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Research Article

Oxygen dependence of nitric oxide-mediated signaling[☆]

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ARTICLE INFO

Article history:

Received 21 November 2012

Accepted 29 November 2012

Keywords:

Nitric oxide
Nitric oxide synthase
Oxygen
Autooxidation
sGC
p53

ABSTRACT

Nitric oxide (\bullet NO) is a biologically important short-lived free radical signaling molecule. Both the enzymatic synthesis and the predominant forms of cellular metabolism of \bullet NO are oxygen-dependent. For these reasons, changes in local oxygen concentrations can have a profound influence on steady-state \bullet NO concentrations. Many proteins are regulated by \bullet NO in a concentration-dependent manner, but their responses are elicited at different thresholds. Using soluble guanylyl cyclase (sGC) and p53 as model \bullet NO-sensitive proteins, we demonstrate that their concentration-dependent responses to \bullet NO are a function of the O_2 concentration. p53 requires relatively high steady-state \bullet NO concentrations (> 600 nM) to induce its phosphorylation (P-ser-15), whereas sGC responds to low \bullet NO concentrations (< 100 nM). At a constant rate of \bullet NO production (liberation from \bullet NO-donors), decreasing the O_2 concentration (1%) lowers the rate of \bullet NO metabolism. This raises steady-state \bullet NO concentrations and allows p53 activation at lower doses of the \bullet NO donor. Enzymatic \bullet NO production, however, requires O_2 as a substrate such that decreasing the O_2 concentration below the K_m for O_2 for nitric oxide synthase (NOS) will decrease the production of \bullet NO. We demonstrate that the amount of \bullet NO produced by RAW 264.7 macrophages is a function of the O_2 concentration. Differences in rates of \bullet NO production and \bullet NO metabolism result in differential sGC activation that is not linear with respect to O_2 . There is an optimal O_2 concentration (≈ 5 –8%) where a balance between the synthesis and metabolism of \bullet NO is established such that both the \bullet NO concentration and sGC activation are maximal.

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1. Introduction

Nitric oxide (nitrogen monoxide, \bullet NO) is a short-lived signaling molecule involved in regulating numerous physiological and pathological functions. Nitric oxide is synthesized by nitric oxide synthase (NOS) of which there are three main isoforms (iNOS, eNOS, nNOS). The substrates for this enzyme are arginine and oxygen (O_2). Also, NADPH, FMN, BH_4 , and FAD are required as cofactors. Not surprisingly, changes in the availability of any one of

these substrates or cofactors can affect the rate of \bullet NO production. When there is an abundance of cofactors, the rate of \bullet NO synthesis will be a function of both the arginine and O_2 concentrations. Arginine availability can vary based on cellular uptake and competing consumptive pathways (i.e. arginase and the urea cycle). The O_2 availability in tissues is a function of its rate of delivery from the vasculature and the rate at which it is consumed locally via mitochondrial respiration. The K_m 's for arginine and O_2 are different for each NOS isoform, thus changes in substrate concentrations will alter the \bullet NO output in an isoform-dependent manner. The K_m 's for O_2 for eNOS, iNOS and nNOS are 23 μ M, 135 μ M and 350 μ M respectively. These differences indicate that for a given O_2 concentration the rate of \bullet NO synthesis from nNOS will be dramatically affected while production from eNOS will remain comparatively constant [1]. Since many phenotypic responses to \bullet NO occur in a concentration-dependent manner [2], the cellular response to \bullet NO produced in a relatively hypoxic environment could be substantially different than in tissues which are well oxygenated even if they express an equivalent amount and type of NOS.

The steady-state concentration of \bullet NO ($[\bullet NO_{ss}]$) is determined by the balance between its rate of production and its rate of disappearance. Although several studies have attempted to elucidate the mechanism(s) of cellular \bullet NO metabolism, to date the dominant pathway(s) for this process remains undetermined.

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Abbreviations: BH_4 , tetrahydrobiopterin; cGMP, cyclic guanosine monophosphate; DETA/NO, (Z)-1-[N-(2-aminoethyl)-N-(2-ammonioethyl)amino]diazene-1-ium-1,2-diolate; eNOS, endothelial nitric oxide synthase; FAD, flavin adenine dinucleotide; FMN, flavin mononucleotide; iNOS, inducible nitric oxide synthase; K_m , Michaelis constant; LPS, lipopolysaccharide; NADPH, nicotinamide adenine dinucleotide phosphate, reduced; nNOS, neuronal nitric oxide synthase; NO_3^- , nitrate; NO_2^- , nitrite; \bullet NO, nitric oxide; ODQ, 1H-[1,2,4]Oxadiazolo[4,3-a]quinoxalin-1-one; O_2 , oxygen; P-Ser-15, phospho-serine 15; sGC, soluble guanylyl cyclase; Sper/NO, (Z)-1-[N-[3-aminopropyl]-N-[4-(3-aminopropylammonio)butyl]-amino]diazene-1-ium-1,2-diolate

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We and others have shown, however, that metabolism of •NO by non-erythroid cells is an O₂-dependant process. Kinetic studies indicated that, although •NO metabolism requires O₂, it is not a direct reaction of •NO with O₂ (autooxidation) [3]. Herein, we demonstrate that the local O₂ concentration determines [\bullet NO]_{ss} which differentially regulates signaling responses to •NO.

2. Materials and methods

2.1. Chemicals

(Z)-1-[N-(2-aminoethyl)-N-(2-ammonioethyl)amino]diazene-1,2-diolate (DETA/NO) and (Z)-1-[N-[3-aminopropyl]-N-[4-(3-aminopropylammonio)butyl]-amino]diazene-1,2-diolate (Sper/NO) were generous gifts from Dr. Joseph Hrabie (National Institutes of Health, NCI). 1H-[1,2,4]Oxadiazolo [4,3-a]quinoxalin-1-one (ODQ), lipopolysaccharides (LPS) and aminoguanidine hydrochloride were purchased from Sigma. All cell culture reagents were purchased from Invitrogen.

2.2. Cell culture

MCF-7 human breast carcinoma cells and RAW 264.7 murine macrophages were cultured in DMEM supplemented with 10% fetal bovine serum and 1% penicillin/streptomycin. All cells were grown to 85% confluence and MCF-7 cells were serum-starved overnight prior to treatment. To achieve a hypoxic environment (< 21% O₂), cells were grown in a hypoxic chamber glove-box (Coy Scientific) at 37 °C and 5% CO₂. Oxygen concentration was constantly monitored and regulated by purging with nitrogen gas. RAW 264.7 cells were activated using LPS (1 μg/ml) to induce iNOS expression. For coculture experiments, activated RAW 264.7 cells were trypsinized, counted and added at indicated densities to MCF-7 cells in serum free media in the presence or absence of aminoguanidine (1 mM), an iNOS inhibitor. To inhibit sGC, MCF-7 cells were pretreated with 10 μM ODQ for 30 minutes prior to addition of RAW 264.7 cells.

2.3. Western Blotting

Whole cell lysates were collected using CellLytic™ M Cell Lysis Reagent (Sigma), supplemented with 1% protease inhibitor cocktail (Calbiochem) and 1 mM PMSF (Sigma). Proteins were transferred to polyvinylidene difluoride (PVDF) membranes using the iBlot transfer system (Invitrogen). The membranes were blocked with 5% nonfat dry milk in Tris-glycine-SDS containing 0.1% Tween 20 for 1 h at room temperature. They were then incubated overnight at 4 °C with suitable primary antibodies: p53 P-Ser-15, p53, β-actin (Cell Signaling Technology, Beverly, MA, USA) or iNOS (Santacruz Biotechnology, Santa Cruz, CA, USA). After incubation with horseradish peroxidase-conjugated secondary anti-rabbit antibody (Cell Signaling Technology, Beverly, MA, USA) for 1 h, the blots were analyzed in a Fluor Chem HD2 imager (Alpha Innotech) using SuperSignal West Femto Maximal Sensitivity Substrate (Thermo Scientific).

2.4. cGMP Measurements

MCF-7 cells were grown to 50% confluency in 96-well microtiter plates. Cells were serum-starved overnight and treated with various concentrations of Sper/NO or cocultured with activated RAW 264.7 macrophages at different densities. After the removal of •NO source, MCF-7 cells were lysed at the indicated time points, and cell extracts were assayed for cGMP by enzyme immunoassay (Amersham Biosciences).

2.5. Cellular •NO metabolism

Rates of disappearance of •NO were measured as described previously [4]. RAW 264.7 cells were trypsinized and put into a reaction chamber at 6 × 10⁶ cells/ml of serum-free media. Cell suspensions were constantly stirred in a sealed, water-jacketed, temperature-controlled (37 °C) chamber. 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, response time of < 0.2 s, sensitivity of 25 nM; Innovative Instruments, Tampa, FL, USA).

2.6. Real-time •NO measurements

MCF-7 cells were grown in 10 cm culture dishes and an •NO-selective electrode (amiNO-700, Innovative Instruments) connected to an Apollo 4000 free radical analyzer (World Precision Instruments) was positioned ~1 mm above, and perpendicular to the cellular monolayer. After equilibration for 2 h at 37 °C, •NO donor compounds were added and steady-state •NO concentrations were measured over time.

2.7. NO₂⁻/NO₃⁻ measurements

Accumulation of •NO metabolites in the media was measured by chemiluminescence with a Sievers nitric oxide analyzer 280i. 25 μl aliquots of media were injected into a reaction chamber containing vanadium chloride in HCl as described previously [4].

2.8. Statistics

Significance was determined using either t-test or one way ANOVA with Bonferoni post-hoc analysis. Error bars indicate SEM.

3. Results

3.1. Proteins respond to •NO in a concentration-dependent manner

We and others have shown that the tumor suppressor protein p53 is phosphorylated at the serine 15 residue (p53 P-Ser-15) in response to relatively high steady-state concentrations of •NO (> 600 nM [\bullet NO]_{ss}) [5]. Conversely, relatively low concentrations of •NO (< 100 nM [\bullet NO]_{ss}) activate soluble guanylyl cyclase (sGC) [6]. The differing sensitivities of these proteins to •NO makes them, in effect, biological •NO dosimeters. We treated MCF-7 cells with increasing concentrations of the •NO-donor Sper/NO and measured changes in p53 P-Ser-15 and cGMP production. When the Sper/NO concentration was increased, the [\bullet NO]_{ss} concentration rose in a corresponding manner (Fig. 1A). As predicted, p53 was phosphorylated at the highest concentration of Sper/NO which produced [\bullet NO]_{ss} concentrations approaching 1 μM (Fig. 1B). Next, cGMP was measured in these cells as an indicator of sGC activation (Fig. 1C). Very low •NO concentrations were sufficient for maximal sGC activation (≈ 5 μM Sper/NO, < 50 nM [\bullet NO]_{ss}).

3.2. The impact of O₂ concentration on [\bullet NO]_{ss}

We have previously shown that, for a given cell type and density, the rate of cellular •NO metabolism displays first-order kinetics

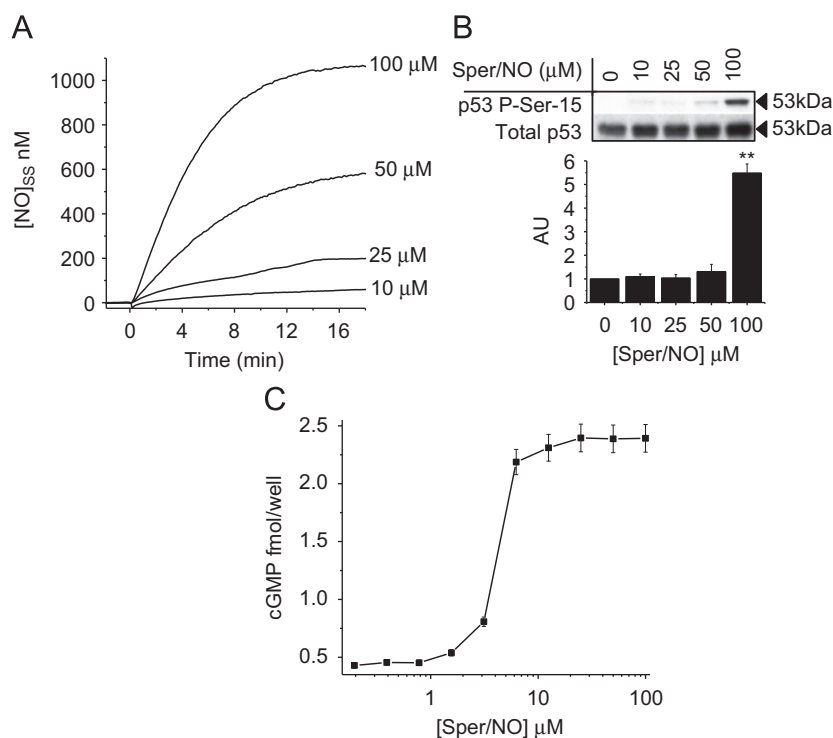


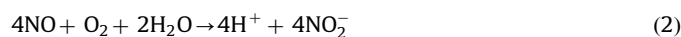
Fig. 1. Signaling effects of •NO are a function of its steady-state concentration. MCF-7 cells were treated with increasing concentrations of Sper/NO (0–100 μM) at 21% O₂. (A) Measurement of [•NO]_{ss} in media (•NO selective electrode, ~1 mm above monolayer). (B) Immunoblot demonstrating p53 P-Ser-15 accumulation (*t*=2 h). (C) Measurement of cGMP production as an indicator of sGC activation (*t*=30 min). *n*=3, ** indicates *p* < 0.01 with respect to untreated controls which are arbitrarily set to 1.0.

which is a function of the both the •NO and O₂ concentrations (Eq. 1) [3].

$$-\frac{d[\text{NO}]}{dt} = k_{\text{obs}}[\text{O}_2][\text{NO}][\text{Cells}] \quad (1)$$

Using an •NO-selective electrode, we measured differences in •NO metabolism after either bolus addition or continuous exposure to •NO at 1% and 21% O₂. For the purposes of this study, metabolism is defined as the disappearance of •NO in the presence of cells. Fig. 2A demonstrates that when cells in suspension received a 1000 nM •NO bolus, •NO metabolism was slower at 1% than at 21% O₂ (*t*_{1/2}=260s vs. 45 s). We also measured [•NO]_{ss} in the culture media of cells treated with the •NO-donor DETA/NO at either 1% or 21% O₂. The rate of •NO liberation from diazeniumdiolates such as DETA/NO is a function of pH and temperature and it is not affected by differences in O₂ concentration [7]. For an equivalent concentration of DETA/NO (500 μM), the [•NO]_{ss} was greater at 1% O₂ than it was at 21% O₂ (Fig. 2B). These observed differences in [•NO]_{ss} at 1% vs. 21% O₂ can be partially attributed to differences in the rates of •NO metabolism which decreases as the O₂ concentration is lowered.

For cell culture experiments, the volume of media far exceeds the volume of cells. Therefore, in addition to cellular metabolism, the disappearance of •NO can result from other consumptive processes such chemical reactions in the media. The predominant reaction of •NO in the media is with dissolved O₂ through the autooxidation reaction [8]. This reaction is first-order with respect to the O₂ concentration (Eq. 3) and, therefore, its rate will vary based on differences in O₂. We calculated the contribution of this reaction to the observed differences in the rates of •NO disappearance at both O₂ concentrations. Autooxidation occurs with the following overall stoichiometry [8]:



The rate of disappearance of •NO through this reaction is given by

$$-\frac{d[\text{NO}]}{dt} = 4k[\text{NO}]^2[\text{O}_2] \quad (3)$$

where $k = 1.8\text{--}6.6 \times 10^6 \text{ M}^{-2}\text{s}^{-1}$.

Calculations of •NO disappearance via the autooxidation reaction at 1% and 21% O₂ are shown in Fig. 2C and D. These illustrate the well-known fact that the half-life of •NO is inversely correlated to the •NO concentration and the O₂ concentration. At 1% O₂, the half-life of •NO is on the order of hours to days at 1000 nM •NO (Fig. 2C). At 21% O₂, the half-life of •NO is reduced to minutes or hours (Fig. 2D). These calculations reveal that, at 1% O₂, 1000 nM •NO disappears ≈ 58 times slower via the autooxidation reaction than in the presence of cells (Fig. 2A) and ≈ 15 times slower at 21% O₂. The lower the •NO concentration, however, the less autooxidation contributes to the observed differences in the rates of •NO disappearance in the presence of cells because autooxidation is second order in •NO. This is exemplified in Fig. 2E, which demonstrates that the disappearance of 200 nM •NO is negligible in the absence of cells.

In addition to cellular consumption and autooxidation, there are other means by which •NO can disappear in cell culture experiments. These include volatilization from the culture flask and reactions with other components in the media [7]. In comparison to the reaction of •NO with cells or O₂, the disappearance of •NO via other mechanisms should be constant at any O₂ concentration. As a result, these other consumptive pathways should not be a confounding factor when observing differences in the rates of •NO disappearance at different O₂ concentrations. Our data indicate that the dominant mechanism of •NO loss is via cellular metabolism, which prevails over other processes including autooxidation.

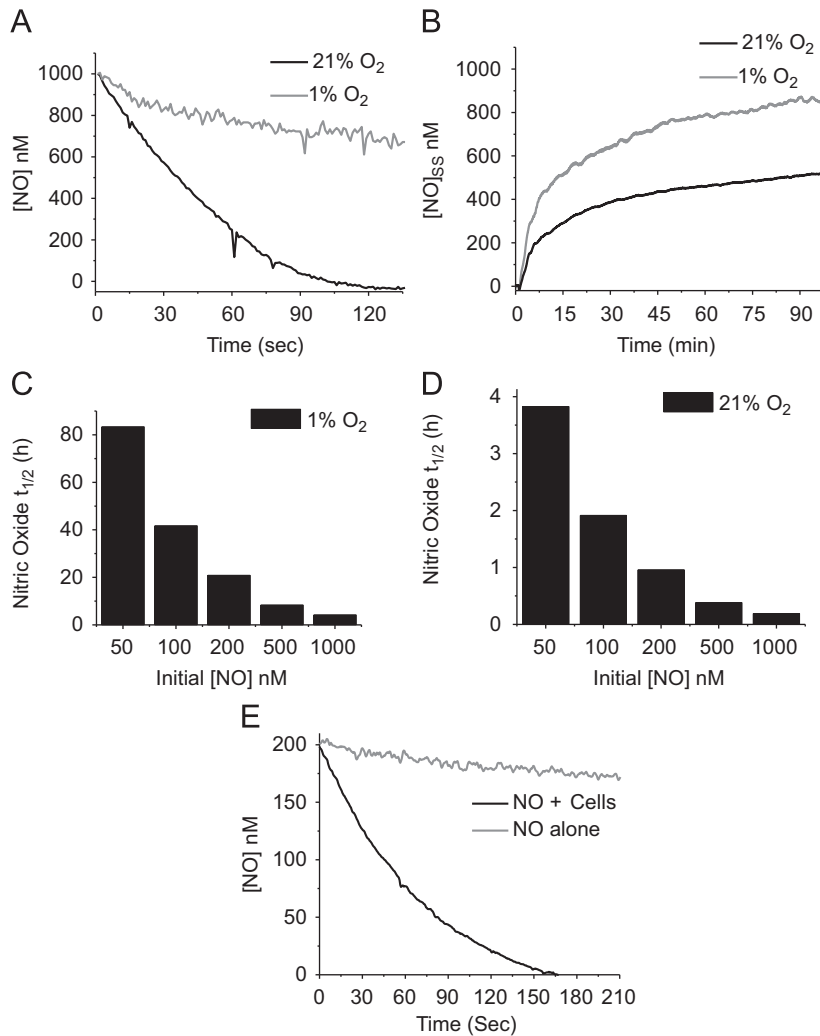


Fig. 2. Oxygen concentration influences •NO consumptive process. (A) MCF-7 cells were placed in suspension and stirred in a reaction vessel at 37 °C with an •NO-selective electrode. •NO metabolism (disappearance) was measured after the additions of bolus amounts of an •NO-saturated stock solution at 21% O₂ or 1% O₂. (B) Measurement of [•NO]_{ss} in the media of cells treated with 500 μM of the •NO-donor DETA/NO (•NO-selective electrode, ~1 mm above monolayer) at 21% O₂ or 1% O₂. (C,D) Calculated half-life of •NO via autooxidation from 50–1000 nM starting concentrations at 1% and 21% O₂. (E) Disappearance of 200 nM •NO at 21% O₂ in the presence or absence of cells.

3.3. Oxygen modulates •NO signaling processes

At a constant rate of •NO production (i.e. from •NO-donor compounds), the [•NO]_{ss} is predominantly a function of the O₂-dependant rate at which it is metabolized by cells (Fig. 2B). We set out to determine if the effect of O₂ on [•NO]_{ss} would differentially affect •NO-regulated signaling. To test this, we measured the accumulation of p53 Ser-15 phosphorylation in MCF-7 cells treated with increasing concentrations of the •NO-donor Sper/NO at either 1% O₂ or 21% O₂ for 2 h. Fig. 3 demonstrates that, at 1% O₂, the concentration of Sper/NO necessary to induce p53 P-Ser-15 is only one fourth of the amount required at 21% O₂. For a given Sper/NO concentration, the rate of •NO production is the same at both O₂ concentrations, yet the [•NO]_{ss} is greater at 1% O₂ due to differences in the rates of cellular •NO metabolism. Since p53 phosphorylation requires a specific threshold concentration of •NO (Fig. 1B), this implies that the [•NO]_{ss} from 25 μM Sper/NO at 1% O₂ is roughly equivalent to what is achieved at 21% O₂ from 100 μM Sper/NO. This demonstrates that, at a constant rate of •NO production, it is the O₂ concentration that ultimately determines the extent of •NO signaling.

3.4. Oxygen determines the rate of enzymatic •NO synthesis

Unlike the liberation of •NO from synthetic donors, the biological production of •NO is enzymatic and requires molecular O₂ as a substrate. For a given NOS isoform, changes in O₂ concentrations will affect enzymatic •NO production. To demonstrate this, we cultured RAW 264.7 macrophages over a range of O₂ concentrations and stimulated them with LPS to induce iNOS expression and •NO production. The total amount of enzymatic •NO synthesis was determined by measuring the accumulation of NO₃⁻/NO₂⁻ in the media over time. In Fig. 4A it can be seen that, as expected, there was a temporal and linear increase in •NO production as the O₂ concentration rose from 1–8%. More importantly, however, above 8% O₂ no further increase in the rate of •NO production was observed. This indicates that O₂ is no longer a limiting factor for •NO synthesis at concentrations greater than 8%.

Hypoxia is known to affect the magnitude of iNOS expression which could potentially influence the overall amount of •NO synthesis [4]. We examined this possibility by culturing RAW 264.7 cells over a range of O₂ concentrations and measuring the levels of iNOS expression subsequent to LPS stimulation. Fig. 4B

demonstrates that decreasing the O₂ concentration increased the magnitude of iNOS expression. Next, we set out to compare the effects of enzyme concentration vs. O₂ concentration on the amount of •NO produced. Activated RAW 264.7 cells were cultured at 1% and 21% O₂ for 16 or 24 h and the amount of NO₂⁻/NO₃⁻ was measured (Fig. 4C, columns 1,2,4,5). At both time points, the amount of •NO synthesis was considerably higher at 21% O₂ than at 1% O₂ as seen in Fig. 4A.

To understand the effects of differences in enzyme expression on •NO synthesis, we cultured RAW 264.7 cells for 16 h at 1% O₂. This resulted in a substantial increase in cellular iNOS protein expression (Fig. 4B). After 16 h these cells were moved to 21% O₂ for an additional 8 h (24 h total) and NO₂⁻/NO₃⁻ was measured (Fig. 4C, column 3). This resulted in a dramatic increase in •NO production at 21% O₂ that we attributed to the hypoxia-mediated increase in iNOS expression (Fig. 4C, compare columns 1, 2 and 3).

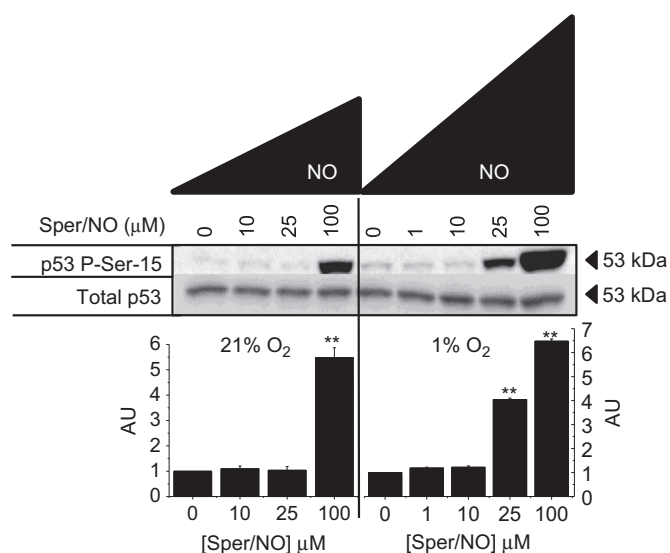


Fig. 3. Low O₂ decreases the dose of •NO-donor compound required for p53 activation. MCF-7 cells were treated with increasing concentrations of Sper/NO (0–100 μM) at 21% O₂ or 1% O₂ for 2 h. Immunoblot demonstrating accumulation of p53-Ser-15 phosphorylation. *n*=3, **indicates *p*<0.01 with respect to untreated controls which are arbitrarily set to 1.0.

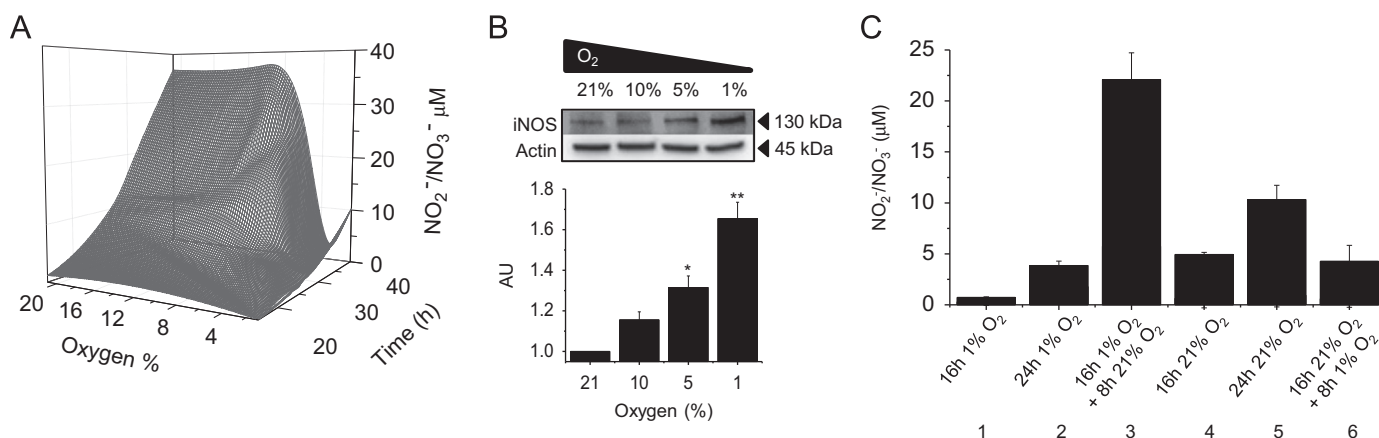


Fig. 4. The rate of •NO synthesis is a function of both the NOS and O₂ concentrations. RAW 264.7 cells were cultured at different O₂ concentrations and stimulated with LPS (*t*=0 h). The rate of •NO synthesis was measured by quantifying NO₂⁻/NO₃⁻ accumulation in the media by chemiluminescence. (A) Contour plot demonstrating the temporal relationship between changing O₂ concentrations and rates of •NO synthesis. (B) Immunoblot demonstrating increased iNOS expression as a function of decreasing O₂ concentrations (*t*=10 h). (C) (columns 1, 2, 4, & 5) Cells were grown at 21% O₂ or 1% O₂ and NO₂⁻/NO₃⁻ was measured at 16 h and 24 h. (columns 3 & 6) Cells were grown at 21% O₂ or 1% O₂. After 16 h the cells at 21% O₂ were moved to 1% O₂ and the cells at 1% O₂ were moved to 21% O₂. Both sets of cells were incubated for an additional eight hours (24 h total) and NO₂⁻/NO₃⁻ was measured. Values given at 24 h are the addition of the 16 h and additional 8 h reads. *n*=3, * indicates 0.01 < *p* < 0.05; **indicates *p* < 0.01 with respect to untreated controls which are arbitrarily set to 1.0.

This experiment was repeated, but the order of oxygen exposure was reversed; 21% O₂ for 16 h followed by 8 h at 1% O₂ (Fig. 4C, column 6). Under these conditions, total •NO synthesis was substantially diminished (Fig. 4C, compare columns 3,6). Together these results indicate that at 21% O₂ enzyme concentration is the major limiting factor for •NO synthesis whereas at 1% O₂, oxygen is the major limiting factor. Therefore, the oxygen concentration can influence •NO production by changing the amount of iNOS expressed or as a limiting substrate for this enzyme.

3.5. Oxygen regulates the magnitude of •NO signaling

Having determined that changes in O₂ concentration could influence both the rate of enzymatic •NO production and •NO metabolism, we went on to examine the impact of these processes on •NO signaling. For these experiments we developed a co-culture model in which MCF-7 cells were cultured in the presence of •NO-producing RAW 264.7 cells. After isolation of the MCF-7 cells, sGC activation was measured by monitoring the cGMP production. Fig. 5A demonstrates that sGC in the target MCF-7 cells was activated in response to RAW 264.7-derived •NO. Cyclic GMP production in the MCF-7 cells was not detected when cells were treated with ODQ, co-cultured with non-activated RAW 264.7 cells or activated RAW cells treated with the iNOS inhibitor aminoguanidine (Fig. 5A). Next, we altered oxygen levels by culturing the cells at different O₂ concentrations (1–21%) and modulated the •NO concentration by changing the ratio of RAW 264.7/MCF-7 cells. We then repeated the co-culture experiment over a range of O₂ and •NO concentrations. Fig. 5B demonstrates that sGC activation was dependent on both the O₂ concentration and the number of •NO-producing cells. Interestingly, the optimal O₂ concentration for maximal cGMP production was at ≈ 5–8% O₂. This suggests that at this O₂ concentration •NO production and •NO consumptive processes are optimized to maximize steady-state •NO concentration and sGC activation.

4. Discussion

Nitric oxide is a small, uncharged, freely diffusible gas that under biological conditions reacts with other free radicals (including oxygen) and metals. Its unique biochemical properties allow it

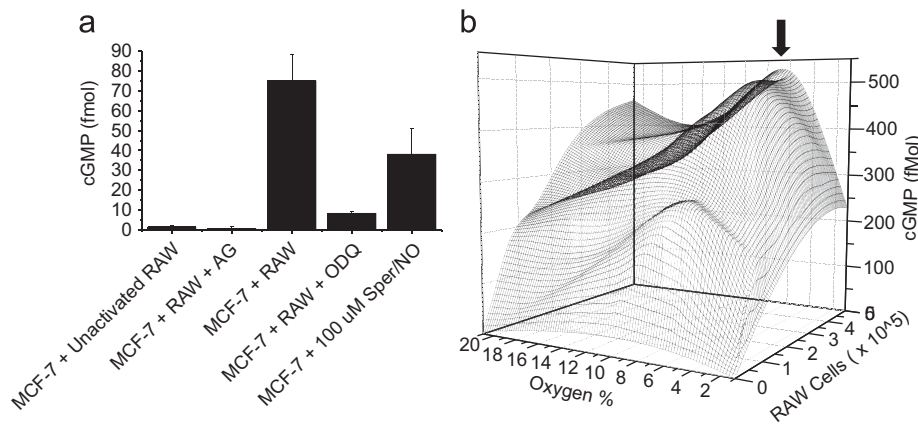


Fig. 5. Signaling effects of endogenously produced \bullet NO are affected by O_2 concentration. (A) Activated RAW 264.7 cells were cocultured in serum free media with MCF-7 cells at 21% O_2 . Coculture with unactivated RAW cells; activated RAW cells \pm the iNOS inhibitor aminoguanidine (AG); or the sGC inhibitor ODQ. After 30 min the RAW 264.7 cells were removed and cGMP production was measured in the MCF-7 cells by ELISA. (B) Increasing amounts of activated RAW 264.7 cells were cocultured in serum free media with MCF-7 cells over a range of O_2 concentrations (0.5–21%). After 30 min the RAW 264.7 cells were removed and cGMP production was measured in the MCF-7 cells by ELISA. Contour plot demonstrating the relationship between cGMP production, O_2 concentration, and different densities of \bullet NO-producing RAW 264.7 cells.

regulate, and sometimes disrupt, numerous aspects of cell physiology. In addition to \bullet NO's ability to activate signal transduction cascades, it can also directly inhibit respiration and sequester cellular iron. Unfortunately, the same properties that confer this diversity of function also make it extremely challenging to predict its exact $[\bullet\text{NO}]_{ss}$ in cells and tissues. It is well-known that both \bullet NO synthesis and \bullet NO consumption are O_2 -dependent, and thus the oxygen concentration will influence the rates of both of these processes. We investigated the overall ability of O_2 to modulate \bullet NO signaling events. If the rate of \bullet NO synthesis is greater than the rate of cellular \bullet NO consumption, $[\bullet\text{NO}]_{ss}$ concentrations will increase. Conversely, if the rate of \bullet NO consumptive processes predominate over its rate of synthesis, $[\bullet\text{NO}]_{ss}$ levels will decrease. Oxygen largely dictates the $[\bullet\text{NO}]_{ss}$ and ultimately the magnitude of \bullet NO signaling. We and others have demonstrated previously that the $[\bullet\text{NO}]_{ss}$ is a significant determinant of cell signaling effects; the various \bullet NO-mediated signaling pathways require different threshold concentrations of \bullet NO for activation. Although many of these concepts have been described previously, our results link \bullet NO concentration, O_2 concentration, \bullet NO synthesis, and \bullet NO metabolism into a unified theory.

Many *in vivo* studies use the expression levels of NOS as justification for \bullet NO-attributable responses [9]. It is often not appreciated that the overall amount of synthesized \bullet NO cannot easily be inferred solely by examining the levels of NOS protein. Our results demonstrate that both the O_2 concentration and the amount of NOS protein are equally important determinants of the amount of \bullet NO synthesized. The relative contribution of both of these parameters varies with the O_2 concentration. At high O_2 , \bullet NO synthesis is dependent on the amount of enzyme present, while at low O_2 , oxygen itself becomes the limiting factor.

We used a \bullet NO-selective electrode to measure the $[\bullet\text{NO}]_{ss}$ in the \bullet NO-donor treated media of cells grown in monolayer. Since \bullet NO is freely diffusible into cells, its steady-state concentration in the media is in equilibrium with its concentration in the cells. This allows for a precise measurement of $[\bullet\text{NO}]_{ss}$ and the identification of signaling cascades that correspond to specific \bullet NO concentrations. As we have previously reported, \bullet NO-mediated cellular signaling is stratified with numerous signaling pathways being activated by differential $[\bullet\text{NO}]_{ss}$ [2]. Our current study builds on these findings by demonstrating how \bullet NO signaling can be significantly modified by the O_2 concentration. In many ways these results are expected given the ability of O_2 to modulate \bullet NO concentrations. However, our results demonstrate

that changes in \bullet NO signaling do not always change linearly with changing O_2 concentrations.

When the rate of \bullet NO production is constant, its steady-state concentration is predominantly a function of its O_2 -dependent rate of metabolism. We demonstrate that, at lower O_2 concentrations, \bullet NO metabolism is considerably slower, and thus $[\bullet\text{NO}]_{ss}$ is higher. This inverse relationship has a direct effect on \bullet NO signaling. Activation of p53 Ser-15 phosphorylation at 1% O_2 , for example, is possible with one-fourth the dose of Sper/NO necessary to achieve the same effect at 21% O_2 . This indicates that, for a given rate of \bullet NO production (synthesis), lowering the O_2 concentration will enable activation of signaling pathways that are not obtainable at higher O_2 concentrations.

In addition to \bullet NO consumptive pathways, we show that O_2 concentrations influence $[\bullet\text{NO}]_{ss}$ by affecting its rate of enzymatic synthesis. Oxygen is a necessary substrate for NOS, and the oxygen gradient itself regulates iNOS expression [4]. Our data reveals that, in \bullet NO-producing RAW 264.7 cells, the rate of \bullet NO synthesis increases in a linear fashion from 1–8% O_2 . Above this amount of O_2 , further increases in the rate of \bullet NO synthesis were not observed. When sGC activation by \bullet NO was examined under changing O_2 concentrations, we noted maximal activation at 5–8% O_2 . At O_2 concentrations higher or lower than these amounts, there was a decrease in sGC activation. We suggest that the balance between O_2 availability and enzyme expression at 5–8% O_2 is optimized such that the rate of \bullet NO production versus its rate of metabolism synergize to achieve maximal $[\bullet\text{NO}]_{ss}$. At higher O_2 concentrations, \bullet NO production is maximal but so is its metabolism. At lower O_2 concentrations, \bullet NO metabolism is slowest but so is its enzymatic synthesis.

5. Conclusions

These findings highlight the importance and dependence of O_2 on biological responses to \bullet NO. The principals discussed in this study are important to consider when assigning a functional role to \bullet NO in the context of numerous normal and disease conditions where oxygen gradients exist. The tissue microenvironment will drive $[\bullet\text{NO}]_{ss}$, thereby modifying which signal transduction cascades are triggered to ultimately influence cellular phenotype. Some of the disparities seen in \bullet NO signaling are thus the result of O_2 mediated changes in $[\bullet\text{NO}]_{ss}$. These are important factors to consider when trying to elucidate the biochemical responses to

•NO synthesis and also the effects of therapeutic •NO donors especially in the context of changing oxygen environments.

Acknowledgments

Research reported in this publication was supported by the National Institute of General Medical Sciences of the National Institutes of Health under award number R01GM085232. The content is solely the responsibility of the authors and does not necessarily represent the official views of the National Institutes of Health. The authors also acknowledge ongoing support from the UIC Cancer Center.

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