueling metastasis [47–50]. However, cells which have completely shifted to a mesenchymal phenotype (extreme EMT) with only mesenchymal markers maintain their highly invasive phenotypes but are less metastatic due to their lack of plasticity and inability to reactivate proliferation after reaching distant organs [47].

3.2. Disseminating Tumor Cells

In addition to acquiring migratory and invasive characteristics, DTCs also need to navigate inhospitable, changing microenvironments and host factors, and thus, most do not survive [51,52]. While metastatic efficiency can vary by cancer type, organ, and additional factors, estimates from the experimental and clinical literature suggest that less than 0.01–0.02% of all DTCs eventually outgrow into overt metastases [53]. Most either die, remain dormant, or are eliminated by host defenses, meaning that the vast majority of DTCs never successfully colonize distant organs [54–57].

For tumors to grow past 1–2 mm³, the threshold of nutrient and oxygen diffusion, they need vascular support [58,59]. Tumor cells produce stimulating factors to hijack the normal biological process of angiogenesis, recruiting endothelial cells from existing blood vessels to form new vessels into the tumor, allowing for sustained oxygen and nutrient delivery. However, these new vessels are leaky, primarily due to abnormal and incomplete maturation, allowing for actively invading tumor cells to more easily make their way into the bloodstream [60,61]. In some aggressive cancers, the tumor cells themselves acquire the ability to form channel-like structures that conduct blood or other fluids. This phenomenon is called vascular mimicry and occurs independently of endothelial cell recruitment from pre-existing blood vessels [62]. Since these mimicked channels may either transport blood or directly connect with endothelial-lined vessels, tumor cells lining these structures are highly exposed to circulating blood flow. As a result, they are more likely to detach and enter the circulation [62–64]. Thus, mosaic vessels, where both tumor cells and endothelial cells form parts of the blood vessel wall, can act as direct bridges, allowing tumor cells to transition from tumor tissue into circulation, effectively enhancing tumor cell shedding into the bloodstream.

Even small tumors can shed millions of cells into the bloodstream daily. Based on experimental measurements across different tumor models, a general estimate of the number of cells able to enter the bloodstream from a 1 cm³ (~1 g) small, vascularized solid tumor is approximately 1 million cells per day, although shedding rates can vary by tumor type, location, vascularity, and microenvironment [65,66]. Thus, metastatic spread may occur simultaneously and in parallel to primary tumor growth, not as a later, progressive event [67]. Strong evidence from clinical, genomic, and mathematical modeling studies all support the fact that metastasis can, and typically does, occur before a primary tumor is detectable by conventional imaging techniques [68,69].

3.3. Circulating Tumor Cells

Circulating tumor cells (CTCs) are DTCs that have entered the bloodstream or lymphatics. From experimental modeling, mouse and human data, the lifetime of a CTC in the bloodstream is limited to only minutes to a few hours [70]. Besides CTCs surviving anoikis, a specific type of apoptosis triggered by extracellular matrix detachment, CTCs need to survive immune cell clearance, blood shear stress, and fragmentation due to size constraints from smaller capillary vessels [71]. Approximately only 0.01–0.1% of CTCs survive this hostile microenvironment to form tumors at a secondary site, and this is thus considered the rate-limiting step of metastasis [72,73].

Most CTCs found in patient or mouse models are found as single cells, but CTC clusters have been detected. CTC clusters, sometimes referred to as circulating tumor microemboli, circulating micrometastases or circulating tumor cell aggregates, are defined by having two or more cells grouped together in a “cluster” [74–76]. Clusters are of clinical importance since CTC clusters are 50–100 times more likely than individual cells to survive dissemination and cause metastatic outgrowth [75,77,78]. CTC clusters in the bloodstream may be either homotypic or heterotypic clusters. Homotypic clusters are composed of

only tumor cells, while heterotypic clusters can include tumor cell interactions with surrounding immune or stromal cells in the blood vasculature. Our group has identified microtentacles (McTNs), tubulin-driven membrane protrusions produced from detached cancer cell lines, dissociated tumor cells from surgical samples, and CTCs isolated from patients' blood [79,80] (Figure 3). McTNs promote cell clustering, endothelial reattachment and CTC retention in distant tissues during metastasis [81–90] and may be a mechanism for initial heterotypic cell clustering and/or reattachment to the blood vessel wall before extravasation (Figure 2). Interestingly, we have shown that inhibition of McTNs dramatically reduces metastasis [90], indicating that the molecular mechanisms that promote McTNs could serve as a potential therapeutic target to reduce metastasis.

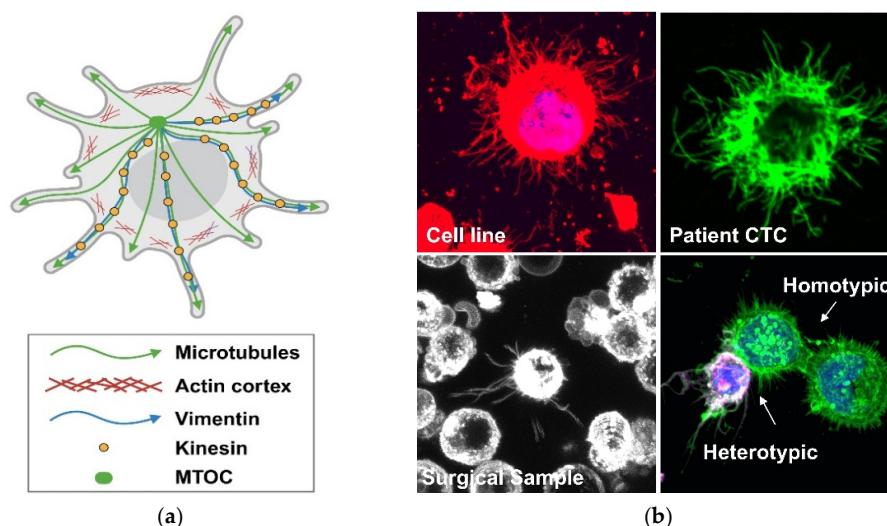


Figure 3. Microtentacles (McTNs) are tubulin-driven protrusions of the membrane on detached cells which aid in cell–cell association. (a) Diagram of a detached cell producing McTNs. McTNs are promoted by tubulin posttranslational modification indicative of microtubule stability, weakening of the actin cortex, and association of vimentin, tau and/or kinesins. Created in BioRender. Ju, J.A. (2026) <https://BioRender.com/c2448457> (accessed on 25 March 2026). (b) I. McTNs produced by the BT549 breast cancer cell line. Suspended cells were immobilized on the TetherChip, fixed and stained with WGA-594 (red, cell membrane) and DAPI (blue, nucleus) The cell body is overexposed to better visualize the McTNs. II. A CTC producing McTNs isolated from the blood of a breast cancer patient immobilized on the TetherChip, fixed and stained with α -tubulin and alexa-488 (green) antibodies. III. Dissociated tumor cells from a patient's biopsy producing McTNs. This image is a representative frame from a 5 min movie of live cells (binary image in black and white). McTNs are dynamic cell protrusions and were moving in and out of the z-stack range. IV. McTNs during homotypic clustering of MDA-MB-231 breast cancer cells expressing GFP-membrane (green) and heterotypic cell clustering between an MDA-MB-231 breast cancer cell and a differentiated neutrophil (white). Clustered cells are immobilized on the TetherChip, fixed and stained. Neutrophil (white) identified with an anti-CD11b antibody (magenta) and WGA-488 (green) to stain the cell membrane. The overlay of the magenta and green channels creates the white cast. DAPI was used to visualize nuclei (blue). Images were taken with a 60 $\times$ objective on either an Olympus IX81 microscope with a Fluoview FV1000 confocal laser scanning system (Olympus Corporation, Center Valley, PA, USA) or a Nikon Ti2-E inverted microscope with a Nikon AX-R confocal system (Nikon Instruments Inc., Tokyo, Japan).

4. Promoting Effect of ROS in Metastasis

Cancer cells exhibit an elevated metabolism to primarily support rapid proliferation, biomass production, and survival in nutrient-poor, hypoxic tumor environments [91]. Their elevated metabolism promotes ROS production through multiple mechanisms due to their reprogramming. Acute-to-moderate hypoxia also led to an increase in ROS due to a redox

imbalance and back-up of electrons in the electron transport chain [92–94]. Beyond initiating DNA damage and primary tumor cell proliferation, oxidative stress and elevated ROS levels can enhance multiple steps within the metastatic cascade. Here, we discuss how NOX activation and hypoxia can induce an EMT and how ROS can alter cell signaling to promote tumor cell migration and invasion during the processes of intra- and extravasation, and we discuss some external sources of ROS from the tumor microenvironment encountered by metastasizing tumor cells.

4.1. NOX Activation and EMT

ROS promote epithelial-to-mesenchymal transition (EMT) through several interconnected mechanisms and signaling pathways. The most described mechanisms involve activation of transcription factors and modulation of key cellular pathways. ROS up-regulate EMT transcription factors such as Snail, Twist, and ZEB1/2, particularly via NF- κ B-dependent signaling [95,96]. Under normoxic conditions (21% O₂), NOX1 activation produces H₂O₂ to promote a Snail-induced EMT via NF- κ B pathway activation, which can be blocked by the ROS scavenger N-acetyl-cysteine (NAC) [97,98]. In A549 lung cancer cells, NF- κ B transcriptionally drove NOX4 expression in response to TGF- β . NOX4 up-regulation increased ROS to promote a Snail-induced EMT. Either an NF- κ B inhibitor or NOX inhibitor was able to inhibit the EMT [99]. In glioblastoma, however, the TGF- β -induced NOX4 is mediated via SMAD3 to induce metabolic reprogramming. While an EMT shift was determined by an increase in the mesenchymal markers N-cadherin and vimentin and phenotypic behaviors such as migration and invasion, the status of any EMT transcription factor was not examined [100]. In both nontumorigenic mammary epithelial MCF-10A cells and metastatic triple-negative breast cancer MDA-MB-231 cells, TGF- β treatment upregulated NOX4 and the production of subsequent extracellular superoxide to induce an EMT (Figure 4) [101,102]. Thus, ROS production via NOX expression and/or activation can drive an EMT shift.

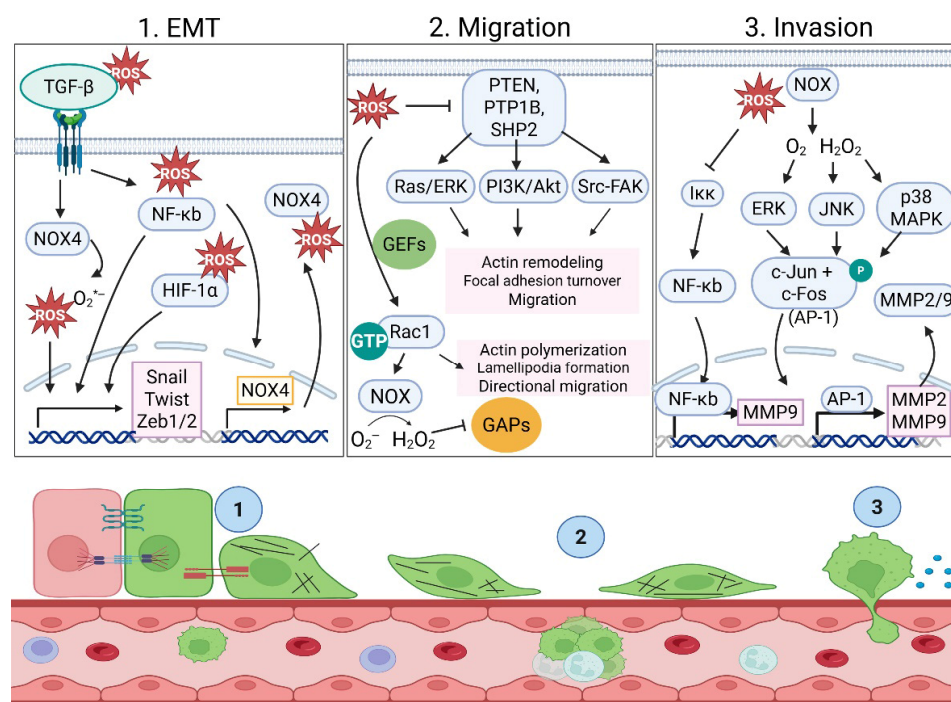


Figure 4. ROS promote EMT, cell migration and cell invasion via multiple mechanisms. Cell signaling pathways stimulated by ROS to promote EMT (1), cell migration (2), and cell invasion (3). As the tumor

cells undergo an EMT (pink to green), their morphology changes and they become less adherent to neighboring cells. During an EMT (1), the tumor cells acquire enhanced migratory (2) and invasive capabilities (3) for movement away from the primary tumor and intravasation into the bloodstream. ROS can directly and indirectly regulate TGF- β , NF- κ B, and HIF-1 α . ROS directly regulate TGF- β by activating its latent form or indirectly by inducing its mRNA and protein expression via signaling pathways such as NF- κ B. Directly, nuclear ROS can modify NF- κ B subunits and reduce their ability to bind DNA, or ROS can indirectly activate the NF- κ B pathway by upstream IKK complex oxidation or via phosphatase inhibition, resulting in prolonged pathway activation. ROS directly regulate HIF-1 α via inhibition of PHD activity or via post-translational modification that prevents HIF-1 α degradation. ROS can also indirectly regulate HIF-1 α via activation of signaling pathways such as PI3K/Akt and MAPK/ERK or via NOX enzymes in a positive feedback loop. ROS directly regulate PTPs (PTEN, PTP1B, SHP2) via active-site cysteine oxidation. NOX4 is considered constitutively active, so it produces ROS without stimuli. NOX4-derived ROS can lead to increased mitochondrial ROS and activate signaling pathways, such as PI3K/AKT or NF- κ B, which, in turn, can further upregulate NOX4 expression, creating a self-perpetuating cycle. Created in BioRender. Ju, J.A. (2026) <https://BioRender.com/c2448457> (accessed on 25 March 2026).

4.2. Hypoxia and EMT

Hypoxia-induced oxidative stress is mainly due to increasing mitochondrial and non-mitochondrial ROS production while simultaneously impairing antioxidant defenses. ROS generated from acute-to-moderate hypoxia leads to the stabilization of Hypoxia-Inducible Factor 1 alpha (HIF-1 α) by inhibiting prolyl hydroxylases (PHDs), which mark HIF-1 α for proteasomal degradation [103]. As HIF-1 α levels decline and the cellular response shifts from acute adaptation to chronic hypoxia management, HIF-2 α and/or HIF-3 α may increase. While HIF-1 α is the best-characterized HIF isoform in EMT, there is some evidence to suggest that HIF-2 α and HIF-3 α may complement HIF-1 α in promoting an EMT [104–106]. However, HIF-1 α is considered the master regulator protein that controls the expression of genes involved in an acute hypoxic response and directly binds to hypoxia response elements (HRE, 5'-RCGTG-3') in the promoters of *Twist*, *Snail* and *ZEB1* [107,108]. *Twist* and *Snail* are most robustly transcriptionally upregulated by HIF-1 α in epithelial and cancer cells, but *ZEB1* also contains HRE sites for HIF-1 α binding (Figure 4) [108–111]. HIF-1 α does not directly bind to the *Slug* or *ZEB2* promoters. Instead, these EMT-driving transcription factors are indirectly induced via NF- κ B and Twist networks [112]. While hypoxia acts as a powerful trigger for oxidative stress, creating conditions where ROS accumulate despite low oxygen levels, hypoxia can also induce an EMT via transcriptional activation of EMT-promoting factors.

4.3. Intravasation and Extravasation

4.3.1. Tumor Cell Migration

ROS promote tumor cell migration through multiple interconnected mechanisms that remodel the cytoskeleton, alter adhesion dynamics, and drive pro-migratory gene expression. ROS can amplify kinase signaling by inactivating key phosphatases controlling the intensity or duration of cellular signals. The best-studied mechanism involves H₂O₂ generated from superoxide (O₂^{•−}) produced by NOX or mitochondria to oxidize phosphatases [113]. By targeting a catalytic cysteine residue in active sites of specific phosphatases, through a series of reversible or irreversible modifications, ROS inactivate these enzymes. Examples of ROS-targeted phosphatases are PTEN, PTP1B, SHP2, and LMW-PTPs (Figure 4) [114–118]. Their inactivation leads to overactive PI3K/AKT, Ras/ERK, and/or Src-focal adhesion kinase (FAK) signaling, thus enhancing actin remodeling, focal adhesion turnover, and tumor cell migration [119,120]. In addition to phosphatase inactivation, ROS can activate Rac1 and other Rho-family GTPases to coordinate actin polymerization, lamellipodia formation, and directional migration by enhancing guanine

exchange factor (GEF) activity or inhibition relief, thereby increasing Rac1-GTP levels [121]. In cancer cells, growth factor or integrin signaling generates NOX-derived ROS to amplify Rac1 activation by stabilizing or recruiting GEFs to the plasma membrane, promoting lamellipodial protrusion [122]. Rac1-GTP can bind NOX subunits directly, activating NOX complexes and generating superoxide, which is rapidly converted to H_2O_2 . This NOX-derived H_2O_2 can oxidize and inhibit Rac1-GTPase-activating proteins (GAPs) or phosphatases that would otherwise terminate Rac1 activity. Thus, Rac1 and ROS can engage in a positive feedback loop where Rac1 stimulates ROS production and ROS, in turn, sustain Rac1 activation [123]. Inhibiting these negative regulators (Rac1-GAPs or phosphatases) shifts the balance toward Rac1-dependent actin polymerization at the leading edge, supporting persistent lamellipodia and forward movement [121,124].

Alternatively, an increase in or chronic ROS contributes to aberrant cell migration by modulating cytoskeletal proteins and dynamics [125–129]. NOX-generated ROS can promote actin polymerization in the front of the cell by targeting specific cysteine residues (Cys139 and Cys147) in cofilin, forming intramolecular disulfide bonds that inactivate its actin-severing activity, promoting actin polymerization [127,128]. However, ROS have also been shown to indirectly promote cofilin binding to F-actin by oxidizing 14-3-3, which allows the cofilin phosphatase SSH-1L to be released and dephosphorylate cofilin [126]. LIM-Kinase (LIMK) is responsible for phosphorylation and cofilin inactivation. While ROS do not typically target LIMK directly, they tune LIMK activity by modulating upstream GTPases, kinases, and phosphatases in the Rho/ROCK–LIMK–cofilin and Rac/Cdc42–PAK–LIMK–cofilin pathways. ROS are also necessary for the establishment of cell polarity during PDGF-stimulated directional migration. Depleting ROS by NOX1 removal disrupts the phosphatase PP2A, leading to aberrant phosphorylation and activation of the Par3/aPKC/Tiam polarity complex, which promotes multiple small lamellipodia-like protrusions from the periphery of the cell instead of the appropriate formation of a single, polarized lamellipodia. The re-expression of NOX1 or exogenous H_2O_2 reverses this phenotype [130]. Thus NOX1-induced ROS can control aPKC and Par3 phosphorylation by regulating PP2A activity, leading to unique and functional lamellipodia. Another study shows that superoxide production by NOX5 is stimulated by actin effector molecules, and knockdown of NOX5 impairs pancreatic cancer cell migration [131]. L-plastin (LPL) is an actin-bundling protein that, when oxidized via H_2O_2 , has diminished actin-bundling capacity, leading to inhibited tumor cell spreading and reduced migration [129]. While multiple cysteines in actin have been reported to undergo oxidation, there is not yet an example of localized oxidation of actin during cell migration [125]. These studies begin to elucidate how ROS affect multiple points in signaling pathways to control actin dynamics and polarity to affect cell migration.

ROS can also alter adhesion dynamics by modulating cell–cell junctions, integrin signaling, and focal adhesions through redox-sensitive kinases, phosphatases, and cytoskeletal regulators. One mechanism by which ROS mediates the weakening of the cell–cell junctions is by inhibiting PTPs responsible for dephosphorylating β -catenin. Phosphorylation of β -catenin disrupts association with E-cadherin, indicating that ROS can help convert junctional disengagement into a pro-migratory program [132,133]. In head-and-neck squamous cell carcinoma (HNSCC), loss of E-cadherin-mediated cell adhesion and cell–cell interactions triggered Rac1–NOX–ROS, which activated the nonreceptor tyrosine kinase Src and STAT3 to drive migration [134]. Integrins connect the extracellular matrix (ECM) to the intracellular cytoskeleton and are the point of internal focal adhesion (FA) complex assembly. ROS can enhance integrin clustering and activation and modify FA components, influencing their dynamics and impacting cell adhesion, migration, and downstream signal transduction [135]. Multiple redox-sensitive pathways tune the lifetime and size

of FA, favoring smaller, more dynamic adhesions that support faster, directional migration [136]. In addition to integrin clustering, ROS can oxidize regulatory cysteines in Src, with Cys245 in the SH2 domain and Cys487 in the kinase domain being the main redox-sensitive residues [137]. Oxidation favors a conformational change, opening the protein for autophosphorylation of Tyr416 in the activation loop to then activating FAK, which enhances focal adhesion turnover and actin remodeling to promote cell migration. As previously mentioned, ROS can also reversibly oxidize cysteine residues in PTPs that normally dephosphorylate Src and FAK, prolonging their activity [118]. Sustained Src-FAK signaling increases phosphorylation of paxillin and talin, promoting FA turnover at the leading edge, which is required for persistent migration [117,135]. Supporting the importance of Src and FAK in metastasis is the fact that an increased Src or FAK expression or activity has been identified in many types of cancer and is correlated with a poor prognosis [138–141]. These studies identify the importance of ROS in remodeling the cytoskeleton and in altering adhesion dynamics to induce a pro-migratory phenotype.

4.3.2. Tumor Cell Invasion: ROS Regulation of MMPs

Matrix metalloproteinases (MMPs) are a family of proteolytic enzymes that degrade components of the extracellular matrix to facilitate cancer cell invasion, with MMP2 and MMP9 (gelatinases) particularly implicated in cancer spread. There are multiple examples of ROS promoting cancer cell invasion via MMP2/MMP9 expression or activity, while antioxidants decrease MMP2-/MMP9-stimulated invasion [142–148]. ROS, particularly H₂O₂ and mitochondrial/NOX-derived superoxide, activate the ERK, JNK, and p38 MAPK pathways, leading to phosphorylation of c-Jun and c-Fos and formation of the AP-1 complex, which binds to the promoters of *MMP2* and *MMP9* to increase their mRNA and protein expression (Figure 4) [149–155]. Although *MMP2* is often described as less dependent on AP-1 than *MMP9*, perhaps in part because it is basally active, functional AP-1 sites exist within its promoter and contribute to its transcription in several cell types [156,157]. ROS can upregulate *MMP2* and *MMP9* via other transcription factors, including ETS factors, HIF-1 α , NF- κ B, or ATF-2 [158–163]. HIF-1 α , stabilized by ROS, can directly transactivate the *MMP9* promoter or indirectly transactivate the *MMP2* promoter via additional factors or intermediate regulators to increase mRNA and protein [164–166]. NF- κ B can be activated by ROS oxidizing and inactivating I κ B or by modulating upstream kinases such as IKK, leading to NF- κ B nuclear translocation and MMP-9 upregulation [167,168]. ROS can also stabilize *MMP9* mRNA. Mori et al. determined that targeted reduction of hydrogen peroxide-inducible clone-5 (HIC-5) promoted lung metastasis in mice after tail vein injection without affecting primary tumor growth. The mechanism for enhanced metastasis followed a HIC-5-NOX4-mtROS-MMP9 axis [169]. In addition to ROS promoting transcriptional upregulation of *MMP2* and/or *MMP9* or stabilizing *MMP9* mRNA, ROS can post-translationally activate *MMP2* and *MMP9*. MMPs are latent proproteins held in an inactive conformation by the cysteine in the prodomain interacting with Zn²⁺ bound to the catalytic domain. Hydrogen peroxide, peroxyxynitrate, nitric oxide and the hydroxyl radical can oxidize the cysteine thiol in the prodomain to disrupt its interaction with Zn²⁺, causing a “cysteine switch” and enzyme activation [170]. The versatility of ROS in activating MMPs at multiple levels underscores their role in metastasis.

4.4. CTC Survival and Immune Cell Evasion

The most important parameter that determines the successful metastasis of tumor cells is CTC survival [171]. One of the survival challenges CTCs must overcome is apoptotic cell death caused by ECM detachment, also known as anoikis. When epithelial-originating tumor cells detach from ECM, they experience a loss in pro-survival signaling pathways,

resulting in disinhibition of the apoptotic pathways. Cancer cells have developed abrogated metabolic mechanisms that alter redox balance to circumvent cell death pathways and ultimately become resistant to anoikis [172]. NOX enzymes have been identified as contributors to ROS-dependent anoikis resistance. NOX4 was identified as an oncogene that is overexpressed in tumorigenic breast cell lines, primary breast tumors, and primary ovarian tumors [173]. NOX4-overexpressing MCF-10A cells demonstrated resistance to Etoposide-induced apoptosis, in addition to induction of other tumorigenic phenotypes, including anchorage-independent growth, invasion and cell division. In gastric and lung cancer cells, NOX4 is upregulated in suspended cells relative to attached cells and confers anoikis resistance [174,175]. NOX4-mediated anoikis resistance occurred via elevation of ROS levels, leading to oxidation and activation of Src and EGFR, which sustained suspended cell survival through the activation of the PI3K/AKT and ERK/MAPK signaling pathways (Figure 5) [174,175]. Furthermore, NOX4 overexpression and ROS treatment increased EGFR levels to promote anoikis resistance, which was attenuated upon NOX4 knockdown and depletion of ROS. An in vivo xenograft colorectal cancer (CRC) model revealed NOX4 inhibition reduced tumor growth and lung metastasis [176]. Although this study did not investigate suspended CRC cells, it is clear that NOX4 plays an essential role in regulation of apoptotic cell death in multiple solid tumors [177]. The overexpression of NOX4 in both CRC and oral tongue squamous cell carcinoma was shown to be associated with poor overall survival [176,178]. Beyond the tumor-intrinsic contributions of NOX4-generated ROS, tumor-extrinsic NOX4 expression in cancer-associated fibroblasts exert tumor-promoting functions [179]. To demonstrate the effect of NOX4 in the stroma, mouse mammary tumor cells were implanted into wild-type mice and mice with a constitutive deletion of NOX4 (*Nox4*^{−/−}). The mice with *Nox4* deleted had a reduced tumor volume and a significant decrease in distant metastases compared to the wild-type control mice, suggesting that the oxidative tumor microenvironment promotes tumor cell survival via paracrine signaling pathways. Overexpression of NOX4 is associated with poor patient survival, and its inhibition reduces tumor growth and metastasis, identifying it as a key therapeutic target.

Isolation of CTC clusters from breast cancer patients revealed the majority (85.5–91.7%) of CTC-associated white blood cells to be Ly-6G positive cells with neutrophil nuclear morphology, suggesting CTC–neutrophil clusters to be the primary CTC–cell interaction in circulation aside from other tumor cells [180]. We have recently shown that, in addition to cancer cells, neutrophil-differentiated HL-60 cells and primary human neutrophils form McTNs to aid in heterotypic tumor cell clustering [181]. Circulating granulocytic peripheral mononuclear-myeloid-derived suppressor cells (PMN-MDSCs) are often referred to as “pathologically activated neutrophils” and have been found in heterotypic clusters with CTCs isolated from melanoma and breast cancer patients. When a combination of patient-isolated PMN-MDSCs and brain metastatic breast cancer cells (MDA-MB-231BR) were introduced into mice via intracardiac injection, the combination enhanced tumor cell dissemination [182]. PMN-MDSCs facilitated CTC survival through ROS generation, which activates the NRF2-ARE axis and induces Notch1 gene expression in the associated CTCs (Figure 5). Suspended lung and breast cancer cell lines induce NOX4-mediated upregulation in fibronectin and desmosomal proteins to facilitate anoikis resistance through cell aggregate formation (Figure 5) [183]. It is possible that McTN-mediated cell clustering brings tumor cells within close proximity to peripheral blood immune cells that can support this ROS-dependent CTC survival.

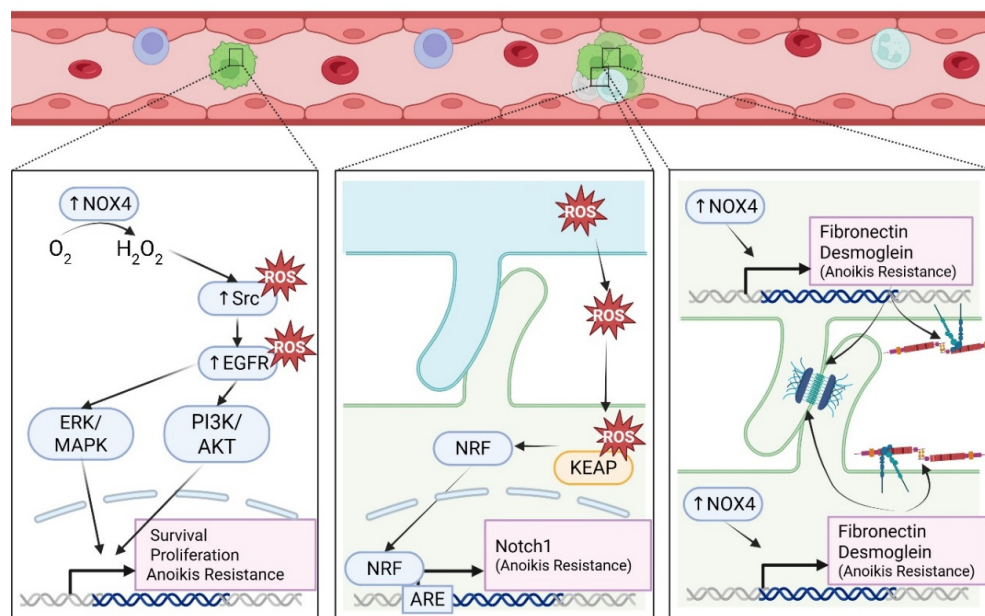


Figure 5. ROS can aid in CTC survival. Upregulation of NOX4 (vertical arrow pointing up) can enhance anoikis resistance and CTC survival via upregulation of the MAPK and PI3K pathways. NOX4-derived ROS can inhibit Src activity via direct oxidation of a cysteine residue, acting as a “redox” switch to limit Src activity during high oxidative stress, and ROS can also indirectly activate Src by inhibiting PTPs that normally turn it off. For EGFR, ROS can directly oxidize cysteine in EGFR’s kinase domain to enhance tyrosine kinase activity and indirectly activate signaling intermediary proteins by inactivating negative regulators such as PTPs, driving EGFR phosphorylation and downstream signaling. Peripheral mononuclear-myeloid-derived suppressor cells (PMN-MDSCs) and tumor cells form heterotypic clusters. The ROS generated from PMN-MDSCs, primarily from high activity of NOX2, act on associated CTCs to directly inhibit KEAP1 binding to NRF2 and activate the NRF2-ARE axis to induce Notch1 gene expression and promote anoikis resistance. Homotypic cell clustering induces a NOX4-mediated upregulation in fibronectin and desmosomal proteins to promote cancer cell aggregation and facilitate anoikis resistance. Created in BioRender. Gilchrist, D.E. (2026) <https://BioRender.com/c2448457> (accessed on 25 March 2026).

4.5. Redox-Dependent Metabolic Reprogramming: DTC Dormancy

Driven by stresses at the secondary site, DTCs can enter into dormancy, a state of reversible cell cycle arrest, as a protective survival mechanism to avoid apoptosis. The new microenvironment is typically one of high oxidative stress. Additionally, ROS can damage components of the electron transport chain and inhibit ROS-sensitive enzymes like aconitase, both reducing the forward (oxidative) flux of the TCA cycle and making oxidative metabolism less efficient. Pushing the cells to metabolically adapt, the cells shift away from glycolysis towards mitochondrial oxidative phosphorylation (OXPHOS) and fatty acid oxidation [184]. Nuclear factor erythroid 2-related factor 2 (NRF2) is known to be a master regulator in response to oxidative stress [185], and multiple studies have identified NRF2 target genes that regulate glycolysis, glycogen metabolism, one-carbon metabolism, nucleotide metabolism, fatty acid metabolism, glutamine metabolism, the pentose phosphate pathway and glutathione metabolism [186,187]. While studying metabolic changes in dormant breast cancer cells, Fox et al. determined that the dormant cells upregulated NRF2 to manage increased ROS cells and indeed induced an NRF2 antioxidant transcriptional program [187]. In the absence of oxidative stress, NRF2 is bound by Kelch-like ECH-associated protein 1 (KEAP1) and targeted for degradation, but ROS can oxidize critical cysteine residues on KEAP1 to induce a conformational change, preventing NRF2 binding. In the absence of KEAP1 binding, NRF2 can then accumulate, translocate into the

nucleus, and bind to antioxidant elements (AREs) in gene promoters [188]. High NRF2 levels not only sustain dormancy, but direct manipulation of NRF2 activity was shown to also facilitate the metabolic reprogramming required for dormant breast cancer cells to start growing again. However, persistent ROS were determined not to be the cause of the elevated levels of NRF2 in the recurrent tumors. Instead, NRF2 elevation was possibly due to a noncanonical mechanism such as impaired protein degradation [187].

As an alternative to glutamine metabolism and a way to maintain biosynthesis, dormant cells can switch to reductive carboxylation, which is basically a reversal of a part of the TCA cycle. During reductive carboxylation, α -ketoglutarate, a primary glutamine metabolite, is converted to isocitrate and then citrate using NADPH. The produced citrate is transported into the mitochondria to detoxify ROS via glutathione and related antioxidant systems. Reductive carboxylation typically increases under conditions of low oxygen; however, cells grown as anchorage-independent spheroids increase reductive carboxylation to detoxify ROS under normal oxygen levels [189]. This suppression of conventional metabolism and induced reductive carboxylation was determined to be dependent on isocitrate dehydrogenase 1 (IDH1). Depending on the type of cancer, DTCs can persist in a dormant state for many years until being “reawakened” to proliferate by appropriate stimuli [190].

5. Challenges/Controversies/Remaining Gaps

5.1. The ROS Seesaw Effect

Thus far, we have highlighted the metastasis-promoting roles of ROS; however, the sensitive redox homeostasis of a tumor cell can tip the cell from pro-metastatic to anti-metastatic. High ROS levels can be detrimental to the survival of cancer cells by overwhelming their upregulated antioxidant systems and damaging essential proteins and nucleic acids via oxidation. Upon loss of matrix attachment, MCF-10A breast epithelial cells experienced reduced cellular ATP levels and elevated ROS levels [191]. NAC and Trolox antioxidants eliminated detachment-induced ROS and rescued cellular ATP levels to support anchorage-independent cell survival and luminal filling of cells during MCF-10A acini formation. Thus, cellular adaptation to remove excess ROS and reduce oxidative stress could actually support CTC survival. This was further validated by the expression or knockdown of catalase enzyme to demonstrate its critical role to support anchorage-independent growth in MCF-10A, MDA-MB-231, and T47D breast cells [192]. Amongst a screen of small molecules that could be used for successful ex vivo expansion of single CTCs isolated from breast cancer patients, NAC proved to be the best compound across four primary CTC lines. NAC treatment was sufficient to rescue CTC proliferation, likely through metabolic alterations that mitigate oxidative stress [193].

While low or moderate ROS levels can act as signaling messengers that can promote cell movement, high levels cause excessive protein oxidation, DNA damage, and cell death, overriding pro-migratory and pro-invasion pathways. Extremely high levels of ROS induce mitochondrial dysfunction, leading to both caspase-dependent and -independent apoptosis [31]. As mentioned, ROS can promote actin polymerization to enhance migration, but levels that are not spatially regulated and/or too high can restrict the cell's ability to move by altering its mechanical properties [194]. High ROS can trigger F-actin polymerization specifically at the actin cortex, which makes the cell more rigid, significantly slowing migration rates, an effect that can be reversed by NAC treatment. Amino acids within microtubules and actin are susceptible to oxidation, and thus excess ROS can interfere with cell movement by direct oxidation of the cytoskeleton. High ROS can cause actin filament severing and reduced microtubule polymerization, effectively freezing the cell's internal transport and structural remodeling. As mentioned, spatially controlled NOX-induced ROS

can oxidize and inactivate cofilin to promote actin polymerization at the leading edge of the cell. However, excess ROS can lead to uncontrolled cofilin inactivation, where, although cofilin may be able to bind actin, it loses its ability to sever or depolymerize the filaments, completely halting actin turnover [195]. Additionally, ROS can deplete energy that is essential for migration by activating poly (ADP-ribose) polymerase (PARP-1) upon DNA damage, which consumes NAD⁺ and inhibits glycolysis, leading to ATP depletion [196]. Excessive ROS can also inhibit PTPs by irreversible oxidation rather than reversible signaling, altering the phosphorylation balance necessary for focal adhesion turnover.

5.2. Novel ROS Detection Approaches

ROS play a dynamic pathophysiological role throughout the metastatic cascade that is determined by the levels of ROS (high or low), the source of ROS and the stage of metastasis (EMT, invasion, migration, intravasation/extravasation, and circulation). As a result, biomolecular markers for real-time detection methods to assess ROS activity in various stages of the metastatic cascade are critical to understand the metastatic propensity of tumor cells. The rapid redox fluctuations, short half-lives, and high reactivity of ROS enhance the difficulty of their direct, real-time measurement. However, novel approaches to measure ROS in vitro and in vivo are continually being developed. Most in vitro methods to measure ROS are commercial kits with fluorescent or chemiluminescent probes that interact with all ROS species or are species-selective [197]. Oxidation products have also been used as a measure of cellular ROS activity. Integration of these probes with poly-HEMA-coated tissue culture plates enabled the assessment of ROS roles in extracellular matrix-detached cancer cells [198]. DCF-DA, CellROX, and MitoSOX fluorescent ROS indicators were applied to detached cells to elucidate pro-survival roles of antioxidant activity in suspended cells, representative of circulating tumor cells (Figure 6) [191]. Our group patented a novel optically clear cell tethering nanosurface technology, TetherChip, to spatially immobilize nonadherent cells. TetherChip technology engages hydrophobic interactions with the cell plasma membrane to spatially immobilize cells while simultaneously preventing the formation of protein-based adhesions [79]. TetherChip has enabled the elucidation of key biomolecular mechanisms, including tubulin-driven protrusions (McTNs) that support metastatic phenotypes in nonadherent cells, which were identified with intracellular immunofluorescence approaches [79,81–90,181,199–204]. It is possible TetherChip technology can be incorporated with ROS detection probes in live tumor cells to advance our understanding of the pro-survival or pro-death roles of ROS in matrix-detached states.

Beyond these effective in vitro ROS detection methods, in vivo detection and quantification of ROS will enable minimally invasive, real-time assessment of patient tumors. A novel photoacoustic and fluorescent probe, JW41, was developed for H₂O₂-selective detection and quantification, proving to be effective in vitro and in vivo [205]. The capped near-infrared probe (JW41) irreversibly reacts with supraphysiological levels of H₂O₂, which forms an uncapped derivative (JW35) that experiences a shift in the absorption spectrum to be detected by photoacoustic and fluorescent imaging. This probe was evaluated to be stable in plasma at 37 °C, nontoxic to MCF-7 and MDA-MB-231 breast cancer cells and taken up efficiently via glucose transporters. In vivo testing with JW41 demonstrated specific accumulation and retention in tumor tissue with spectral signals observed from conversion to the JW35 derivative up to 24 h. A chemoselective bioluminescent probe, Peroxy Caged Luciferin-1 (PCL-1), afforded real-time detection of H₂O₂ in androgen-sensitive prostate tumor (LNCaP) xenograft mice models [206]. Interaction with increasing concentrations of H₂O₂ releases firefly luciferin from PCL-1 to produce increasing total photon flux upon luciferase exposure. This probe was able to detect real-time ROS fluctuations in LNCaP tumors following testosterone stimulation or NAC treatment of the tumors.

To detect early signs of doxorubicin-induced cardiotoxicity in cancer patients, a novel positron emission tomography (PET) tracer, ^{18}F -DHMT, was used to identify elevated ROS production [207]. ^{18}F -DHMT radiotracer was introduced in rats to demonstrate sensitive detection of early increases in myocardial superoxide levels to establish an effective non-invasive detection method. While this work focused on usage in chemotherapy-induced cardiotoxicity, it is possible that ^{18}F -DHMT could be applied with the standard of care imaging platform, PET, as a detection method for elevated ROS in primary or circulating tumor cells to indicate EMT, cell survival, and other pro-metastatic phenotypes.

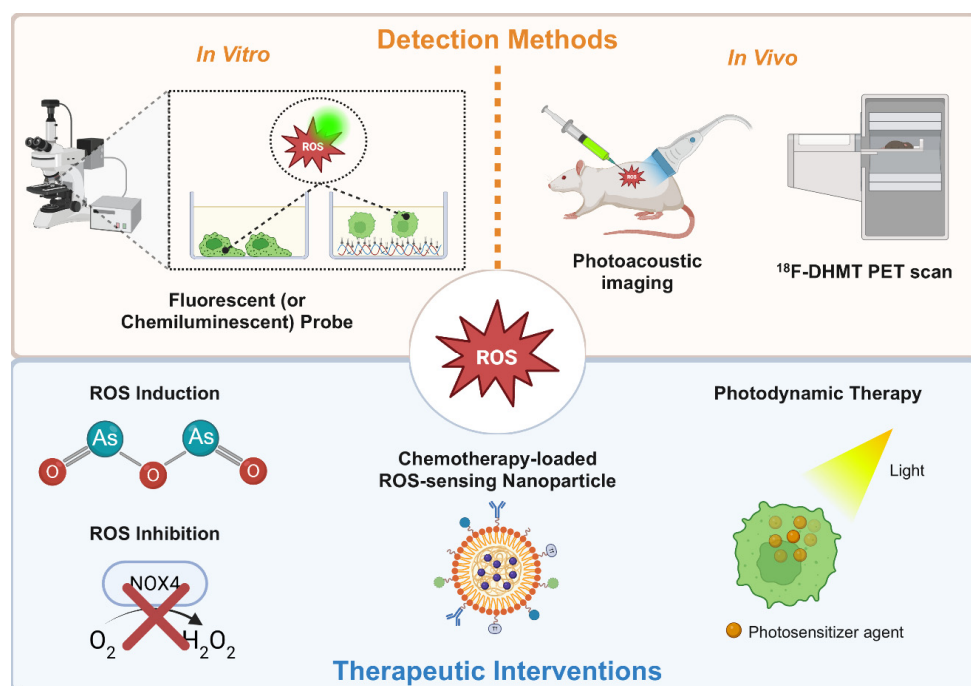


Figure 6. ROS detection methods and therapeutic interventions. In vitro and in vivo ROS detection methods and ROS-centered therapeutic strategies aiming to alter metabolic state of tumor cells by manipulating ROS levels. Created in BioRender. Glichrist, D.E. (2026) <https://BioRender.com/c2448457> (accessed on 25 March 2026).

5.3. Translational Considerations and Redox-Modulating Agents

Tumor cells take advantage of ROS signaling to exert pro-metastatic phenotypes that contribute to poor patient outcomes. However, we also highlighted that tumor cells can leverage antioxidants to reduce oxidative stress burden, thus maintaining an optimal level of elevated ROS to be pro-tumorigenic without deleterious effects. As a result, current ROS-centered therapeutic strategies aim to disrupt that altered metabolic state of tumor cells by increasing ROS beyond manageable levels or reducing ROS below pro-tumorigenic/metastatic levels (Figure 6). Over the years, multiple ROS-modulating drugs have been tested in clinical trials. Antioxidant inhibitors such as auranofin, buthionine sulfoximine (BSO), PX-12, disulfiram and ATN-224 selectively disable protective antioxidant systems (Table 2). The anti-rheumatic drug auranofin inhibits thioredoxin reductase and promotes the accumulation of ROS by disrupting the thioredoxin antioxidant system and is being repurposed for cancer treatment. Combined auranofin and sirolimus for advanced lung cancer (NCT01737502) was generally tolerable but produced only modest clinical benefit, with a median overall survival of 4.4 months. While this specific combination did not support a new standard of care, auranofin is advancing into other trials combined with immune checkpoint inhibitors, targeting different cancers such as recurrent glioblastoma and ovarian cancer. BSO has been tested in multiple Phase 1 clinical trials but almost exclusively

in combination with chemotherapy agents to overcome drug resistance by depleting glutathione in cancer cells [208]. Similarly, PX-12 showed promise when used in combination with chemotherapy to overcome drug resistance, but these combinations largely remain at the research stage rather than progressing into clinical trials. Disulfiram in the presence of copper forms a potent anticancer complex (CuET) that increases intracellular ROS and inhibits the NF- κ B pathway and ubiquitin proteasome system. Unfortunately, disulfiram has proven unsuccessful thus far in multiple clinical trials due to poor stability, rapid metabolism, and/or short half-life within the plasma, which reduces the accumulation of necessary CuET concentrations in the tumor microenvironment [209]. Finally, ATN-224 is a copper chelator that leads to the inhibition of the copper-/zinc-dependent enzyme SOD1, increasing intracellular superoxide and decreasing hydrogen peroxide levels, and it has shown limited-to-moderate success as a cancer treatment [210].

Table 2. Redox-modulating drugs tested as cancer treatments.

Drug Category	Drug Name	Target Enzyme/Mechanism	Cancer Type
Antioxidant Inhibitors	Auranofin	Thioredoxin Reductase (TrxR)	Lung Cancer (NSCLC and SCLC) Ovarian Cancer, Chronic Lymphocytic Leukemia (CLL)
	Buthionine Sulfoximine (BSO)	γ -Glutamylcysteine ligase (GCL)	Neuroblastoma, Melanoma, Advanced Solid Tumors
	PX-12	Thioredoxin-1 (Trx-1)	Advanced or Metastatic Cancer
	Disulfiram	ALDH; GSH/GSSG balance	NSCLC, Prostate Cancer, Metastatic Melanoma, Metastatic Breast Cancer
	ATN-224	Superoxide Dismutase 1 (SOD1)	Prostate Cancer, NSCLC, Breast Cancer, Hepatocellular Carcinoma, Hematological Malignancies
Prooxidant Agents	Elesclomol	Mitochondrial ROS generation	Metastatic Melanoma
	Arsenic Trioxide	Mitochondrial ROS/TrxR	Acute Promyelocytic Leukemia
	High-dose Vitamin C	Hydrogen Peroxide	Pancreatic Cancer, Glioblastoma, NSCLC, Ovarian and Colorectal Cancers
	Artesunate	Iron-mediated ROS generation	Colorectal and Pre-cancerous Cervical Neoplasia
Redox Catalysts	Setanaxib (GKT137831)	NOX1/NOX4 Inhibitor	Head and Neck Cancer
	APX3330 (E3330)	APE/Ref-1 redox signaling	Advanced Solid Tumors

Prooxidant agents such as Elesclomol, arsenic trioxide, high-dose vitamin C and artesunate directly generate harmful ROS to promote cell death. Unfortunately, a Phase 3 clinical trial (NCT00522834) using Elesclomol in combination with paclitaxel for treatment of metastatic melanoma was stopped early due to an imbalance in deaths [211]. However, the results from a 2025 completed Phase 3 clinical trial (NCT02688140) supports the use of the FDA-approved arsenic trioxide (ATO, As₂O₃) in combination with all-trans retinoic acid (ATRA) for the treatment of acute promyelocytic leukemia (APL) [212]. Intravenous, high-dose vitamin C in combination with chemotherapy and radiation has shown promising results in Phase 2 and 3 clinical trials [213]. For metastatic pancreatic cancer, adding high-dose vitamin C to standard chemotherapy doubled the median overall survival for patients from 8 to 16 months (NCT02905578) [214]. Preliminary findings from Phase 1 (NCT01752491) and Phase 2 (NCT02344355) trials testing the addition of intravenous vitamin C to the treatment regime of glioblastoma patients demonstrated a 5-month increased

survival time. Artesunate, primarily used as an antimalarial, is being tested for its ability to promote ferroptosis and apoptosis in cancer cells. Results from an early Phase 1 foundational pilot study showed it to be well tolerated while extending survival times for the patients receiving artesunate compared to the placebo (NCT02353026) [215]. These promising results helped to launch a current larger, multicenter Phase 2 trial to test artesunate as a neoadjuvant treatment for colorectal cancer (NCT07095309). Artesunate is also being evaluated in Phase 2 trials for its safety and effectiveness in preventing the progression of pre-cancerous cervical intra-epithelial neoplasia to invasive cancer (NCT04098744 and NCT07095478). Although artesunate is in testing as a monotherapy, most redox modulators show minimal clinical activity alone and are increasingly investigated for their ability to sensitize tumors to radiation or standard chemotherapy.

As discussed, ROS can serve as promoters of tumor growth and metastasis, and the direct reduction in ROS or inhibition of ROS-generating enzymes may pose as beneficial therapeutic approaches. While NOX enzymes are a plausible anticancer redox target, most evidence testing NOX-targeted treatment are still in the preclinical or early clinical stages. However, Setanaxib (GKT137831) is an oral, first-in-class inhibitor of NOX1/4 and has received the FDA orphan drug designation (ODD) for treatment against systemic sclerosis, idiopathic pulmonary fibrosis, and primary biliary cholangitis (NCT03865927, NCT05014672) [216]. Considering the elucidated role of NOX4 in tumor cell survival that supports metastatic propensity, Setanaxib would be an interesting drug to test against solid tumors. Thus far, a Phase 2 clinical trial combining Setanaxib with Pembrolizumab in patients with recurrent or metastatic squamous cell carcinoma of head and neck cancer showed significant improvements in progression-free survival (PFS) and overall survival (OS) (NCT05323656). A novel approach to overcome resistance to radiotherapy against tumor cells encapsulated Setanaxib into a bioactive and CD44-targeted hyaluronic acid nanoparticle (HANP) [217]. The nanoparticle payload was systemically delivered to breast cancer patient-derived xenograft (PDX) models along with low-dose local radiotherapy and demonstrated tumor growth inhibition and elevated cell death. As of April 2026, the general trend shows a clear split between the described redox modulators being used in specific combinations and those which have stalled due to efficacy and toxicity issues. Studies are shifting away from BSO, PX-12, ATN-224, and Elesclomol for the treatment of cancers, but Auranofin, vitamin C, artesunate, Setanaxib, and APX3330 show promise and are moving forward into specialized or combination trials.

The FDA has also approved photodynamic therapy (PDT) using light-activated photosensitizers to generate ROS and kill cancer cells (Figure 6). PDT has been used to treat multiple pre-cancerous and cancerous solid tumors, including esophageal cancer, non-small-cell lung cancer, and skin cancer [218]. Current clinical trials often focus on intraoperative or endoscopic delivery to target residual cells that surgery might miss. Though beneficial for reducing tumors that exist specifically at the irradiated site, PDT is not as effective at reducing CTCs or disseminated metastases undetected by current imaging modalities. While not an FDA-approved therapy, Wang et al. demonstrated that their docetaxel-loaded pH/ROS dual-responsive nanoparticle shows potential for preventing tumor metastasis. The intracellular acidic environment and endogenous ROS accelerate the degradation of the nanomaterial, and a component from the degraded material induces more ROS through mitochondrial damage to amplify the docetaxel payload release [219]. This nanoparticle demonstrated tumor-selective capabilities due to the agent-release mechanisms and efficacy to reduce in vivo tumor growth and metastasis. Although our group showed that Taxol agents can induce metastatic phenotypes, this nanoparticle platform may serve as a strong candidate to carry anti-tumor and anti-metastatic therapeutic payloads that induce tumor cell death through oxidative stress mechanisms [220]. These novel approaches to challenge

tumor cells with reduced ROS or increased ROS to induce cell death along each stage of the metastatic cascade raise excitement for the advancement of cancer therapies.

6. Conclusions

In this review, we have highlighted the dynamic, bifunctional nature of reactive oxygen species (ROS) in tumor metastasis. ROS species are produced through redox reactions and ROS-producing enzymes or as byproducts of normal metabolic activity. ROS levels can overwhelm normal cellular antioxidant systems to cause damage and even cell death. On the other hand, elevated ROS in malignant tumor cells can exert pro-metastatic phenotypes in a stage-dependent manner. Hypoxia-induced and NOX-derived ROS facilitate the epithelial–mesenchymal transition to increase cell motility, invasive capacity, and resilience against internal or external stressors. Once tumor cells have entered the vasculature, ROS activate intracellular and paracrine pro-survival, anti-apoptotic signaling mechanisms to support CTCs during the harsh conditions experienced in circulation. Although substantial evidence demonstrates ROS as pro-metastatic factors, we have discussed in this review mechanisms by which ROS can be damaging to cells. Therefore, there is evidence showing that reduced ROS, via antioxidant upregulation, can also promote metastatic phenotypes. The bifunctional and context-dependent nature of ROS presents a need to develop novel detection methods to investigate ROS at different stages in vitro and in vivo with fluorescent probes and medical imaging modalities. Advancements in our understanding of the multifaceted roles of ROS in metastasis, and other pathological diseases, have led to clinical trials and FDA approval of ROS-centered therapeutics, including ROS inducers, inhibitors, and drug delivery agents. Our ongoing efforts are to elucidate the roles of ROS in each stage of metastasis, to develop novel technologies for ROS detection, and to manufacture or repurpose ROS-sensing, -targeting, or -generating agents to reduce metastasis and improve patient outcomes.

Supplementary Materials: The following supporting information can be downloaded at: <https://www.mdpi.com/article/10.3390/antiox15050529/s1>, Table S1: Characteristics of Additional ROS Species.

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microtentacles and microfluidic cell tethering, it did not bias the interpretation of this work. To maintain impartiality, blinded raw data was subject to a standardized image analysis by a third party.

Abbreviations

The following abbreviations are used in this manuscript:

^{18}F -DHMT	^{18}F -6-(4-((1-(2-fluoroethyl)-1H-1,2,3-triazol-4-yl)methoxy)phenyl)-5-methyl-5,6-dihydrophenanthridine-3,8-diamine
aPKC	Atypical protein kinase C
ALDH	Aldehyde dehydrogenase
APE	Apurinic/apyrimidinic endonuclease 1
APL	Acute promyelocytic leukemia
ARE	Antioxidant elements
ATF2	Activating transcription factor 2
ATO	Arsenic trioxide
ATP	Adenosine triphosphate
ATRA	All-trans retinoic acid
CAT	Catalases
CellROX	Cellular reactive oxygen species indicator probe
c-Fos	Cellular FBJ murine osteogenic sarcoma virus
cGMP	Cyclic guanosine monophosphate
c-Jun	Cellular Jun protein
CLL	Chronic lymphocytic leukemia
CRC	Colorectal cancer
CTC	Circulating tumor cells
DAPI	4',6-diamidino-2-phenylindole
DCF-DA	2',7'-dichlorodihydrofluorescein diacetate
DNA	Deoxyribonucleic acid
DTC	Disseminating/disseminated tumor cell
DUOX	Dual oxidase
ECM	Extracellular matrix
EGFR	Epidermal growth factor receptor
EMT	Epithelial-to-mesenchymal transition
eNOS	Endothelial nitric oxide synthase
ER	Endoplasmic reticulum
ERK	Extracellular signal-regulated kinase
ETC	Electron transport chain
ETS	Erythroblast transformation-specific
FA	Focal adhesion
FAD	Flavin adenine dinucleotide
FAK	Focal adhesion kinase
Fas	FS-7-associated surface antigen
FDA	Food and Drug Administration
GAP	GTPase-activating protein
GCL	γ -Glutamylcysteine ligase
GEF	Guanine nucleotide exchange factor
GFP	Green fluorescent protein
GPx	Glutathione peroxidases
GSH	Reduced glutathione
GSSG	Oxidized glutathione
GTP	Guanosine triphosphate
HANP	Hyaluronic acid nanoparticle

HEMA	2-Hydroxyethyl methacrylate
HIC-5	Hydrogen peroxide-inducible clone-5
HIF-1 α	Hypoxia-inducible factor 1 alpha
HRE	Hypoxia response elements
HSNCC	Head-and-neck squamous cell carcinoma
IDH1	isocitrate dehydrogenase 1
IKK	Inhibitor of kappa beta kinase
iNOS	inducible nitric oxide synthase
JNK	c-Jun N-terminal kinase
KEAP1	Kelch-like ECH-associated protein 1
KGDHC	α -Ketoglutarate dehydrogenase complex
LDL	Low-density lipoprotein
LIMK	LIM domain kinase
LMW-PTP	Low-molecular-weight protein tyrosine phosphatase
LPL	L-plastin
Ly-6G	Lymphocyte antigen 6 complex, locus G
MAPK	Mitogen-activated protein kinase
McTNs	Microtentacles
MitoSOX	Mitochondrial superoxide indicator
MMPs	Matrix metalloproteinases
mRNA	Messenger ribonucleic acid
mtROS	Mitochondrial reactive oxygen species
NAC	N-acetylcysteine
NADH	Nicotinamide adenine dinucleotide
NADPH	Nicotinamide adenine dinucleotide phosphate
NCT#####	National clinical trial (registered on ClinicalTrials.gov) [221]
NF- κ B	Nuclear factor kappa-light-chain-enhancer of activated B cells
nNOS	Neuronal nitric oxide synthase
NOS	Nitric oxide synthase
NOX	NADPH oxidases
NRF2	Nuclear factor erythroid 2-related factor 2
NSCLC	Non-small-cell lung cancer
ODD	Orphan drug designation
OS	Overall survival
OXPHOS	Oxidative phosphorylation
PAK	p21-activated kinases
Par3	Partitioning-defective 3
PARP	Poly (ADP-ribose) polymerase
PCL-1	Peroxy caged luciferin-1
PDGF	Platelet-derived growth factor
PDT	Photodynamic therapy
PET	Positron emission tomography
PFS	Progression-free survival
PHD	Prolyl hydroxylase
PI3K	Phosphoinositide 3-kinase
PMN-MDSC	Peripheral mononuclear-myeloid-derived suppressor cells
PP2A	Protein phosphatase 2A
PPARs	Peroxisome-proliferator-activated receptors
PTEN	Phosphatase and tensin homolog
PTP1B	Protein tyrosine phosphatase 1B
PTPs	Protein tyrosine phosphatase
Ref-1	Redox factor-1
RES	Reactive electrophile species
RET	Reverse electron transfer

RNS	Reactive nitrogen species
ROCK	Rho-associated coiled-coil-containing protein kinase
ROS	Reactive oxygen species
RSS	Reactive sulfur species
SH2	Src homology 2
SHP2	Src homology 2 domain-containing protein tyrosine phosphatase 2
SMAD	Suppressor of mothers against decapentaplegic homolog
SOD	Superoxide dismutase
SSH-1L	Slingshot homolog-1 long
STAT	Signal transducer and activator of transcription
TCA	Tricarboxylic acid
TGF- β	Transforming growth factor-beta
Tiam	T-lymphoma invasion and metastasis-inducing protein
Trolox	6-Hydroxy-2,5,7,8-tetramethylchroman-2-carboxylic acid
Trx-1	Thioredoxin-1
TrxR	Thioredoxin reductase
WGA	Wheat germ agglutinin
ZEB1/2	Zinc finger E-box-binding homeobox 1/2

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