

# PAPP-A ENHANCES THE ANTIOXIDATIVE EFFECTS OF IGF-1 DURING BOVINE IN VITRO EMBRYO PRODUCTION

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## ABSTRACT

We investigated whether exogenous pregnancy-associated plasma protein-A (PAPP-A) enhances the antioxidant role of insulin-like growth factor-1 (IGF-1) in bovine *in vitro* embryo production (IVP). We performed standard *in vitro* maturation (IVM) and *in vitro* culture (IVC) or added menadione to promote an oxidative stressed microenvironment and evaluated the antioxidant effect of IGF-1 alone or in combination with PAPP-A (IGF-1/PAPP-A). In IVM, the treatments did not affect oocyte nuclear development, total GSH content, *cumulus* cell gene expression, and blastocyst yield. Nevertheless, IGF-1/PAPP-A treatment prevented an increase in reactive oxygen species (ROS) and mitochondrial membrane potential (MMP) levels. In IVC, the treatments did not affect the total GSH content on blastocysts and IVC media, but IGF-1 and IGF-1/PAPP-A treatments increased blastocyst yield compared to the menadione group. In addition, IGF-1/PAPP-A treatment

had lower ROS levels and regulated genes related to embryonic quality compared to the control and menadione groups. Overall, we showed that PAPP-A could enhance the antioxidant role of IGF-1 during IVP in cattle by avoiding higher ROS levels in oocytes and blastocysts and modulating the transcriptional abundance of genes involved in oxidative protection and embryonic quality.

**Keywords:** Pappalysin, oxidative stress, ROS, cattle, insulin-like growth factor-1, gene expression.

## 1. INTRODUCTION

Reactive oxygen species (ROS) are produced as natural products of aerobic metabolism, even when cells are under basal conditions. Most ROS production is a natural result of the oxidative phosphorylation process that occurs in the mitochondrial electron transport chain to generate adenosine triphosphate (ATP)(Reviewed by Guerin et al., 2001). However, excessive ROS levels have been associated with oxidative stress, which can cause lipid peroxidation, mitochondrial function damage, and DNA structure alterations. The latter has been demonstrated by events of base pair loss or fragmentation of the DNA core (Birben et al., 2012; Nasr-Esfahani et al., 1990; Noda et al., 1991; Richter et al., 1995; Roth, 2018). Some specific cells and/or conditions require higher ATP production, such as the oocytes and embryos during the early development stages (Chappel, 2013). Under physiological conditions, these cells rely on an antioxidant system that assures reduced and/or absence of any cell damage caused by increased ROS levels production. However, high ROS levels have been described in bovine oocytes submitted to *in vitro* maturation (IVM) and bovine blastocysts produced by *in vitro* embryo production (IVEP) (Dalvi et al., 2005; Morado et al., 2009), such high ROS levels

in *in vitro* conditions even have led to embryo development arrest at the 2-cell stage in mice (Nasr-Esfahani et al., 1990).

Supporting the fact that an increase in ROS levels has harmful effects, two studies demonstrated that bovine embryos when cultured with menadione, a pro-oxidant stressor inducer of ROS production when under oxygen presence (Monks et al., 1992; Nutter et al., 1992), indeed resulted in decreased the blastocyst yield production and increased apoptotic cells number (de Assis et al., 2015; Moss et al., 2009). On the contrary, the insulin-like growth factor-1 (IGF-1) has shown a systemic antioxidant effect. (Higashi et al., 2010; Sukhanov et al., 2007). Corroborating with that, it has been demonstrated that IGF-1 media supplementation prevented menadione deleterious effects (Monks et al., 1992; Nutter et al., 1992), although such effect seems to be independent of ROS concentrations in bovine embryo culture (Moss et al., 2009).

The IGF-1 bioavailability system is regulated by IGF-binding proteins (IGFBPs), which decrease IGF-1 activity. Oppositely, the pregnancy-associated plasma protein-A (PAPP-A) increases IGF-1 availability and facilitates IGF-1 binding to its receptors (Froesch et al., 1985; Jones & Clemmons, 1995; Lawrence et al., 1999; LeRoith et al., 1992). PAPP-A is mainly secreted by granulosa cells (Conover et al., 1999). It acts by cleaving IGFBP-2, IGFBP-4, and IGFBP-5 in many species' tissues, including cattle (Laursen et al., 2007; Laursen et al., 2001; Mazerbourg et al., 2003; Mazerbourg & Monget, 2018; Monget et al., 2003; Rivera & Fortune, 2003). The relevance of its role in reproductive tissues has been demonstrated by producing knockout mice for PAPP-A presence, which resulted in reduced animal fertility, lower *cumulus*-oocyte complexes (COCs) quantification, and lower progesterone and estradiol levels during ovarian superovulation cycles (Laursen et al., 2001; Nyegaard et al., 2010). Because of that, PAPP-A has been used as a dominant follicle selection marker. Moreover, its

contributions to the reproductive field have not been limited to *in vivo* conditions. Under *in vitro* conditions, the presence of PAPP-A increased the IGF-1 availability in the COCs *in vitro* maturation media, which later in the blastocyst production was associated with modulation on genes involved with lipid metabolism, mitochondrial functionality, and blastocyst competence (Dellaqua et al., 2023; Giroto et al., 2019).

Collectively, due to its physiological relevance, many studies have evaluated the effect of IGF-1 supplementation on ROS production during *in vitro* embryo production (IVEP), however, the results of these *in vitro* studies are controversial. Taking into consideration the IGF-1 bioavailability system, we hypothesize that the effect of IGF-1 supplementation in the IVEP system might be enhanced by its association with PAPP-A. Therefore, our objective was to evaluate the antioxidant effect of IGF-1 associated with PAPP-A on the bovine embryo production system, regarding the oocyte and blastocyst quality. To the best of our knowledge, the synergist of PAPP-A and IGF-1 has not yet been investigated.

## 2. MATERIALS AND METHODS

Unless otherwise mentioned, all products were purchased from Sigma-Aldrich Co. (St. Louis, MO, USA) with high purity.

### 2.1. Experimental design

To expand our understanding of the IGF-1/PAPP-A system involvement with oxidative stress, ROS production was induced in oocytes and embryos samples with menadione treatment (5  $\mu$ M), as previously described (Cavallari et al., 2019; Chen et al., 2017; de Assis et al., 2015; Moss et al., 2009). In a pilot experiment, the effect of PAPP-A alone on ROS concentration was investigated, however, no modulation was

observed in this parameter (data not shown) Therefore, this present study followed with PAPP-A associated with IGF-1 (Figure 1).

## FIGURE 1

### 2.2. *Cumulus*-oocyte complex collection

Ovaries from crossbred female cattle (*Bos taurus indicus*) were obtained in a slaughterhouse and transported to the laboratory in sodium chloride 0.9% at 36 °C. Follicles with a diameter of 2-8 mm were aspirated with an 18-gage needle and collected in a 15 mL conical tube. After sedimentation, COCs were collected and selected using a stereomicroscope. Only COCs with homogeneous cytoplasm and a compact multilayer of grade 1 and 2 *cumulus* were used for the experiments, as described by Stojkovic et al. (Stojkovic et al., 2001).

### 2.3. In vitro maturation

The selected COCs were washed in buffered wash medium and randomly transferred to four-well dishes containing 10 µL/COC basic IVM medium, consisting of TCM199 with bicarbonate and Earle's salts, recombinant follicular stimulating hormone (0.1 IU/mL solution; Gonal-f; Merck Serono, Bari, Italy), sodium pyruvate (22 µg/mL), amikacin (75 µg/mL), and fatty acid-free bovine serum albumin (BSA; 4 mg/mL). The basic IVM medium was used from the control group (C group). In addition, three other groups consisted of IVM medium supplemented with menadione (5 µM): only menadione (M group), menadione with IGF-1 (100 ng/mL, IGF-1 recombinant human; Life Technologies, Carlsbad, CA, USA; M/I group), or menadione with IGF-1 and PAPP-A (100 ng/mL, recombinant human Pappalysin-1; R&D Systems Inc., Minneapolis, MN, USA; M/I-P group) according to the experimental design (Figure 1a). After 24 hours at

38.5 °C in humidified air with 5.5% CO<sub>2</sub>, the COCs were submitted to *cumulus* cell removal, and oocyte and cumulus cell samples were collected separately for analysis.

## 2.4. In vitro fertilization

The presumptive matured COCs (not treated, basic IVM media) were washed three times with in vitro fertilization (IVF) medium and placed in plates with 90 µL of IVF medium (25 COCs/droplet) and covered with silicone oil (Quimesp Química, Guarulhos, São Paulo, Brazil). The IVF medium contained thyroid albumin-lactate-pyruvate supplemented with BSA (6 mg/mL), pyruvate (22 µg/mL), amikacin (75 µg/mL), heparin (30 µg/mL), and penicillamine–hypotaurine–epinephrine solution (20 µM penicillamine, 10 µM hypotaurine, and 1 µM epinephrine). Nelore bull semen from a single ejaculation (CRV Lagoa, Sertãozinho, São Paulo, Brazil) was thawed at 37 °C for 30 s and spermatozoa were washed in a discontinuous gradient with BotuFIV Select SPERM<sup>®</sup> (Botupharma, Botucatu, Brazil) prepared by adding 45% to 90% BotuFIV Select SPERM<sup>®</sup> in a 1.5 mL centrifuge tube. The sperm samples were added to the top of the gradient and centrifuged at 3000 × g for 5 minutes (MiniSpin<sup>®</sup>; Eppendorf, Hamburg, Germany). The supernatant was then removed, and the sperm pellet obtained was counted in a Neubauer chamber. Sperm were resuspended in IVF medium to reach a final concentration of 1×10<sup>6</sup> cells/mL, and 7 µL were added to each IVF droplet and incubated for 18 hours at 38.5 °C in a saturated humidity atmosphere containing 5.5% CO<sub>2</sub>. Subsequently, the presumptive zygotes (PZ) were denuded of the remaining surrounding *cumulus* cells by vortexing and washed three times in Synthetic Oviduct Fluid (SOF) medium (Holm et al., 1999).

## 2.5. In vitro culture

Embryo culture was performed in four-well dishes with 10  $\mu$ L/PZ of IVC medium (SOF medium, sodium pyruvate (0.2 mM), BSA (5 mg/mL), and 2.5% v/v fetal bovine serum). On day 4 post-IVF, all structures were transferred to a new culture medium as previously performed by Moss et al. (2009). All structures were placed in a new four dish containing IVC medium without supplementation (control, C group) or in wells supplemented with menadione (5  $\mu$ M; M group), menadione with IGF-1 (100 ng/mL; M/I group) or menadione with IGF-1 and PAPP-A (100 ng/mL; M/I+P group), as shown in the experimental design (Figure 1b). On day 5, the structures were washed three times in IVC medium and transferred to a new four-well dish with IVC medium (without treatments) until day 7. All cultures were performed in a humidified and controlled atmosphere (5% CO<sub>2</sub>, 5% O<sub>2</sub>, and 90% N<sub>2</sub>) at 38.5 °C. Blastocyst yield was determined on day 7 after IVF and embryos were collected for analysis (Figure 1b).

## 2.6. Transcript profile of *cumulus* cells and blastocysts

Three out of six replicates were randomly selected for gene expression analysis of *cumulus* cell samples (n=3 samples/group). For blastocyst analysis, a sample was defined as a pool of three blastocysts (n=5 samples/group).

Total RNA was extracted using the PicoPure™ RNA Isolation Kit (Applied Biosystem®, Foster City, CA, USA) according to the manufacturer's protocol. After purification, the RNA samples were eluted in 12  $\mu$ L of RNase-free water. Total RNA (100 ng/reaction for *cumulus* cell samples and total RNA samples for embryos) was incubated with DNase I (1 IU/ $\mu$ g; Invitrogen, São Paulo, Brazil) and then reverse transcribed with random primers according to the protocol of the High-Capacity Kit (Applied Biosystem®, Foster City, CA, USA).

Subsequently, the mRNA abundance of 96 genes (see Supplementary Table S1) was assessed in the *cumulus* cells and blastocysts samples. All analyzes were performed using the Biomark HD assay (Fluidigm, South San Francisco, CA, USA) as previously described (Dellaqua et al., 2023; Fontes et al., 2020; Franchi et al., 2019; Giroto et al., 2019). The relative expression values for each gene were calculated with the  $\Delta\text{Ct}$  method and the data were normalized using the geometric mean values of the most stable reference genes for each cell type (NormFinder software tested). The geometric mean of the mRNA abundance of *ACTB*, *GAPDH*, *PPIA*, and *RPL30* was used to normalize the blastocyst samples data, while *ACTB*, *B2M*, *PPIA*, and *RPL30* were used for the *cumulus* cells data.

## **2.7. Assessment of nuclear progression, mitochondrial activity, and intracellular reactive oxygen species in oocytes and embryos**

A combined fluorescent staining technique using Hoechst 33342, MitoTracker<sup>®</sup> Red CMXRos, and CellROX<sup>™</sup> Green Reagent (Invitrogen) was used to determine the nuclear maturation stage, indirect mitochondrial inner membrane potential (MMP), and ROS concentration in oocytes, respectively. In the blastocyst samples, only the ROS concentrations of the non-expanded blastocysts were analyzed.

After IVM, oocytes were denuded by vortexing and washed in phosphate-buffered saline (PBS; pH 7.2) containing 1 mg/mL polyvinylpyrrolidone (PBS-PVP). Next, oocytes were incubated with Hoechst (1  $\mu\text{g/mL}$ ), CellROX<sup>™</sup> Green (5  $\mu\text{M}$ ), and MitoTracker<sup>®</sup> Red (0.5  $\mu\text{M}$ ). For blastocyst samples, CellROX<sup>™</sup> Green (5  $\mu\text{M}$ ) and Hoechst (1  $\mu\text{g/mL}$ ) were used. In both sample types, the incubations were done for 30 minutes at 5.5%  $\text{CO}_2$  and 38.5  $^{\circ}\text{C}$ . After, samples were washed three times in PBS-PVP and fixed with 4% paraformaldehyde for 15 minutes. Last, samples were washed three

times in PBS-PVP, placed in glycerol microdroplets on a microscope slide, covered with a coverslip, and analyzed with a Leica DM4500 epifluorescence microscope (Leica Camera AG, Wetzlar, Germany) to determine DNA nuclear progression and ROS and MMP fluorescence intensities. Images were acquired with LAS software using a camera attached to the microscope. Quantification of average pixel intensity was performed using Image J software (NIH, Bethesda, MD, USA) for ROS and MMP analysis.

## 2.8. Glutathione total content determination

Total GSH content was quantified using a recycling assay previously as described (Anderson, 1985). Two sample types were analyzed, cells and culture media samples. The cell samples were embryos (n=5 pools of 5 embryos/pool/group), denuded oocytes (n=5 pools of 10 oocytes/pool/group), and *cumulus* cells (n=5 pools of *cumulus* cells from 10 oocytes/pool/group) were stored in trichloroacetic acid (TCA) at -80 °C. The culture medium samples were IVM media, IVC media (Day 5), and IVC media (Day 7), all samples consisted of 20 µL volume (n=5 samples/group), which were stored at -80 °C and TCA was added immediately before the GSH assay. Each sample (except the medium samples) was vortexed and centrifuged at 10000 ×g for 20 s. Standards containing 1.56 to 100 nmol/mL reduced GSH were prepared simultaneously with the samples in TCA. The samples and standards were stored on ice until they were loaded onto the microtiter plate. Then, 20 µL of each sample or standard was added to a 96-well microtiter plate, followed by 140 µL of NADPH (0.3 mM), 5 µL of sodium phosphate buffer (143 mM) with EDTA (6.3 mM), and 20 µL of DTNB (6 mM). All reagents were diluted in sodium phosphate buffer (143 mM) with EDTA (6.3 mM). The plate was incubated at 30 °C for 5 minutes and then 15 µL of GSH reductase (4.0 IU/mL) was added. Immediately, the

plate was analyzed at 412 nm in a microplate reader (BioTek Instruments, Inc., Winooski, VT, USA) with 10 replicate measurements.

## 2.9. Statistical analysis

Meiosis progression and blastocyst rates were calculated as percentages and arcsine-transformed. All data were tested for normality using the Shapiro-Wilk test. The data of gene expression, total GSH content, MMP, and ROS fluorescent intensity, which showed a non-normal distribution, were log-transformed to achieve a normal distribution. For parametric data, analysis of variance (ANOVA) followed by the Tukey test was performed to compare means, and for non-parametric data, the Kruskal-Wallis test followed by the Wilcoxon test was performed to compare means. All data are expressed as mean  $\pm$  standard error of the mean, and differences were considered significant when  $P \leq 0.05$ .

## 3. RESULTS

### 3.1. Effects of IGF-1/PAPP-A treatment during bovine IVM

When the treatments were performed during the IVM, no difference was observed regarding the percentage of oocytes that reached metaphase II and the blastocyst yield (Table 1). The total GSH content of the COCs, embryos, and IVM medium was also unaffected (Table 2). On the other hand, when analyzing oocyte samples, similar MMP levels were detected in the IGF-1/PAPP-A group (M/I+P) and the control group (C), which was lower compared to the menadione (M) and IGF-1 (M/I) groups (Figure 2b). In addition, the ROS levels in the oocyte samples from IGF-1/PAPP-A (M/I+P) were lower than menadione (M) and IGF-1 (M/I) groups, although, higher than the control group (C, Figure 2b). Concerning the transcripts abundance analysis, in the *cumulus* cells samples,

the *VNN1* gene, a stress marker, was remarkably lower in the IGF-1 alone (M/I group) and it combined with PAPP-A (M/I+P group) when compared to the menadione group (M) (Figure 3). While, in the blastocyst samples, no difference was observed among the groups, although *CDX2* appears to be more abundant in the control (C) and IGF-1/PAPP-A (M/I+P) groups compared to the M and M/I groups (P=0.0847) and *FOXO2* tended to be more abundant in the M/I+P group (P=0.0647; Figure S1).

TABLE 1

TABLE 2

FIGURE 2

FIGURE 3

### 3.2. Effects of IGF-1/PAPP-A treatment during bovine IVC

When the treatments were performed during the IVC, the blastocyst yield was higher in the IGF-1 alone (M/I) and it combined with PAPP-A (M/I+P) groups compared to the menadione (M) group (Figure 4b). Moreover, lower ROS levels were detected in the IGF-1/PAPP-A (M/I+P) group compared to control (C) and menadione (M) groups (Figure 4). Furthermore, the transcripts abundance of *ACSL3*, *EGFR*, *HMOX1*, *IMPDH2*, and *SLC2A5* were higher in the IGF-1/PAPP-A group (M/I+P) compared to the menadione group (M; Figure 5). While no alteration was observed in the total GSH quantification in the IVC (D5 and D7) medium or the blastocysts samples (Table 3).

FIGURE 4

FIGURE 5

TABLE 3

## 4. DISCUSSION

In this study, we proposed a new strategy for the use of recombinant IGF-1 in the *in vitro* embryo production technology. For the first time, the association of recombinant PAPP-A with IGF-1 enhanced the antioxidant role of the latter when under the *in vitro* stressed microenvironment. The results show that exogenous IGF-1/PAPP-A, when added to the IVM media, prevents ROS and MMP levels increase in the oocyte after maturation. In addition, blastocysts derived from these oocytes had lower ROS levels. Moreover, when IGF-1/PAPP-A was added to the IVC media, it demonstrated an impressive modulation in the blastocyst transcriptional profile.

Previous studies have presented conflicting results regarding the effects of ROS or IGF-1 on the oocyte nuclear stage (Ascari et al., 2017; Chen et al., 2017; Gomez et al., 1993; LaRosa & Downs, 2006; Lorenzo et al., 1994; Reed et al., 1993; Rieger et al., 1998; Takami et al., 1999; Tamura et al., 2008; Yang et al., 2022). A study in rodents has shown a reduced progression in the oocyte nuclear maturation when they are under a stressful environment induced by menadione. In the same study, the authors demonstrated that the oocyte maturation progress until the MII stage could be restored by using an antioxidant agent, such as melatonin (Chen et al., 2017). In disagreement with this data, in our study, the menadione treatment did not impair the oocyte nuclear maturation (Table 1). This same absence of menadione effect on meiosis progression was observed in murine COCs, which was demonstrated to be due to the protection effect of the *cumulus* cells (LaRosa & Downs, 2006). Therefore, the presence of *cumulus* cells surrounding the oocytes might be an explanation for the lack of effect of the menadione treatment in our study, although higher ROS levels were observed in these oocytes.

Corroborating with this latter result, other studies also observed higher ROS levels in oocytes and embryos exposed to menadione treatment. And again, treatments with antioxidants, such as melatonin, were able to prevent it (Ascari et al., 2017; Cavallari et

al., 2019; Chen et al., 2017; de Assis et al., 2015; Moss et al., 2009). Although IGF-1 is classified as an antioxidant, a previous study demonstrated no effect of IGF-1 in modulating ROS levels (Yang et al., 2022). In the same way, our study showed no modulation of ROS levels by the IGF-1 group (M/I), however, when IGF-1 was associated with PAPP-A (M/I+P), the ROS levels in oocytes were significantly lower than in the menadione group.

Interestingly, the quantification of ROS levels in blastocysts showed a similar pattern in the menadione-treated and control group, which might be due to the *in vitro* condition already being a stressful environment. Under these conditions, the IGF-1/PAPP-A combination (M/I+P) resulted in lower ROS levels compared to the control and menadione groups. IGF-1 has been shown to lower systemic and vascular ROS concentrations while increasing total serum antioxidant activities and capacity, thereby reducing oxidative damage in the brain and liver of rodents (Garcia-Fernandez et al., 2008; Higashi et al., 2010; Sukhanov et al., 2007). The discrepancy in the outcomes of using IGF-1 alone or combined with PAPP-A might be the key factor. As mentioned before, the IGFBPs decrease IGF-1 activity, and their expression is known to occur in oocytes and embryos during their development (Armstrong et al., 2002; Satrapa et al., 2013; Sawai et al., 2005; Winger et al., 1997). Moreover, BSA and/or fetal bovine serum are often added to IVM and IVC media, and both supplements contain IGFBPs (Slaaby et al., 2008; Zheng et al., 2006). As proof of that, it has been shown that BSA presence in culture media reduces IGF-1/IGF-1R binding. In contrast, PAPP-A addition in bovine IVM media increases IGF-1-free concentrations, even with BSA supplementation (Giroto et al., 2019). Therefore, we suggest that PAPP-A presence is required to increase IGF-1 activity to reduce ROS levels in bovine IVP systems.

Moreover, the increase in ROS levels has also been correlated to mitochondrial damage, which typically leads to lower MMP levels (Hao et al., 2011; Loor et al., 2010). On the other hand, MMP is generated by protons moving during ATP synthesis, which means that higher MMP has been associated with greater energy production (Cottet-Rousselle et al., 2011). However, under unbalanced energy demand, the exponential increase in MMP levels leads to high ROS production (Lambert & Brand, 2009; Murphy, 2009; Starkov, 2008), which even impairs ATP synthesis and results in detrimental effects in the cells (Berruyer et al., 2004; Miwa & Brand, 2003; Ramzan et al., 2010; Rottenberg et al., 2009; Starkov & Fiskum, 2003). Therefore, it is expected from normal cell function maintenance of MMP levels (Kaim & Dimroth, 1999; Korshunov et al., 1997; Ramzan et al., 2010; Rottenberg et al., 2009; Starkov & Fiskum, 2003). Regarding this correlation, it has been proposed that controlling the MMP might be a mechanism for mitochondrial ROS regulation. For instance, a 10 mV reduction in MMP was sufficient to eliminate approximately 70% of the ROS production by complex I (Miwa & Brand, 2003). It was observed in the present study high ROS levels in oocytes from M and M/I groups, which were associated with high MMP levels. This pattern might indicate a disbalance in the cells from these groups. However, higher ROS levels were detected in the M/P+I group compared to the control group, while the MMP level was similar in the M/P+I and C groups. Based on this, we suggest that the decrease of MMP levels by the IGF-1/PAPP-A association (M/I+P) could be a mechanism to reduce ROS levels in oocytes.

Regarding the relative transcripts abundance analyzis, it appears that neither IGF-1 (M/I) nor PAPP-A (M/I+P) affected *cumulus* cells compared to control or menadione. Except by *VNNI*, whose relative mRNA abundance was downregulated by IGF-1 (M/I) alone and in combination with PAPP-A (M/I+P) compared to the menadione group (M). *VNNI* is a stress marker that modulates the release of glutathione (GSH) when cells are

exposed to oxidative stress (Berruyer et al., 2004; Nivet et al., 2013). GSH appears to be the most important protection against oxidative stress in oocytes and embryos (Feuerstein et al., 2007; Gardiner & Reed, 1994; Takahashi et al., 1993), and indeed studies have observed that the use of antioxidant substances during IVM or IVC leads to an inverse correlation between GSH and ROS levels (Kere et al., 2013; Kwak et al., 2012; Wang et al., 2014). However, the GSH levels analyzed in the present study were unaffected by the experimental conditions when quantified in the COCs, embryos, IVM, and IVC media. Interestingly, Sovernigo, et al. (2017) found that only some antioxidants in cattle show this inverse correlation between ROS and GSH levels, which might explain why our results showed that the IGF-1 is not directly related to the GSH pathway.

Concerning the relative transcripts abundance in blastocysts, few genes were upregulated in the M/I+P compared to the M group. The modulated genes were *IMPDH2* and *EGFR*, which are related to cell replication, proliferation, and differentiation (Adamson & Wiley, 1997; Jackson et al., 1975; Kozhevnikova et al., 2012; Zimmermann et al., 1998); *ACSL3*, which plays a key role in lipid biosynthesis and has been shown to restore survival rates after vitrification of low-quality embryos (Mashek et al., 2007; Soupene & Kuypers, 2008; Valente et al., 2019); *HMOX1* is reported to activate the transcriptional machinery that drives the induction of antioxidant genes (Lin et al., 2007), and *SLC2A5* that is a membrane transporter which has a high specificity for fructose (Burant et al., 1992; Mueckler & Thorens, 2013). It is also known that *SLC2A5* transcription began at the time of bovine embryonic genome activation, suggesting that the early embryo is capable of transporting this energy substrate and that it may be associated, for example, with the production of ribose 5-phosphate, an essential precursor for nucleotide synthesis (Augustin et al., 2001).

#### GRAPHICAL ABSTRACT

## 5. CONCLUSION

Taking it all together, we suggest that PAPP-A might be a crucial factor for the antioxidant role of IGF-1 in bovine *in vitro* production. Only associated with PAPP-A that IGF-1 prevented an increase in the ROS levels in oocytes and embryos under menadione-induced stress. Furthermore, the combination of these proteins appears to activate pathways associated with repair processes, which we demonstrated to result in higher embryo quantity and quality. We conclude that IGF-1 needs PAPP-A association to act as an antioxidant factor in bovine oocytes and embryos produced *in vitro*.

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## FIGURE LEGEND

**Figure 1.** Experimental design. Experiment 1: (a) Insulin-like growth factor-1 (IGF-1) and pregnancy associated plasma protein-A (PAPP-A) addition in bovine in vitro maturation (IVM). Immature *cumulus*-oocyte complexes (COCs) (50 COCs/group) were selected and randomly distributed among the groups: C (IVM medium), M (IVM medium + menadione), M/I (IVM medium + menadione + IGF-1), and M/I+P (IVM medium + menadione + IGF-1 + PAPP-A). The COCs were matured for 24 h, and then the oocytes were separated to *cumulus* cells mechanically or followed to in vitro fertilization (IVF) and culture (IVC) until day 7. The IVM medium was collected to assess the total glutathione (GSH) content. (b) IGF-1 and PAPP-A addition in bovine IVC. Experiment 2: immature COCs (50 COCs/well) were selected and matured for 24 h, followed by IVF and IVC for 3 days in IVC medium. On day 4, all the structures were distributed among the groups: C (IVC medium), M (IVM medium + menadione), M/I (IVM medium + menadione + IGF-1), and M/I+P (IVM medium + menadione + IGF-1 + PAPP-A) and incubated for 24 h. On day 5, embryos were washed thrice in IVC medium and replaced in a four-well dish with the control IVC medium (without any treatment) until day 7. The IVC medium was collected at days 5 and 7 to assess the total GSH content.

**Figure 2.** Effects of insulin-like growth factor-1 (IGF-1) with or without pregnancy-associated plasma protein A (PAPP-A) on bovine *cumulus*-oocyte complex in vitro matured in a stressed environment induced by menadione presence. The symbols + and - indicate presence and absence, respectively. (a) Representative image of oocytes stained by MitoTracker® Red CMXRos dye (red) and CellROX™ Green dye (green). (b) Pixels intensity of mitochondrial membrane potential (MMP) of oocytes (n=77-88 oocytes/group; 5 replicates; p=0.0001) detected by MitoTracker® Red CMXRos dye and reactive oxygen species (ROS) (n=123-132 oocytes/group; 5 replicates; P= 0.0001) detected using CellROX™ Green dye. The symbols + and - indicate presence and absence, respectively. Data are presented as mean  $\pm$  standard error of the mean. Differences were considered significant when  $P \leq 0.05$ . Different letters indicate statistical differences. C: Control; M: Menadione; M/I: Menadione + IGF-1; M/I+P: Menadione + IGF-1 + PAPP-A.

**Figure 3.** mRNA relative abundance of genes differently expressed in the *cumulus* cells. The symbols + and - indicate presence and absence, respectively. Target genes were normalized with the reference genes (*ACTB*, *B2M*, *PPIA*, and *RPL30*) geometric mean using the  $\Delta\Delta C_t$  method. Data are presented as fold change [ $2^{(-\Delta\Delta C_t)}$ ]  $\pm$  standard error of the mean. Differences were considered significant when  $P \leq 0.05$ . Different letters indicate statistical differences. C: Control; M: Menadione; M/I: Menadione + IGF-1; M/I+P: Menadione + IGF-1 + PAPP-A.

**Figure 4.** Effects of insulin-like growth factor-1 (IGF-1) with or without pregnancy associated plasma protein-A (PAPP-A) during bovine in vitro culture on embryo development of blastocysts submitted to the stressed environment induced by menadione presence on day 4 post-in vitro fertilization for 24 h. The symbols + and - indicate presence and absence, respectively. (A) Representative image of blastocysts stained by hoechst 33342 (blue) and CellROX™ Green dye (green) (B) Percentage rate of blastocysts produced at day 7 (4 replicates) and pixels intensity of reactive oxygen species (ROS) detected using CellROX™ Green dye in the blastocysts (n=20-30/blastocysts/group; 4 replicates). Data are presented as mean  $\pm$  standard error of the mean, and differences were considered significant when  $P \leq 0.05$ . Different letters

indicate statistical differences. C: Control; M: Menadione; M/I: Menadione + IGF-1; M/I+P: Menadione + IGF-1 + PAPP-A.

**Figure 5.** mRNA relative abundance of genes differently expressed in the blastocysts. The symbols + and - indicate presence and absence, respectively. Target genes were normalized with the reference genes (*ACTB*, *GAPDH*, *PPIA*, and *RPL30*) geometric mean using the  $2^{-\Delta Ct}$  method. Data are presented as mean  $\pm$  standard error of the mean. Differences were considered significant when  $P \leq 0.05$ . Different letters indicate statistical differences. C: Control; M: Menadione; M/I: Menadione + IGF-1; M/I+P: Menadione + IGF-1 + PAPP-A.

**Figure 6.** The proposed system with PAPP-A, a metalloprotease that cleaves the IGFBPs, as an enhancer of IGF-1 antioxidative role in IVP. The addition of IGF-1/PAPP-A in bovine IVM of COCs exposed to the stressed environment induced by menadione presence enhances the IGF-1 antioxidative role due to avoid the increased ROS concentration and controlling MMP levels in oocytes besides the downregulation of VNN1, a stress markers in *cumulus* cells. The IGF-1/PAPP-A association also showed to enhance the IGF-1 antioxidative role when applied in IVC. The proteins association increased the blastocyst yield and avoided the high ROS concentration in blastocysts besides modulated transcripts encoding proteins involved with antioxidative protection, lipid metabolism, and embryonic quality showing the PAPP-A is a feasible enhancer to IGF-1 antioxidative role. Red arrows: downregulation/decrease ( $P < 0.05$ ); blue arrow: upregulation/increase ( $P > 0.05$ ); green arrow: upregulation ( $0.1 > P > 0.05$ ).

## TABLE LEGEND

**Table 1.** Oocytes' nuclear stage and embryo development from COCs exposed to the stressed environment during IVM for 24 h with no treatment (M), IGF-1 (M/I), IGF-1/PAPP-A (M/I+P), and control (C).

C: control; IVM medium.

M: IVM medium + menadione (5  $\mu$ M);

M/I: IVM medium + menadione + IGF-1 (100 ng/mL);

M/I+P: IVM medium + menadione + IGF-1 + PAPP-A (100 ng/mL).

Blastocyst yield: total of blastocysts/total of oocytes in IVM.

Abbreviations: TII, telophase II; MII, metaphase II.

**Table 2.** Total GSH content (mean  $\pm$  SEM) in IVM medium and COCs submitted for 24h to the stressed environment with no treatment (M), IGF-1 (M/I), IGF-1/PAPP-A (M/I+P), and control (C).

C: control; IVM medium.

M: IVM medium + menadione (5  $\mu$ M);

M/I: IVM medium + menadione + IGF-1 (100 ng/mL);

M/I+P: IVM medium + menadione + IGF-1 + PAPP-A (100 ng/mL).

771 *Cumulus* cells pool: *cumulus* cells from 10 COCs; n = 5 pools/group.

772 Medium samples: n = 5 samples of 20  $\mu$ L.

773 Oocytes pool: 10 oocytes; n = 5 pools/group.

774 **Table 3.** Total GSH content (mean  $\pm$  SEM) in IVC medium on days 5 and 7 and blastocyst  
775 submitted for 24h to the stressed environment with no treatment (M), IGF-1 (M/I), IGF-  
776 1/PAPP-A (M/I+P) and control (C).

777

778 C: control; IVC medium.

779 M: IVC medium + menadione (5  $\mu$ M);

780 M/I: IVC medium + menadione + IGF-1 (100 ng/mL);

781 M/I+P: IVC medium + menadione + IGF-1 + PAPP-A (100 ng/mL);

782 Blastocyst pool: 5 blastocysts; n = 5 pools/group.

783 Medium samples: n = 5 samples of 20  $\mu$ L.

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