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## Research Article

## Oxygen dependence of nitric oxide-mediated signaling



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## 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 ( $\approx 5$ –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  $\mu$ M, 135  $\mu$ M and 350  $\mu$ 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.

**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; ODO, 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 O<sub>2</sub>-dependant process. Kinetic studies indicated that, although •NO metabolism requires O<sub>2</sub>, it is not a direct reaction of •NO with O<sub>2</sub> (autooxidation) [3]. Herein, we demonstrate that the local O<sub>2</sub> concentration determines [•NO]<sub>ss</sub> 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% O<sub>2</sub>), cells were grown in a hypoxic chamber glove-box (Coy Scientific) at 37 °C and 5% CO<sub>2</sub>. 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 × 10<sup>6</sup> 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 O<sub>2</sub> 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. NO<sub>2</sub><sup>-</sup>/NO<sub>3</sub><sup>-</sup> 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.

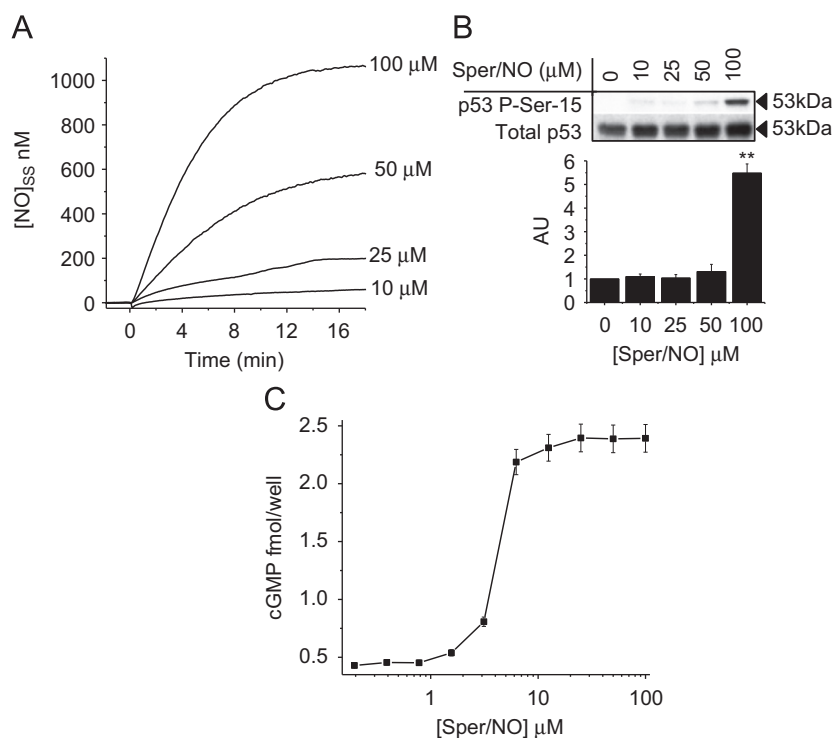
## 3. Results

### 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 [•NO]<sub>ss</sub>) [5]. Conversely, relatively low concentrations of •NO (< 100 nM [•NO]<sub>ss</sub>) 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 [•NO]<sub>ss</sub> concentration rose in a corresponding manner (Fig. 1A). As predicted, p53 was phosphorylated at the highest concentration of Sper/NO which produced [•NO]<sub>ss</sub> 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 [•NO]<sub>ss</sub>).

### 3.2. The impact of O<sub>2</sub> concentration on [•NO]<sub>ss</sub>

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



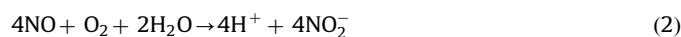
**Fig. 1.** Signaling effects of •NO are a function of its steady-state concentration. MCF-7 cells were treated with increasing concentrations of Sper/NO (0–100 μM) at 21% O<sub>2</sub>. (A) Measurement of [•NO]<sub>ss</sub> in media (•NO selective electrode, ~1 mm above monolayer). (B) Immunoblot demonstrating 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 O<sub>2</sub> 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% O<sub>2</sub>. 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% O<sub>2</sub> (*t*<sub>1/2</sub>=260s vs. 45 s). We also measured [•NO]<sub>ss</sub> in the culture media of cells treated with the •NO-donor DETA/NO at either 1% or 21% O<sub>2</sub>. 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 O<sub>2</sub> concentration [7]. For an equivalent concentration of DETA/NO (500 μM), the [•NO]<sub>ss</sub> was greater at 1% O<sub>2</sub> than it was at 21% O<sub>2</sub> (Fig. 2B). These observed differences in [•NO]<sub>ss</sub> at 1% vs. 21% O<sub>2</sub> can be partially attributed to differences in the rates of •NO metabolism which decreases as the O<sub>2</sub> 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 O<sub>2</sub> through the autooxidation reaction [8]. This reaction is first-order with respect to the O<sub>2</sub> concentration (Eq. 3) and, therefore, its rate will vary based on differences in O<sub>2</sub>. We calculated the contribution of this reaction to the observed differences in the rates of •NO disappearance at both O<sub>2</sub> concentrations. Autooxidation occurs with the following overall stoichiometry [8]:



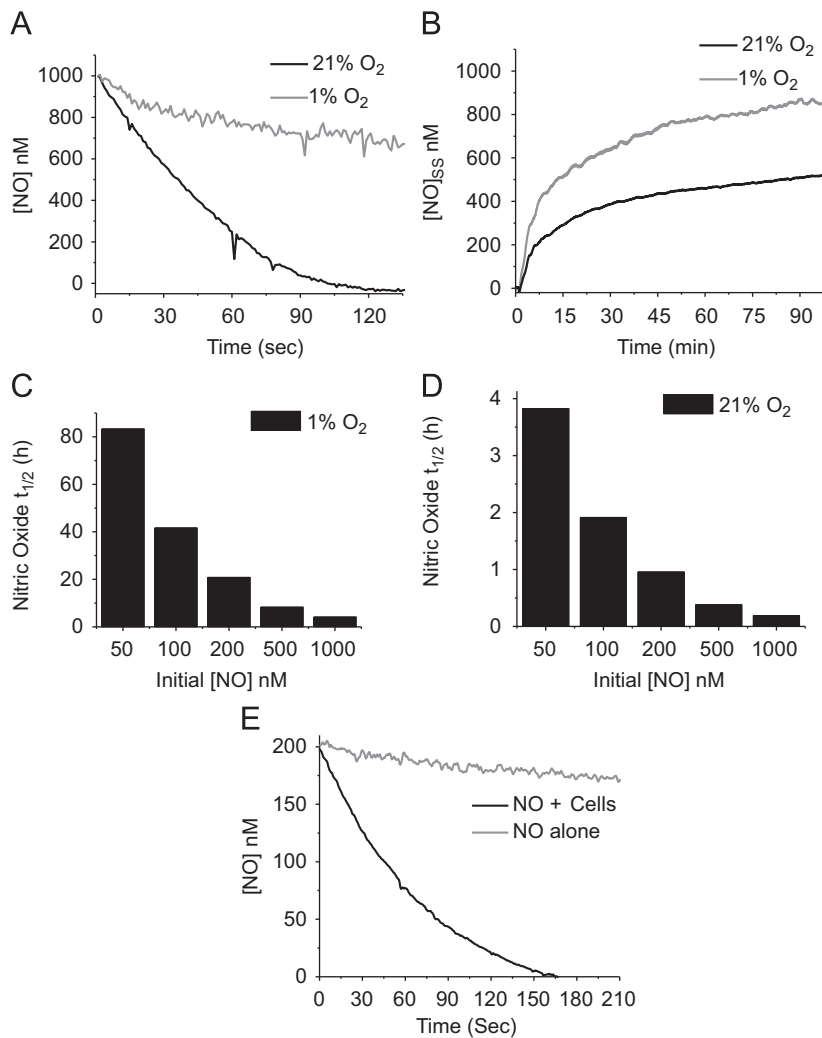
The rate of disappearance of •NO through this reaction is given by

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

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% O<sub>2</sub> 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 O<sub>2</sub> concentration. At 1% O<sub>2</sub>, the half-life of •NO is on the order of hours to days at 1000 nM •NO (Fig. 2C). At 21% O<sub>2</sub>, the half-life of •NO is reduced to minutes or hours (Fig. 2D). These calculations reveal that, at 1% O<sub>2</sub>, 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% O<sub>2</sub>. 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 O<sub>2</sub>, the disappearance of •NO via other mechanisms should be constant at any O<sub>2</sub> 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 O<sub>2</sub> 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% O<sub>2</sub> or 1% O<sub>2</sub>. (B) Measurement of [•NO]<sub>ss</sub> in the media of cells treated with 500 μM of the •NO-donor DETA/NO (•NO-selective electrode, ~1 mm above monolayer) at 21% O<sub>2</sub> or 1% O<sub>2</sub>. (C,D) Calculated half-life of •NO via autooxidation from 50–1000 nM starting concentrations at 1% and 21% O<sub>2</sub>. (E) Disappearance of 200 nM •NO at 21% O<sub>2</sub> 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]<sub>ss</sub> is predominantly a function of the O<sub>2</sub>-dependant rate at which it is metabolized by cells (Fig. 2B). We set out to determine if the effect of O<sub>2</sub> on [•NO]<sub>ss</sub> 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% O<sub>2</sub> or 21% O<sub>2</sub> for 2 h. Fig. 3 demonstrates that, at 1% O<sub>2</sub>, the concentration of Sper/NO necessary to induce p53 P-Ser-15 is only one fourth of the amount required at 21% O<sub>2</sub>. For a given Sper/NO concentration, the rate of •NO production is the same at both O<sub>2</sub> concentrations, yet the [•NO]<sub>ss</sub> is greater at 1% O<sub>2</sub> 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]<sub>ss</sub> from 25 μM Sper/NO at 1% O<sub>2</sub> is roughly equivalent to what is achieved at 21% O<sub>2</sub> from 100 μM Sper/NO. This demonstrates that, at a constant rate of •NO production, it is the O<sub>2</sub> concentration that ultimately determines the extent of •NO signaling.

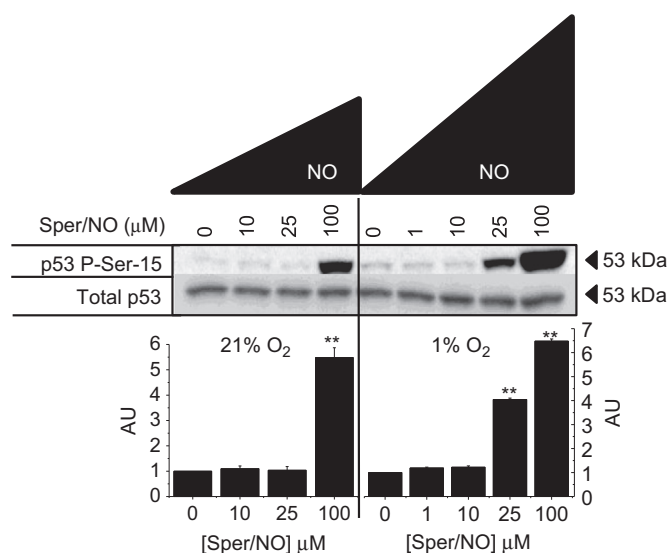
### 3.4. Oxygen determines the rate of enzymatic •NO synthesis

Unlike the liberation of •NO from synthetic donors, the biological production of •NO is enzymatic and requires molecular O<sub>2</sub> as a substrate. For a given NOS isoform, changes in O<sub>2</sub> concentrations will affect enzymatic •NO production. To demonstrate this, we cultured RAW 264.7 macrophages over a range of O<sub>2</sub> 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 NO<sub>3</sub><sup>-</sup>/NO<sub>2</sub><sup>-</sup> 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 O<sub>2</sub> concentration rose from 1–8%. More importantly, however, above 8% O<sub>2</sub> no further increase in the rate of •NO production was observed. This indicates that O<sub>2</sub> is no longer a limiting factor for •NO synthesis at concentrations greater than 8%.

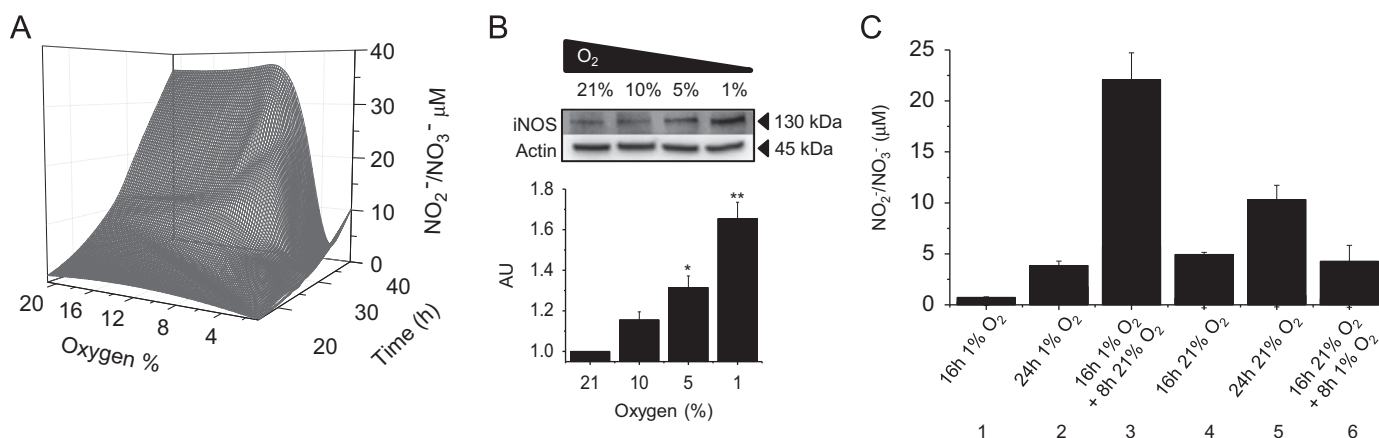
Hypoxia is known to affect the magnitude of iNOS expression which could potentially influence the overall amount of •NO synthesis [4]. We examined this possibility by culturing RAW 264.7 cells over a range of O<sub>2</sub> concentrations and measuring the levels of iNOS expression subsequent to LPS stimulation. Fig. 4B

demonstrates that decreasing the O<sub>2</sub> concentration increased the magnitude of iNOS expression. Next, we set out to compare the effects of enzyme concentration vs. O<sub>2</sub> concentration on the amount of •NO produced. Activated RAW 264.7 cells were cultured at 1% and 21% O<sub>2</sub> for 16 or 24 h and the amount of NO<sub>2</sub><sup>-</sup>/NO<sub>3</sub><sup>-</sup> was measured (Fig. 4C, columns 1,2,4,5). At both time points, the amount of •NO synthesis was considerably higher at 21% O<sub>2</sub> than at 1% O<sub>2</sub> 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% O<sub>2</sub>. This resulted in a substantial increase in cellular iNOS protein expression (Fig. 4B). After 16 h these cells were moved to 21% O<sub>2</sub> for an additional 8 h (24 h total) and NO<sub>2</sub><sup>-</sup>/NO<sub>3</sub><sup>-</sup> was measured (Fig. 4C, column 3). This resulted in a dramatic increase in •NO production at 21% O<sub>2</sub> that we attributed to the hypoxia-mediated increase in iNOS expression (Fig. 4C, compare columns 1, 2 and 3).



**Fig. 3.** Low O<sub>2</sub> 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% O<sub>2</sub> or 1% O<sub>2</sub> 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 O<sub>2</sub> concentrations. RAW 264.7 cells were cultured at different O<sub>2</sub> concentrations and stimulated with LPS (*t*=0 h). The rate of •NO synthesis was measured by quantifying NO<sub>2</sub><sup>-</sup>/NO<sub>3</sub><sup>-</sup> accumulation in the media by chemiluminescence. (A) Contour plot demonstrating the temporal relationship between changing O<sub>2</sub> concentrations and rates of •NO synthesis. (B) Immunoblot demonstrating increased iNOS expression as a function of decreasing O<sub>2</sub> concentrations (*t*=10 h). (C) (columns 1, 2, 4, & 5) Cells were grown at 21% O<sub>2</sub> or 1% O<sub>2</sub> and NO<sub>2</sub><sup>-</sup>/NO<sub>3</sub><sup>-</sup> was measured at 16 h and 24 h. (columns 3 & 6) Cells were grown at 21% O<sub>2</sub> or 1% O<sub>2</sub>. After 16 h the cells at 21% O<sub>2</sub> were moved to 1% O<sub>2</sub> and the cells at 1% O<sub>2</sub> were moved to 21% O<sub>2</sub>. Both sets of cells were incubated for an additional eight hours (24 h total) and NO<sub>2</sub><sup>-</sup>/NO<sub>3</sub><sup>-</sup> 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.

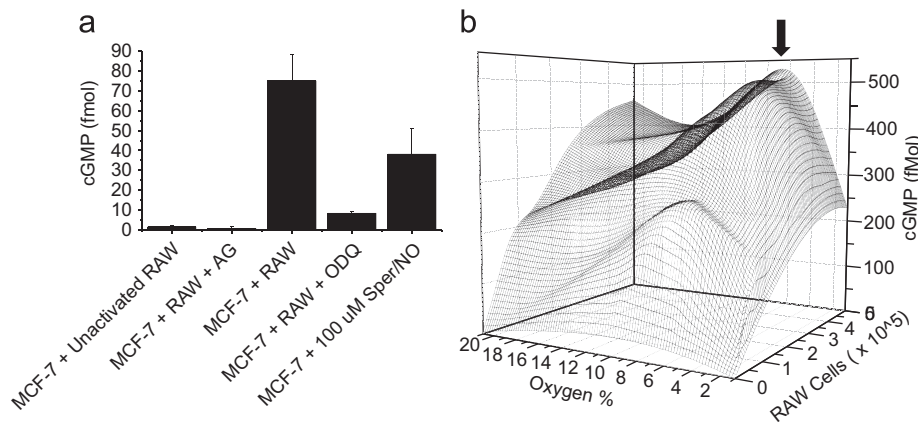
This experiment was repeated, but the order of oxygen exposure was reversed; 21% O<sub>2</sub> for 16 h followed by 8 h at 1% O<sub>2</sub> (Fig. 4C, column 6). Under these conditions, total •NO synthesis was substantially diminished (Fig. 4C, compare columns 3,6). Together these results indicate that at 21% O<sub>2</sub> enzyme concentration is the major limiting factor for •NO synthesis whereas at 1% O<sub>2</sub>, 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 O<sub>2</sub> 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 O<sub>2</sub> 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 O<sub>2</sub> and •NO concentrations. Fig. 5B demonstrates that sGC activation was dependent on both the O<sub>2</sub> concentration and the number of •NO-producing cells. Interestingly, the optimal O<sub>2</sub> concentration for maximal cGMP production was at ≈ 5–8% O<sub>2</sub>. This suggests that at this O<sub>2</sub> 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<sub>2</sub> concentration. (A) Activated RAW 264.7 cells were cocultured in serum free media with MCF-7 cells at 21% O<sub>2</sub>. 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<sub>2</sub> 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<sub>2</sub> 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  $[\bullet\text{NO}]_{\text{ss}}$  in cells and tissues. It is well-known that both  $\bullet$ NO synthesis and  $\bullet$ NO consumption are O<sub>2</sub>-dependent, and thus the oxygen concentration will influence the rates of both of these processes. We investigated the overall ability of O<sub>2</sub> to modulate  $\bullet$ NO signaling events. If the rate of  $\bullet$ NO synthesis is greater than the rate of cellular  $\bullet$ NO consumption,  $[\bullet\text{NO}]_{\text{ss}}$  concentrations will increase. Conversely, if the rate of  $\bullet$ NO consumptive processes predominate over its rate of synthesis,  $[\bullet\text{NO}]_{\text{ss}}$  levels will decrease. Oxygen largely dictates the  $[\bullet\text{NO}]_{\text{ss}}$  and ultimately the magnitude of  $\bullet$ NO signaling. We and others have demonstrated previously that the  $[\bullet\text{NO}]_{\text{ss}}$  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<sub>2</sub> 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<sub>2</sub> 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<sub>2</sub> concentration. At high O<sub>2</sub>,  $\bullet$ NO synthesis is dependent on the amount of enzyme present, while at low O<sub>2</sub>, oxygen itself becomes the limiting factor.

We used a  $\bullet$ NO-selective electrode to measure the  $[\bullet\text{NO}]_{\text{ss}}$  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  $[\bullet\text{NO}]_{\text{ss}}$  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  $[\bullet\text{NO}]_{\text{ss}}$  [2]. Our current study builds on these findings by demonstrating how  $\bullet$ NO signaling can be significantly modified by the O<sub>2</sub> concentration. In many ways these results are expected given the ability of O<sub>2</sub> to modulate  $\bullet$ NO concentrations. However, our results demonstrate

that changes in  $\bullet$ NO signaling do not always change linearly with changing O<sub>2</sub> concentrations.

When the rate of  $\bullet$ NO production is constant, its steady-state concentration is predominantly a function of its O<sub>2</sub>-dependent rate of metabolism. We demonstrate that, at lower O<sub>2</sub> concentrations,  $\bullet$ NO metabolism is considerably slower, and thus  $[\bullet\text{NO}]_{\text{ss}}$  is higher. This inverse relationship has a direct effect on  $\bullet$ NO signaling. Activation of p53 Ser-15 phosphorylation at 1% O<sub>2</sub>, for example, is possible with one-fourth the dose of Sper/NO necessary to achieve the same effect at 21% O<sub>2</sub>. This indicates that, for a given rate of  $\bullet$ NO production (synthesis), lowering the O<sub>2</sub> concentration will enable activation of signaling pathways that are not obtainable at higher O<sub>2</sub> concentrations.

In addition to  $\bullet$ NO consumptive pathways, we show that O<sub>2</sub> concentrations influence  $[\bullet\text{NO}]_{\text{ss}}$  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<sub>2</sub>. Above this amount of O<sub>2</sub>, further increases in the rate of  $\bullet$ NO synthesis were not observed. When sGC activation by  $\bullet$ NO was examined under changing O<sub>2</sub> concentrations, we noted maximal activation at 5–8% O<sub>2</sub>. At O<sub>2</sub> concentrations higher or lower than these amounts, there was a decrease in sGC activation. We suggest that the balance between O<sub>2</sub> availability and enzyme expression at 5–8% O<sub>2</sub> is optimized such that the rate of  $\bullet$ NO production versus its rate of metabolism synergize to achieve maximal  $[\bullet\text{NO}]_{\text{ss}}$ . At higher O<sub>2</sub> concentrations,  $\bullet$ NO production is maximal but so is its metabolism. At lower O<sub>2</sub> concentrations,  $\bullet$ NO metabolism is slowest but so is its enzymatic synthesis.

## 5. Conclusions

These findings highlight the importance and dependence of O<sub>2</sub> 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  $[\bullet\text{NO}]_{\text{ss}}$ , 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<sub>2</sub> mediated changes in  $[\bullet\text{NO}]_{\text{ss}}$ . 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.

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