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Review Article

Breathing new life into nitric oxide signaling: A brief overview of the interplay between oxygen and nitric oxide[☆]



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ABSTRACT

Nitric oxide ($\bullet\text{NO}$, nitrogen monoxide) is one of the most unique biological signaling molecules associated with a multitude of physiologic and pathological conditions. In order to fully appreciate its numerous roles, it is essential to understand its basic biochemical properties. Most signaling effector molecules such as steroids or proteins have a significant life-span and function through classical receptor–ligand interactions. $\bullet\text{NO}$, however, is a short-lived free-radical gas that only reacts with two types of molecules under biological conditions; metals and other free radicals. These simple interactions can lead to a myriad of complex intermediates which in turn have their own phenotypic effects. For these reasons, responses to $\bullet\text{NO}$ often appear to be random or contradictory when outcomes are compared across various experimental settings. This article will serve as a brief overview of the chemical, biological, and microenvironmental factors that dictate $\bullet\text{NO}$ signaling with an emphasis on $\bullet\text{NO}$ metabolism. The prominent role that oxygen (dioxygen, O_2) plays in $\bullet\text{NO}$ metabolism and how it influences the biological effects of $\bullet\text{NO}$ will be highlighted. This information and these concepts are intended to help students and investigators think about the interpretation of data from experiments where biological effects of $\bullet\text{NO}$ are being elucidated.

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Introduction to nitric oxide

Nitric oxide is a free radical, which is any molecule with an unpaired electron. Before $\bullet\text{NO}$ was found to be endogenously synthesized in humans in the 1980s, most free radicals were largely thought to have deleterious biological effects [1]. For example

oxygen-, nitrogen-, and carbon-centered radicals are strong oxidants that can indiscriminately damage a host of macromolecules including DNA, lipids, and proteins. The most notable examples include the hydroxyl radical ($\bullet\text{OH}$), superoxide (O_2^-), nitrogen dioxide ($\bullet\text{NO}_2$), and carbonate anion radical ($\bullet\text{CO}_2^-$) [2]. Nitric oxide, on the other hand, is considered a relatively stable free radical and under biological conditions it only reacts with two types of molecules: metals and other free radicals [3]. In this context, $\bullet\text{NO}$ is designated to be stable because it does not spontaneously decay or react with itself (dimerize) [4]. Although $\bullet\text{NO}$

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rapidly reacts with other free radicals, the reaction of •NO with another •NO molecule to form N₂O₂ does not occur under biological conditions based largely on entropic grounds [1].

Unlike other biologically relevant free radicals, •NO does not directly undergo one-electron oxidation/reduction reactions to form the nitrosonium cation (NO⁺) or nitroxyl anion (NO⁻), or its conjugate acid (HNO), respectively [5,6]. Although these species can be formed via redox reactions between •NO and metals or O₂, a significant thermodynamic barrier precludes their direct formation. For these reasons, •NO is not considered a particularly good oxidant or reductant. Nevertheless •NO does rapidly react with numerous cellular targets that contribute to its short biological life-time, which in most circumstances is less than 2 s [7]. When compared to other signaling molecules, this is extremely short. However, in relation to other free radicals, whose life-span is often on the order of milliseconds, this is a relatively long time. Being composed of only one atom of nitrogen and one atom of oxygen, •NO is one of the smallest known signaling molecules. Although it is a free radical with an unpaired electron, it does not possess an electrical charge. These are important properties because they allow •NO to readily cross biological membranes by passive diffusion. In addition, •NO is soluble in both hydrophobic and hydrophilic environments. In fact, it is more soluble in hydrophobic environments by a factor of about 10, which turns out to be an important determinant of its biological actions and reactivity [8].

Biological targets of •NO

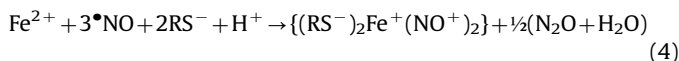
Iron

As mentioned above, in a cellular setting, •NO will only react with transition metals and other free radicals. The majority of •NO/metal reactions involve ferrous iron (Fe²⁺) and to a lesser extent copper (Cu⁺) [3]. There are several forms of iron in cells and •NO is known to react with all of them, albeit to varying degrees. A large proportion of iron exists in the form of heme (iron protoporphyrin), where it is used for enzyme catalysis or to transport O₂. In erythrocytes and myocytes, the prominent heme-containing proteins are hemoglobin and myoglobin. The main function of these proteins is to transport dioxygen (O₂), which binds directly to the ferrous heme. In these proteins, •NO can bind reversibly to the ferrous heme in the absence of O₂, or it can react with an O₂ bound to the heme [9]. The reaction and binding of •NO to reduced heme leads to the formation of iron-nitrosyls via a process called nitrosylation Eq. (1). In the absence of O₂, this is a relatively stable bond. On the other hand, in the presence of O₂, its reaction with the •NO-bound iron center converts it to nitrate (NO₃⁻) Eq. (2). Conversely, •NO can react with oxyHb, which also forms NO₃⁻ Eq. (3). Reactions (2) and (3) are extremely fast, diffusion limited reactions, and result in the rapid scavenging of •NO. This is one of the main consumptive mechanisms of •NO in the vasculature [4].



In non-erythroid cells, the reactions of •NO with heme-proteins have important signaling consequences as opposed to merely scavenging bioavailable •NO. One of the most important examples is the reaction of •NO with soluble guanylyl cyclase (sGC). This protein converts GTP to cGMP and binding of •NO to its heme center leads to a 100-fold increase in its catalytic activity [10].

Another key population of cellular iron is “chelatable iron” or the chelatable iron pool (CIP). This is iron in transition between cellular uptake and incorporation into iron storage protein (ferritin) or the catalytic site of enzymes. The CIP is methodologically defined as being the population of cellular iron that is accessible to chemical iron chelators [11]. It is mostly cytosolic but also has mitochondrial and nuclear components. The CIP is redox-active and represents a small but chemically significant population of total cellular iron (< 3%). The rapid reaction of •NO with the CIP and cellular thiols results in the formation of dinitrosyliron complexes (DNIC, (RS⁻)₂Fe⁺(NO⁺)₂) Eq. (4) [12]. In fact, 100% of the CIP is quantitatively converted into DNIC upon cellular exposure to •NO [13]. Moreover, DNIC are the most abundant population of •NO-derived cellular adducts, much more than S-nitrosothiols, heme nitrosyls, or nitrotyrosine [14]. The consequences of DNIC formation can range from iron sequestration and signaling to participating in S-nitrosothiol (RSNO) formation [15–20].



Cellular iron is also used for the activity of non-heme iron oxygenases, which are important enzymes that catalyze a wide variety of degradatory and biosynthetic reactions. The source of iron for these enzymes is the CIP [21–23]. Nitric oxide can bind reversibly to the catalytic iron and inhibit a variety of these enzymes to varying degrees. As O₂ is the normal substrate for these enzymes, the reaction of •NO at the iron site is competitive and it will depend on the relative concentrations of the two gases. As a general rule, any protein that will bind O₂ will also bind •NO (and CO) with either greater or lesser affinity. This will depend on the iron coordination and other bound ligands as well as the oxidation state. Lastly, there are iron-sulfur proteins which are used for redox and electron transfer reactions in a diverse set of proteins. The interaction of •NO with iron-sulfur proteins is usually destructive but they may also be involved in the formation of protein-bound DNIC [24,25].

Free radicals

Paired electrons are stable and nearly 100% of all electrons in our bodies are paired. Just as •NO reacts with iron and other transition metals because they have unpaired electrons (in their *d* orbitals), •NO reacts with free radicals because they possess unpaired electrons as well. In addition to •NO, there are several free radicals of biological interest. Superoxide (O₂⁻) is probably the most notable. Superoxide is the one electron reduction product of O₂ and it is formed as a natural byproduct of oxidative metabolism. It is also the substrate for several types of enzymes as well as the product; superoxide dismutase and xanthine oxidase, respectively. The reaction of •NO with O₂⁻ is near diffusion limited and the product is peroxynitrite (ONOO⁻) [26]. Peroxynitrite can go on to react with numerous other molecules including •NO and O₂⁻ [27]. In general, ONOO⁻ performs strong oxidative chemistry. Depending on the relative flux rates of •NO and O₂⁻, however, the net effect of ONOO⁻ formation is a decrease in the concentrations of both species. The most significant effect of ONOO⁻ may simply be attributed to scavenging of •NO and O₂⁻, thereby diminishing their signaling capabilities [28]. There are other radical-radical reactions with •NO that have biological consequences (for detailed reviews on oxidative stress see and •NO reactivity see [3,6,9,29,30]).

Nitric oxide production

In mammalian systems the dominant mode of •NO production is via enzymatic synthesis from one of three isoforms of nitric

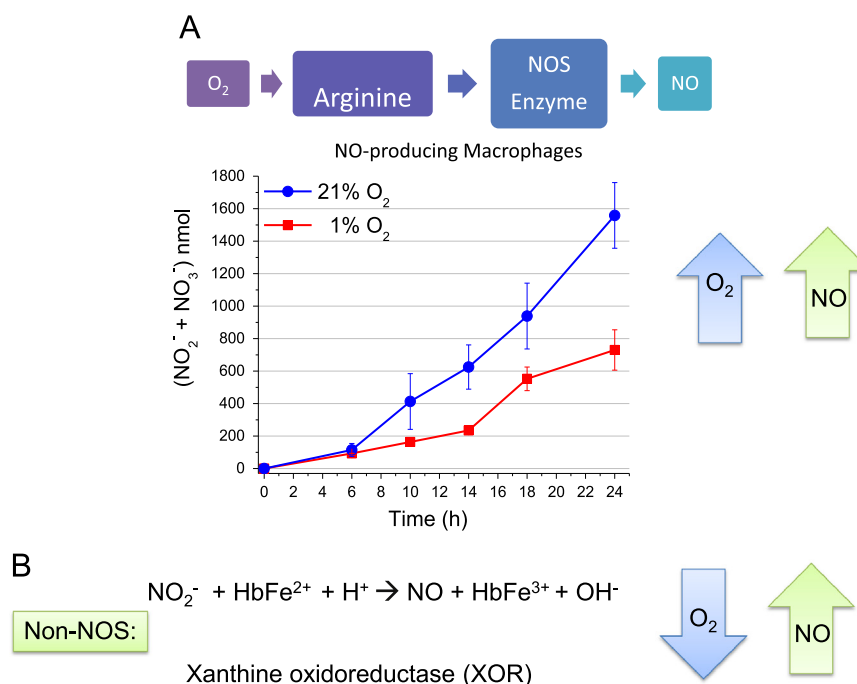
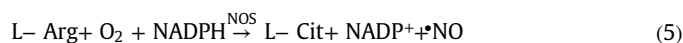


Fig. 1. Oxygen determines the rate of •NO production. (A) Enzymatic synthesis of •NO is a function of O₂ concentration. RAW 264.7 murine macrophages were cultured at 1% or 21% O₂ and stimulated with LPS (*t*=0). Total •NO synthesis was assessed by measuring NO₂⁻/NO₃⁻ in the media at the indicated time points by chemiluminescence. (B) NOS independent mechanisms of •NO synthesis.

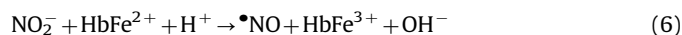
oxide synthase (neuronal nNOS, inducible iNOS, and endothelial eNOS, or NOSI, NOSII, and NOSIII, respectively). The differences between each NOS are based on a variety of factors including their tissue distributions as well as their modes of expression and regulation. nNOS and eNOS are constitutively expressed and generally produce lower amounts of •NO (nM) for short periods of time. iNOS however is “inducible” and it becomes upregulated in response to various stimuli including cytokines and bacterial endotoxins. It is capable of producing greater amounts of •NO for prolonged periods of time. The NOS enzymes are flavoproteins that transfer electrons via NADH, FAD, FMN, and Fe²⁺. They also require the cofactor tetrahydrobiopterin (BH₄). Substrates for NOS enzymes are dioxygen (O₂) and the amino acid L-arginine (Arg), with the products being •NO and the amino acid L-citrulline Eq. (5) [31,32].



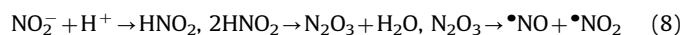
The amount of •NO being synthesized from NOS depends on a variety of factors including substrate and cofactor availability. Posttranslational modifications, such as phosphorylation, as well as tetrahydrobiopterin (BH₄) availability can both influence the rate of •NO synthesis. For example, BH₄ deficiency can result in uncoupling of the enzyme and favor the generation of O₂⁻ [33]. Arginine is usually not a limiting substrate except in disease states but other enzymes such as arginase can compete with NOS for its availability. Under conditions where there is an abundance of substrates and cofactors, the most significant determinant of the rate/amount of •NO synthesis is the local O₂ concentration. The K_M for O₂ for each of the NOS isoforms varies widely (eNOS=23 μM, iNOS=135 μM, and nNOS=350 μM) [32]. At O₂ concentrations below the K_M for O₂, the rate of •NO synthesized from each NOS will be proportional to the O₂ concentration and it will increase linearly as the O₂ concentration raises. For example, eNOS, with the lowest K_M, will maximally produce •NO over a range of physiologic O₂ concentrations. For nNOS, however, the production of •NO will increase over a wide range of O₂ concentrations. Fig. 1A

demonstrates how differences in O₂ concentrations affect total •NO production from iNOS in cultured macrophages [34]. At 24 h, total •NO synthesis at 1% O₂ is about half the amount that is produced at 21% O₂. This is what would be predicted as 1% O₂ is well below the K_M for O₂ for iNOS.

In addition to •NO synthesis from NOS enzymes, there are other biological mechanisms to generate •NO. These include nitrite reduction by heme proteins such as hemoglobin Eq. (6) as well as enzymatic reduction of nitrite by non-NOS enzymes such as xanthine oxidoreductase Eq. (7) (Fig. 1B) [35–37].



Unlike NOS-mediated •NO synthesis which requires O₂, the generation of •NO by the above reactions is enhanced at low O₂. They are thought to serve as compensatory mechanisms for •NO synthesis under hypoxic conditions. Nitric oxide can also be generated by the acidification of nitrite in tissue compartments or cellular organelles where the pH is low Eq. (8) [38]. This reaction is known to occur in the low pH environment of the stomach from dietary nitrite.



As far as total body •NO synthesis, the above 3 mechanisms appear to be minor contributors. However, they do account for the physiologic and therapeutic effects of dietary and pharmacologic nitrite. Although nitrate and nitrite were once thought to be carcinogenic, it is now becoming increasingly clear that the physiologic benefits derived from their ability to generate •NO far outweigh their deleterious properties [39].

Nitric oxide consumption

Although it is well understood how •NO is produced under biological conditions, much less is known about its mode of catabolism. The half-life of •NO is short (0.1–2 s), which is a function

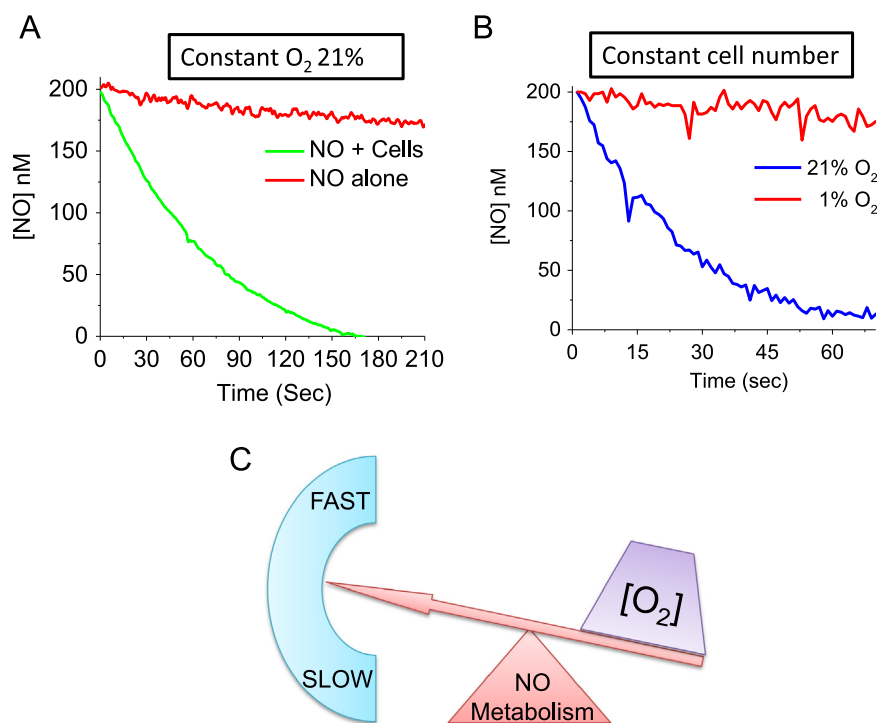


Fig. 2. Oxygen determines the rate of •NO catabolism. RAW 264.7 cells were trypsinized and suspended into a sealed, water-jacketed, temperature-controlled (37 °C) reaction chamber with constant stirring. The reaction chamber was equipped with both •NO and O₂ electrodes connected to an Apollo 4000 free radical analyzer (World Precision Instruments, Sarasota, FL, USA). Headspace in the vessel was negligible compared to the vessel volume to ensure that the rate of •NO volatilization was insignificant compared with its reaction in solution. Reactions were initiated by injection of a saturated •NO solution with a gas-tight syringe, and •NO metabolism was measured using an •NO-selective electrode (amiNO-700, Innovative Instruments, Tampa, FL, USA). (A) At time “0”, the cells were treated with a bolus of •NO (200 nM) and the disappearance of •NO was measured over time. Or •NO added to the reaction chamber in the absence of cell. (B) Disappearance of •NO was measured in cells treated with 200 nM •NO (at 21% O₂) or the cells were allowed to consume O₂ in the reaction chamber until a level of 1% O₂ was achieved before the addition of •NO. (C) The direct relationship between the O₂ concentration and the rate of •NO metabolism.

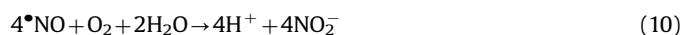
of the mechanisms by which •NO is consumed by cells [7]. These consumptive mechanisms are determined by the rate at which •NO reacts with various cellular targets. Although •NO only reacts with transition metals and other free radicals including O₂, there are a multitude of potential reactants that will differ based on the cell type and local redox environment. Regardless of what the cellular reactants are, cellular consumption of •NO most likely is not simply the result of stoichiometric reactions with cellular targets. Continuous exposure to •NO demonstrated that cells consume •NO at a constant rate which did not appear to be saturable [7]. This indicates that the reactants for •NO are either continuously produced or rapidly recycled. As mentioned above one well-known reaction of •NO is with oxyhemoglobin to form nitrate (NO₃⁻) (Eqs. (2) and (9)). This is an extremely rapid reaction and is the major consumptive mechanism of •NO in the vasculature.



In non-erythroid cells, however, there are a multitude of other potential reactants for •NO including radicals like superoxide (O₂⁻), which forms nitrate (NO₃⁻), or hypervalent metal oxo species (Fe⁴⁺=O), which form nitrite [9]. Nitric oxide will also react with many heme and non-heme iron proteins as well as chelatable iron, ultimately being oxidized to nitrate and nitrite. It is important to point out that the exact mechanism(s) by which •NO is metabolized by cells is not precisely known. This is because •NO has numerous cellular targets and therefore the means by which it is consumed is the sum of several dominant reactions. The contribution of each reaction toward the rate of •NO consumption is a function of the rate of each reaction and the concentration of each reactant. Fig. 2A demonstrates that when a bolus of 200 nM •NO is added to a suspension of cells, it disappears over time via its

reactions with various cellular targets. It can be seen that the rate of •NO disappearance in the presence of cells is much more rapid than its disappearance in the absence of cells, which is known to occur via a direct reaction with O₂ in solution [7,34].

Although the types of specific cellular reactants for •NO may differ between cell types, research has shown that the predominant mechanisms of •NO metabolism require O₂ [7]. This is illustrated in Fig. 2B, which demonstrates the rates of •NO disappearance in the presence of cells at two different O₂ concentrations (21% and 1%). For the same number of cells, •NO is metabolized much more rapidly at 21% O₂ than at 1% O₂. Therefore, there is a direct relationship between the O₂ concentration and the rate of cellular •NO metabolism (Fig. 2C). The discovery that •NO is metabolized by cells in an O₂-dependent manner may not at first seem surprising as it is well-known that •NO directly reacts with O₂ (autoxidation, Eq. (10)).



$$\text{Rate law: } -d[\bullet\text{NO}]/dt = 4k_{\text{aq}}[\bullet\text{NO}]^2[\text{O}_2]$$

There are two critical pieces of evidence that rule out autoxidation as a significant means of •NO metabolism. First, the autoxidation reaction is second-order in •NO concentration whereas the metabolism of •NO by cells follows first-order kinetics Eq. (11) [7,8].

$$-d[\bullet\text{NO}]/dt = k_{\text{obs}}[\text{O}_2][\bullet\text{NO}][\text{Cell}] \quad (11)$$

$$k_{\text{obs}} = 5.38 \pm 0.3 \times 10^{-4} \text{ M}^{-1} \text{ s}^{-1} (\text{cell}/\text{mL})^{-1}$$

Second, if •NO were reacting directly with O₂, its rate of disappearance would be exceedingly slow at physiologic •NO concentrations. Because the rate of •NO autoxidation is second-order with respect to •NO, it means that at high •NO concentrations, the

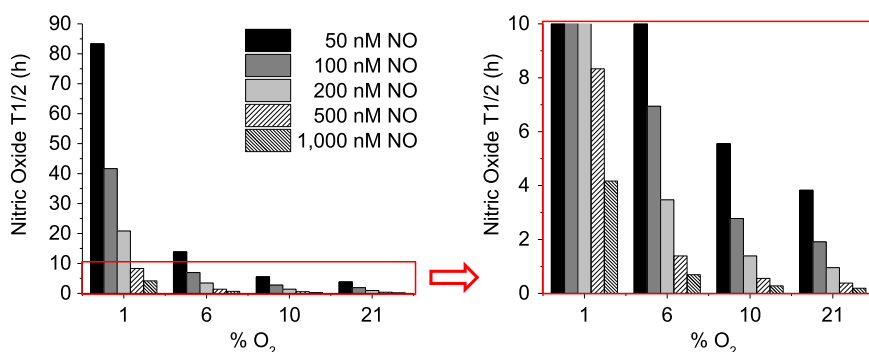


Fig. 3. Concentrations of $\bullet\text{NO}$ and O_2 determine the rate autoxidation. Calculated theoretical half-lives of $\bullet\text{NO}$ via autoxidation at various O_2 concentrations based on initial starting $\bullet\text{NO}$ concentrations of 50, 100, 200, 500, and 1000 nM. The half-life of a reaction is the amount of time it takes for the concentration of a reactant ($\bullet\text{NO}$) to decrease to one-half of its initial value. These values represent the *first* half-life of $\bullet\text{NO}$. Since the autoxidation reaction is a second-order reaction (with respect to the $\bullet\text{NO}$ concentrations), the half-life will change over time as the concentration of $\bullet\text{NO}$ changes.

autoxidation reaction is much faster than at low concentrations. Fig. 3 illustrates how the half-life of $\bullet\text{NO}$ via autoxidation changes over a range of physiologic $\bullet\text{NO}$ and O_2 concentrations. What should be emphasized is that if the dominant reaction of $\bullet\text{NO}$ within cells was directly with O_2 (autoxidation), the half-life of $\bullet\text{NO}$ would be on the order of hours to days at physiologic $\bullet\text{NO}$ and O_2 concentration. Even when the O_2 concentration is high (21%), $\bullet\text{NO}$ could last for hours. This is not to say that $\bullet\text{NO}$ autoxidation does not occur under biological conditions. Experiments have shown that this reaction preferentially occurs in hydrophobic environments such as cell membranes and may be an important source of reactive nitrogen intermediates (N_2O_2 , $\bullet\text{NO}_2$) [8,40]. We know, however, that the biological half-life of $\bullet\text{NO}$ is short (< 2 s), therefore, autoxidation cannot quantitatively account for the loss of a significant proportion of total $\bullet\text{NO}$.

Oxygen determines the steady-state concentration of $\bullet\text{NO}$

Oxygen determines the rate of $\bullet\text{NO}$ synthesis by acting as a substrate for NOS and it also determines the rate of $\bullet\text{NO}$ metabolism. Therefore, the local O_2 concentration will play a significant role in determining the steady-state concentration of $\bullet\text{NO}$ (Fig. 4A). If we think of the steady-state concentration of $\bullet\text{NO}$ as water in a bathtub, then the water level will be determined by the rate of water flow into the tub ($\bullet\text{NO}$ synthesis) relative to its rate of flow out of the drain ($\bullet\text{NO}$ metabolism). Since O_2 differentially controls the rates of both $\bullet\text{NO}$ synthesis and metabolism, the steady-state concentration of $\bullet\text{NO}$ will be a function of the relative differences in their respective rates. At both the macroscopic and microscopic level, O_2 gradients exist within the human body. Differences in O_2 concentrations can vary widely based on the organ, tissue, cell type, and even the intercellular location. What determines the microenvironmental O_2 concentration depends on the rate of O_2 delivery from the vasculature as well as the rate of mitochondrial O_2 consumption. Fig. 4B illustrates differences in the half-life of $\bullet\text{NO}$ calculated from average tissue O_2 concentrations. This model, however, does not take into account the effects of O_2 on $\bullet\text{NO}$ synthesis only the effect of O_2 on the rate of $\bullet\text{NO}$ disappearance [7]. It does, however, emphasize how over a range of physiologic O_2 concentrations the biological half-life of $\bullet\text{NO}$ can vary significantly.

Why is the half-life of $\bullet\text{NO}$ important? The broad answer is that it ultimately determines the magnitude of $\bullet\text{NO}$ signaling. To properly understand this, it is important to appreciate how $\bullet\text{NO}$ moves within the cellular environment, i.e., by diffusion. Nitric oxide is small and uncharged and is soluble in both hydrophobic and hydrophilic environments. For these reasons, it is freely diffusible and cell membranes pose no barrier to its movement. With

a relatively large diffusion coefficient ($\approx 3800 \mu\text{m}^2/\text{s}$), $\bullet\text{NO}$ can travel considerable distances in a short period of time depending on the local environment (≈ 1 cell length/25 ms) [41]. Once $\bullet\text{NO}$ is synthesized, it moves away from its point of origin by random diffusion (Fig. 5A). Movement of $\bullet\text{NO}$ in any direction is equally likely at any point in time, however, net movement of molecules in one direction will occur if there is a spatial concentration gradient. The movement will always occur from a region of higher $\bullet\text{NO}$ concentration, such as in an $\bullet\text{NO}$ -producing cell, to a region of lower $\bullet\text{NO}$ concentration, an adjacent cell (Fig. 5B) [42].

The distance $\bullet\text{NO}$ will diffuse is proportional to its concentration and half-life. This brings up the importance of O_2 . The O_2 concentration is an important determinant of the $\bullet\text{NO}$ concentration, which is a factor in determining how far $\bullet\text{NO}$ diffuses. At high O_2 concentrations, the half-life of $\bullet\text{NO}$ is short and so is the distance it can diffuse. The opposite is true for low O_2 concentrations (Fig. 6A). Just as the half-life of $\bullet\text{NO}$ can be calculated as a function of O_2 concentration, so can its diffusional distance

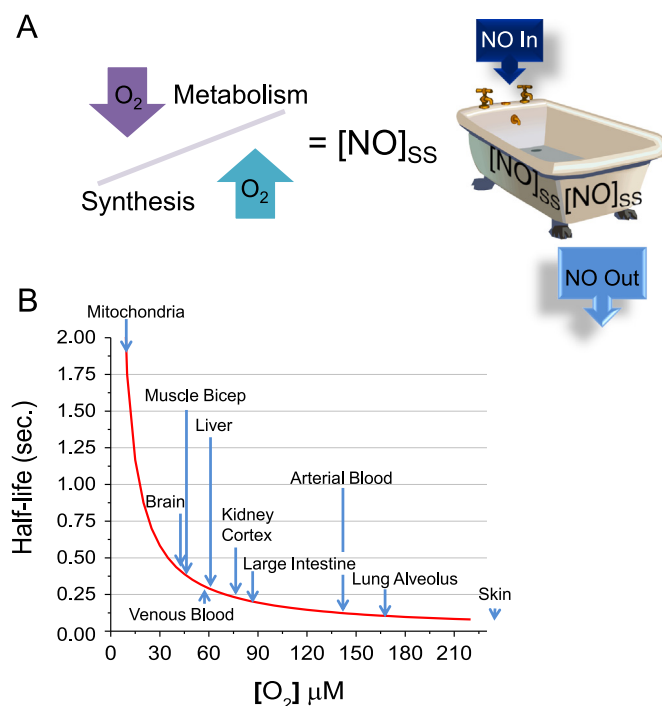


Fig. 4. Oxygen determines the concentration of $\bullet\text{NO}$ and its half-life. (A) The steady-state concentration of $\bullet\text{NO}$ is a function of its rate of synthesis relative to its rate of degradation. (B) Different tissues have different average O_2 concentrations. The half-life of $\bullet\text{NO}$ changes with respect to the O_2 concentration. Adapted from Ref. [7].

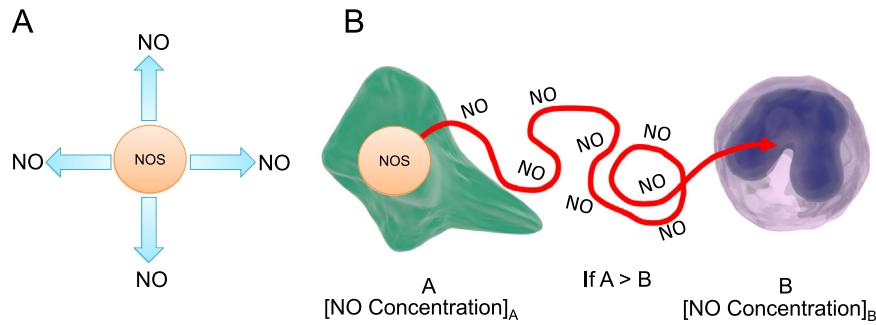


Fig. 5. Nitric oxide moves by random diffusion. (A) Once $\bullet\text{NO}$ is synthesized by NOS, it diffuses away in all directions. (B) Net movement of $\bullet\text{NO}$ occurs from an area of high concentration (a cell “A” where $\bullet\text{NO}$ is being made) to a region of lower concentration (“B” surrounding cells). The movement of $\bullet\text{NO}$ is random but down a concentration gradient.

(Fig. 6B). It can be seen that differences in physiologic O_2 concentrations dramatically affect the diffusional distance of $\bullet\text{NO}$ [7]. This has important implications for predicting the magnitude of $\bullet\text{NO}$ signaling under various O_2 concentrations. At a high pO_2 , for example, the half-life of $\bullet\text{NO}$ is short and so is the distance it will diffuse. At low pO_2 , $\bullet\text{NO}$ will travel significantly longer distances and potentially exert its influence many cell lengths away (Fig. 6B). What this means in terms of cell signaling is that the size of the diffusional sphere surrounding a $\bullet\text{NO}$ -producing cell will change in response to changes in the O_2 concentration. This will ultimately determine the amount and types of cellular targets by changing the number of neighboring cells being affected in a spatially heterogeneous tissue environment.

Although this discussion has focused on the impact of O_2 on $\bullet\text{NO}$ bioavailability it would not be complete without mentioning how $\bullet\text{NO}$ can affect local O_2 concentrations. One of the earlier discoveries about the biological functions of $\bullet\text{NO}$ was the observation that it could inhibit cellular respiration [43]. It was determined that $\bullet\text{NO}$ could regulate mitochondrial function and metabolism through its ability to interact at several sites in the respiratory chain. This is another example that emphasizes the importance of $\bullet\text{NO}$ /metal interactions. At Complex IV (cytochrome c oxidase) subnanomolar amounts of $\bullet\text{NO}$ competitively inhibit respiration. Here $\bullet\text{NO}$ binds to the ferrous heme-iron or oxidized copper, but not both simultaneously at the heme iron:copper binuclear center (a(3)/Cu(B)) [44]. At Complex I greater amounts of $\bullet\text{NO}$ were shown to be inhibitory but this was attributed to $\bullet\text{NO}$ -mediated oxidation or S-nitrosation of specific thiols [45]. The majority of total body O_2 utilization is via its four-electron

reduction to water at mitochondrial complex IV. Inhibition of mitochondrial respiration by $\bullet\text{NO}$, therefore, prevents O_2 consumption which extends the half-life of O_2 . This mechanism is thought to participate in O_2 homeostasis by increasing local O_2 delivery from the vascular and extending its delivery into tissues. Therefore, there is a reciprocal relationship between $\bullet\text{NO}$ and O_2 whereby each molecule plays a role in regulating the concentration of the other [46].

Oxygen and $\bullet\text{NO}$ signaling

As we have seen, the O_2 concentration is a key determinant of both the half-life of $\bullet\text{NO}$ as well as its diffusional distance. These parameters are important because they influence signaling responses to $\bullet\text{NO}$. Most cellular targets for $\bullet\text{NO}$ respond in a concentration and time dependent manner [47–49]. Numerous examples have demonstrated that the activation of specific proteins or the induction of signal transduction cascades respond to distinct concentration thresholds of $\bullet\text{NO}$. For example, soluble guanylyl cyclase (sGC) is one of the most important biological targets for $\bullet\text{NO}$. It is activated by very low steady-state amounts of $\bullet\text{NO}$ ($[0.5\text{--}5\text{ nM}]_{\text{SS}}$). ERK is another protein that becomes phosphorylated in the presence of $\bullet\text{NO}$ but it requires slightly higher amounts ($> [100\text{ nM}]_{\text{SS}}$). Another class of proteins, non-heme iron oxygenases, are also targets for $\bullet\text{NO}$. These include the HIF prolyl hydroxylases and the lysine histone demethylase (KDM) enzymes [28,50]. These are inhibited by intermediate concentrations of $\bullet\text{NO}$ ($[100\text{--}400\text{ nM}]_{\text{SS}}$). Proteins like p53 are regulated by $\bullet\text{NO}$ at the upper limits of physiologic concentrations. Phosphorylation of this

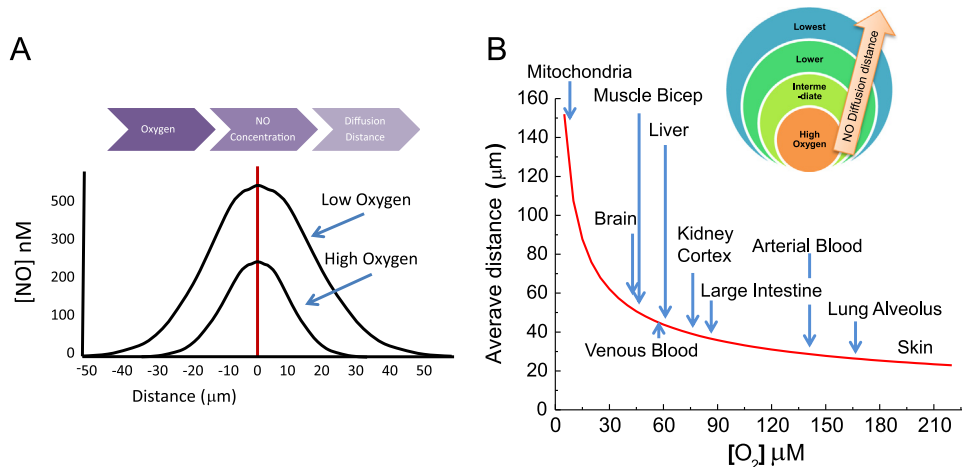


Fig. 6. Oxygen determines the concentration of $\bullet\text{NO}$ and its diffusional distance. (A) At a constant rate of $\bullet\text{NO}$ synthesis, the lower the O_2 concentration, the slower $\bullet\text{NO}$ is metabolized and the greater its steady-state concentration is. At high O_2 concentrations, the steady-state $\bullet\text{NO}$ concentration is lower. The $\bullet\text{NO}$ concentration will determine its half-life as well as its diffusional distance. (B) The O_2 concentration determines the half-life of $\bullet\text{NO}$, which influences how far it will diffuse.

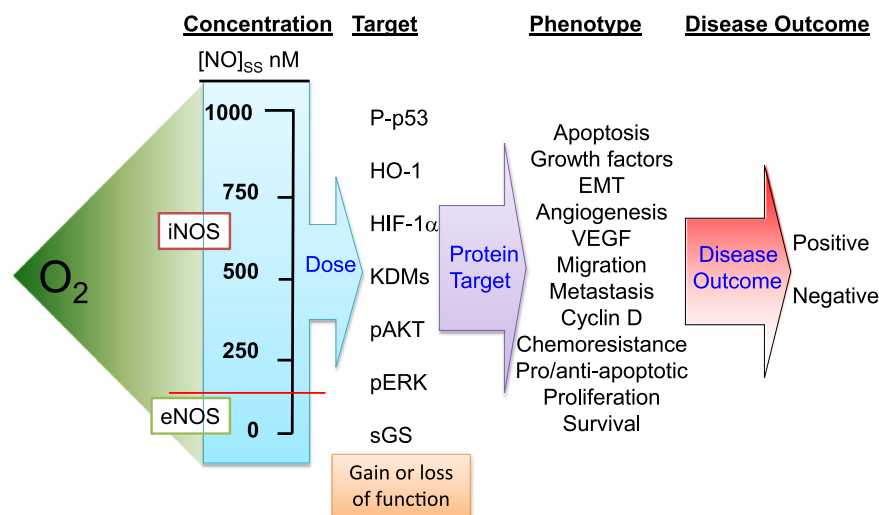


Fig. 7. Oxygen is an upstream driving force of cell phenotype by regulating the \bullet NO concentration. Many proteins differ in the threshold amounts of \bullet NO necessary for regulation. Since O_2 determines the concentration (dose) of \bullet NO, it will indirectly influence proteins regulated by \bullet NO in a concentration-dependent manner. \bullet NO-mediated changes in the protein profile will result in altered cellular phenotypes. This will ultimately have a positive or negative effect on disease outcome.

protein occurs in the presence of high concentrations of \bullet NO ($> [500 \text{ nM}]_{ss}$) (Fig. 7).

A long-standing challenge for researchers in the \bullet NO field has been to provide plausible mechanisms to explain opposite biological responses to \bullet NO under seemingly similar circumstances. There are many physiologic and pathological examples where \bullet NO has been shown to have a positive signaling effect and an equal number of situations where \bullet NO has been shown to have a negative effect in the same system. For example, \bullet NO has numerous purported roles in cancer etiology [49,51,52]. In some cases, \bullet NO is beneficial in that it decreases tumor size, is cytotoxic, induces apoptosis, and is an antioxidant. There are other reports, however, where \bullet NO is a negative prognostic indicator by virtue of its ability to increase angiogenesis, stimulate migration and invasion, and induce DNA damage. The question that is often raised, “is \bullet NO good or bad?”. The answer is “both”. To appreciate how \bullet NO can have such dissimilar and opposing responses, one has to understand the concentration and temporal-dependent effects of \bullet NO signaling. Differences in tissue O_2 concentrations will partially determine the steady-state concentration of \bullet NO. The concentration of \bullet NO will dictate what cellular targets it interacts with. The types and distributions of cellular targets will determine cell phenotype, ultimately leading to positive or negative effects on disease outcome (Fig. 7).

High O_2 concentrations will maximize \bullet NO production from NOS, however, it will also increase the rate of \bullet NO metabolism. Conversely, low O_2 concentrations will reduce enzymatic \bullet NO synthesis while also decreasing the rate of \bullet NO metabolism. Is there an optimal O_2 concentration where a balance between \bullet NO production and consumption is achieved to maximize steady-state \bullet NO concentrations? One study showed that the amount of cGMP produced by the activation of sGC in \bullet NO-producing cells was biphasic over a range of O_2 concentrations [34]. The maximal amount of cGMP was measured at an O_2 concentration of $\approx 8\%$. The interpretation was that this O_2 concentration resulted in the highest steady-state \bullet NO concentration and therefore maximal sGC activation.

Just as the concentration of \bullet NO is a key determinant of cell phenotype, so is the duration of \bullet NO exposure. Not only do the three NOS isoforms produce different amounts of \bullet NO, they can also differ in the duration of synthesis. This becomes important because some proteins are activated (or inhibited) immediately by \bullet NO whereas others require continuous and sometimes prolonged \bullet NO exposure [47]. Also, the effect of \bullet NO on certain proteins is

reversible such that they become inactivated (or reactivated) when the \bullet NO source is removed. Yet in other cases, prolonged activation is achieved long after \bullet NO synthesis has ceased. This indicates that some proteins are real-time \bullet NO sensors whose regulation parallels the duration of \bullet NO synthesis. In cases where the effects of \bullet NO result in prolonged protein regulation, even short bursts of \bullet NO exposure could have long-lasting phenotypic effects.

Spatial constraints are also contributing factors that influence phenotypic responses to \bullet NO. As described above, the concentration of \bullet NO will diminish at further distances from its point of synthesis (Fig. 8). Since most proteins respond to \bullet NO in a concentration-dependent manner, distance will ultimately dictate the population of targets being regulated by \bullet NO. Proteins, with different concentration threshold requirements for activation by \bullet NO, will be differentially regulated depending on their proximity to the \bullet NO source. For example, a protein that requires small amounts of \bullet NO like sGC may be fully activated at locations both near and far from the point of \bullet NO synthesis (Fig. 8, green protein). Whereas a different protein requiring intermediate amounts of \bullet NO (HIF-1 α) will be fully activated when it is close to the point of \bullet NO synthesis. At further distances where the steady-state \bullet NO concentration has diminished, this protein may not be activated at all (Fig. 8, orange protein). Proteins with the highest \bullet NO-concentration requirements such as p53 may not respond to \bullet NO at all regardless of whether they are proximal or distal to the \bullet NO source (Fig. 8, red protein). These examples demonstrate how spatial differences in the location of target proteins can dictate phenotypic outcomes based solely on the differences in their concentration thresholds for activation by \bullet NO.

Fig. 9 illustrates the complexities surrounding the interpretation of phenotypic data from cultured cells treated with \bullet NO when there are differences in both \bullet NO concentration and exposure time. In this simple experiment, cancer cells in culture were exposed to different concentrations of \bullet NO for 48 h and cell viability was measured at 3 separate time points (24, 30, and 48 h). If the experiment was terminated at 24 h, the interpretation would be that \bullet NO stimulates tumor cell growth (\bullet NO is bad). If endpoint measurements were not made until 30 h, the interpretation might be that \bullet NO has no effect on cell viability. However, if the cells were exposed to \bullet NO for a full 48 h before measuring cell viability, the logical conclusion would be that \bullet NO is cytotoxic to tumor cells (\bullet NO is good). Furthermore, if we only looked at low concentrations of \bullet NO (5 μ M DETA/NO), the outcome would be that \bullet NO has no effect on tumor cell viability from 0 to 48 h.

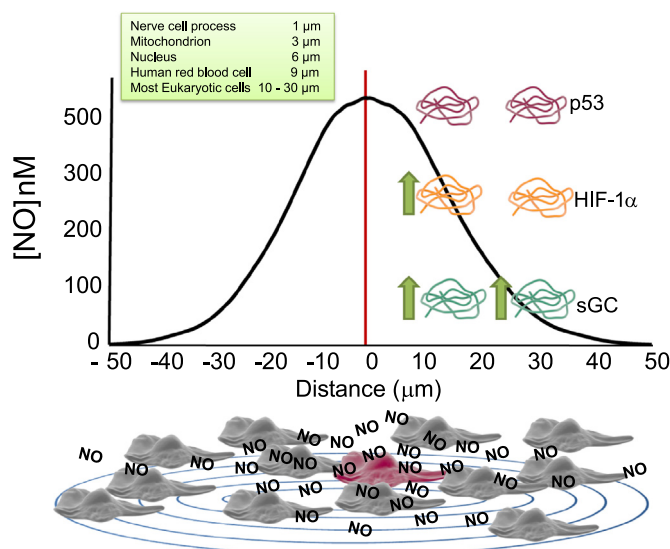


Fig. 8. The location of target proteins relative to the \bullet NO source will influence their degree of regulation. The concentration of \bullet NO diminishes at further distances from an \bullet NO-producing cell (red cell). Proteins requiring small amounts of \bullet NO will be regulated even at distances far away from the \bullet NO source (green protein). Proteins with higher concentration requirements for \bullet NO must be closer to the \bullet NO source (orange protein). Proteins requiring high steady-state amounts of \bullet NO may not be regulated at any location within the radius of \bullet NO diffusion if the threshold \bullet NO concentration requirement is not achieved (red protein). Green arrows indicate the regulation, activation, or deactivation by \bullet NO (relative sensitivity to \bullet NO green > orange > red protein).

However, if we only looked at higher concentrations of \bullet NO, the interpretation would be completely different at every time point.

Conclusion

The concepts that have been presented herein are undoubtedly an oversimplification of numerous interrelated complex biochemical and signaling mechanisms of \bullet NO. The aim of this short review, however, was to provide a framework for experimental

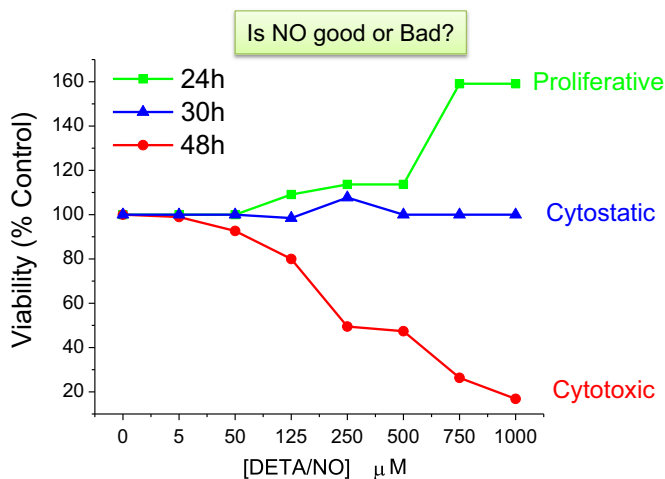


Fig. 9. Both the concentration and the duration of \bullet NO exposure determine the phenotypic outcome. MDA-MB-231 breast cancer cells were exposed to a range of physiologic \bullet NO concentrations using the \bullet NO-donor DETA/NO. DETA/NO has a long half-life ($t_{1/2}$ = 22 h), which enables prolonged and continuous \bullet NO exposure. Real-time measurements of cell proliferation was conducted using the xCELLigence[®] DP system (ACEA Biosciences, Inc.). MDA-MB-231 cells were seeded in E-plates containing 10% serum and allowed to adhere for 12 h before the addition of \bullet NO (5–1000 μM DETA/NO). Cell proliferation was measured continuously for 48 h following \bullet NO treatment.

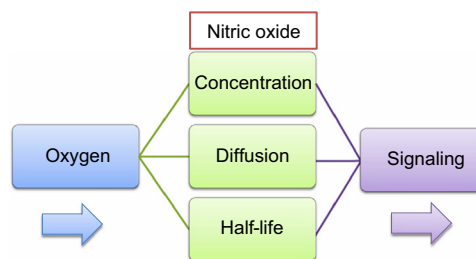


Fig. 10. Oxygen is a dominant and vital regulator of \bullet NO chemical biology.

design and data interpretation by taking into consideration these numerous confounding parameters. Although \bullet NO may appear to have contradictory effects under seemingly similar biological situations, a careful look at the environmental conditions will often reveal logical explanations for these differences. By understanding the importance of O_2 , target location, \bullet NO concentration, and exposure time on \bullet NO signaling responses, it becomes much easier to comprehend the sheer magnitude of potential outcomes (Fig. 10). Phenotypic responses attributed to \bullet NO suddenly do not appear to be random at all but rather predictable outcomes that are simply a reflection of the local redox environment. Nitric oxide, being a free radical, is unique among signaling molecules. It does not act indiscriminately and it follows the same rules of chemistry and physics as all molecules. With a better appreciation of its complex chemical biology, meaningful information will continue to be discovered about this fascinating and important molecule.

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