

# Nitric oxide regulates steroid synthesis by bovine antral granulosa cells in a chemically defined medium

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## Abstract

Nitric oxide (NO) in bovine ovary has been characterized as one of the controllers of granulosa cells' (GC) steroidogenesis and apoptosis. One of the pathways used by NO to have these effects is cGMP. The objectives of the present study were to verify the effect of sodium nitroprusside (SNP), a NO donor, on steroidogenesis, cell viability (mitochondrial activity) and GC cell cycle distribution and if this effect occurs by the NO-cGMP signaling pathway with the addition of SNP with or without 1H-[1,2,3] oxadiazolo[4,3a]quinoxaline-1-one (ODQ), a selective soluble guanylate cyclase inhibitor. The antral GC from 3 to 5 mm diameter cattle follicles was cultured without treatment (control), with ODQ ( $10^{-4}$  M) and  $10^{-5}$ ,  $10^{-3}$  and  $10^{-1}$  M SNP with or without ODQ for 24 h. Nitrate/nitrite ( $\text{NO}_3^-/\text{NO}_2^-$ ) concentrations were evaluated by Griess method, progesterone ( $\text{P}_4$ ) and  $17\beta$ -estradiol ( $\text{E}_2$ ) concentrations by chemiluminescence, viability and cell cycle stage by MTT method (3-[4,5-dimethylthiazol-2-yl]-2,3 dipheniltetrazolium bromide) and flow cytometry, respectively. Nitrate/nitrite concentration in culture medium increased ( $P < 0.05$ ) in a dose-dependent manner according to SNP concentration added to the culture medium. The GC cultured without treatment, with ODQ and with  $10^{-5}$  M SNP in the presence or absence of ODQ developed into cell aggregates and did not vary in cell viability ( $P > 0.05$ ), while GC cultured with  $10^{-3}$  and  $10^{-1}$  M SNP with or without ODQ presented disorganized GC aggregates or did not develop into cell aggregates and also had substantially decreased cell

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viability (mitochondrial activity inhibition) and steroids synthesis ( $P < 0.05$ ), and effects were not reversed with us of ODQ. Most GC cultured without treatment (control) or with ODQ,  $10^{-5}$  and  $10^{-3}$  M SNP with or without ODQ were in the G0/G1 (80–75%) stage and in a lesser proportion (20–25%) in the S + G2/M stage of the cell cycle, while the  $10^{-1}$  M SNP treatment resulted in GC in G1 phase arrest. The treatment with  $10^{-5}$  M SNP increased ( $P < 0.05$ )  $E_2$  synthesis and inhibited ( $P < 0.05$ ) progesterone synthesis. The addition of ODQ reversed ( $P < 0.05$ ) the stimulatory effect of  $10^{-5}$  M SNP treatment on  $E_2$ , but not on  $P_4$  synthesis ( $P > 0.05$ ). These results demonstrated that  $E_2$  synthesis by antral GC from small follicles is modulated by lesser NO concentrations via the cGMP pathway, but not  $P_4$  while steroids inhibition cGMP pathway independent, mitochondrial damage and the interference on cell cycle progression caused by greater NO concentration can lead to cell death.

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## 1. Introduction

$17\beta$ -Estradiol ( $E_2$ ) has an essential role to establish and maintain the reproductive system. Estradiol synthesis by granulosa cells (GC) depends on enzyme P450 aromatase (P450arom), which metabolizes androgens into  $E_2$ . Its increase is induced by FSH, and when bound to the receptor attaches to G protein, activates adenylyl cyclase and stimulates the production of the second messenger cAMP (Richards et al., 1995). Besides FSH, there are other hormones, growth factors, proteins, peptides, intercellular and intracellular factors that modulate GC functions (Chun et al., 1995; Lapolt and Hong, 1995).

Nitric oxide (NO) is a colorless gas at room temperature and pressure. The maximum solubility of NO in water at room temperature and pressure is slightly greater than the solubility of dioxygen ( $O_2$ ). Similar to  $O_2$ , NO is somewhat lipophilic and is 6–8 times more soluble in nonpolar solvents and lipid membranes than in water. Thus, NO reaction rates can be dramatically greater in a hydrophobic environment than in water due to its concentration increase (Liu et al., 1998), because it has a greater solubility in biological membranes.

NO is produced from L-arginine in a reaction catalyzed by the enzyme nitric oxide synthase (NOS) that exists in three isoforms: endothelial (eNOS) and neuronal (nNOS), which are  $Ca^{2+}$ /CaM-dependent and inducible (iNOS), regulated by cytokines (Snyder et al., 1996). The presence of eNOS and iNOS as a source of NO in ovarian compartments and its function as a steroidogenesis regulator in GC has previously been demonstrated in mice (Janblonka-Shariff and Olson, 1997; Matsumi et al., 1998; Hurwitz et al., 2002) and cattle (Basini et al., 1998). However, there are still inconsistencies with regard to the mechanism by which NO effects GC.

NO is a negative regulator of steroids synthesis. NO exerts its effects by binding to the prosthetic heme group of enzymes. Thus, NO may directly bind to the enzyme, P450 aromatase, and inhibit gonadotropin-stimulated steroidogenesis (Van Voorhis et al., 1994; Hanke et al., 1998).

Another mechanism of action mediated by NO is the activation of soluble guanylate cyclase that converts GTP (guanosine triphosphate) to cGMP and inorganic pyrophosphate (Murard, 1999; Hanafy et al., 2001) in a heme-dependent manner (Craven and DeRubertis, 1978). It has been suggested that cGMP increases phosphodiesterase 2 (PDE2) activity which in turn increases cAMP hydrolysis, the FSH intracellular second messenger.

In cultured GC from mice (Ishimaru et al., 2001) and pigs (Grasselli et al., 2001) the NO/cGMP pathway has been suggested as one of the mechanisms used by NO to inhibit steroidogenesis.

The use of cGMP analogs in cows, however, did not efficiently inhibit steroidogenesis (Basini et al., 2000).

These results are conflicting, probably due to variations among culture systems, presence of blood cells, differences among species and variation in GC differentiation stages of cells that are used. BSA is normally used in most culture systems, including those previously reported, and is contaminated with steroids, cholesterol, peptides and other yet to be recognized molecules (Wang et al., 1997; Mingoti et al., 2002). For this reason, culture medium becomes semi-defined possibly causing interference in the results. Moreover, the presence of blood cells can also alter GC steroidogenesis and NO synthesis/concentration (Beckmann et al., 1991; Shakil and Whitehead, 1994) in culture medium. Therefore, in the present experiment, BSA was replaced by polyvinyl alcohol (PVA) in a culture system which maintains GC P450 $\alpha$  activity without undergoing luteinization (Piccinato et al., 2000). Methods to extract red blood cells and macrophages that are accidentally collected during ovarian follicle aspiration were also performed. Using this culture system, the objectives were: (1) to verify NO effect using sodium nitroprusside (SNP), a NO donor, on antral GC steroid synthesis, viability and cell cycle distribution obtained from small follicles (3–5 mm) of cattle, because these have greater steroidogenic activity (Rouillier et al., 1996, 1998) and are not yet differentiated having greater proliferative activity (Gutierrez et al., 1997); (2) to verify if these effects are modulated by cGMP pathway using a treatment with 1H-[1,2,3]oxadiazolo[4,3a]quinoxaline-1-one (ODQ), a selective soluble guanylate cyclase inhibitor.

## 2. Materials and methods

### 2.1. Reagents

Most reagents were obtained from Sigma (St. Louis, MO, USA). Reagents from other laboratories are specified.

### 2.2. Experimental design

GC were cultured with 0,  $10^{-5}$ ,  $10^{-3}$ ,  $10^{-1}$  M sodium nitroprusside without or with  $10^{-4}$  M ODQ (Basini et al., 2000). The synthesis of  $\text{NO}_3^-/\text{NO}_2^-$ , progesterone ( $\text{P}_4$ ) and  $\text{E}_2$  by GC and GC viability and cell cycle status were evaluated at the end of 24 h of culture.

### 2.3. GC collection

Ovaries were collected postmortem at a local abattoir from mixed-age cows in the luteal and follicular phase of the estrous cycle and the absence of palpable products of pregnancy (Ireland et al., 1980) and were immediately transported (30 °C) to the laboratory in sterile saline solution (NaCl 0.9%) supplemented with antibiotics (100 IU/ml penicillin and 100  $\mu\text{g}/\text{ml}$  streptomycin sulphate). Excess tissue was trimmed. Ovaries were washed with saline solution containing antibiotics and were briefly rinsed with 70% ethanol and then washed (three times) with saline solution supplemented with antibiotics.

Small well-vascularized follicles (3–5 mm) having more than 70% of clear exposed surface (Grimes et al., 1987) were punctured. Pool of antral GC were then aspirated (Rouillier et al., 1998) and placed in tubes of 15 ml containing 2 ml base medium [Dulbecco's Modification of Eagle's Medium/Ham's F-12 containing HEPES 15 mM (Gibco-BRL Life Technologies), supplemented

with 10 mM sodium bicarbonate; 100 UI/ml penicillin G and 100 µg/ml streptomycin sulphate (Merck Indústrias SA, Rio de Janeiro, Brazil)] and 50 UI heparin to prevent cell aggregates.

GC suspension was filtered through nylon mesh to extract oocytes and cellular debris and then centrifuged ( $300 \times g/10$  min) at room temperature. Supernatant was decanted and GC were washed (twice) using 1 ml base medium. Regarding the fact that red blood cells can bind to NO (McMahon, 2002), that macrophages can synthesize NO (Whitehead and Lacey, 1996) and that during follicle puncture blood may also be aspirated, two procedures were accomplished. GC pellet was treated with 300 µl 0.9% prewarmed ammonium chloride solution at 37 °C and incubated for 1 min to extract red blood cells. Following, GC isotonicity was recovered by adding base medium up to 12 ml into the tubes containing GC. GC were centrifuged ( $300 \times g/10$  min) and the pellet was washed with base medium. The GC pellet was re-suspended in 1 ml base medium and GC were seeded in a 32 mm tissue culture dish (Corning Incorporated, Corning, NY, USA) containing 2 ml base medium. GC were incubated for 2 h at 37.5 °C in a 5% CO<sub>2</sub> atmosphere to remove macrophages, because these cells can easily adhere to culture plate (Beckmann et al., 1991).

Remaining GC (not adhered) were collected from culture plate, centrifuged ( $300 \times g/10$  min) and re-suspended in 2 ml culture medium [Dulbecco's Modification of Eagle's Medium/Ham's F-12 containing HEPES 15 mM, 10 mM sodium bicarbonate, 100 U/ml penicillin G, and 100 µg/ml streptomycin sulphate (Merck Indústrias SA);  $10^{-7}$  M androstenedione; 0.1% PVA; 1% MEM nonessential amino acid mixture 100% and 1% solution ITS 100% (1.4 ng/ml sodium selenite, 5.0 µg/ml transferrin, 10 ng/ml insulin)].

Viability and cell number were estimated under an inverted microscope (Telaval 31, Zeiss) at 400× magnification using a hemocytometer and Trypan blue (0.4%) exclusion method. Cellular viability average was 70%.

Smears of the cellular suspension were realized before and after red blood cells and macrophages removal. After staining with hematoxyline, 100 cells were counted in 10 replicates (1000 total cells). For each 100 cells counted before the treatment an average of  $39.8 \pm 7.7\%$  red blood cells and  $8.5 \pm 3\%$  macrophages were found. After treatment  $9 \pm 4\%$  red blood cells and  $0.4 \pm 0.5\%$  macrophages were found.

#### 2.4. GC culture

$5.0 \times 10^5$  antral GC of follicles of 3–5 mm diameter were seeded in 24-well plates with 1000 µl culture medium containing the treatments. GC were then incubated at 37.5 °C in a 5% CO<sub>2</sub> atmosphere.

#### 2.5. Cell viability determination

The NO effect on cell viability was evaluated by MTT method. This assay indirectly determines cell viability by measuring the activity of mitochondrial enzymes. Yellow tetrazolium salt 3-(4,5-dimethylthiazol-2-yl)-2,3 diphenyltetrazolium bromide (MTT) is metabolized by mitochondrial dehydrogenases to form the final product blue formazan (metyltriazole) (Mosmann, 1983). At the end of 24 h of culture period, after the removal of 700 µl culture medium for hormone and NO analysis, 30 µl MTT (5 mg/ml) were added to each well and GC were held in incubation for 3 h at 37 °C in CO<sub>2</sub>:95% air. After incubation 300 µl isopropanol acidified with HCl (0.04 N) containing 10% Triton X 100 were added to dissolve formed crystals and remained 14–16 h at room temperature. After this period, 200 µl blue formazan/mixture were transferred to 96-

well plates and analyses were determined by a spectrophotometer (Multiskan EX Primary EIA V 2.1–0) at wavelength of 570 nm, with background subtraction (620 nm). The relationship between absorbance and cell number was determined by the previous incubation of a known cell number with MTT to establish a standard curve. A linear relationship between cell number and absorbance values ( $R^2 = 0.99$ ;  $P < 0.05$ ) was found.

## 2.6. Cell cycle

To determine the effect of different treatments on GCs cell cycle distribution the flow cytometry determination was used. For that reason, the method described by Blondin et al. (1996) was used with some modifications. GC were collected after incubation with 1 ml PBS  $\text{Ca}^{2+}/\text{Mg}^{2+}$ -free and EDTA (125 mg/l) for 10 min and then centrifuged at  $400 \times g$  at  $4^\circ\text{C}$  for 5 min. The supernatant was decanted and the pellet was re-suspended in 500  $\mu\text{l}$  PBS  $\text{Ca}^{2+}/\text{Mg}^{2+}$ -free. GC were fixed with 1 ml absolute ethanol during agitation in vortex for at least 1 h at  $4^\circ\text{C}$ . Afterwards, GC were centrifuged and the pellet re-suspended with 500  $\mu\text{l}$  ethanol 70% and stored at  $-20^\circ\text{C}$  until flow cytometry analysis. Prior to staining GC were centrifuged, the supernatant was decanted and cells were washed twice with PBS ( $400 \times g$  at  $4^\circ\text{C}$  for 5 min). Fixed GC were re-suspended with 1 ml PBS containing RNase at 50  $\mu\text{g}/\text{ml}$  (Merck Indústrias SA), propidium iodide (50  $\mu\text{g}/\text{ml}$ ) and incubated for 4 h at  $4^\circ\text{C}$ . Samples were filtered through 40  $\mu\text{m}$  nylon mesh. The percentage of cells with degraded DNA and cell cycle distributions for single cells was determined using a ploidy analyzer (PARTEC PA) with UV filter. Data derived from DNA histogram were analyzed by flow cytometer consisting of GC percentage in stage G0/G1 (diploid cells), S + G2/M (DNA synthesis and mitosis).

## 2.7. Nitrate/nitrite and steroids dosages

After a 24 h period, culture medium was collected and stored at  $-20^\circ\text{C}$  to dose nitrate/nitrite ( $\text{NO}_3^-/\text{NO}_2^-$ ),  $\text{P}_4$  and  $\text{E}_2$ .

### 2.7.1. $\text{NO}_3^-/\text{NO}_2^-$

Nitrate/nitrite concentrations were determined by a method based on the colorimetric reaction of Griess (Ricart-Jané et al., 2002). Because very little or no  $\text{NO}_2^-$  is normally found in GC cultures, no attempt was made to differentiate between the amounts of  $\text{NO}_3^-$  and  $\text{NO}_2^-$  and therefore, the results are reported as NO. Griess reagent is composed of a mixture of sulphanilamide 2% and *N*-(1-naphthyl) ethylene-diamine 0.2% in deionized water. The first reacts with the nitrite in the sample to form diazonium salt, which in turn reacts with the second reagent to give a purple-azo-dye product with a peak absorbance at 540 nm. To transform  $\text{NO}_3^-$  into  $\text{NO}_2^-$  samples (40  $\mu\text{l}$ ) were incubated in 96-well plates with the reduction solution (100  $\mu\text{l}$  of 10 UI  $\text{NO}_3^-$  reductase enzyme diluted in deionized water +900  $\mu\text{l}$  deionized water), 1000  $\mu\text{l}$  cofactor NADPH (5 mg/ml) diluted in deionized water, 1000  $\mu\text{l}$  phosphate potassium buffer (0.5 M) at  $37^\circ\text{C}$  for 14–16 h. Afterwards, 80  $\mu\text{l}$  Griess reagent were added to samples. The standard curve of  $\text{NaNO}_3$  was diluted in DMEM/Ham's-F12 ranging from 0.5 to 100  $\mu\text{M}$ . The analysis was made in a spectrophotometer (Multiskan EX Primary EIA V 2.1-0). A dispersion graph was generated with the absorbance values. There was a linear ( $R^2 = 0.98$ ,  $P < 0.05$ ) relation between absorbance and NO concentration. All solutions were protected from light.

### 2.7.2. Steroids

Progesterone and E<sub>2</sub> concentrations were determined by an automated chemiluminescence system (ACS:180<sup>®</sup>, Bayer Corporation, Tarrytown, NY, USA) which is a competitive immunoassay that uses direct chemiluminescent technology (progesterone and estradiol derivative are labeled with acridinium ester). The monoclonal mouse anti-rabbit antibodies are coupled to paramagnetic particles and are highly specific, possesses a cross-reactivity of less than 0.95% with other compounds. Progesterone and E<sub>2</sub> concentrations ranged from 0.86 to 29.80 ng/ml and 88 to 785 pg/ml, respectively. Samples containing 0 and 10<sup>-5</sup> M SNP with or without ODQ were diluted (1/400), however, culture medium samples with 10<sup>-3</sup> and 10<sup>-1</sup> M SNP with or without ODQ were not. The average intra-assay coefficient of variation was 1.6% and 3.0% for P<sub>4</sub> and E<sub>2</sub>, respectively.

### 2.8. Statistical analysis

Each experiment assessing cell viability, cell cycle, steroidogenesis and NO concentration was repeated two times independently and in each experiment treatment two replicates culture wells. Data are expressed as mean ± S.E.M. Statistical differences between treatments and interactions were calculated with multifactorial ANOVA. When significant differences were found, means were compared by Tukey test considered significant at  $P < 0.05$ . SAS (1996) (Statistical Analysis System) software was used.

## 3. Results

### 3.1. Morphological characteristics of GC

At the end of 24 h, GC cultured with 0 (control), 10<sup>-5</sup> M SNP with or without ODQ (Fig. 1, panels a–d) maintained the typical round shape of steroidogenically active GC and formed cell groups, while GC cultured with 10<sup>-3</sup> M SNP with or without ODQ (Fig. 1, panels e and f) formed disorganized groups, had dark color with little adherence to the culture plate and the cultured with 10<sup>-1</sup> M SNP with or without ODQ (Fig. 1, panels g and h) did not form groups, had dark color and were flattened in the culture plate.

### 3.2. NO<sub>3</sub><sup>-</sup>/NO<sub>2</sub><sup>-</sup> concentration

NO concentration in culture medium increased ( $P < 0.05$ ) in a dose-dependent manner according to the SNP concentration added in culture medium: SNP 10<sup>-5</sup> M with or without ODQ (6.7 ± 2.27, 6.6 ± 2.3 μM, respectively), SNP 10<sup>-3</sup> M with or without ODQ (48.6 ± 11, 50.9 ± 2.6 μM, respectively) and SNP 10<sup>-1</sup> M with or without ODQ (65.6 ± 2.9, 68.3 ± 11.7 μM, respectively) (Fig. 2a).

### 3.3. Cell viability

Cell viability was not altered ( $P > 0.05$ ) among control group (6.8 × 10<sup>5</sup> ± 0.27 viable cells) and groups cultured with ODQ (6.6 × 10<sup>5</sup> ± 0.75 viable cells) and 10<sup>-5</sup> M SNP with ODQ (7.0 × 10<sup>5</sup> ± 0.5 viable cells) or without ODQ (7.6 × 10<sup>5</sup> ± 0.8 viable cell). However, after the addition of 10<sup>-3</sup> and 10<sup>-1</sup> M SNP with or without ODQ, 100% GC ( $P < 0.05$ ) became non-viable (Fig. 2b).

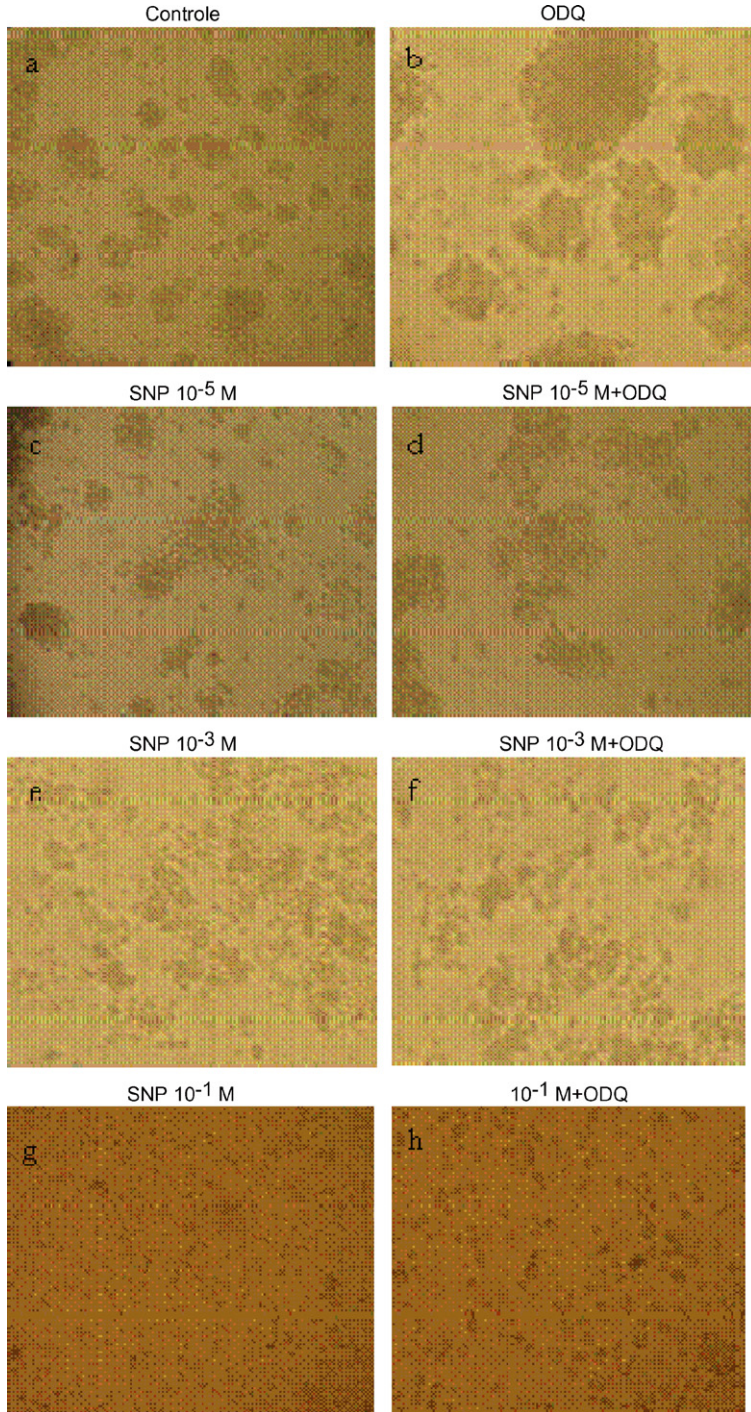


Fig. 1. Granulosa cells cultured without SNP and ODQ (control) (a) or with  $10^{-4}$  M ODQ (b);  $10^{-5}$  M SNP (c);  $10^{-5}$  M SNP +  $10^{-4}$  M ODQ (d);  $10^{-3}$  M SNP (e);  $10^{-3}$  M SNP +  $10^{-4}$  M ODQ (f);  $10^{-1}$  M SNP (g);  $10^{-1}$  M SNP +  $10^{-4}$  M ODQ (h) (at 400 $\times$  magnification).

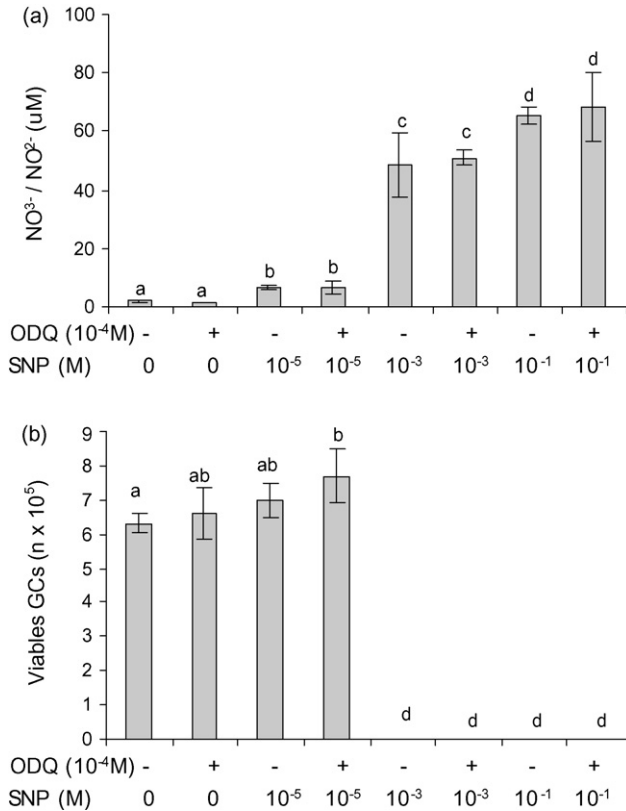


Fig. 2. Effect of the addition of SNP (0, 10<sup>-5</sup>, 10<sup>-3</sup> and 10<sup>-1</sup> M) with or without ODQ (10<sup>-4</sup> M) in NO<sub>3</sub><sup>-</sup>/NO<sub>2</sub><sup>-</sup> concentration in culture medium and in cell viability (number of viable GCs × 10<sup>5</sup>) after 24 h culture. Values with different superscripts (a–d) are different ( $P < 0.05$ ) by Tukey test. Each value represents mean ± S.E.M. of a total of 4 determinations from two independent experiments.

### 3.4. Cell cycle

Most GC cultured with 0 (control), 10<sup>-5</sup> and 10<sup>-3</sup> M SNP with or without ODQ were in G0/G1 (80–70%) stage after 24 h culture, while 20–30% were in the stage S + G2/M. The GC cultured with 10<sup>-1</sup> M SNP, however, had a decreased cell count and a G1 phase arrest (Fig. 3).

### 3.5. Steroidogenesis

SNP at 10<sup>-3</sup> and 10<sup>-5</sup> M inhibited ( $P < 0.05$ ) P<sub>4</sub> synthesis (14.39 ± 0.003; 129.1 ± 0.006 ng/ml, respectively) compared to control (174.7 ± 0.02 ng/ml); 10<sup>-3</sup> SNP had a greater effect ( $P < 0.05$ ) than SNP 10<sup>-5</sup> M. The addition of 10<sup>-4</sup> M ODQ did not influence the negative effect of 10<sup>-3</sup> and 10<sup>-5</sup> M SNP on P<sub>4</sub> synthesis (16.35 ± 0.0007; 126.68 ± 0.044 ng/ml, respectively) and ODQ also inhibited P<sub>4</sub> synthesis (91.3 ± 0.02 ng/ml) (Fig. 4, panel a).

Although the addition of both 10<sup>-4</sup> M ODQ and 10<sup>-5</sup> M SNP stimulated ( $P < 0.05$ ) E<sub>2</sub> synthesis (3.4; 6.8 ± 0.5 ng/ml, respectively) when compared to the control (2.0 ng/ml,  $P < 0.05$ ), E<sub>2</sub> synthesis was greater ( $P < 0.05$ ) for the 10<sup>-5</sup> M SNP than for ODQ treatments. Even though the

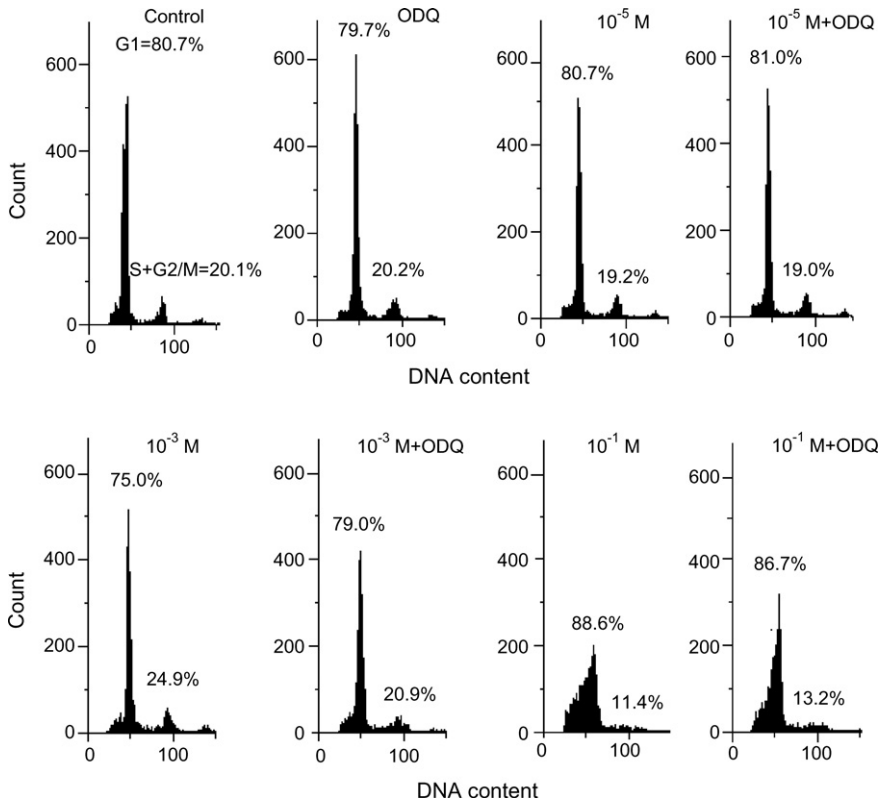


Fig. 3. Effect of SNP addition on cell cycle of GC cultured without SNP and ODQ (control);  $10^{-4}$  M ODQ;  $10^{-5}$  M SNP,  $10^{-5}$  M SNP +  $10^{-4}$  M ODQ;  $10^{-3}$  M SNP;  $10^{-3}$  M SNP +  $10^{-4}$  M ODQ;  $10^{-1}$  M SNP;  $10^{-1}$  M SNP +  $10^{-4}$  M ODQ. G0: subdiploid DNA levels; G0/G1: diploid cells; S and G2/M: DNA synthesis/cell mitosis.

addition of ODQ inhibited ( $P < 0.05$ ) the increase of  $E_2$  synthesis ( $3.3 \pm 0.3$  ng/ml) which was stimulated by the addition of  $10^{-5}$  M SNP, no difference ( $P > 0.05$ ) in  $E_2$  synthesis between cultures containing ODQ and ODQ with  $10^{-5}$  M SNP was observed. The addition of  $10^{-3}$  M SNP inhibited ( $P < 0.05$ )  $E_2$  synthesis ( $0.02 \pm 0.001$  ng/ml) which was not reversed by the addition of ODQ ( $0.02 \pm 0.1$  ng/ml) (Fig. 4, panel b).

Progesterone and  $E_2$  synthesis in cultures containing  $10^{-1}$  M SNP was less than the minimum concentration ( $>0.86$  ng/ml and  $>88.0$  pg/ml, respectively) detected by the method utilized and, therefore, is not included in the figures.

#### 4. Discussion

The current research demonstrated that the culture system containing PVA maintained the viability of GC, allowed cells to progress through the cellular cycle, and maintained the morphological characteristics of steroidogenically active GC described by Gutierrez et al. (1997), such as a round shape and cell groups adhered to the plate by fibroblast-like cells. In addition, the treatment to remove red blood cells and macrophages was efficient. Blood cells may interfere in steroidogenesis and macrophages may synthesize NO (Beckmann et al., 1991; Shakil and

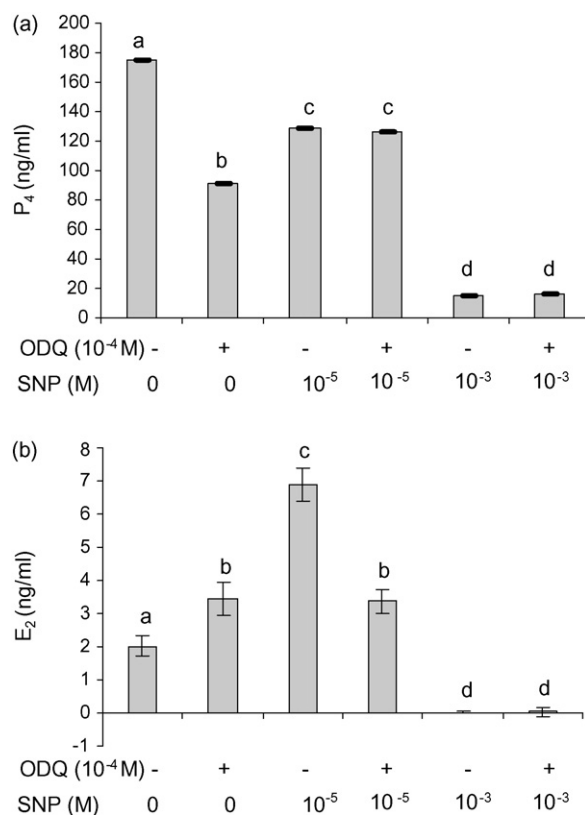


Fig. 4. Effect of SNP addition (0, 10<sup>-5</sup>, 10<sup>-3</sup> M) with or without ODQ (10<sup>-4</sup> M) on P<sub>4</sub> and E<sub>2</sub> synthesis by GC after 24 h culture. Values with different superscripts (a–d) are different ( $P < 0.05$ ) by Tukey test. Each value represents mean  $\pm$  S.E.M. of a total of four determinations from two independent experiments.

Whitehead, 1994). Consequently, this treatment becomes important to verify NO effects in GC steroidogenesis.

An inhibitory effect on E<sub>2</sub> synthesis without the cGMP pathway activation (Basini et al., 2000) in cattle GC cultured after the addition of increasing concentrations of NO donor, *S*-nitroso-*N*-acetyl-penicillamine (SNAP), has been verified. Some studies, however, suggest that the NO/cGMP pathway is a GC steroidogenesis inhibition mechanism in rats (Ellman et al., 1993) and pigs (Grasselli et al., 2001). Although the E<sub>2</sub> synthesis inhibition mechanism that has been proposed is different, they have in common the fact that NO inhibits steroid synthesis. On the contrary, the results of the current experiment indicates that at least during the first 24 h culture NO stimulates E<sub>2</sub> synthesis by antral GC through the activation of soluble nitric oxide-sensitive guanylyl cyclase, because 10<sup>-5</sup> M SNP stimulated E<sub>2</sub> synthesis, which was inhibited by the addition of ODQ. In spite of these inconsistencies, there are reports the increase of nitrite concentration (NO metabolite) is correlated with follicular volume and E<sub>2</sub> concentration, but not with progesterone concentration in human follicular fluid during follicular phase of the reproductive cycle (Anteby et al., 1996). Additionally, Bonello et al. (1996) found a decrease in E<sub>2</sub> synthesis in rats through ovarian perfusion using the inhibitor of an NOS enzyme (N $\omega$ -nitro-L-arginine methyl ester, L-NAME), suggesting that NO may positively regulate E<sub>2</sub> synthesis. The present results

agree with these latter results suggesting that NO may take part in the positive regulation of E<sub>2</sub> synthesis by GC of cattle.

Most NO biological effects occur by the soluble guanylyl cyclase activation that catalyzes the conversion of GTP into cGMP (Denninger and Marletta, 1999). cGMP causes cAMP concentrations to decrease by the phosphodiesterase-cAMP (PDE-cAMP) activation that cleaves cAMP (MacFarland et al., 1991). However, there are two PDE categories: PDE 2 activated by NO which cause cAMP hydrolysis (MacFarland et al., 1991), and PDE 3 (Conti et al., 1995) inhibited by NO and cGMP, resulting in cAMP increase and in the activation of the camp-dependent pathway (Draijer et al., 1995; Kurtz et al., 1998). Oocyte PDE 3 gene expression and GC PDE 4 gene expression has been reported (Tsafiriri et al., 1996). Thus, it becomes clear that other components of the cGMP signaling pathway must be elucidated to comprehend NO actions on cGMP and cAMP pathway.

Besides the previously mentioned considerations, the culture system used in the present study was different from that in previous studies because of GC type and differentiation stage, and also because of the different supplementations added to the culture (Basini et al., 2000; Grasselli et al., 2001). Cultures supplemented with BSA become undefined because BSA contains cholesterol, testosterone, progesterone, various growth factors and other yet to be determined components. Moreover, there are variations in the component concentrations, because these are biological materials (Wang et al., 1997; Maurer, 1992; Keskintepe and Brackett, 1996; Mingoti et al., 2002). Mingoti et al. (2002) also demonstrated a decrease on estradiol synthesis by *cumulus* cells when bovine *cumulus*–oocyte complexes (COC) were matured in culture medium supplemented with BSA and testosterone, suggesting that BSA increases androgen concentrations in the medium, causing a negative effect on estradiol synthesis, and hence influencing the results obtained in such studies. Additionally, the culture supplementation with insulin, not verified in the culture system utilized by Basini et al. (2000), might have resulted in the differences observed in results regarding the NO role on steroid synthesis by GC of cattle, because insulin maintains GC steroidogenic activity (Gutierrez et al., 1997).

Although 10<sup>-5</sup> M SNP stimulated E<sub>2</sub> synthesis, it inhibited progesterone without soluble nitric oxide-sensitive guanylyl cyclase activation, in accordance with Basini et al. (2000). Besides the cGMP pathway, NO can activate enzymes containing heme groups such as cyclooxygenases (Rettori et al., 1992) and cytochrome P450 (Wink et al., 1993). Snyder et al. (1996) suggested that NO regulates the steroidogenesis by the direct inhibition of cytochrome P450 enzymes. Accordingly, the results for the present experiment showed that NO inhibited progesterone synthesis by GC through cGMP independent mechanisms, but it can also positively regulate E<sub>2</sub> synthesis by cGMP in the first 24 h culture.

It was also shown that the NO independent cGMP pathway is involved with steroids synthesis, because of the cGMP pathway inhibition by the addition of ODQ without SNP stimulated estradiol synthesis in comparison with the control. Therefore, the 50% estradiol increase was due to the 10<sup>-5</sup> M SNP addition that stimulated estradiol synthesis by cGMP pathway, while the 20% estradiol increase was due to cGMP pathway inhibition which is stimulated by factors not elucidated in this experiment. It was also observed that there are other factors regulating progesterone synthesis through the cGMP pathway, because its inhibition decreased progesterone synthesis, however, no negative effect was detected on progesterone synthesis when ODQ was added to 10<sup>-5</sup> M SNP treatment. Even though NO is the major guanylyl cyclase physiological activator (Ignarro, 1991) there are other physiological regulators involved in its activation such as calcium (Sitaramayya, 2002) and protoporphyrin IX (Mingone et al., 2006). Little is known about guanylyl cyclase regulators in ovaries (Lapolt et al., 2002), and thus research in this field must be done

to verify other cGMP pathway regulators which will provide a greater understanding about the events that control ovarian function.

Results from the present study also showed that greater SNP concentrations ( $10^{-3}$  and  $10^{-1}$  M) inhibit both progesterone and estradiol synthesis by independent cGMP pathway mechanisms as previously detected by Basini et al. (2000). Basini et al. (1998) showed that NO in greater concentrations ( $10^{-3}$  M SNAP) protects GC from apoptosis. Results of the present study also did not demonstrate that GC apoptosis was induced by greater concentrations of SNP ( $10^{-3}$  and  $10^{-1}$  M), but it did indicate that greater NO concentrations did not protect GC from cellular damage, on the contrary, these cells had a lesser viability as verified by substantial mitochondrial damage and altered the morphological characteristics of steroidogenically active GC which were previously described by Gutierrez et al. (1997), and had disorganized GC aggregates ( $10^{-3}$  M) or did not develop into cell aggregates ( $10^{-1}$  M).

NO in greater concentrations induces apoptosis and/or necrosis in rat ovaries (Ellman et al., 1993), human epithelial-endometrial cells (Li et al., 2001) and cattle embryos (Orsi, 2006). NO in greater concentrations compete with oxygen and binds to enzyme heme groups from the respiratory chain, as well as cytochrome oxidase (Clementi et al., 1999; Sarti et al., 1999). After several hours of NO treatment, cytochrome oxidase enzyme can be irreversibly inhibited due to NO conversion into nitrogen reactive species (peroxynitrite), which inhibits various cellular respiratory sites (Clementi et al., 1999; Sarti et al., 1999). This mechanism triggers calcium release (Richter, 1993). Disturbances in mitochondrial calcium concentrations can kill cells through excessive calcium cycling across the inner mitochondrial membrane which can have serious consequences such as disruption of mitochondrial membrane potential, ATP depletion and cell death (Richter, 1993). Therefore, the stable maintenance of mitochondrial membrane potential can inhibit the onset of apoptosis. Thus, the present data suggest that mitochondrial damage caused by toxic effect of greater SNP concentration perturbs the Krebs cycle which in normal physiologic conditions generated nicotinamide adenine dinucleotide phosphate (NADPH) utilized by P450<sub>scc</sub> and P450<sub>arom</sub> to P<sub>4</sub> and E<sub>2</sub> synthesis, respectively (Strauss et al., 1981; Lieberman et al., 1984). The possibility cannot be ignored that NO may also bind with the heme group of P450<sub>scc</sub> and P450<sub>arom</sub> as an inhibition mechanism of steroid synthesis (Van Voorhis et al., 1994; Snyder et al., 1996; Hanke et al., 1998).

Although  $10^{-3}$  M SNP concentrations kept most GC in the G0/G1 stage, these cells progressed through the cell cycle similarly to control and treatments containing  $10^{-5}$  M SNP with or without ODQ while  $10^{-1}$  M SNP induced G1 phase arrest as demonstrated by flow cytometry. NO inhibitory effects on cell proliferation were demonstrated in vascular smooth muscle cells (VSMCS) (Tanner et al., 2000), and T cells (Sato et al., 2007). NO is involved on over-expression of p21 (cyclin-dependent kinase inhibitors) which induces G1 arrest in VSMCS (Tanner et al., 2000), and this elevated concentration is consistent with the anti-proliferative effect of NO (Yang et al., 1996). A disruption of DNA synthesis can interfere with cell replication and lead to cell death (Burney et al., 1997). The present results demonstrated that greater NO concentrations inhibited GC progression through the cell cycle suggesting a cytostatic effect.

The results from the current experiment, that utilized the culture system previously described, demonstrated that lesser NO concentrations can negatively regulate P<sub>4</sub> synthesis by a cGMP independent pathway; however, this pathway is involved in the negative regulation of progesterone synthesis by factors not elucidated. The NO/cGMP pathway is, however, involved with synthesis, but this pathway also seems to be involved in inhibition of estradiol synthesis by independent NO mechanisms. NO may, therefore, be involved in steroidogenesis regulation, GC differentiation by independent or dependent cGMP mechanisms. In addition, cytotoxic effects verified by greater

NO concentration treatments in the present experiment likely initiated with mitochondrial function inactivation and subsequently with interference on cell cycle progression and P<sub>4</sub> and E<sub>2</sub> synthesis inhibition. Probably this NO toxicity that was induced could lead a cell death by apoptosis or necrosis. Therefore, these findings have important implications regarding NO concentrations in GC function.

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