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Hydrogen Peroxide Induces Nitric Oxide and Proteasome Activity in Endothelial Cells: A Bell-Shaped Signaling Response

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Abstract

We investigated nitric oxide ($\cdot\text{NO}$)-mediated proteasomal activation in bovine aortic endothelial cells (BAEC) treated with varying fluxes of hydrogen peroxide (H_2O_2) generated from glucose/glucose oxidase (Glu/GO). Results revealed a bell-shaped $\cdot\text{NO}$ signaling response in BAEC treated with Glu/GO (2–20 mU/ml). GO treatment (2 mU/ml) enhanced eNOS phosphorylation and $\cdot\text{NO}$ release in BAEC. With increasing GO concentrations, phospho eNOS and $\cdot\text{NO}$ levels decreased. A bell-shaped response in proteasomal function and $\cdot\text{NO}$ induction was observed in BAEC treated with varying levels of GO (2–10 mU/ml). Proteasomal activation induced in GO-treated BAEC was inhibited by L-NAME pretreatment, suggesting that $\cdot\text{NO}$ mediates proteasomal activation. Intracellular $\cdot\text{NO}$ induced by H_2O_2 was detected by isolating the DAF-2/ $\cdot\text{NO}/\text{O}_2$ -derived “green fluorescent product” using the HPLC-fluorescence technique, a more rigorous and quantitative methodology for detecting the DAF-2/ $\cdot\text{NO}/\text{O}_2$ reaction product. Finally, the relationship between H_2O_2 flux, proteasomal activation/inactivation and endothelial cell survival and apoptosis is discussed.

Keywords

hydrogen peroxide; nitric oxide; proteasome activation; apoptosis; diaminofluorescein; DAF-2; HPLC

Introduction

The chemical biology of the reactive oxygen species (ROS) (e.g., superoxide, hydrogen peroxide) and their signal transduction mechanisms in physiological and pathophysiological processes are complex, intriguing and continuously evolving [1–5]. Hydrogen peroxide (H_2O_2) is generated in most mammalian cells as a byproduct of oxygen metabolism. Low levels of ROS generation can stimulate cell growth and cell proliferation [6–8]. However, excessive

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generation of ROS under pathophysiological conditions (inflammation, cardiovascular and neurodegenerative disorders) can induce cell injury due to oxidative damage to lipid membranes, proteins and DNA. Thus, there is a great deal of interest in research pertaining to the regulation of antioxidant enzymes (superoxide dismutases and peroxidases) and other processes (proteasomal activation and ubiquitinylation) that are responsible for repairing oxidatively-damaged lipids, protein, and DNA [9,10]. H₂O₂ has been suggested to be a “second messenger” in redox-signaling events [11,12]. Recent research suggests that H₂O₂ can stimulate calcium, *NO, and cGMP/cAMP signaling in vascular endothelial and smooth muscle cells [13,14]. Emerging research indicate that *NO/cGMP/cAMP pathway is responsible for upregulating the proteasomal signaling mechanism in endothelial cells [15]. The proteasomal machinery serves an important antioxidant function in repairing oxidatively damaged proteins and lipids [10]. Thus, it is important to understand oxidant-induced regulation of the proteasomal function.

Endothelial injury is an early oxidative injury in several vascular diseases [16–18]. Endothelial cells are exposed to H₂O₂ and other lipid peroxides originating from shear stress and from leukocytes and macrophages [19,20]. The mechanism(s) by which peroxides induce endothelial dysfunction are not fully understood [3,21]. H₂O₂ induces transcriptional activation of eNOS in endothelial cells [5,13,22,23]. The cytoprotective effects of *NO against H₂O₂-mediated cellular toxicity have been attributed to iron chelation, radical scavenging or restoration of mitochondrial respiration [24–26]. At higher concentrations of H₂O₂, the “uncoupling” of eNOS was reported to stimulate additional superoxide formation [27]. To more fully define the peroxide-mediated downstream signaling pathway, we investigated the dose-response relationship between H₂O₂, *NO formation and proteasomal activation in BAEC treated with glucose/glucose oxidase (Glu/GO).

The present results suggest that H₂O₂ stimulates a bell-shaped, *NO -mediated proteasomal signaling response in endothelial cells. At low levels of H₂O₂, there was an increase in *NO signaling. This signaling mechanism was abolished in cells treated with higher concentrations of H₂O₂. The present data reveal a new mechanistic perspective in the oxidant-induced regulation of signal transduction processes involved in cell survival and cell death. In addition, we have described in this study a HPLC-fluorescence methodology to detect and unambiguously quantitate the DAF-2/*NO/O₂-derived fluorescent product. The formation of DAF-2 triazole product, as detected by HPLC-fluorescence, is more sensitive to NOS inhibitors as compared to optical microscope technique.

Materials and Methods

Materials

Glucose oxidase, clasto-lactacystin- β -lactone, N_ω-Nitro-L-arginine-methyl ester (L-NAME) and N^G-monomethyl-L-arginine monoacetate (L-NMMA) were obtained from Sigma Chemical Co (St. Louis, MO). 4,5-Diaminofluorescein (DAF-2), DAF-2 diacetate (DAF-2DA) and DAF-2-triazole (DAF-2T) were purchased from Calbiochem (San Diego CA). The polyclonal antibodies directed against phospho-eNOS (Ser1177), Akt, and phospho-Akt (Ser473) were purchased from Cell Signaling Technology (Beverly, MA). Cell culture reagents including L-glutamine, penicillin, streptomycin and fetal bovine serum (FBS) were obtained from Invitrogen (Carlsbad, CA). All other chemicals used were of analytical grade.

Endothelial cell culture

Bovine aortic endothelial cells (BAEC) were obtained from Clonetics (San Diego, CA) and maintained (37°C, 5% CO₂) in Dulbecco’s modified Eagle’s medium (DMEM) containing 10% FBS, L-glutamine (4 mM), penicillin (100 units/ml), streptomycin (100 μ g/ml) and

glucose (25 mM). Cells were obtained between passages 2 and 10. One day before treatment the medium was replaced with DMEM containing 2% FBS. GO (1–20 mU/ml) was added to cells in the medium containing 25 mM Glu. H₂O₂ (1 μM/min) was generated from Glu/GO (20 mU/ml) in 10 ml DMEM containing 2% FBS, and this was measured by using a YSI Model 25 oxidase meter (Yellow Springs Instruments). Nitric oxide synthase inhibitors [L-NAME (2 mM) and L-NMMA (200 μM)] and proteasome inhibitor [lactacystin-β-lactone (10 μM)], were added 2 h before GO treatment.

Intracellular conversion of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) to formazan was used as an indicator of cell viability ($\lambda_{\max} = 562 \text{ nm}$). After treatment with H₂O₂, the culture media were removed, and cells were washed twice with Dulbecco's phosphate-buffered saline (DPBS). Cells were incubated with the culture medium (DMEM) alone without FBS containing 0.25 mg/ml MTT for 1 h at 37°C. Following this, the medium was removed. Cells were washed again with DPBS and finally dissolved in dimethyl sulfoxide (DMSO). The amount of formazan formed was measured spectrophotometrically as described previously [28].

Measurement of caspase-3 activity

The cytosolic enzymatic activity of caspase-3 was measured as described previously [29]. Briefly, cells were washed twice with DPBS following treatment with H₂O₂ and then lysed with 50 mM HEPES buffer (pH 7.4) containing 5 mM CHAPS and 5 mM dithiothreitol. After removing the cytosolic fraction by centrifugation at 12,000 × *g* for 30 min, the activity of caspase-3 was measured using the substrates of ac-DEVD-*p*NA (acetyl-Asp-Glu-Val-Asp-*p*-nitroanilide). The absorbance at 405 nm of the released *p*NA was monitored in a spectrophotometer and quantitated using a *p*NA standard.

Measurement of apoptosis by TUNEL assay

The terminal deoxynucleotidyl transferase-mediated nick-end labeling (TUNEL) assay was used for microscopic detection of apoptosis [30]. This assay is based on labeling of 3'-free hydroxyl ends of the fragmented DNA with fluorescein-dUTP catalyzed by terminal deoxynucleotidyl transferase. Procedures were followed according to a commercially available kit (ApoAlert) from Clontech. Apoptotic cells exhibit a strong nuclear green fluorescence that can be detected using a standard fluorescein filter (520 nm). All cells stained with propidium iodide exhibit a strong red cytoplasmic fluorescence at 620 nm. The areas of apoptotic cells were detected by fluorescence microscopy equipped with rhodamine and FITC filters. The quantification of apoptosis was performed using the Metamorph image analysis package.

Western blot analysis

BAEC were washed with ice-cold phosphate buffered saline (PBS) and homogenized in 100 μl of radioimmunoprecipitation assay buffer (20 mM Tris-HCl, pH 7.4/2.5 mM EDTA/1% Triton X-100/1% sodium deoxycholate/1% SDS/100 mM NaCl/100 mM sodium fluoride) containing 1 mM sodium *ortho*-vanadate and a mixture of protease inhibitors. The homogenate was centrifuged at 750 × *g* for 10 min at 4°C to pellet out the nuclei. The remaining supernatant was centrifuged for 30 min at 12,000 × *g*. Proteins were resolved on 8% for TrR, eNOS, phospho eNOS, Akt and phospho Akt by SDS/PAGE gels and blotted onto nitrocellulose membranes. Membranes were probed with primary antibodies and then incubated with horseradish peroxidase-conjugated rabbit anti-mouse IgG secondary antibody. Protein bands were detected by using the ECL method (Amersham Pharmacia).

Measurement of intracellular \bullet NO

Intracellular \bullet NO levels were monitored using the DAF-2 fluorescence probe [31,32]. The treated cells were washed with DPBS and incubated in 2 ml of fresh culture medium without FBS. DAF-2 was added at a final concentration of 5 μ M, and the cells were incubated for 1 hr. The cells were washed twice with DPBS and maintained in 1 ml of culture medium. Fluorescence was monitored using a Nikon fluorescence microscope (excitation, 488 nm; emission, 610 nm) equipped with an FITC filter. The values of fluorescent intensity were calculated using the Metamorph software.

HPLC method

The HPLC method to detect intracellular \bullet NO is more quantitative and specific, as DAF-2T (the product of \bullet NO/O₂ and DAF-2) can be readily isolated and its concentration determined by comparison with an authentic standard. BAEC were treated with different concentrations of GO before incubation with DAF-2DA. Cells were subsequently washed twice with DPBS and treated with 5 μ M DAF-2DA for 1 hr in DMEM medium containing 2% FBS. The medium was removed and cells were washed twice with DPBS and then harvested with DPBS. After centrifugation (5 min \times 4,000 rpm) and removal of supernatant liquid, the pellets were stored at -80° C until the day of analysis by HPLC. On the day of analysis, the pellets were thawed to an ambient temperature and cells were lysed with 250 μ l of phosphate buffer (10 mM, pH 7.5) containing 100 μ M diethylenetriaminepentaacetic acid (DTPA) and 0.1% Triton X 100. Lysates (5 μ l) were taken for protein measurement. To the remaining solution, 1-butanol (0.5 ml) was added and the mixture was vortexed for 10 min and then centrifuged (2 min \times 5,000 rpm). An aliquot (450 μ l) from n-butanol phase (upper layer) was separated and dried in a MultiVap analytical evaporator (Organomation) at 30° C using air flow. The dried samples were reconstituted in 50 μ l of phosphate buffer (10 mM) containing 100 μ M DTPA and used for HPLC analysis.

DAF-2, DAF-2T and other DAF-2DA - derived products (Fig. 1S in Supplementary Information) were separated on an Agilent 1100 HPLC system equipped with fluorescence and UV-Vis absorption detectors. Typically, 10 μ l of sample was injected into the HPLC system with a C₁₈ column (Agilent, Zorbax 80 \AA , Extend-C18, 4.6 \times 250 mm, 5 μ m) equipped with a guard column. The compounds of interest were separated by isocratic elution using an aqueous mobile phase consisting of potassium phosphate buffer pH 7.5 (10 mM) and acetonitrile (5% by volume) at a flow rate of 1 ml/min. Fluorescence detection at 490 nm (excitation) and 515 nm (emission) and absorbance changes at 220, 250, 280, 320, 490 nm were used to monitor the products.

Nitrate measurement

Nitrate, an oxidative metabolite of \bullet NO, was measured by chemiluminescence according to a modified method [33]. Briefly, after treatment of BAEC with GO, the culture medium was collected, and the released nitrate was detected using the \bullet NO analyzer. Protein concentration in each well was measured according to the Bradford method. For nitrate measurement, a solution containing vanadium and HCl was used. \bullet NO generated from sodium nitrate was used as standard.

26S proteasome activity

Proteasome function was measured as reported [34,35]. Briefly, cells were washed with buffer I (50 mM Tris, pH 7.4/2 mM DTT/5 mM MgCl₂/2 mM ATP) after treatment, and homogenized with buffer I containing 250 mM sucrose. Twenty micrograms of 10,000 \times g supernatant were diluted with buffer I to a final volume of 900 μ l. The fluorogenic proteasome substrates SucLLVY-AMC (chymotrypsin-like) and Z-Leu-Leu-Lys-AMC (trypsin-like) were added in

a final concentration of 80 μM . Proteolytic activity was measured by monitoring the release of the fluorescent group 7-amido-4-methylcoumarin (excitation 380 nm, emission 460 nm).

20S proteasome activity

Activity of the 20S proteasome was determined according to Grune *et al.* (36). Cells were lysed in PBS containing 0.1% Triton X-100 and 0.5 mM DTT. The assay mixture contained 50 μl of buffer (50 mM Tris-HCl, pH 7.8/20 mM KCl/5 mM MgCl_2 /0.1 mM DTT), 250 μM sLLVY-MCA, and 50 μl of cell lysate (15 μg of protein). After 30 min at 37°C, the reaction was stopped by adding 1 ml of 0.2 M glycine buffer, pH 10, and the fluorescence of the liberated 7-amido-4-methylcoumarin was measured using the excitation and emission wavelengths at 365 and 460 nm, respectively.

Statistical analysis

Data are presented as mean \pm SD. Comparisons between different treatments were performed using a Student's t-test. p values less than 0.05 were considered to be significant.

Results

H_2O_2 induces a bell-shaped nitric oxide signaling response

Endothelial cells were treated with different concentrations of GO (1–20 mU/ml). The rate of generation of H_2O_2 under these conditions varied from 0.05 $\mu\text{M}/\text{min}$ to 1 $\mu\text{M}/\text{min}$ [21]. Figure 1A shows the Western blots of phospho eNOS (phosphorylation at Ser1179), eNOS protein, phospho Akt and Akt induced in BAEC treated with a range of GO concentrations. Exposure to lower levels of GO (1–2 mU/ml) increased eNOS phosphorylation at Ser1179 (and not at Thr497) in BAEC, followed by a gradual decline in phospho eNOS induction in cells treated with increasing levels of GO (10–20 mU/ml). A similar trend was also noticed with total eNOS protein expression. These results are in agreement with the published report showing eNOS activation by a PI_3 -kinase dependent phosphorylation at Ser1177 [5]. The present data indicate that H_2O_2 regulates Akt phosphorylation in a dose-dependent manner. As reported previously [5], we observed an increase in phosphoAkt (phosphorylated at Ser473) in BAEC treated with H_2O_2 generated at a lower rate, followed by a decrease in phosphoAkt expression in cells exposed to a higher rate of H_2O_2 generation. There was no detectable change in the total Akt levels under these conditions (Fig. 1A). These results suggest that treatment of endothelial cells with low levels of peroxide activate a unique signal transduction mechanism that was absent in cells treated with high peroxide levels.

Next we investigated whether changes in the phosphorylation status of eNOS protein parallel the increase in its activity. To this end, $\bullet\text{NO}$ levels were examined, using different techniques, in cells exposed to varying H_2O_2 fluxes. Treatment with low GO levels (1–5 mU/ml) enhanced $\bullet\text{NO}$ release in BAEC, as monitored by nitrate measurements in the medium (Fig. 1B), intracellular DAF-2-derived “green” fluorescence (Fig. 1C and D) and by HPLC analysis of intracellular DAF-2T product (Fig. 1E). All of these indicators of $\bullet\text{NO}$ decreased considerably in cells treated with 20 mU/ml. Previously, it had been shown that DAF-2 forms a fluorescent triazole-type product in the presence of an oxidant derived from $\bullet\text{NO}$ and oxygen interaction [37,38].

To investigate the effect of L-NAME (a non-specific inhibitor of all forms of NOS enzymes) on $\bullet\text{NO}$ formation, BAEC were pretreated with L-NAME for 2 h and then treated with GO (2 mU/ml) for 4 h. As shown in Figure 2A and B, L-NAME pretreatment decreased the green fluorescence by about 20% in BAEC treated with 2 mU/ml of GO. In contrast, L-NAME pretreatment dramatically lowered the intracellular formation of DAF-2T, the characteristic marker product of DAF-2, $\bullet\text{NO}$, and oxygen (Fig. 2C). Figure 2E shows the actual HPLC

chromatograms of the standards DAF-2 and DAF-2T and an unknown peak. The advantage of monitoring $\cdot\text{NO}/\text{DAF-2}/\text{O}_2$ product by HPLC is that it enables us to monitor the intracellular uptake and metabolism of DAF-2DA. Although DAF-2 exhibits a much weaker fluorescent intensity (nearly three orders of magnitude less than DAF-2T), DAF-2 concentration in cells is nearly three orders of magnitude higher than that of DAF-2T. Thus, DAF-2 can contribute significantly to the intracellular fluorescence measured by microscopy. This may explain the much lower sensitivity to L-NAME of the fluorescence microscopy technique as compared to the HPLC assay. On the other hand the peak height due to DAF-2 is nearly identical during the various treatment conditions (Fig. 2E). This suggests that the observed changes in DAF-2T are not due to the differential cellular uptake of DAF-2DA. Figure 2D shows the levels of nitrate measured in the extracellular media under various conditions (Fig. 2A). We used the NOS inhibitor, L-NMMA instead of L-NAME, because L-NAME interfered with nitrate measurement by chemiluminescence. L-NMMA decreased the DAF-2 green fluorescence, albeit to a moderate extent (data not shown). These results suggest that DAF-2 green fluorescence, as monitored by fluorescence microscopy, could also originate by another oxidative route that is independent of $\cdot\text{NO}$ formation under these conditions. As shown in Figure 2E, another green fluorescent product appeared at 4 min in the presence of 20 mU/ml GO. In contrast, DAF-2T measurement by HPLC in cells is more sensitive to changes in intracellular $\cdot\text{NO}$ than the fluorescence microscopic technique.

H₂O₂ induces a bell-shaped proteolytic signaling response-Intermediacy of $\cdot\text{NO}$

The proteasomal pathway is a major clearing route for damaged proteins, and failure of its function increases the accumulation of abnormal proteins [39]. The proteasomal function was assessed by measuring the activities of two of the most important proteasomal enzymes, chymotrypsin and trypsin. Both 26S and 20S proteasomal functions were measured in BAEC treated with a range of GO concentrations, as described previously [40]. GO treatment at concentrations ranging from 1–5 mU/ml significantly increased the proteasomal function (both 26 and 20S), whereas at a higher concentration (20 mU/ml) the proteasomal function significantly decreased (Figure 3A–B and 3C–D). Proteasomal activities were inhibited in the presence of L-NMMA or lactacystin. It was found that all these interventions could significantly prevent the proteasomal activation (26S and 20S) induced by 2mU/ml GO (Figure 4A–B and 4C–D). We independently confirmed this result by monitoring the proteasomal degradation of TfR in endothelial cells. Immunoblot studies using the antibody directed against TfR showed an accumulation of this protein in cells treated with a higher concentration of GO (20 mU/ml) and that this accumulation was not detected in cells treated with lower levels of GO (Figure 3E).

Lactacystin, a specific inhibitor of proteasome, was shown to induce eNOS expression in cells [41], and thus the effect of this inhibitor was investigated in cells exposed to 2 mU/ml of GO. As shown in Figure 2S in Supplementary Information, eNOS expression increased in the presence of 10 μM lactacystin and further increased with both lactacystin and 2 mU/mL GO. Functional $\cdot\text{NO}$ was also detected under these conditions by measuring the DAF-2T product and nitrate release (Figure 2S).

Time-dependent $\cdot\text{NO}$ and proteosomal activation in BAEC treated with GO

BAEC were treated with both low and high (2 mU/ml and 20 mU/ml) concentrations of GO for different time points ranging from 2–24 h, and $\cdot\text{NO}$ and proteasome activities were measured. As shown in Figure 5, nitrate (A), DAF-2T (B), and eNOS (E) levels were elevated up to 8 h and maintained this level even at 24 h in BAEC exposed to GO (2 mU/ml), and nitrate and DAF-2T but not eNOS protein decreased at 24 h. In contrast, there was a significant decrease after 2 h in eNOS expression, nitrate and DAF-2T levels in cells treated with 20 mU/ml of GO during the same time period. Both 26S and 20S proteasomal activities were increased

in BAEC treated with 2 mU at 2 h time point, and this increase was maintained for up to 24 h (Fig. 6A–D).

Effect of low versus high concentrations of H₂O₂ on endothelial cell viability and apoptosis

BAEC were treated with H₂O₂ at concentrations ranging from 1 to 20 mU/ml GO and the cell viability was assessed by the MTT assay. The cell viability was virtually not affected in cells exposed to low levels of GO (1–10 mU/ml), whereas almost 70% of the cells were not viable in the presence of a higher concentration of GO (20 mU/ml) (Fig. 7B). Under these conditions, the caspase-3 activity was measured, which was significantly elevated in cells treated with 20 mU/ml of GO (Fig. 7A). Apoptosis was further confirmed by measuring DNA fragmentation using the TUNEL assay (Fig. 7C and D).

Discussion

In this work, we report that the exposure of endothelial cells to low concentrations of H₂O₂ stimulates •NO-mediated proteasomal signaling. There was, however, a decrease in •NO and proteasomal activation in cells treated with higher H₂O₂ fluxes. The fact that H₂O₂ induces a bell-shaped signaling response has important implications in endothelial cell survival and endothelial apoptosis induced by peroxides (Fig. 8).

Peroxide-induced endothelial NOS signaling

Previous studies have shown that exogenously-added H₂O₂ is a potent stimulator of •NO production in endothelial cells [5]. It was shown that both Ca²⁺ and calmodulin kinase II were involved in the signaling cascade linking H₂O₂ to eNOS induction [23]. More recently, endogenous levels of H₂O₂ generated in response to pathophysiological stimulus, angiotensin II, was reported to be sufficient to stimulate endothelial •NO formation [42]. The NADPH oxidase was shown to be responsible for angiotensin II-mediated H₂O₂ formation. Induction of eNOS by H₂O₂ was postulated as an important response to vascular oxidative stress [42].

H₂O₂-induced proteasomal function: Role of •NO

Recently, we reported that the •NO/cGMP/cAMP pathway can directly upregulate immunoproteasomal subunits (LMP2 and LMP7), thereby stimulating the proteasomal activities in endothelial cells [15]. Proteasomal activities were increased in BAEC treated with an •NO donor or with cell-permeable analogs of cGMP or cAMP. This increase in proteasomal activity was inhibited by specific adenylyl and guanylyl cyclase inhibitors [15]. Cells pretreated with protein kinase G and protein kinase A inhibitors markedly decreased •NO-dependent proteasome activation. The •NO/cGMP/cAMP signaling mechanism enhanced the phosphorylation of the transcription factor cAMP-response element-binding (CREB) protein, the cAMP-response element promoter activity and the expression of the immunoproteasomal subunits. In the present work, low levels of H₂O₂ induce •NO-dependent proteasomal activation. The regulatory mechanisms by which low levels of H₂O₂ upregulate proteasomal activities are proposed to be similar to the •NO-mediated signaling pathway. •NO as a stimulator of proteasomal function is an emerging concept [15]. The antioxidative and cytoprotective effects of •NO were attributed to proteasomal activation, as proteasomal inhibitors counteracted •NO-induced proteasomal activation [15]. Previous studies have shown that pretreatment of cells with antioxidants increased •NO stimulation and proteasomal function [43]. This is attributed to a decrease in ROS (e.g., H₂O₂) that in turn stimulates NOS signaling.

Previously, we reported that treatment of endothelial cells with a higher flux of H₂O₂ using GO (20 mU/ml) increased the expression of TfR levels leading to enhanced iron uptake and cellular apoptosis [21]. We also reported that this oxidant-induced iron signaling was effectively antagonized by exogenously-added •NO donors [40]. This effect was attributed to

an increase in proteasomal activation and the subsequent degradation of TfR levels on the cell surface. The present data further strengthens our previous suggestion [40], in that a lower flux of H₂O₂ enhanced eNOS protein phosphorylation and •NO-mediated proteasomal activation. This process, in turn, regulates the accumulation of the oxidatively-modified proteins (e.g., TfR). However, in the presence of a higher flux of H₂O₂ the proteasomal activities were decreased, leading to increased accumulation of modified proteins and cell death.

Although the exact mechanism by which higher flux of H₂O₂ inhibits proteasome is not known, it is likely that this may be due to an increase in phosphatases activity resulting in eNOS dephosphorylation. Preliminary data indicate that okadaic acid reversed the proteasomal inactivation observed at higher flux of H₂O₂ (data not shown). Additional experiments are needed to fully explain these effects.

A bell-shaped response in redox biology

There exist several examples in oxidative pathophysiology where a bell-shaped dose-response was observed with respect to changes in the levels of antioxidant enzymes and antioxidant enzyme mimetics [44–46]. Early on, it was shown that MnSOD exhibited a bell-shaped dose-response in its protection against ischemic injury in isolated rabbit heart [44,45]. The bell-shaped dose-response behavior of SOD was invoked to explain the paradoxical overproduction of protein carbonyls in cells with varying amounts of SOD [47]. As H₂O₂ is formed as an end-product in most pathophysiological processes, the bell-shaped cell signaling response reported for H₂O₂ could have major implications in understanding the cytotoxic and cytoprotective mechanisms of antioxidant enzymes and antioxidant enzyme mimetics in disease models. Biphasic redox-signaling was demonstrated in endothelial cells treated with ceramides [43]. Ceramides, a group of naturally occurring sphingolipid second messenger molecules, induce both •NO and ROS in endothelial cells [43]. Lower levels of ceramide (5–20 μM) induced •NO generation, whereas exposure to higher concentrations (>20 μM) induced ROS in endothelial cells. •NO stimulation in endothelial cells exposed to lower concentrations of ceramide was attributed to Ca²⁺ activation and translocation of eNOS [14]. Antioxidants such as Mito-Q enhanced •NO-mediated effects (i.e., proteasomal function) in BAEC treated with lower concentrations of ceramide [43]. Under these conditions, it is likely that ceramide effects were mediated through a biphasic ROS signaling [43]. More recently, superoxide was shown to counteract the signaling effects mediated by •NO [48].

Finally, we report in this study that although DAF-2T (the authentic product of the reaction between DAF-2, •NO, and molecular oxygen) exhibits a characteristic green fluorescence, DAF-2 is also oxidatively converted to other green fluorescent products *via* a •NO-independent mechanism. In addition, DAF-2 itself can contribute significantly to the intracellular green fluorescence (despite its much lower fluorescence quantum yield) due to its enhanced intracellular concentration as compared to DAF-2T. Consequently, the HPLC-fluorescence is a more reliable method for identification and quantitation of DAF-2T, the product of •NO/O₂ reaction with DAF-2.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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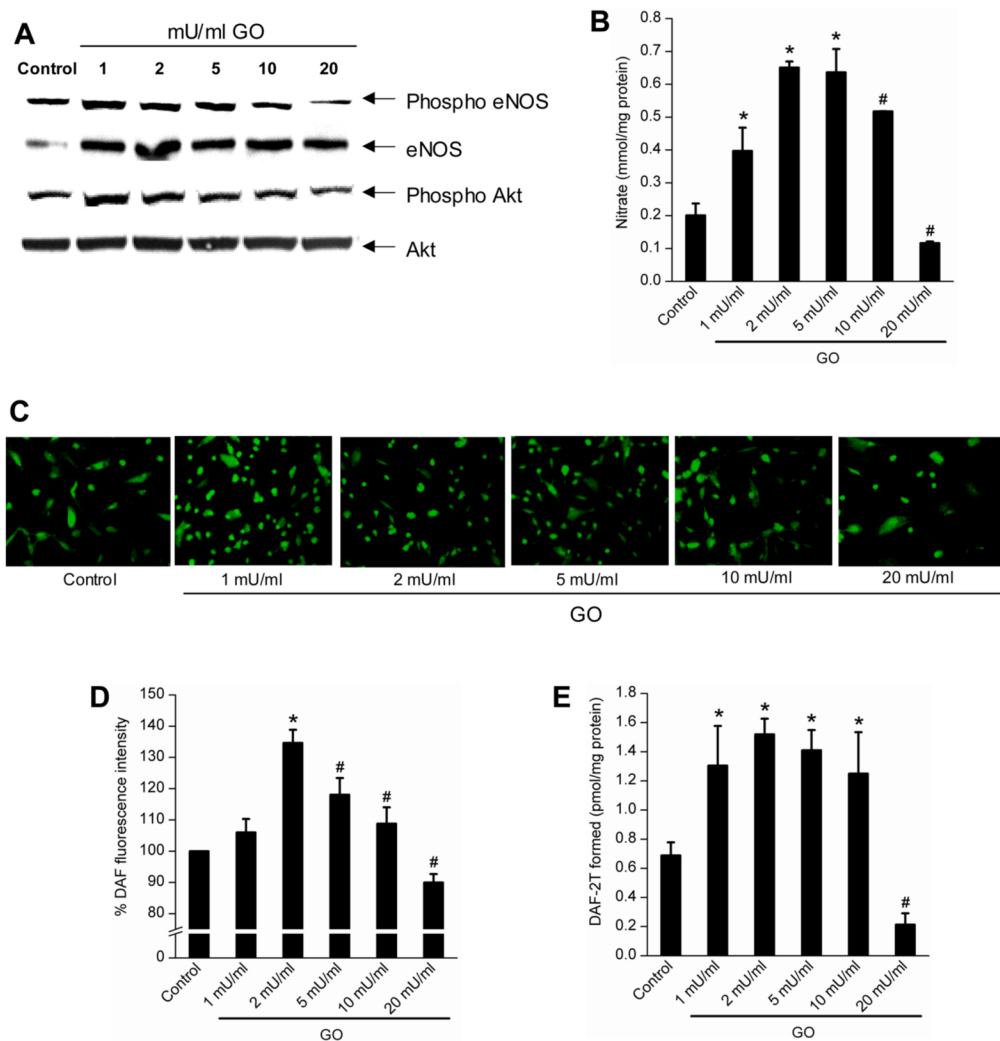
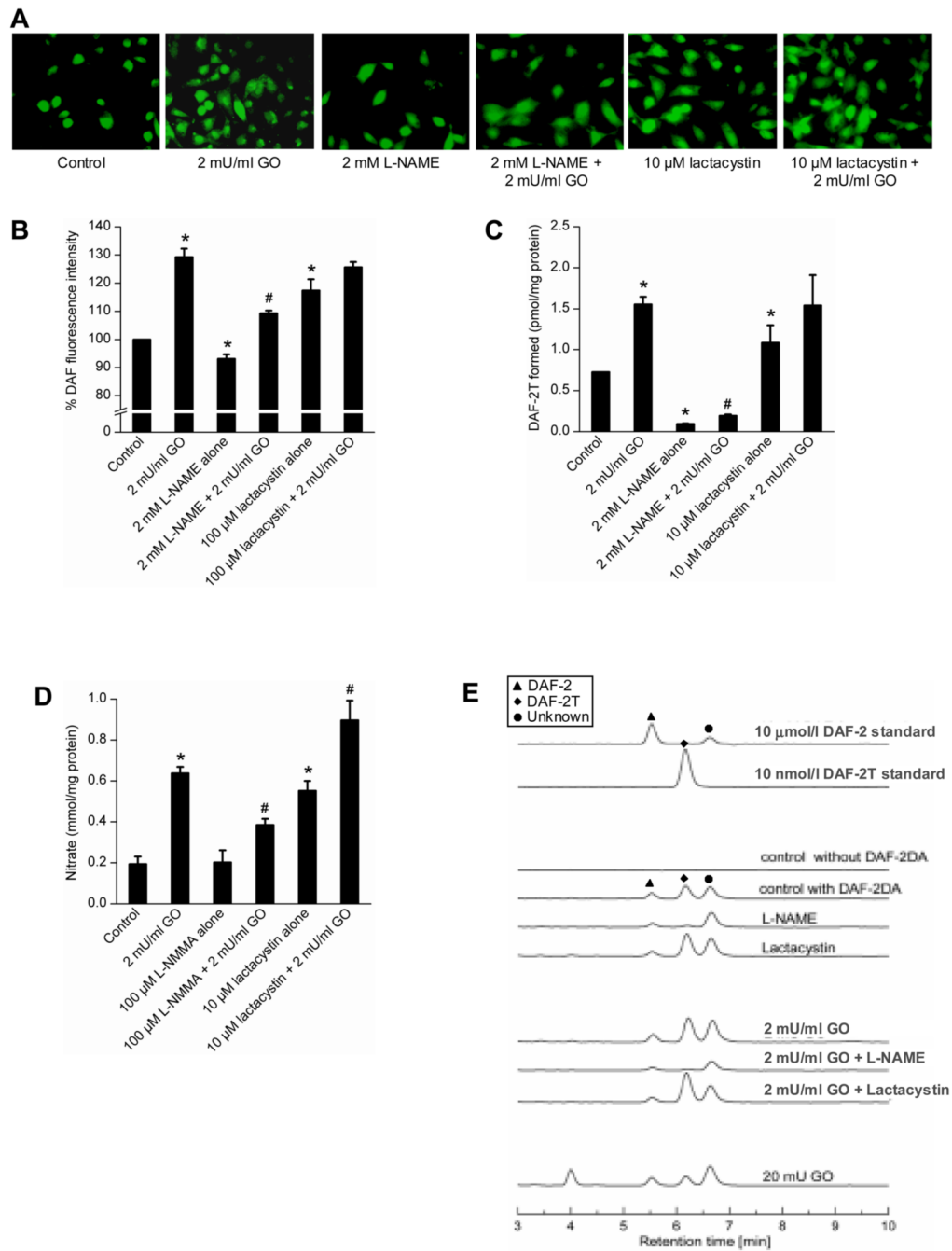


Figure 1. Dose-dependent effects of H₂O₂ on eNOS and Akt phosphorylations and *NO generation in BAEC treated with Glu/GO. **A**, BAEC were treated with different concentrations of GO for 4 h, and cell lysates probed with anti-phospho eNOS, eNOS, phospho Akt, and total Akt antibodies as described in *Materials and Methods*. **B**, Cells were treated as in **A** and the culture media analyzed for nitrate levels. **C**, Fluorescence micrographs obtained from cells treated as in **A** followed by a treatment with 5 μ M DAF-2DA as described in *Materials and Methods*. **D**, Densitometric analysis of the fluorescence data shown in **C**. The control fluorescence intensity was normalized to 100%. **E**, DAF-2-triazole extracted from BAEC treated with varying concentrations of GO as in **C**. DAF-2-triazole was measured by HPLC as described in *Materials and Methods*. * $p < 0.05$ as compared to control; # $p < 0.05$ as compared to 2 mU/ml GO sample.

**Figure 2.**

The effect of L-NAME and lactacystin on H₂O₂-induced DAF-2 fluorescence and DAF-2T formation. **A**, BAEC were treated with GO (2 μM/ml) for 4 h with and without L-NAME and lactacystin pretreatment and fluorescent micrographs obtained after treatment with DAF-2DA. **B**, Same as **A** except that the graph shows the densitometric analysis of the fluorescence data shown in **A**. The control fluorescence intensity was taken as 100%. **C**, DAF-2T formation as measured by HPLC from cells treated under conditions shown in **A**. **D**, Graph showing nitrate levels measured in the cell culture media following treatment of cells as in **A** but in the absence of DAF-2DA. **E**, HPLC traces of DAF-2 and DAF-2T standards, n-butanol extracts of control and GO-treated cells with and without pretreatment of L-NAME and lactacystin as shown in

A. Symbols marked in the HPLC traces represent the species, as shown in the inset. * $p < 0.05$ as compared to control; # $p < 0.05$ as compared to 2 mU/ml GO sample..

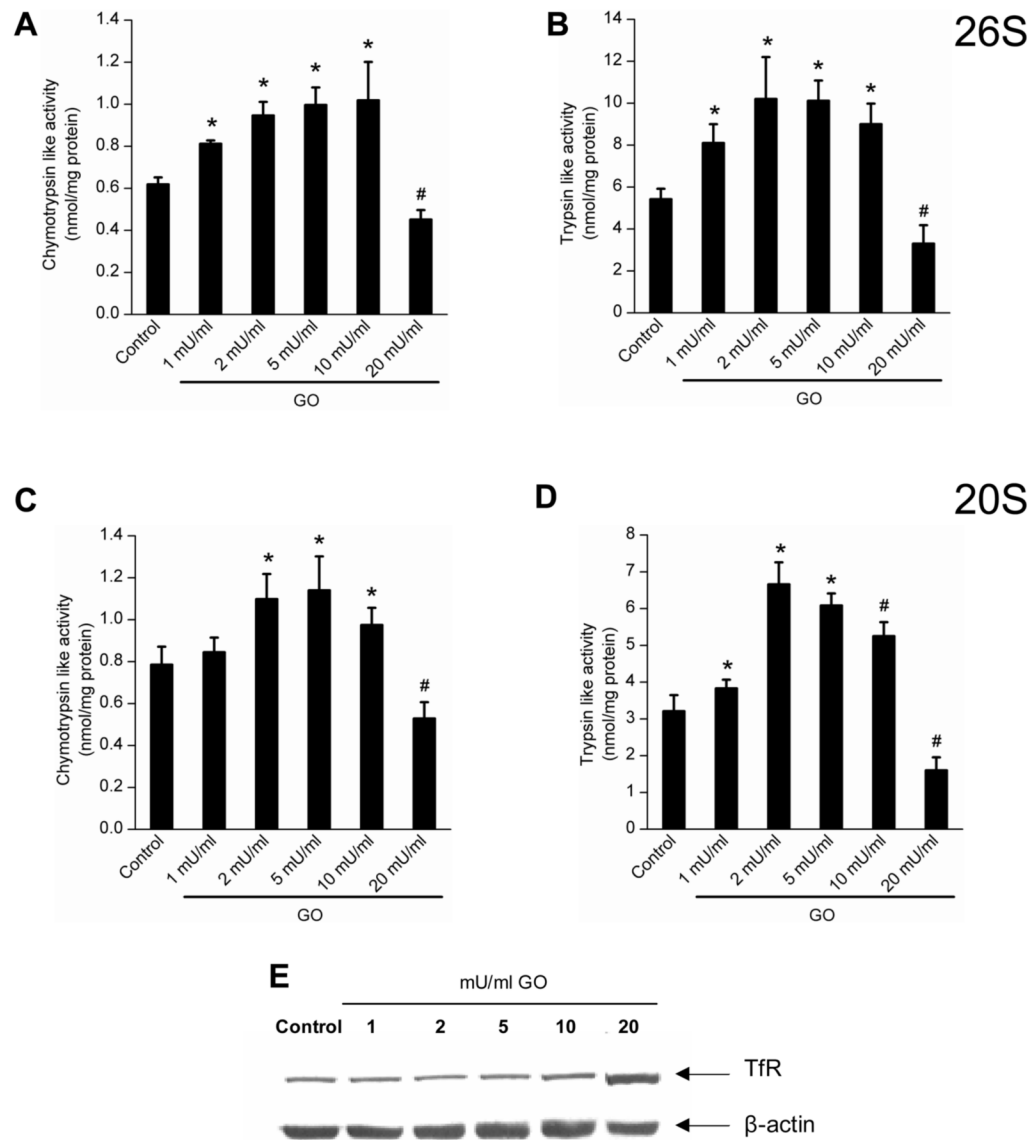
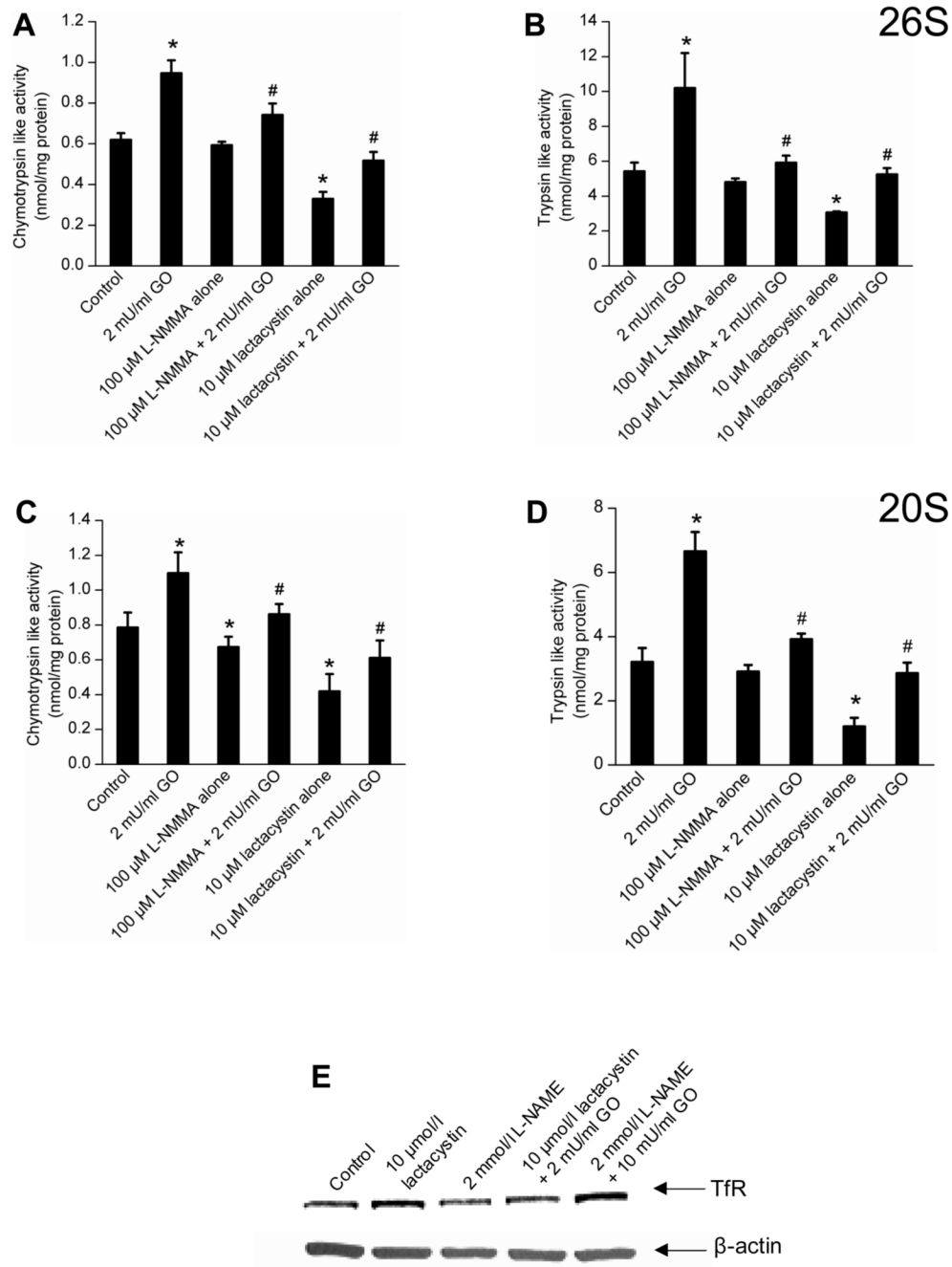


Figure 3. Dose-dependent effects of GO on proteasomal function. **A** and **B**, BAEC were treated with different concentrations of GO (1–20 mU/ml) for 4 h, and chymotrypsin- and trypsin-like activities of 26S proteasome measured as described in *Materials and Methods*. **C** and **D**, Treatment conditions same as above and the chymotrypsin and trypsin-like activities of 20S proteasome were measured. **E**, Western blot of transferrin receptor protein levels monitored in cell lysates of BAEC after treatment with GO (1–20 mU/ml). * $p < 0.05$ as compared to control; # $p < 0.05$ as compared to 2 mU/ml GO sample.

**Figure 4.**

The effects of NOS and proteasome inhibitors on proteasomal activation in BAECs treated with low levels of GO. **A** and **B**, BAEC were treated for 4 h with GO (2 mU/ml) before and after pretreatment with L-NMMA and lactacystin for 2 h. The graphs show the chymotrypsin- and trypsin-like activities of 26S proteasome measured under different treatment conditions. **C** and **D**, Same as above, except the chymotrypsin- and trypsin-like activities of 20S proteasome were measured. **E**, The Western blot of TfR protein levels was measured following the same treatment conditions as described above. * $p < 0.05$ as compared to control; # $p < 0.05$ as compared to 2 mU/ml GO sample.

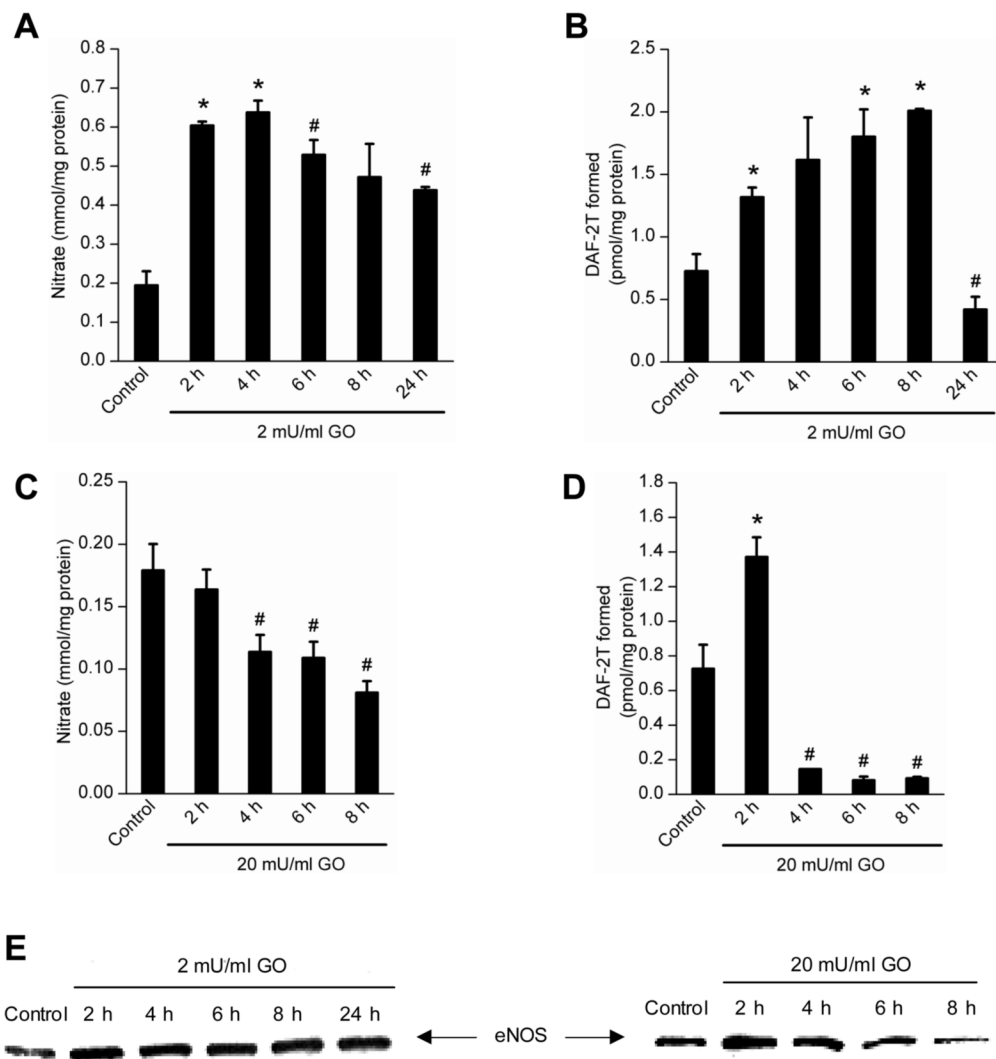


Figure 5. Measurements of nitrate and DAF-2T in cells treated with low and high levels of GO. **A** and **B**, BAEC were treated with 2 mU/ml of GO for different time periods and nitrate release (**A**) and intracellular DAF-2T levels (**B**) were measured. **C** and **D**, Same as above, except that cells were treated with GO (20 mU/ml) and nitrate (**C**) and DAF-2T (**D**) measured as described in *Materials and Methods*. **E**, Immunoblots of eNOS protein measured in cell lysates after treatment with 2 and 20 mU/ml of GO. * $p < 0.05$ as compared to control; # $p < 0.05$ as compared to 4 h (**A**), 6 h (**B**) and 2 h (**C** and **D**) incubation with GO.

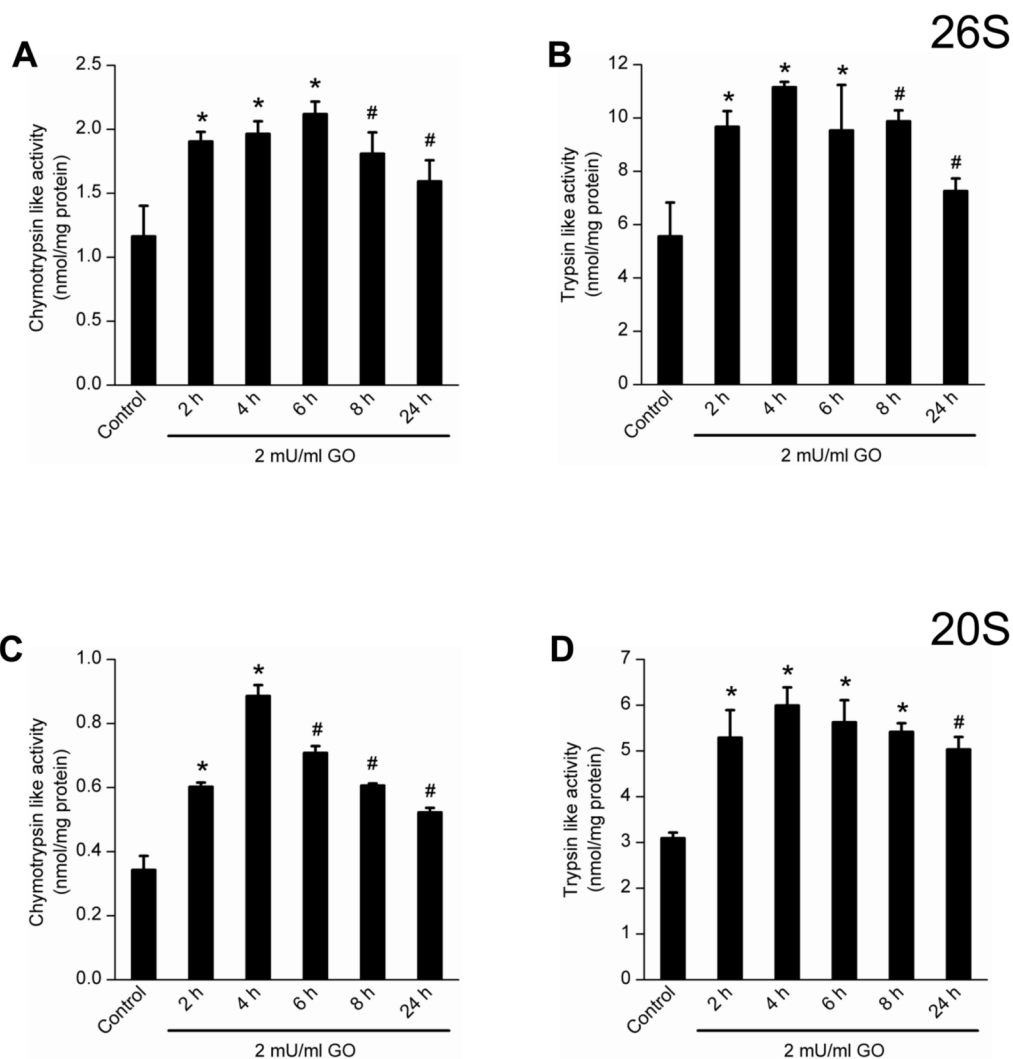


Figure 6. Time-dependent changes in proteasome activity in BAEC treated with low dose of GO. **A** and **B**, BAEC were treated with 2 mU/ml for different time periods as shown, and the chymotrypsin- and trypsin-like activities of 26S proteasome measured in cell lysates. **C** and **D**, Same as above, except that the chymotrypsin- and trypsin-like activities of 20S proteasome were measured in cell lysates. * $p < 0.05$ as compared to control; # $p < 0.05$ as compared to 6 h (**A**), and 4 h (**B**, **C** and **D**) incubation with GO.

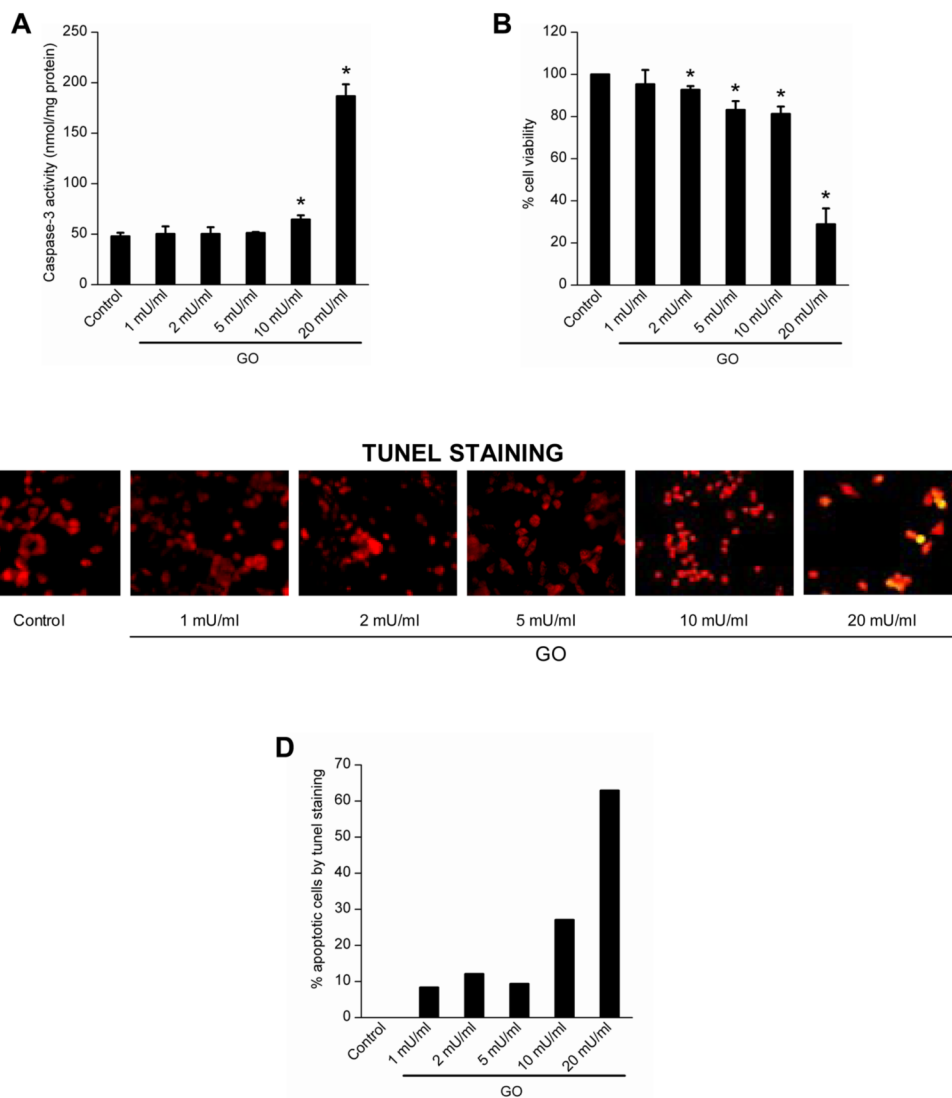


Figure 7. Dose-dependent effects of GO on endothelial apoptosis. **A**, BAEC were treated with GO (1–20 μU/ml) for 8 h, and the caspase-3 activity measured by monitoring the release of p-nitroanilide. **B**, Treatment conditions same as above and the cell viability was measured by the MTT assay. **C**, Treatment conditions same as in **A** and cells stained for TUNEL-positive cells and examined by fluorescence microscopy. Photographs are overlaid images of propidium iodide and FITC-stained cells. Apoptotic and non-apoptotic cells are denoted by yellow and red colors. **D**, Treatment conditions same as **A** and the graph shows the percent of TUNEL-positive cells in response to different treatment conditions. * $p < 0.05$ as compared to control.

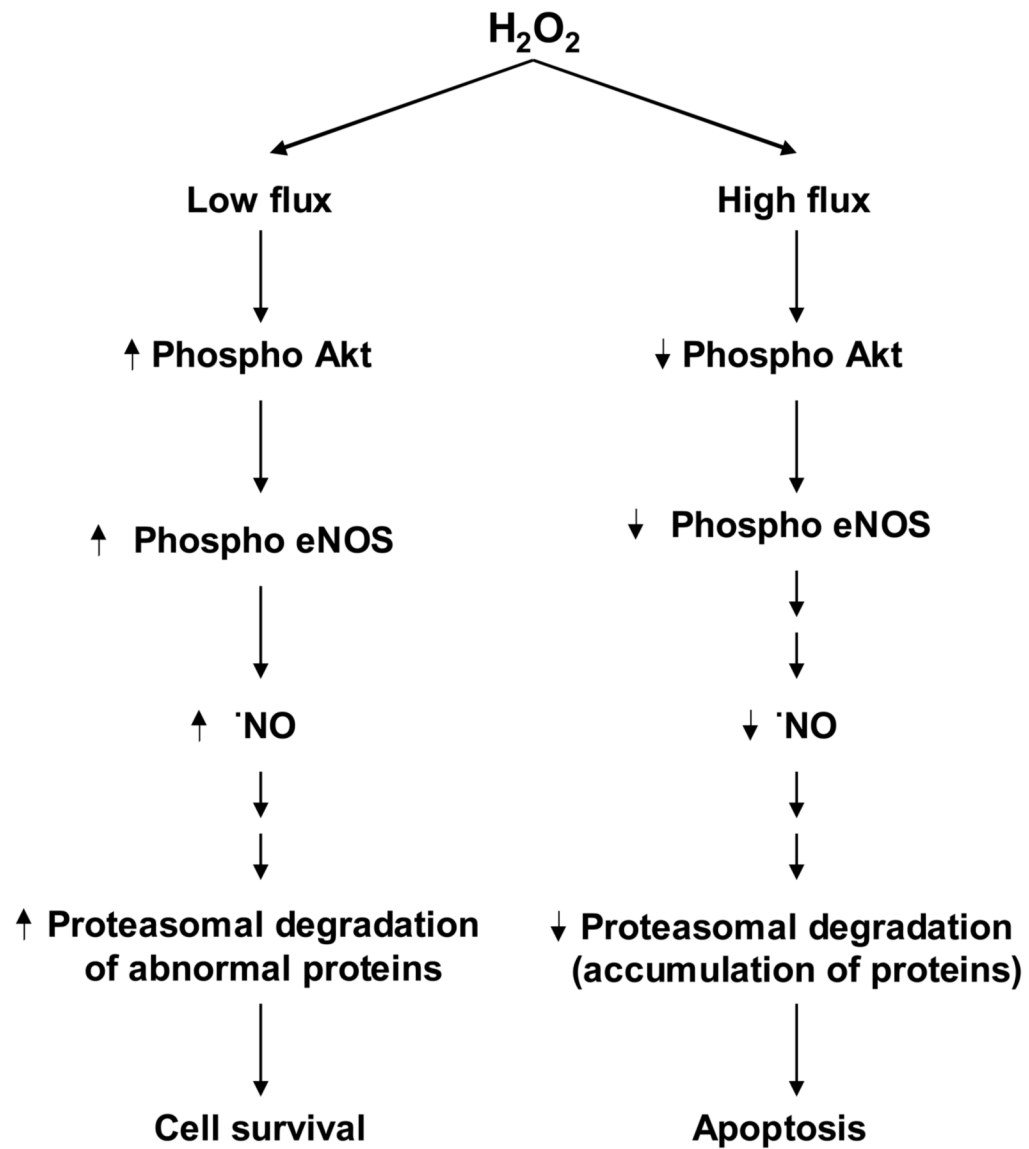


Figure 8. Nitric oxide mediated regulation of proteasomal function in response to H₂O₂-induced endothelial cell survival and cell death.