

## Kinetic model of the inhibition of respiration by endogenous nitric oxide in intact cells

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### ABSTRACT

Nitric oxide (NO) inhibits mitochondrial respiration by decreasing the apparent affinity of cytochrome *c* oxidase (CcO) for oxygen. Using iNOS-transfected HEK 293 cells to achieve regulated intracellular NO production, we determined NO and O<sub>2</sub> concentrations and mitochondrial O<sub>2</sub> consumption by high-resolution respirometry over a range of O<sub>2</sub> concentrations down to nanomolar. Inhibition of respiration by NO was reversible, and complete NO removal recovered cell respiration above its routine reference values. Respiration was observed even at high NO concentrations, and the dependence of IC<sub>50</sub> on [O<sub>2</sub>] exhibits a characteristic but puzzling parabolic shape; both these features imply that CcO is protected from complete inactivation by NO and are likely to be physiologically relevant. We present a kinetic model of CcO inhibition by NO that efficiently predicts experimentally determined respiration at physiological O<sub>2</sub> and NO concentrations and under hypoxia, and accurately predicts the respiratory responses under hyperoxia. The model invokes competitive and uncompetitive inhibition by binding of NO to the reduced and oxidized forms of CcO, respectively, and suggests that dissociation of NO from reduced CcO may involve its O<sub>2</sub>-dependent oxidation. It also explains the non-linear dependence of IC<sub>50</sub> on O<sub>2</sub> concentration, and the hyperbolic increase of c<sub>50</sub> as a function of NO concentration.

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### 1. Introduction

Nitric oxide (NO) is a fundamental cell messenger produced in cells by NO synthases (NOS). Many physiological actions of NO are mediated through activation of soluble guanylate cyclase and subsequent production of cGMP [1,2]. NO is also an effective inhibitor of mitochondrial respiration [3–5]: it reversibly inhibits cytochrome *c* oxidase (CcO), the terminal electron acceptor of the mitochondrial respiratory system, in a process which occurs in competition with oxygen (O<sub>2</sub>). NO, like other inhibitors that raise the *K'*<sub>m</sub> of CcO, induces O<sub>2</sub> limitation under apparently normoxic conditions. The inhibition of mitochondrial respiration by NO has been implicated in a wide range of physiological processes, including the regulation of the

affinity of mitochondrial respiration for O<sub>2</sub> [6–10], the control of mitochondrial generation of superoxide [11] and hydrogen peroxide [12], the activation of hypoxia-inducible factor [13], the activation of AMPK in astrocytes [14], the modulation of O<sub>2</sub> delivery to tissues [15], the modulation of calcium-mediated cell signaling in the brain [16] and the maintenance of constant cerebral O<sub>2</sub> consumption at varying blood flow [17]. The interaction between CcO and NO has been implicated in various pathophysiological situations [18,19].

The molecular mechanism by which NO inhibits CcO has not been fully defined (for a review see [20]). The enzyme belongs to the superfamily of heme-copper oxidases and contains a highly conserved bimetallic active site composed of a high-spin heme *a*<sub>3</sub> and a copper ion Cu<sub>B</sub>. The binuclear center is the binding site of the physiological substrate O<sub>2</sub> and of other ligands such as CN<sup>-</sup>, CO, and NO [21]. O<sub>2</sub>-binding demands the complete reduction of the active site: two electrons donated by cytochrome *c* enter CcO at Cu<sub>A</sub>, from which they are transferred, via cytochrome *a*, to the binuclear center. Conflicting hypotheses have been proposed to explain the mechanism of CcO inhibition by NO [22–24]; however, recent studies demonstrate that NO interactions with CcO cannot be adequately described by a simple competitive model and that only one NO molecule binds per binuclear center [25]. The enzyme can bind NO either at the reduced heme *a*<sub>3</sub> iron (competitive with O<sub>2</sub>) or at the oxidized Cu<sub>B</sub> (non-competitive with O<sub>2</sub>)

**Abbreviations:** NO, nitric oxide; NOS, NO synthase; CcO, cytochrome *c* oxidase; O<sub>2</sub>, oxygen; CN<sup>-</sup>, cyanide; CO, carbon monoxide; iNOS, inducible NOS; DMEM, Dulbecco's modified Eagle's medium; Tet-iNOS 293, tetracycline-inducible iNOS-expressing HEK 293 cells; S-EITU, S-ethylisothiourea; FCCP, carbonylcyanide *p*-(trifluoromethoxy) phenylhydrazone; HBSS, Hanks balanced salt solution; HbO<sub>2</sub>, oxyhemoglobin

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[25–28]. NO binds to reduced heme  $a_3$  very quickly and with high affinity, to yield a  $\text{Fe}^{2+}$ -NO nitrosyl-adduct. Through binding to oxidized  $\text{Cu}_B$ , donation of an electron to the metal and subsequent hydration, NO is oxidized to nitrite. The predominance of one inhibitory mechanism over the other is controlled by the electron flux through the enzyme [29]. NO and its derivative peroxynitrite have also been reported to irreversibly decrease the affinity of CcO for  $\text{O}_2$  [30], although this effect is unlikely to occur at physiological NO levels.

The complexity of the cellular environment provides the ideal setting for studying how the interplay between  $\text{O}_2$  and NO concentrations determines the extent of respiratory inhibition and the  $\text{O}_2$ -dependence of iNOS-mediated NO production. However, because of technical limitations, previous studies have not provided accurate kinetic data at low  $[\text{O}_2]$ . Additionally, in many studies sensitivity of  $\text{O}_2$  consumption to NO donors has been examined at  $[\text{O}_2]$  much higher than those likely to exist *in vivo*. High-resolution respirometry allows accurate determination of respiration at low  $\text{O}_2$  concentrations and during transition to anoxia [31]. Here we present a detailed study of the  $\text{O}_2$  kinetics of cellular respiration in intact cells producing controlled amounts of NO [32]. Insertion of an NO sensor into the Oxygraph-2k respirometer chamber allowed simultaneous recording of NO production and mitochondrial respiration over an extended  $[\text{O}_2]$  range. Based on our results, we present a pseudo-equilibrium kinetic model of inhibition of respiration by NO that accounts for both competitive and uncompetitive inhibition of CcO and has predictive value.

## 2. Materials and methods

### 2.1. Cell culture and reagents

Tetracycline-inducible HEK 293 cells stably expressing human iNOS (Tet-iNOS 293) were generated as described [32]. Cells were cultured in DMEM (Invitrogen, Barcelona, Spain) containing 4.5 g/l D-glucose, 10% (v/v) fetal-calf serum, 200  $\mu\text{g}/\text{ml}$  hygromycin B and 15  $\mu\text{g}/\text{ml}$  blasticidin, at 37 °C in a humidified atmosphere with 5%  $\text{CO}_2$ . S-ethylisothiourea hydrobromide (S-EITU; NOS inhibitor), L-arginine, oligomycin and FCCP were from Sigma-Aldrich (St. Louis, MO). Hygromycin B and blasticidin were from Invitrogen. Tetracycline was from Calbiochem (Darmstadt, Germany).

### 2.2. Induction of endogenous NO production

Expression of iNOS was induced as previously described [33]. Briefly, cells were incubated overnight in a complete growth medium (without selection antibiotics) containing tetracycline (5–50 ng/ml) and 500  $\mu\text{M}$  S-EITU, a potent inhibitor of iNOS activity. Cells were washed with L-arginine-free DMEM supplemented with 1% dialysed fetal-calf serum to eliminate any traces of L-arginine. Cells were incubated in L-arginine-free medium for 1 h to completely eliminate S-EITU and tetracycline without activating NO production. Cells were trypsinized and resuspended at  $1 \times 10^7$  cells/ml in HBSS containing 25 mM Hepes. Endogenous NO production was triggered by addition of L-arginine (1 mM).

### 2.3. Immunoblot analysis of iNOS expression

For preparation of total protein extracts, cells were trypsinized and resuspended at  $1 \times 10^7$  cells/ml in HBSS containing 25 mM Hepes. Cells were spun (300 g, 5 min, 4 °C) and the pellet was resuspended in 60  $\mu\text{l}$  ice-cold lysis buffer (20 mM Hepes, pH 7.5, 400 mM NaCl, 20% (v/v) glycerol, 0.1 mM EDTA, 10 mM NaF, 10  $\mu\text{M}$   $\text{Na}_2\text{MoO}_4$ , 1 mM  $\text{NaVO}_3$ , 10 mM PNPP (*p*-nitrophenyl phosphate), and 10 mM  $\beta$ -glycerophosphate) supplemented with 1 mM dithiothreitol, 1 mM pepablock, pH 7.4, and a protease-inhibitor cocktail (Roche Diagnostics, Barcelona, Spain). After 15 min on ice, cleared whole cell extracts were obtained by recovering the supernatant after centrifugation

(16,000 g, 15 min, 4 °C). Protein samples (200  $\mu\text{g}$ ) were resolved by SDS/7.5%-(w/v)-PAGE and transferred to nitrocellulose membranes. Membranes were blocked with 5% (w/v) non-fat dry milk in TBS-T (20 mM Tris/HCl, pH 7.2, 150 mM NaCl, 0.1% Tween 20) and incubated overnight with polyclonal anti-iNOS antibody (1:2000, Transduction Laboratories, Erembodegem, Belgium) in a blocking solution at 4 °C. Protein bands were detected by incubation for 1 h with horseradish peroxidase-coupled goat anti-rabbit IgG (1:5000; Vector Laboratories, Burlingame, CA) in blocking solution at room temperature, followed by enhanced chemiluminescence (GE Healthcare, Amersham, UK).

### 2.4. Simultaneous measurement of $\text{O}_2$ consumption and NO production

$\text{O}_2$  consumption at physiological tissue  $[\text{O}_2]$  was determined by high-resolution respirometry (Oxygraph-2k; Oroboros Instruments, Innsbruck, Austria). Cells were trypsinized after overnight treatment with tetracycline and resuspended in HBSS containing 25 mM Hepes at  $1 \times 10^7$  cells/ml. The instrumental background flux was calculated as a linear function of  $[\text{O}_2]$  and the experimental data were corrected for the whole range of  $[\text{O}_2]$  using DatLab software (Oroboros Instruments) [31]. To separate mitochondrial and residual  $\text{O}_2$  consumption, all results were corrected for residual  $\text{O}_2$  consumption of 0.5  $\text{pmol O}_2 \cdot \text{s}^{-1} \cdot 10^{-6}$  cells, measured at 35  $\mu\text{M O}_2$  (Table 1). Compared with CcO, residual oxygen uptake has a lower affinity for oxygen [34]. This was accounted for by assigning a 5-fold higher  $c_{50}$  to residual respiration as a basis for application of the residual  $\text{O}_2$  consumption-correction over the entire experimental  $[\text{O}_2]$  range. At air saturation, 37 °C and local barometric pressure (92.6 kPa), the  $[\text{O}_2]$  in culture medium (HBSS) was 175.7  $\mu\text{M}$  ( $\text{O}_2$  solubility factor 0.92). Measurements were taken in cell suspensions (2 ml) gently agitated at 37 °C in parallel Oxygraph-2k chambers; NO production was initiated in one chamber by addition of 1 mM L-arginine; a control run (in cells treated with the same amount of tetracycline) was simultaneously performed in the other chamber. Because light reverses NO-induced inhibition of respiration [35–37]  $\text{O}_2$  consumption was measured in the dark, except for analysis of inhibition reversibility, when cell suspensions were illuminated with cold light (Intralux 5000-1; Volpi, Switzerland). Reversibility was also studied by adding  $\text{HbO}_2$  to quench NO or S-EITU (1 mM) to inhibit iNOS activity.

For the simultaneous measurement of endogenous NO production, an NO sensor (ISO-NOP; World Precision Instruments, Stevenage, UK) was inserted into the Oxygraph-2k chamber. For this, a second capillary was drilled into the PVDF stopper to tightly fit the 2 mm diameter sleeve of the NO sensor. The opening of the second capillary did not increase  $\text{O}_2$  backdiffusion into the chamber at low  $[\text{O}_2]$ . The NO sensor was calibrated by addition of known  $\text{NaNO}_2$  concentrations under reducing conditions ( $\text{KI}/\text{H}_2\text{SO}_4$ ) at 37 °C. To initiate NO production, 1 mM L-arginine was added to the chamber at 60  $\mu\text{M O}_2$ . This  $[\text{O}_2]$  is sufficient to sustain iNOS activity [38,39]. NO concentrations were monitored until  $\text{O}_2$  was exhausted and the NO signal returned to basal. The NO signal

**Table 1**

Routine respiration, electron transport capacity, and resting and residual respiration in Tet-iNOS 293 cells. Respiration of control cells in the absence of NO (cells were not treated with tetracycline and no arginine was added). Routine: basal respiration using endogenous substrates. Resting: addition of oligomycin (80 ng/ml) to measure respiration in the absence of ATP synthesis. Uncoupled: addition of optimum FCCP concentration (0.35  $\mu\text{M}$ ) for maximum flow to measure the capacity of the electron transport system. Corrections were made for residual  $\text{O}_2$  consumption (1 mM KCN at 30 to 40  $\mu\text{M O}_2$ ). Measurements were made at 37 °C at  $1 \times 10^7$  cells/ml in the chambers of a high-resolution respirometer. Values are means  $\pm$  SD (number of independent experiments in parentheses).

| Respiratory state<br>endogenous substrates | Respiration<br>[ $\text{pmol O}_2 \cdot \text{s}^{-1} \cdot 10^{-6}$ cells] | Flux control ratio<br>per uncoupled rate |
|--|---|--|
| Routine                                    | 14.5 $\pm$ 1.9 (8)  | 0.31 $\pm$ 0.03                          |
| Resting (oligomycin)                       | 4.2 $\pm$ 0.4 (3)   | 0.089 $\pm$ 0.004                        |
| Uncoupled (FCCP)                           | 47.4 $\pm$ 7.1 (8)  |  |
| Residual (KCN)                             | 0.51 $\pm$ 0.08 (5)   | 0.011 $\pm$ 0.003                        |

before addition of L-arginine was used for internal zero calibration, and non-linear zero drift was observed before addition of L-arginine and after zero  $O_2$  was reached. A polynomial interpolation of these sections of the zero signal was used to construct a zero-baseline, which was subtracted from the total signal.

### 3. Results

#### 3.1. Routine, uncoupled and oligomycin-inhibited respiration

Cell respiration was measured in intact Tet-iNOS 293 cells, which express iNOS upon treatment with tetracycline [32] (Fig. 2E). Cell respiration with endogenous substrates was determined in cells not expressing iNOS, suspended in HBSS containing 25 mM Hepes. Routine  $O_2$  consumption was  $14.5 \pm 1.9 \text{ pmol } O_2 \cdot \text{s}^{-1} \cdot 10^{-6} \text{ cells}$ , corrected for residual  $O_2$  consumption after inhibition of CcO. Uncoupled respiration was three-fold higher than routine respiration (Table 1), indicating that endogenous respiration has a large potential for activation and is not substrate-limited. Upon addition of oligomycin to inhibit ATP synthase, respiration declined to around 10% of uncoupled or 30% of routine respiration, showing that a high proportion of routine respiration is coupled to ATP synthesis. Residual  $O_2$  consumption, measured after addition of 1 mM KCN to inhibit CcO, was as low as 3% of routine respiration.

#### 3.2. $O_2$ kinetics in the absence of NO

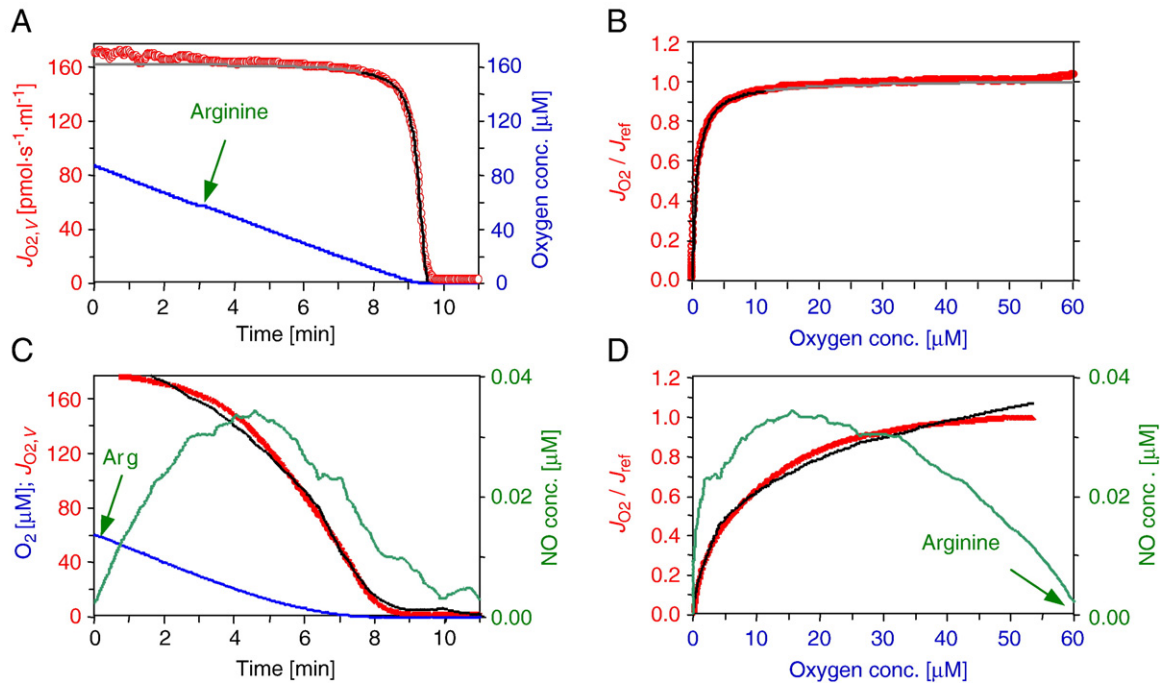
In the absence of NO, routine respiration followed a strictly hyperbolic function, fitted in the low- $O_2$  range ( $<11 \mu\text{M}$ ; Fig. 1A and B). The  $[O_2]$  at half-maximum flux,  $c_{50}$  (equivalent to the apparent  $K_{m,O_2}$ ) [40], was  $0.65 \mu\text{M} \pm 0.04 \text{ SD}$  ( $n=5$ ) for routine respiration in control cells (not expressing iNOS) and was unaffected by addition of 1 mM L-arginine (Fig. 1A and B). The  $c_{50}$  agrees with the results obtained in human umbilical vein endothelial cells [41] and fibroblasts [42], and is comparable to the  $c_{50}$  in isolated mitochondria [40,43].

#### 3.3. Endogenous NO production and inhibition of respiration by NO in the low- $O_2$ range

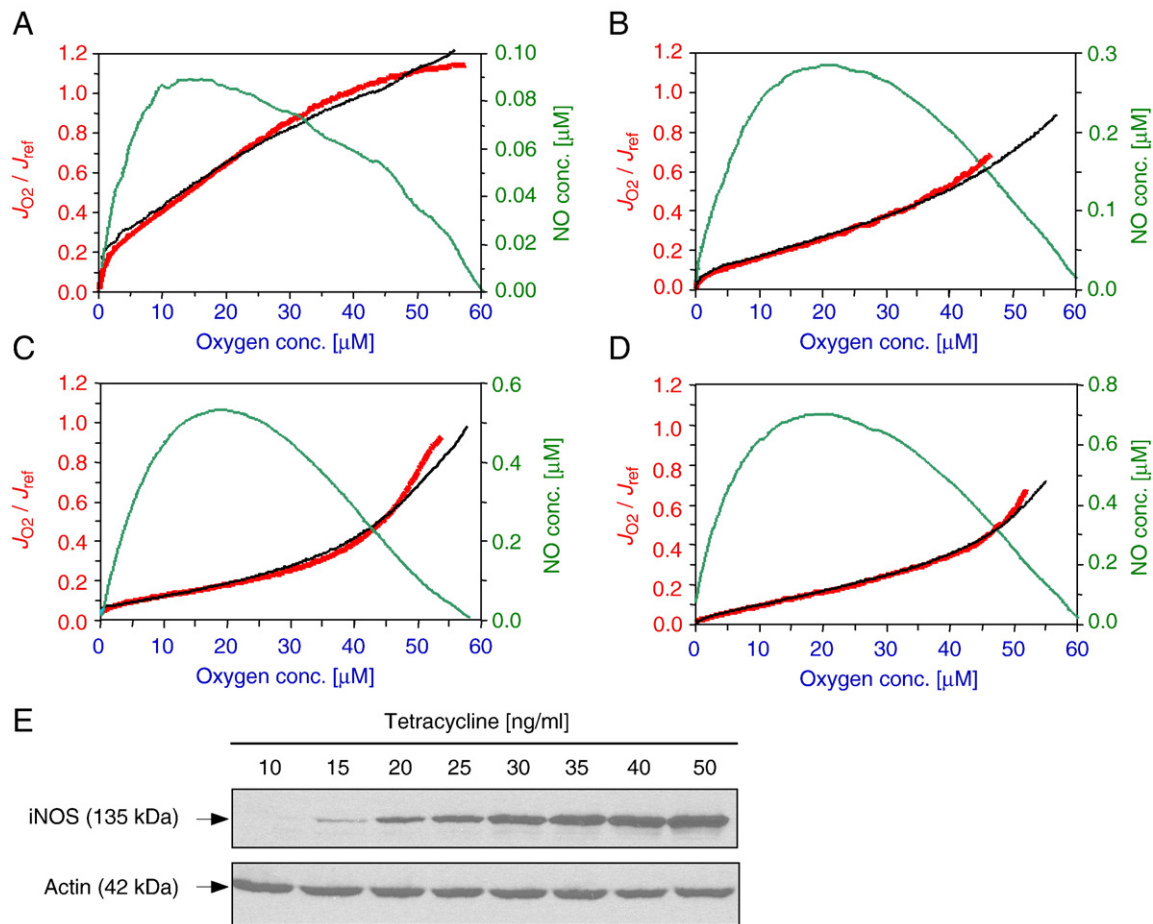
Overnight incubation of cells with increasing concentrations of tetracycline (10–50 ng/ml) induced a concentration-dependent expression of iNOS, as determined by western blot (Fig. 2E). Addition of the iNOS substrate L-arginine (1 mM) at  $60 \mu\text{M } O_2$  immediately triggered NO production, as shown by a steady increase of [NO] (Fig. 1C). The amount of NO produced by cells after addition of L-arginine was proportional to the level of iNOS expression. In the physiological  $O_2$  range, even low [NO] (30 nM) caused a significant right-shift in the  $O_2$  kinetics of cell respiration (Fig. 1D; compare with Fig. 1B). When  $[O_2]$  had dropped to  $\sim 15 \mu\text{M}$ , NO degradation balanced NO production. iNOS activity is progressively limited as  $[O_2]$  decreases (iNOS has a high  $K_m$  for  $O_2$ ; [38,39]). At low  $[O_2]$ , however, mitochondria are capable of producing NO from

#### 2.5. Data analysis

Data were analyzed by standard least squares non-linear minimization routines, written with the Matlab program (MathWorks, Natick, MA).



**Fig. 1.** Cell respiration in the absence or presence of low levels of NO production.  $O_2$  consumption and NO production were measured in cells without (A,B) and with (C,D) iNOS induction (20 ng/ml tetracycline). (A,C) Oxygraph-2k traces of  $[O_2]$  (blue),  $O_2$  flux (red) and [NO] (green) as a function of time, with data points plotted at 1 s intervals. (B,D) Kinetic plots of  $O_2$  flux as a function of  $[O_2]$  in the low- $O_2$  range ( $<11 \mu\text{M}$ ). (A,B) The hyperbolic fits (black) almost perfectly superimpose the measured data points. (D) The graph shows relative measured flux (red) calculated relative to routine flux recorded at saturating  $[O_2]$  ( $J_{ref}$ ), relative calculated flux from the kinetic model (black), and [NO] (green) as a function of  $[O_2]$  at low levels of NO production (20 ng/ml tetracycline).



**Fig. 2.** Endogenous NO production and NO-induced inhibition of cell respiration. (A–D) Kinetic plots of  $O_2$  flux as a function of  $[O_2]$  in the low- $O_2$  range ( $<11 \mu M$ ), and electrochemical measurement of NO, in cells with iNOS induction. The graphs show relative measured flux (red) calculated relative to routine flux recorded at saturating  $[O_2]$  ( $J_{ref}$ ), relative calculated flux from the kinetic model (black), and  $[NO]$  (green) as a function of  $[O_2]$  at three levels of NO production: (A) low production (20 ng/ml tetracycline); (B,C) intermediate production (25 ng/ml tetracycline), and (D) high production (30 ng/ml tetracycline). Traces are representative of five similar experiments. (E) Western blot showing the expression of iNOS after overnight induction with tetracycline (10–50 ng/ml). Expression of actin was detected as a loading control.

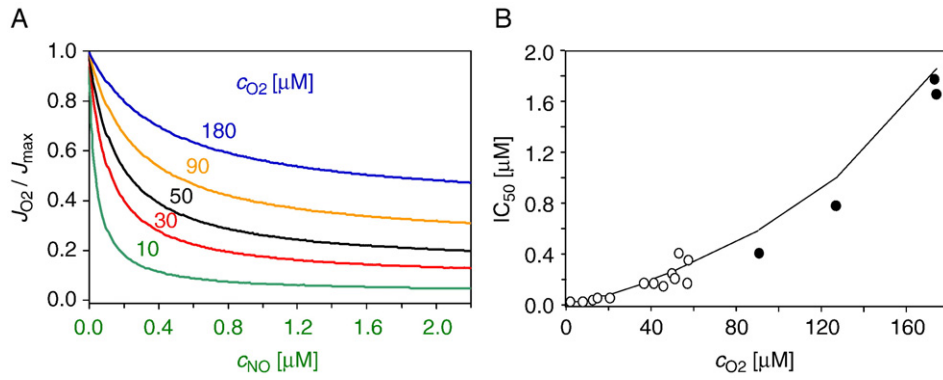
nitrite ( $NO_2^-$ ) [44].  $[NO]$  declined to zero when  $O_2$  was exhausted by cell respiration (Fig. 1C and D). Accordingly, addition of L-arginine at zero  $[O_2]$  did not induce detectable NO production (not shown).

The rate of NO production and the maximum  $[NO]$  obtained in the experimental low- $O_2$  regime were a function of iNOS expression (Fig. 1C and D; Fig. 2A–D). For each experiment, the reference flux ( $J_{ref}$ ), corresponding to routine respiration, was measured when  $[O_2]$  was not limiting and before addition of L-arginine. Respiration was inhibited as a complex function of  $O_2$ -dependent NO production and  $O_2$  limitation of CcO (Fig. 1D; Fig. 2A–D). Two experimental observations were particularly puzzling. First, CcO maintained residual activity up to  $2 \mu M$  NO even at  $[O_2]$  in the low  $\mu M$  range, in spite of the fact that CcO inhibition was already detectable at low-nM  $[NO]$  (see Fig. 2D; Fig. 5B and D). This cannot be explained by  $O_2$  consumption by other enzyme systems and suggests that at least one NO-bound derivative of CcO is active and reduces  $O_2$ . Second, the  $[NO]$  required to inhibit CcO by 50% at any given  $[O_2]$  ( $IC_{50}$ ) has an unusual parabolic dependence on  $[O_2]$  (Fig. 3B; see also [24]). A similar upward parabolic trend was obtained when NO was generated from the NO donor DETA-NONOate (Fig. S1).

The interdependence of NO and  $O_2$  concentrations generally makes it impossible to apply kinetic fitting procedures that depend on variation of substrate ( $O_2$ ) concentration at a constant concentration of the inhibitor (NO). We therefore applied an approach in which kinetic parameters were fitted to all data points of all experiments simultaneously.

### 3.4. Kinetic model of CcO inhibition by NO

Consistent with previous observations [25–28], our data cannot be adequately explained by a simple competitive inhibition model in which NO binds to the reduced heme  $a_3$  of CcO in place of  $O_2$ , especially in relation to the residual activity at high  $[NO]$  and the dependence of  $IC_{50}$  on  $[O_2]$ . These perplexing features would be explained if CcO catalyzed oxidation of NO, at least under some experimental conditions. We therefore tested two minimal chemically-sound, albeit oversimplified, models that allow NO to be oxidized by  $O_2$ . One considers oxidation of NO bound to the reduced heme  $a_3$ ; the other, oxidation of NO bound to the oxidized  $Cu_B$ . Although both models were comparably successful, only the first is described here (see Fig. 4), since it has better independent support [45,46]. In this model, when NO occupies the heme  $a_3$ , effectively competing with  $O_2$ , an alternative catalytic pathway is opened, whereby  $O_2$  transiently binds to reduced  $Cu_B$ , is activated to superoxide and is forced to react with NO to yield peroxynitrite, which spontaneously converts to nitrite ( $NO_2^-$ ) [46].



**Fig. 3.** Relationship between NO concentration and the degree of inhibition of respiration. (A) Effect of [NO] on the relative O<sub>2</sub> flux, calculated from the kinetic parameters at constant [O<sub>2</sub>] of 10 μM (green), 30 μM (red), 50 μM (black), 90 μM (yellow), or 180 μM (blue). (B) IC<sub>50</sub> (μM) as a function of [O<sub>2</sub>]; open symbols are from experiments carried out in the low [O<sub>2</sub>] range; closed symbols from the high [O<sub>2</sub>] range. The line is drawn according to Eq. (4), using the parameters reported in the text.

In the model, both reduced species, CcO<sub>r</sub>NO and CcO<sub>r</sub>, are catalytically active. The sum of the enzyme intermediates predicted at any given concentrations of O<sub>2</sub> and NO is given by the following partition function:

$$[CcO_{tot}] = [CcO_r](1 + [O_2]/K_{m1} + [NO]/K_{icNO} + [O_2][NO]/K_{m2}K_{icNO} + [O_2][NO]/K_{uNO}K_{m1}) \quad (1)$$

where CcO<sub>tot</sub> is the total (all species) CcO; CcO<sub>r</sub> is the reduced CcO; K<sub>m1</sub> is the K<sub>m</sub> of CcO for O<sub>2</sub> in the absence of NO; K<sub>m2</sub> is the K<sub>m</sub> of CcO for O<sub>2</sub> when NO is bound to heme a<sub>3</sub>; K<sub>icNO</sub> is the competitive inhibition constant of NO for CcO, and K<sub>uNO</sub> is the uncompetitive inhibition constant of NO for CcO.

The rate of respiration is described by Eq. (2):

$$V = \frac{[CcO_{tot}]V_{max1}([O_2]K_{uNO}K_{m2}K_{icNO} + r[O_2][NO]K_{m1}K_{uNO})}{K_{m1}K_{m2}K_{icNO}K_{uNO} + [O_2]K_{m2}K_{icNO}K_{uNO} + [O_2][NO]K_{m2}K_{icNO} + [NO]K_{m1}K_{m2}K_{uNO} + [O_2][NO]K_{m1}K_{uNO}} \quad (2)$$

where  $r$  represents the ratio  $V_{max2}/V_{max1}$  (which we estimate at  $\sim 1.35$ ). Notice that the oxidative dissociation of NO bound to Cu<sub>B</sub> as nitrite ion governed by  $V_{NO}$  (see Fig. 4) does not involve O<sub>2</sub> consumption and therefore does not appear in Eq. (2).

Analysis by standard least squares non-linear minimization routines yielded the following parameter values:  $K_{m1} = 0.81 \mu\text{M}$ ,  $V_{max1} = 16.5 \text{ pmol O}_2 \cdot \text{s}^{-1} \cdot 10^{-6} \text{ cells}$ ,  $K_{icNO} = 3.63 \text{ nM}$ ,  $K_{m2} = 520 \mu\text{M}$  and  $V_{max2} = 22 \text{ pmol O}_2 \cdot \text{s}^{-1} \cdot 10^{-6} \text{ cells}$ .  $K_{uNO}$  was higher than the usual NO concentrations achieved in this study, thus it was poorly determined and could be dropped from the fit. The best fit to the experimental data requires the affinity of O<sub>2</sub> for CcO to be modulated by enzyme-bound NO, such that the species CcO<sub>r</sub>NO (with NO bound to heme a<sub>3</sub>) has a higher K<sub>m</sub> for O<sub>2</sub> (K<sub>m2</sub>) than the uninhibited enzyme (K<sub>m1</sub>). This is consistent with the inhibitory activity of NO and the lower affinity of O<sub>2</sub> for Cu.

The model was applied to all the data from experiments carried out at low [O<sub>2</sub>] (some of which are presented in Fig. 1C and D and Fig. 2A–D). The kinetic parameters thus obtained were used to simulate the experimental traces collected over a 4-fold extended [O<sub>2</sub>] range, which are shown in Fig. 5, and to generate the lines in Fig. 3A and B. The experiments in the high-O<sub>2</sub> range were designed to test the predictive ability of the model beyond the range over which the parameters were fitted.

In the minimum model of linear competitive inhibition [22], the apparent inhibition constant,  $K'_i$ , determines the proportionality by which the  $c_{50}$  (the [O<sub>2</sub>] required to achieve half  $V_{max}$  at any given [NO]) is increased as a linear function of [NO]. This model fails to explain the non-linear dependence of  $c_{50}$  on [NO] that we observed in intact cells (not shown). A more accurate prediction is given by the following equation, derived from our model:

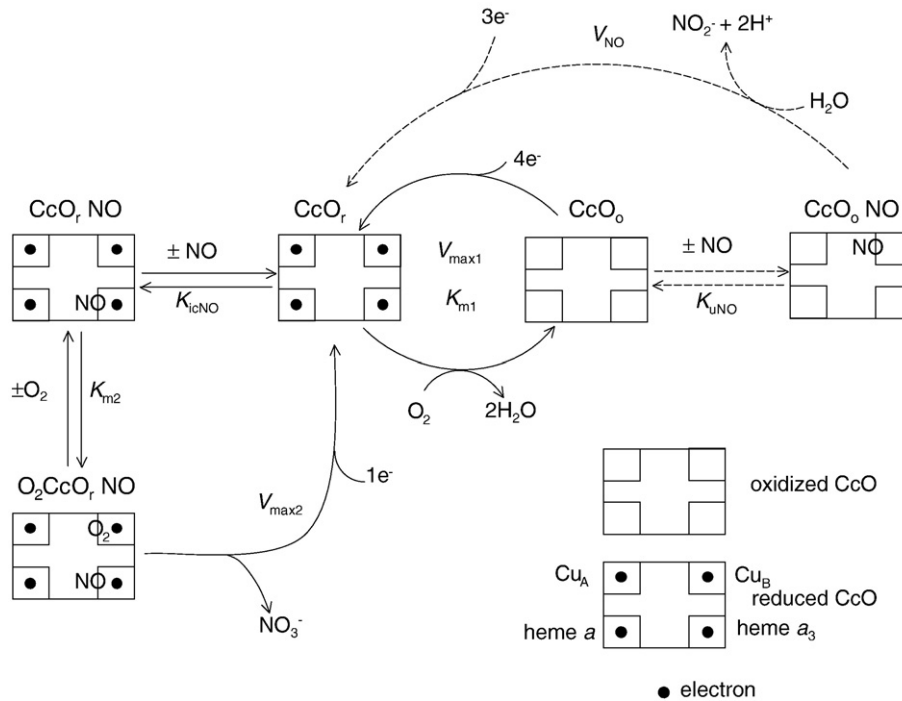
$$c_{50} = \frac{K_{uNO}K_{m1}K_{m2}(K_{icNO} + [NO])}{K_{icNO}K_{uNO}K_{m2} + [NO]K_{uNO}K_{m1} + [NO]K_{icNO}K_{m2}} \quad (3)$$

The IC<sub>50</sub> is defined as the [NO] that inhibits mitochondrial respiration by 50% relative to flux at the same [O<sub>2</sub>] in the absence of NO. In the competitive model, IC<sub>50</sub> is a linear function of [O<sub>2</sub>] [22]; our data in intact cells show that IC<sub>50</sub> is a non-linear function of [O<sub>2</sub>] (Fig. 3B), which our model predicts to be determined as follows:

$$IC_{50} = \frac{K_{uNO}K_{icNO}K_{m2}(K_{m1} + [O_2])}{K_{m1}K_{uNO}K_{m2} + [O_2]K_{m2}K_{icNO} - 2rK_{m1}^2K_{uNO} - (2r-1)[O_2]K_{uNO}K_{m1}} \quad (4)$$

where  $r$  represents the ratio  $V_{max2}/V_{max1}$ .

We are aware of the limitations of the proposed model, which (i) assumes that the cell's reducing pressure is constant and neglects the reduction steps of the catalytic cycle, and (ii) may attribute to CcO roles played by other components of the cell. These limitations, however, do



**Fig. 4.** Scheme representing the proposed model of NO-inhibition of CcO. A simplified catalytic cycle under constant reducing pressure is adopted for CcO in the absence of NO, described by the steady state parameters  $K_{m1}$  and  $V_{max1}$ . NO is assumed to bind to both the fully reduced and the fully oxidized states of the enzyme, with the dissociation constants  $K_{icNO}$  and  $K_{uNO}$  respectively. When NO binds to the reduced heme  $a_3$  dissociation can also occur after oxidation of the gas to nitrate ion, thanks to a molecule of  $O_2$  activated by  $Cu_B$  (see text and Refs. [45,46]); this goes through a complete alternative catalytic cycle described by the steady state parameters  $K_{m2}$  and  $V_{max2}$ . NO bound to  $Cu_B$  can in turn be oxidized by the metal and be released as nitrite ion ( $NO_2^-$ ) in a reaction that does not require  $O_2$  (described by  $V_{NO}$ ). Dashed arrows indicate reactions described in the literature but that occur only to a minimal extent under our experimental conditions and can be safely neglected in the quantitative analysis of our data.  $CcO_r$ , reduced CcO;  $CcO_o$ , oxidized CcO;  $CcO_rNO$ ,  $CcO_r$  with NO bound to heme  $a_3$ ;  $O_2CcO_rNO$ ,  $CcO_r$  with NO bound to heme  $a_3$  and  $O_2$  bound to  $Cu_B$ ;  $CcO_o$ , oxidized CcO;  $CcO_oNO$ ,  $CcO_o$  with NO bound to  $Cu_B$ .

not seem to us to be major drawbacks since the aim of the model is to describe cell respiration under NO stress via a chemically sound, though oversimplified, reaction mechanism.

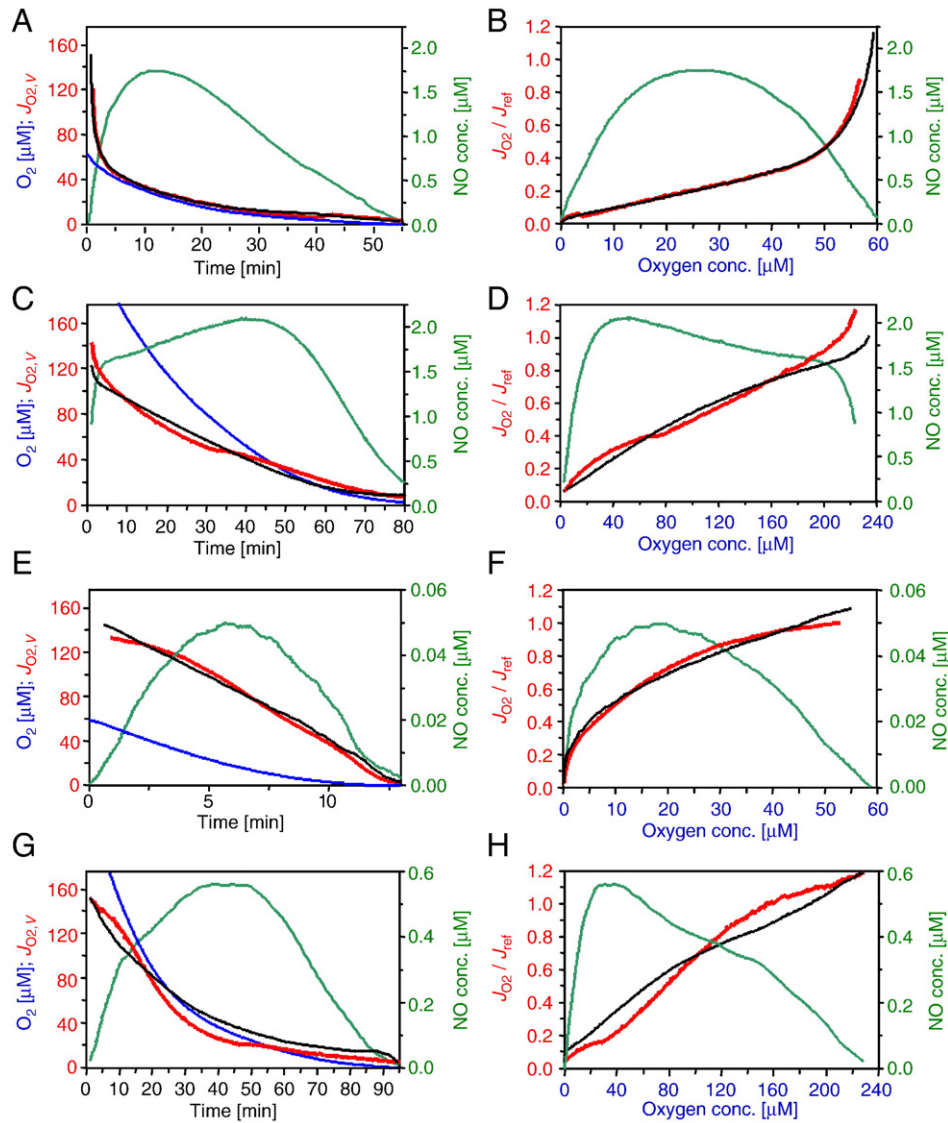
### 3.5. NO dynamics in the low- and high- $O_2$ ranges

At high levels of iNOS expression (50 ng/ml tetracycline) NO production in the low- $O_2$  range is sufficiently fast to reach high, near steady-state NO concentrations of 1.5  $\mu M$  within 10–15 min, when  $O_2$  drops to 30  $\mu M$  and becomes limiting for NO production (Fig. 5A and B). In the high- $O_2$  range, the further increase of [NO] from 10 to 40 min is modest (from 1.5 to 2  $\mu M$ ), indicating a near-balance between NO production and utilization at NO concentrations in the upper pathophysiological range (Fig. 5C and D). In contrast, at low levels of iNOS expression (20 ng/ml tetracycline) in the low- $O_2$  range, respiration is less inhibited and 30  $\mu M$   $O_2$  is reached in less than 5 min (Fig. 5E), at a time when the steep increase in [NO] continues near-linearly at high  $[O_2]$  (Fig. 5G). There is an additional period of approximately 40 min in the high- $O_2$  range, during which net NO production is positive and hence NO accumulates, until again at around 30  $\mu M$   $O_2$  the balance shifts to NO degradation and the [NO] level falls (Fig. 5G and H). A plot of critical  $[O_2]$  for limitation of NO production, against [NO] is shown in Fig. 6. Although the critical  $[O_2]$  is a complex function of iNOS expression levels and  $O_2$  range, it was actually restricted to a surprisingly narrow range under all experimental conditions.

### 3.6. Reversible inhibition and NO-induced activation of the cellular metabolic state

We performed a series of experiments to study the reversibility of the inhibition. Addition of  $HbO_2$  to cells producing high levels of NO (50 ng/ml tetracycline and 1 mM L-arginine) resulted not only in full recovery but in a significant increase in cellular respiration above the routine reference level observed before L-arginine addition (Fig. 7A). Respiration was also recovered after inhibition of iNOS activity with S-EITU (1 mM; not shown). No irreversible effects were observed up to 1.8  $\mu M$  NO. Moreover, when the cell suspension was illuminated with a cold light source at 60  $\mu M$   $O_2$ , the respiration rate immediately increased above the reference level; this reflects the photochemical dissociation of the reduced heme  $a_3$ -NO complex [35–37], again revealing the NO-induced stimulation of cell respiration (Fig. 7B). Light had no effect on respiration in the absence of NO (not shown).

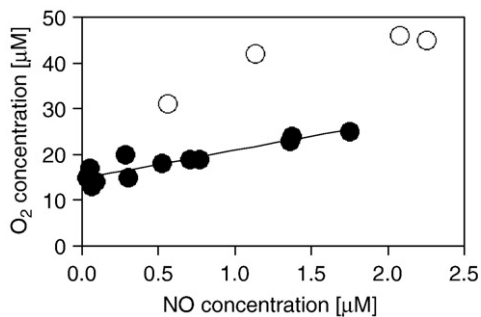
We define the stimulation factor,  $F$ , as the ratio of  $V_{max1}$  after NO removal to  $V_{max1}$  before L-arginine addition (Fig. 7C).  $F$  was usually between 1 and 1.5 and we attribute it to the accumulation of reduced cytochrome  $c$  and a decline in mitochondrial membrane potential caused by CcO inhibition. Stimulation causes  $V_{max1}$  (and  $V_{max2}$ ) to be variable, rather than constant, parameters; however, our model assumes constant reducing pressure and neglects the stimulation effect. Nevertheless, stimulation is significant only at high [NO], where  $V_{max2}$  predominates, while at low [NO] the absence of stimulation is associated with predominance of  $V_{max1}$ . Therefore the stimulation effect is partially reflected in the ratio of  $V_{max2}$  to  $V_{max1}$  ( $r = V_{max2}/V_{max1}$ ). A plot of normalized fluxes predicted from the kinetic model plotted against experimentally determined fluxes is depicted in Fig. 7D.



**Fig. 5.** NO production and NO-induced inhibition of respiration over a standard (A,B,E,F) or extended (C,D,G,H) [O<sub>2</sub>] range. O<sub>2</sub> consumption and NO production were measured at high (50 ng/ml tetracycline; A–D) and low (20 ng/ml tetracycline; E–H) iNOS activities. (A,C,E,G) Oxigraph-2k traces of [O<sub>2</sub>] (blue), O<sub>2</sub> flux (red) and [NO] (green) as a function of time, with data points plotted at 1 s intervals. (B,D,F,H) Relative measured flux (red), estimated flux from the kinetic model (black) and [NO] (green) as a function of [O<sub>2</sub>]. Traces are representative of three similar experiments.

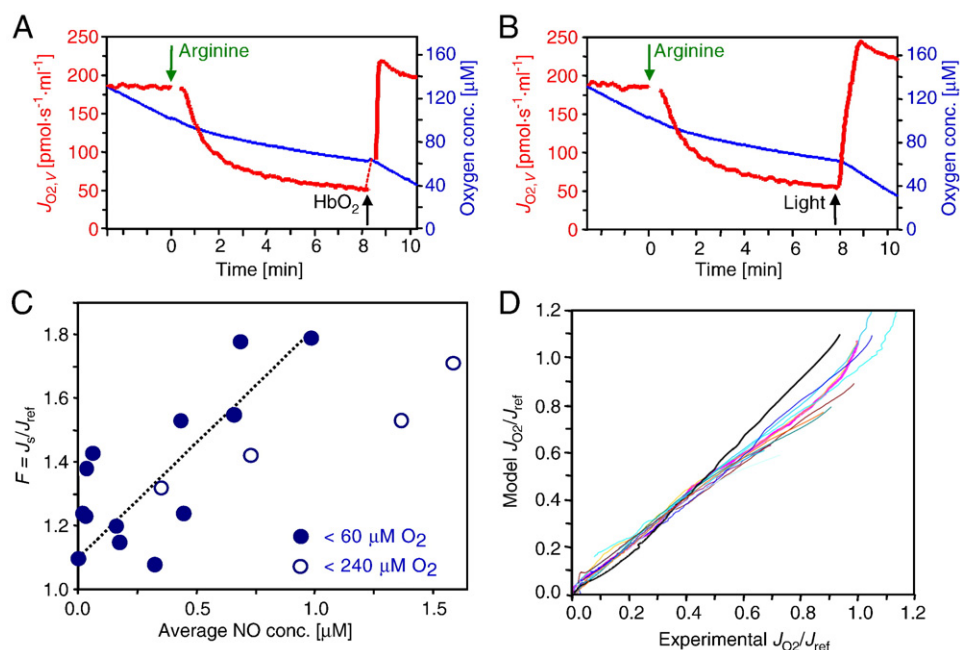
#### 4. Discussion

We present a detailed study of the O<sub>2</sub> kinetics of cellular respiration at varying concentrations of endogenous NO over the pathophysiological



**Fig. 6.** Critical [O<sub>2</sub>] for limitation of NO production. Critical [O<sub>2</sub>] for net NO production, below which [NO] declines. High [NO] corresponds to high iNOS activities in the low- and high-O<sub>2</sub> ranges (full and open circles, respectively).

range. Based on the results, we propose a kinetic model of NO-induced inhibition of cellular respiration that explains the experimental data and has predictive value. The model assumes that the system can be treated under the pseudo-equilibrium approximation and that two cycles, each characterized by its own  $K_m$  and  $V_{max}$ , are possible, one for NO-free and one for NO-bound CcO. The pseudo-equilibrium assumption is unrealistic, but greatly simplifies the analysis at the expense of some parameters becoming apparent, rather than real, kinetic constants. The model invokes both competitive inhibition (at the heme  $a_3$  site) and uncompetitive inhibition (at Cu<sub>B</sub>), as previously described [25–28]. The details of the complex catalytic cycle of CcO are ignored, in particular in relation to the reduction intermediates.  $V_{max1}$  corresponds to routine respiration of the intact cell at saturating [O<sub>2</sub>] in the absence of NO. Routine respiration is regulated at a modest level of activity, significantly below the maximum capacity of the electron transport system, and thus below the true  $V_{max}$  for CcO [41], as shown by the results listed in Table 1 and by the NO-induced stimulation of respiration (Fig. 7A and B). Notably, increases in [NO] that are unable to inhibit cell respiration reduce CcO, and the excess capacity of the enzyme is able to maintain O<sub>2</sub> consumption in the cell [47].



**Fig. 7.** Stimulation of respiration following NO removal. (A,B) Oxygraph traces of [O<sub>2</sub>] (blue) and O<sub>2</sub> flux (red) in cells treated with tetracycline (50 ng/ml), showing the inhibition of respiration induced by the addition of L-arginine (1 mM). Subsequent addition of HbO<sub>2</sub> (A) or illumination with a cold light source (B) stimulated respiration ( $J_g$ ) above the initial reference level. (C) Stimulation factor ( $F = J_g/J_{ref}$ ) as a function of [NO]. (D) Normalized fluxes predicted from the kinetic model (hyperbolic approximation, see Supporting information) plotted against experimentally determined fluxes.

The model is designed to incorporate two important observations from previous studies on purified CcO or its model compounds. First, it allows NO bound to oxidized CcO to act as a slow reductant, rather than as an inhibitor, by donating its unpaired electron. The resulting nitrosonium ion (NO<sup>+</sup>), dissociates *via* conversion to nitrite (NO<sub>2</sub><sup>-</sup>) in a non-redox reaction with a water molecule [27,48]. Second, the model includes an NO-dissociation cycle (for the derivative CcO<sub>2</sub>NO) that consumes O<sub>2</sub>, albeit with a high  $K_m$  ( $K_{m2}$ ). This can account for the low O<sub>2</sub> consumption we observed experimentally at high [NO]. Although this reaction has not been observed unequivocally in isolated CcO, and was not detected in specifically designed experiments [49], it has been suggested by Pearce et al. [45], and has been directly demonstrated for a model compound mimicking the binuclear site of CcO synthesized by Collman et al. [46]. Our data cannot resolve this issue, due to the complexity of the system studied (whole cells, rather than purified CcO). However, we unequivocally observed respiration at NO concentrations that should fully inhibit CcO activity in the absence of a mechanism like the one proposed in Fig. 4 and Refs. [45,46].

Our model fits the experimental data and is compatible with what is known about the interaction between CcO and NO [25,45,46]; however, it may not provide a complete description of the observed O<sub>2</sub> fluxes in intact living cells. Other reactions between O<sub>2</sub> and NO are possible, either spontaneous or catalyzed by cell components other than CcO. These reactions, as well as functions played by other cellular components, would contribute to O<sub>2</sub> consumption and in our model would be wrongly attributed to CcO. Nonetheless, our model provides a chemically sound quantitative description of our data and has predictive value. We also tested a variant model, in which O<sub>2</sub> oxidizes the NO molecule bound to Cu<sub>B</sub>, rather than the one bound to heme a<sub>3</sub> [50]. Although this variant accurately fits the experimental data, it was discarded because it lacks independent support; indeed no such reaction was reported for the synthetic model of CcO [46].

The most important feature of the model presented here is that the relative population of the catalytically active species O<sub>2</sub>CcO<sub>2</sub>NO (represented by the term  $[O_2][NO]/K_{m2}K_{icNO}$  in Eq. (1)) is propor-

tional to the product of the concentrations of NO and O<sub>2</sub>. This provides an explanation for how the enzyme can be active at relatively high inhibitor concentrations and also explains the non-linear IC<sub>50</sub> vs. [O<sub>2</sub>] plot (Fig. 3B). An IC<sub>50</sub> vs. [O<sub>2</sub>] plot similar to ours was presented by Koivisto et al. [24]; however, these authors proposed that two O<sub>2</sub> molecules need to bind to CcO to compete with NO, which is unlikely. Our hypothesis provides a sounder solution to this problem. More recently, using published data mainly obtained with NO donors, an IC<sub>50</sub> vs. [O<sub>2</sub>] plot similar to ours has been reported [51].

In conclusion, we show the dependence of NO production on [O<sub>2</sub>] and the interplay between O<sub>2</sub> and NO concentrations in determining cell respiration. In spite of its complexity, this interplay can be described by a relatively simple and chemically sound kinetic model. The control of respiration by NO regulates O<sub>2</sub> gradients in the cell, redox signaling and apoptosis [52]. Furthermore, interactions of NO with mitochondria have important implications for ischemic biology [33,53] and hypoxic vasodilation [54], and their understanding will help clarify the mechanisms by which NO exerts cardioprotective actions [55].

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## Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.bbabo.2010.01.033.

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