ELISA MARIANO PIOLTINE

USO DE TUDCA NA MODULAÇÃO DO ESTRESSE DO RETÍCULO ENDOPLASMÁTICO EM ETAPAS DA PRODUÇÃO *IN VITRO* DE EMBRIÕES BOVINOS

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Tese apresentada ao Programa de Pósgraduação do Instituto de Biociências de Botucatu, Universidade Estadual Paulista -UNESP, para a obtenção do título de Doutor em Farmacologia e Biotecnologia.

Orientador: Prof. Associado Marcelo Fábio Gouveia Nogueira

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Palavras-chave: Estresse do retículo endoplasmático; Produção *in vitro* de embriões; TUDCA.

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"Na vida, não vale tanto o que temos, nem tanto importa o que somos. Vale o que realizamos com aquilo que possuímos e, acima de tudo, importa o que fazemos de nós."

Chico Xavier

RESUMO

Embora estágios iniciais de embriões de mamíferos tenham aparentemente uma grande plasticidade, eles, assim como o oócito em maturação, são bastante sensíveis a estresses exógenos. Responder a esses estresses é uma parte vital da fisiologia celular. Cada vez mais é aparente que um dos principais mecanismos, para iniciar a resposta celular a uma variedade de estressores exógenos, reside no retículo endoplasmático (RE). O RE é uma importante organela responsável pela síntese, processamento e transporte de proteínas e lipídeos. No entanto e in vitro, a perturbação do microambiente do RE pode causar alterações na estrutura das proteínas levando ao acúmulo de proteínas com o incorreto dobramento, uma condição denominada estresse do RE. Dependendo da intensidade do estresse, o RE pode desencadear a apoptose pela Unfolded Protein Response. Em várias espécies, estudos observaram uma melhoria da competência do oócito e no desenvolvimento do embrião quando foram adicionados inibidores do estresse do RE no meio de maturação e/ou meio de cultivo in vitro (CIV). Dentre os inibidores do estresse do RE, o ácido tauroursodesoxicólico (TUDCA) se mostrou benéfico na produção in vitro de embriões. Entretanto, pouco é conhecido sobre o seu efeito na maturação oocitária e no desenvolvimento do embrião bovino in vitro. Portanto, os objetivos gerais desta Tese foram: Capítulo 1 - investigar o efeito da adição de TUDCA durante a maturação in vitro (MIV) sobre a competência do oócito e do embrião [maturação nuclear, atividade mitocondrial, produção de espécies reativas de oxigênio (ERO), formação de pro núcleo, formação de blastocisto e abundância de transcritos em oócitos e embriões] e Capítulo 2 investigar o efeito da adição de TUDCA durante o CIV sobre a competência embrionária e a criotolerância do embrião (cinética de eclosão e abundância de transcritos de embriões antes e após a vitrificação). A adição de TUDCA na MIV não melhorou a competência oocitária. Entretanto, a adição de 200 µM de TUDCA pareceu aliviar o estresse do RE reduzindo a produção de ERO no oócito e aumentando a abundância de transcritos relacionados positivamente com a defesa antioxidante do oócito e do embrião. Contrariamente, a adição de 1.000 µM de TUDCA na MIV pareceu ser tóxica para o oócito e diminuiu a taxa de maturação nuclear bem como a atividade mitocondrial. Além disso, houve um aumento da abundância de transcritos pró-apoptóticos no oócito. Quando o TUDCA foi suplementado no CIV não houve melhora da competência embrionária. De forma semelhante que na MIV, a adição de 1.000 µM de TUDCA no CIV pareceu ser tóxica para o embrião e diminuiu a taxas de blastocisto eclodido. De modo complementar, houve um aumento da abundância de transcritos relatados negativamente com o estresse do

RE, o estresse oxidativo, o metabolismo lipídico e a sobrevivência celular. Entretanto, a adição de 200 μ M de TUDCA no CIV pareceu aliviar o estresse do RE em embriões submetidos à vitrificação. Com isso, houve o aumento da taxa de embrião eclodido após o aquecimento. Além disso, nos blastocistos eclodidos pós-aquecimento foi observada uma redução da abundância de transcritos relacionados negativamente com o estresse do RE e um aumento daqueles relacionados positivamente com o mecanismo antioxidante e o desenvolvimento embrionário.

Palavras-chave: TUDCA; Oócito; Blastocisto; Estresse do retículo endoplasmático; Vitrificação; Bovino.

ABSTRACT

Although early stages of mammalian embryos appear to retain great plasticity, they and maturing oocytes are quite sensitive to exogenous stresses. Responding to those stresses is a vital part of cellular physiology, and it is increasingly apparent that one of the main mechanisms for initiating that response is based on the endoplasmic reticulum (ER). ER is an important organelle responsible for the synthesis, processing, folding and transport of proteins and lipids. However, in vitro, the disturbance of the ER microenvironment can cause changes in the structure and folding of proteins leading to the accumulation of misfolding proteins, a condition called ER stress. Depending on the intensity of the stress, the ER can trigger apoptosis by Unfolded Protein Response. In several species, studies have reported an improvement in oocyte and embryo competence when ER stress inhibitors have been added to the *in vitro* maturation (IVM) medium and/or *in vitro* culture (IVC) medium. Among the ER stress inhibitors, tauroursodeoxycholic acid (TUDCA) has been shown to be beneficial in in vitro embryo production. However, little is known about TUDCA's effect on oocyte maturation and the development of bovine embryo in vitro. Therefore, the main objectives of this Thesis were: Chapter 1 - to investigate the effect of adding different TUDCA concentrations during IVM on the oocyte and embryo competence [nuclear maturation, mitochondrial activity, reactive oxygen species (ROS) production, pronuclei formation, blastocyst formation and transcripts abundance in oocytes and embryos]; and Chapter 2 - to investigate the effect of adding different TUDCA concentrations during IVC on embryo competence and embryo cryotolerance (hatching kinetics and transcripts abundance in embryo before and after the vitrification). The TUDCA addition to the IVM did not improve oocyte competence. However, the 200 µM of TUDCA appeared to relieve ER stress and reduced the ROS production in the oocyte. Concomitantly, there was an increasing of the transcripts abundance positively reported to the antioxidant defense in the oocyte and embryo. In contrast, the 1,000 µM of TUDCA appeared to be toxic to the oocyte by decreasing the nuclear maturation rate and mitochondrial activity in the oocyte. Moreover, there was an increasing of the proapoptotic transcripts' abundance in the oocyte. When TUDCA was supplemented in IVC there was no improvement in embryo competence. As in IVM, the 1,000 µM of TUDCA appeared to be toxic to the embryo - decreasing the hatched blastocyst rate and increasing the transcripts' abundance negatively reported in ER stress, oxidative stress, lipid metabolism and cell survival. However, the 200 µM of TUDCA addition in the CIV seemed to relieve ER stress in the embryos subjected to vitrification, increasing the

hatched embryo rate post-warming. In addition, in hatched blastocysts post-warming it was observed a reduction of the transcripts' abundance negatively reported to the ER stress and an increasing of those positively reported to the antioxidant mechanism and the embryo development.

Keywords: TUDCA; Oocyte; Blastocyst; Endoplasmic reticulum stress; Vitrification; Cattle.

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PREÂMBULO

Esta Tese para o programa de Pós-graduação em Farmacologia e Biotecnologia se encontra dividida em quatro sessões principais. A primeira, denominada "Introdução", descreve o contexto e a problemática relacionada com este trabalho de maneira mais profunda e detalhada. As duas partes subsequentes, Capítulo 1 -"Modulating the endoplasmic reticulum stress in in vitro maturation of bovine oocytes: Could tauroursodeoxycholic acid improve developmental competence?" e Capítulo 2 - "Molecular and cellular effects of the inclusion of tauroursodeoxycholic acid in in vitro culture before to cryopreserve bovine blastocysts", correspondem a duas propostas de manuscritos a serem submetidos como artigos resultantes do trabalho efetuado. Finalizando, a sessão "Considerações Finais" compreende as conclusões gerais (descrevendo de forma sucinta os resultados obtidos) e os resultados suplementares não inseridos nos artigos.

Conforme normas do programa de Pós-graduação, o presente exemplar consiste na versão definitiva da tese a ser encaminhada à biblioteca e que comporá o conjunto de publicações do Instituto de Biociências de Botucatu da UNESP. Entretanto, os artigos científicos aqui apresentados estão no prelo, mas não comprometidos com a publicação em quaisquer periódicos, estando, portanto, passíveis de correções e modificações de amplo espectro, especialmente por parte da banca de defesa.

INTRODUÇÃO

As biotecnologias da reprodução em bovinos, especialmente as técnicas envolvendo a produção de embriões (*in vivo* e *in vitro*), constituem uma importante ferramenta para aumentar a eficiência da pecuária, proporcionando o aproveitamento racional de algumas áreas destinadas à bovinocultura e contribuindo para o melhoramento genético (Lopes *et al.*, 2012, Oliveira *et al.*, 2014).

Estudos na espécie bovina dos sistemas de produção *in vitro* de embriões (PIVE), foram fundamentais para a compreensão de vários fenômenos e mecanismos biológicos que ocorrem durante a maturação oocitária, a capacitação espermática e a fertilização, além do início do desenvolvimento embrionário em estágios préimplantacionais (Gonçalves *et al.*, 2008). Essa importância é demonstrada - nos registros da Sociedade Internacional de Tecnologia de Embriões (IETS) de 2017 - com o total de embriões produzidos *in vitro* (PIV) que ultrapassou aqueles derivados *in vivo* (992.289 vs. 406.287, respectivamente; Viana *et al.*, 2018).

Contudo, é de conhecimento que as condições *in vitro*, nas quais os oócitos e embriões são expostos, causam uma variedade de estresses celulares que contribuem para a perda da competência no desenvolvimento do embrião (Lane e Gardner *et al.*, 2005). Essas condições incluem tensão de cisalhamento durante o manuseio, temperatura, meio de cultura, pH alterado, alta tensão de O_2 e até incidência luminosa (Lane e Gardner, 2000; Thompson *et al.*, 2002; Fleming *et al.*, 2004; Bell *et al.*, 2009). Como consequências, em gametas e embriões, ocorrem alterações na expressão de alguns genes, nos mecanismos epigenéticos, no metabolismo, na indução da apoptose e, finalmente, na redução do desenvolvimento e da viabilidade embrionária quando comparadas com os embriões derivados *in vivo* (Lane e Gardner, 2000; Thompson *et al.*, 2002; Fong *et al.*, 2007; Bell *et al.*, 2009). Responder ao estresse exógeno é uma parte vital da fisiologia celular. Está cada vez mais aparente que um dos principais mecanismos, para iniciar a resposta celular a uma variedade de estressores exógenos, reside no retículo endoplasmático (RE; Guzel *et al.*, 2017). O RE é uma importante organela responsável pela síntese, processamento e transporte de proteínas e lipídeos (Brodsky *et al.*, 2011). No entanto e *in vitro*, a perturbação do microambiente do RE pode causar alterações na estrutura das proteínas levando ao acúmulo de proteínas com o incorreto dobramento (Latham *et al.*, 2015), uma condição denominada estresse do RE (Tabas *et al.*, 2011). Para combater os efeitos deletérios do estresse do RE, as células desenvolveram várias estratégias, coletivamente conhecidas como resposta a proteínas malformadas ou UPR (*Unfolded Protein Response*). Inicialmente, a UPR restaura a homeostase do RE pelo aumento da produção de chaperonas moleculares envolvidas no dobramento de proteínas e pela degradação de proteínas malformadas (Hussain *et al.*, 2007). Entretanto, o acúmulo excessivo e prolongado de proteínas malformadas no lúmen do RE redireciona a UPR para ativar a sinalização apoptótica (Szegezdi *et al.*, 2006).

Para o correto desenvolvimento do oócito e do embrião, as proteínas devem ser dobradas adequadamente em sua formatação 3D. Assim, a regulação do estresse/homeostase do RE provavelmente é um mecanismo importante durante a PIVE (Zeng *et al.*, 2005; Lockwood *et al.*, 2017).

Em várias espécies, estudos observaram uma melhoria da competência do oócito e no desenvolvimento do embrião quando foram adicionados inibidores do estresse do RE no meio de maturação e/ou meio de cultivo *in vitro* (CIV), reforçando que o dobramento correto das proteínas é essencial para a atividade celular (Song *et al.*, 2014; Lin *et al.*, 2016). Vários inibidores já foram utilizados para reduzir o estresse do RE, tais como o ácido tauroursodesoxicólico (TUDCA), o ácido fenilbutírico (PBA) e o salubrinal (Song *et al.*, 2012; Zhang *et al.*, 2012; Sutton-McDowall *et al.*, 2015).

Com função de chaperona química, o ácido biliar TUDCA tem se mostrado benéfico quando adicionado nos meios de PIVE em mamíferos (Zhang *et al.*, 2012; Kim *et al.*, 2012; Yoon *et al.*, 2014). Na espécie bovina, a adição de TUDCA no CIV aumentou a taxa de blastocisto de embriões clonados por transferência nuclear de célula somática (Song *et al.*, 2011). Além disso, o TUDCA foi eficaz na redução de espécies reativas de oxigênio (ERO) e de transcritos marcadores do estresse do RE em embriões expostos à 20% de O_2 (Yoon *et al.*, 2014). Recentemente, a adição de 10 μ M de TUDCA no CIV demonstrou ser benéfico para a crio tolerância de embriões bovinos (Khatun *et al.*, 2020).

Apesar de estudos relatarem o efeito benéfico de TUDCA na PIVE na espécie bovina, pouco é conhecido sobre os efeitos de TUDCA no alívio do estresse do RE em relação à crio tolerância de embriões produzidos *in vitro*. Além disso, até o momento, nenhum estudo investigou os efeitos da adição de TUDCA na maturação *in vitro* (MIV) de oócitos bovinos.

Dessa forma, é importante uma investigação para aprofundar a compreensão dos mecanismos de ação desse inibidor do estresse do RE em etapas da PIVE, como forma de mitigar os estresses causados pelo ambiente *in vitro* e, potencialmente, aumentar a produção de embriões com maior tolerância à criopreservação.

REVISÃO DA LITERATURA

O retículo endoplasmático e suas vias de ativação de resposta ao estresse celular

O RE é uma organela de células eucarióticas, formado a partir da invaginação da membrana plasmática, constituído por uma rede de túbulos e vesículas achatadas e

interconectadas, que se comunicam com a membrana nuclear. O RE pode ser classificado em dois tipos: retículo endoplasmático rugoso (RER), que possui formato achatado e a presença de ribossomos associados à sua membrana; retículo endoplasmático liso (REL) com uma estrutura mais tubular e sem ribossomos (Pagliassotti, 2012). O RE está envolvido em diversos processos dentro da célula como: síntese, enovelamento, glicosilação e transporte de proteínas; síntese e distribuição de fosfolipídios e esteróis; armazenamento e o controle da liberação de Ca²⁺ para o citoplasma (Walter *et al.*, 2011).

Cerca de um terço do proteoma total é sintetizado no RER, sendo esta organela o principal compartimento subcelular envolvido na dobragem e maturação de proteínas destinada às vias endo/exocíticas. Para tal, as proteínas secretoras e transmembrânicas sintetizadas nos ribossomos são translocadas do citoplasma para o lúmem do RE, onde sofrerão modificações pós-traducionais e a dobragem necessária para ser entregue em seu local de ação (Alberts *et al.*, 2002). Para desempenhar tais funções, o RE possui um ambiente altamente especializado e singular para o enovelamento, a formação de pontes dissulfeto e a glicosilação de proteínas. Este ambiente é mantido pelo alto nível de cálcio, um ambiente oxidativo e pela presença de chaperonas cálcio-dependentes, lectinas, foldases e glucosidases, as quais são responsáveis pelo dobramento e controle de qualidade das proteínas recém-sintetizadas (Walter *et al.*, 2011; Bravo *et al.*, 2013).

Além de garantir a estrutura correta das proteínas, o RE contribui para o armazenamento e a regulação do cálcio, para a síntese e armazenamento de lipídios e para o metabolismo da glicose (Ghemrawi *et al.*, 2018). Dessa forma, o RE é sensível a perturbações na homeostase celular que são desencadeadas por diferentes tipos de estresse, podendo ser de origem endógena ou exógena e incluem, por exemplo, variações no nível de cálcio, inibição da glicosilação, estresse oxidativo, insuficiência

de nutrientes, mutações gênicas e exposição a agentes químicos (Walter et al., 2011; Zhang et al., 2012). Essas perturbações podem causar alterações do equilíbrio redox celular, distúrbios na homeostase de cálcio, falhas nas modificações pós-traducionais e um aumento geral na síntese de proteínas, levando ao acúmulo de proteínas malformadas no RE (Ron & Walter, 2007). Proteínas dobradas erroneamente são potencialmente prejudiciais para a função celular e, portanto, são rigidamente controladas. Contudo, quando esse acúmulo de proteínas é exacerbado ocorre a ativação do estresse do RE (Ghemrawi et al., 2018). Essa condição é detectada por diferentes sensores localizados na membrana do RE, os quais desencadeiam uma via de transdução de sinal que induzem a expressão de chaperonas e foldases, componentes do sistema de degradação associado ao RE ou ERAD (ER-associated Degradation). Juntos, eles atuam para ajudar a célula a lidar com o estresse, atenuando a síntese proteica, degradando as proteínas malformadas e aumentando a capacidade do RE para dobrar proteínas. A ativação coordenada dos sensores constitui uma resposta específica ao estresse do RE, a qual foi denominada de resposta a proteínas malformadas ou UPR (do inglês Unfolded Protein Response; Tabas et al., 2011; Walter et al., 2011; Kroeger et al., 2018).

Vias de sinalização da UPR

A UPR é uma resposta ao estresse celular originada no RE. Nos mamíferos, ela é predominantemente controlada por três sensores principais: IRE1 (*inositol-requiring enzyme 1*), PERK (*protein kinase related endoplasmic reticulum kinase*) e ATF6 (*activating transcription factor 6*; Figura 1). Os domínios luminais dos três sensores de estresse do RE são normalmente ligados a principal e mais abundante chaperona do RE, BiP (*binding protein*), também conhecida como GRP78 (*glucose-regulated protein 78*)

ou HSPA5 (*heat shock 70 kDa protein 5*), mantendo-se em um estado inativo (Kusaczuk *et al.*, 2013; Chipurupalli *et al.*, 2019). Porém, durante o estresse de RE o acúmulo de proteínas malformadas promove a dissociação de BiP dos seus ligantes, para auxiliar no correto dobramento das proteínas e, consequentemente, ocorre a ativação de PERK, IRE1 e ATF6 (Kusaczuk *et al.*, 2013; Chipurupalli *et al.*, 2019; Figura 1).

A IRE1 é uma proteína bifuncional com ação tanto de cinase como de endonuclease. Uma vez ativada, a IRE1 sofre um processo de oligomerização e os domínios cinases. presentes porção citoplasmática, na são justapostos e autofosforilados, deflagrando assim a ativação de sua função endonucleásica no lúmen da cisterna reticular (Ron & Walter, 2007; Korennykh et al., 2011). A ativação do domínio endonucleásico inicia o processamento pós-transcricional do mRNA de alguns genes, como o fator de transcrição XBP1 (X-box binding protein 1). O IRE1 ativado induz uma clivagem seletiva em uXBP1 (unspliced XBP1) e um íntron de 26 nucleotídeos é excisado. As extremidades 5' e 3' do mRNA são religadas, produzindo uma isoforma sXBP1 (spliced XBP1) que é competente para se ligar ao promotor de genes essenciais da UPR, incluindo a expressão de chaperonas, foldases e componentes da via ERAD, a fim de aliviar o estresse do RE e restaurar a homeostase (Lee et al., 2003; Figura 1).

PERK também conhecida como eIF2AK3, compartilha com IRE1 várias características na sua estrutura e na sua ativação, no entanto, falta-lhe o domínio com atividade endorribonuclease. Semelhante a sinalização por IRE1, a dissociação de BiP ligada a PERK permite sua oligomerização e sua ativação por autofosforilação (Harding *et al.*, 2000). O dímero de PERK ativado é capaz de reconhecer e fosforilar a eIF2 α (α -subunit of the eukaryotic initiation factor 2) no resíduo de serina 51, inibindo a troca do

nucleotídeo GDP por GTP feita pelo complexo pentamérico eIF2B e, consequentemente, atenuando a tradução de novas proteínas (Lee *et al.*, 2010). O bloqueio da tradução durante o estresse do RE é importante para reduzir a carga de proteínas na maquinaria de dobragem do RE. Durante esse período, eIF2 α regula positivamente a transcrição de ATF4 (*activating transcription factor-4*). O ATF4 ativo irá induzir genes importantes para combater o estresse oxidativo, bem como à indução da expressão do fator de transcrição pró-apoptótico CHOP (*C/EBP-homologous protein*) e de GADD34 (*growth arrest and DNA damage-inducible protein* 34) (Lee *et al.*, 2010; Figura 1).

O ATF6 é uma proteína transmembrânica tipo II no RE com um domínio aminoterminal que contém um motivo transcricional bZIP situado no citoplasma. Diferente da PERK e IRE1, o ATF6 possui na porção carboxi-terminal (localizada no lúmen do ER) sinais de localização do aparato de Golgi que são mascarados por BiP (Yoshida *et al.*, 2001). Em resposta ao estresse do RE, a liberação de BiP facilita a translocação do ATF6 para o Golgi onde é clivado pelas proteases S1P (*site-1 protease*) e S2P (*site-2 protease*) para produzir um domínio amino-terminal ativo de 50 kDa, o ATF6f. Este é translocado para o núcleo onde induz a síntese de chaperonas e enzimas modificadoras. Além disso, convergindo com as cascatas de sinalização IRE1 e PERK, o ATF6 também pode induzir a expressão de XBP1 e CHOP para aprimorar a sinalização UPR (Yoshida *et al.*, 2001; Shen *et al.*, 2004; Figura 1).



Figura 1. Diagrama esquemático das principais vias de sinalização da unfolded protein response (UPR). O acúmulo de proteínas malformadas no retículo endoplasmático (RE) induz UPR, com a ativação de três receptores: IRE1, PERK e ATF6 (Lin *et al.*, 2019).

Morte celular associada à UPR

A sinalização UPR é extremamente importante para a restauração da homeostase do RE e a manutenção da sua função normal. Quando proteínas malformadas se acumulam no lúmen do RE, a sinalização UPR é ativada como uma via adaptativa para mitigar essa fonte de estresse. No entanto, se o acúmulo de proteínas malformadas for persistente, o estresse não pode ser aliviado e a UPR desencadeia mecanismos de morte celular pela ativação das vias CHOP, JNK (*Jun-N-terminal kinas*e) e caspase 12 (Huang *et al.*, 2017; Figura 2).

O CHOP, também conhecido como GADD153 (growth arrest and DNA damage-inducible protein 153), foi originalmente identificado em resposta a danos no

DNA. No entanto, a indução de CHOP é provavelmente mais sensível às condições de estresse do RE (Szegezdi *et al.*, 2006). Durante o estresse do RE, todos os três sensores da UPR induzem a transcrição de CHOP. Entretanto, para regular positivamente a síntese da proteína CHOP, a via PERK-eIF2α-ATF4 da UPR é essencial. A dissociação de BiP do receptor PERK ativa a transcrição ATF4 que regula positivamente a expressão de CHOP (Bravo *et al.*, 2013). Foi descrito que CHOP induz a apoptose principalmente mediante a regulação negativa da expressão do gene anti-apoptótico BCL2 ou por depleção de glutationa, resultando em aumento da produção de espécies reativas de oxigênio (ERO; Szegezdi *et al.*, 2006; Figura 2).

Diferentemente da via CHOP, a via JNK é ativada apenas pelo sensor IRE1 da UPR. O domínio cinase da IRE1 pode se ligar à proteína adaptadora TRAF2 (*TNF receptor associated factor 2*) o qual estimula a cascata das cinases ASK1 (apoptotic signaling kinase-1)/JNK, e a liberação de Ca²⁺ pelo RE regulada por Bax/Bcl-2 (*Bcl-2 associated X protein/B-cell leukemia/lymphoma 2*) (Van der Kallen *et al.*, 2009). A cinase JNK fosforila alguns membros da família de proteínas supressoras de apoptose, Bcl-2, e induzem a apoptose dependente de Bax (Van der Kallen *et al.*, 2009; Figura 2).

As vias que ligam o estresse do RE à ativação da caspase 12 ainda precisam ser melhor investigadas, porém foi proposto que níveis elevados de cálcio citoplasmático ativam uma protease não caspase (calpaína) que processa proteoliticamente a procaspase 12 (Nakagawa e Yuan, 2000). Ainda, foi sugerido que a caspase 12 possa ser clivada e ativada pela caspase 7 (Rao *et al.*, 2001). Após a sua liberação da membrana do RE, a caspase 12 se associa à caspase 9, o que pode levar à ativação da caspase 3 e, subsequentemente, da apoptose (Shen *et al.*, 2004; Figura 2).



Figura 2. Diagrama esquemático das vias de apoptose induzidas pelo estresse do retículo endoplasmático (RE). Em casos de estresses do RE severos e prolongados a UPR desencadeia mecanismos de morte celular pela ativação das vias CHOP, JNK (*Jun-N-terminal kinas*e) e caspase 12 (Lin *et al.*, 2019).

O estresse do retículo endoplasmático na produção *in vitro* de embriões

O oócito (em sua fase de maturação) e os embriões de mamíferos são notáveis por suas fisiologias celulares únicas e mecanismos singulares de regulação do desenvolvimento. Oócitos e embriões em estágios iniciais carecem de muitos dos mecanismos para desempenhar funções metabólicas e homeostáticas básicas, como sequestradores de radicais livres, transportadores de íons e mecanismos osmorreguladores (Latham *et al.*, 2016). Oócitos e embriões também sofrem eventos celulares únicos, não vistos nas células somáticas (Latham *et al.*, 2016). Essas singularidades de oócitos em maturação e de estágios inciais embrionários criam, igualmente, desafios particulares. De fato, esses desafios podem estar subjacentes à relativa sensibilidade dessas células a insultos exógenos. Embora estágios iniciais de embriões de mamíferos tenham, aparente, uma grande plasticidade – o que permite que eles compensem perturbações dramáticas como a redução celular - o oócito em maturação e o embrião são bastante sensíveis a estresses exógenos (Paczkowski *et al.*, 2014; Takahashi *et al.*, 2012; Sakatani *et al.*, 2004; Paula-Lopes e Hansen, 2002).

Durante a maturação oocitária e o desenvolvimento do embrião pré-implantação, o RE atua como um importante local para a biossíntese de proteínas, lipídios e proteínas secretoras e, portanto, desempenha um papel fundamental à crescente demanda do oócito por novas proteínas (Guzel et al., 2017). Essas proteínas devem ser dobradas adequadamente no RE para manter a maturação e o correto desenvolvimento do embrião. Assim, a regulação do estresse/homeostase do RE provavelmente é um mecanismo importante durante esses processos (Latham et al., 2015; Guzel et al., 2017). O desenvolvimento de gametas e de embriões pode encontrar vários tipos de estresse exógeno no sistema de cultivo in vitro (Paula-Lopes e Hansen, 2002; Sakatani et al., 2004; Takahashi et al., 2012; Paczkowski et al., 2014; Latham et al., 2016). É cada vez mais evidente que alguns desses fatores adversos podem impactar negativamente as funções do RE e a síntese proteica, resultando na ativação do estresse do RE e nas vias de sinalização da UPR (Latham et al., 2016). De fato, o estresse do RE e a sinalização UPR parecem desempenhar papéis críticos durante a retomada meiótica oocitária e o desenvolvimento embrionário pré-implantação (Song et al., 2014; Sutton-McDowall et al., 2015; Latham et al., 2016).

Ativação do estresse do retículo endoplasmático na produção in vitro de embriões

Oócitos e embriões em estágios iniciais são muito sensíveis à variação de diversos fatores exógenos, incluindo a temperatura, o estresse osmótico, o estresse de cisalhamento, o estresse oxidativo, a exposição química etc. Esses fatores induzem o estresse do RE (Latham *et al.*, 2016) e reduzem o potencial desenvolvimento embrionário por meio de alterações na expressão gênica, em mecanismos epigenéticos e no metabolismo (Lane *et al.*, 2005).

Durante a maturação dos oócitos e o desenvolvimento inicial do embrião, várias enzimas e vias metabólicas produzem espécies reativas de oxigênio (ERO) endógenas. A fosforilação oxidativa nas mitocôndrias gera ERO. O próprio estresse do RE pode gerar ERO (Landau *et al.*, 2013). Existe uma conexão íntima entre o estresse do RE e o estresse oxidativo (Yoon *et al.*, 2014). O estresse oxidativo impede o correto dobramento e transporte de proteínas e a homeostase do cálcio, podendo desencadear o estresse do RE (Malhotra e Kaufman, 2007). A cultura embrionária com elevada tensão de oxigênio (ar atmosférico), glicose elevada e outros fatores que alteram a fosforilação oxidativa e a função mitocondrial, podem induzir o estresse oxidativo e, consequentemente, o estresse do RE (Chu *et al.*, 2013).

Os meios de cultura que apresentam uma osmolaridade apropriada demonstraram melhorar a maturação dos oócitos e o desenvolvimento embrionário (Lin *et al.*, 2015). No entanto, a hiperosmolaridade dos meios de cultura diminuiu a capacidade de desenvolvimento embrionário, induzindo o estresse do RE e a apoptose (Zhang *et al.*, 2012).

Da mesma forma, o estresse térmico está associado ao estresse do RE, aumentando os níveis de ERO e a apoptose em oócitos e embriões (Paula-Lopes e Hansen, 2002; Tseng *et al.*, 2006). O estresse por baixas temperaturas associado à

criopreservação, também pode afetar o desenvolvimento embrionário e a regulação dos genes (Saenz-de-Juano *et al.*, 2012). De modo semelhante, a exposição prolongada à temperatura ambiente pode gerar ERO, efeitos negativos nas estruturas intracelulares e levar ao estresse do RE (Hegele-Hartung *et al.*, 1991).

A manipulação de oócitos e de embriões mediante a pipetagem produz tensão de cisalhamento, a qual pode danificar o oócito e afetar negativamente o desenvolvimento embrionário. O estresse de cisalhamento transitório pode não influenciar negativamente os embriões, entretanto, o manuseio prolongado ou repetido pode induzir o estresse do RE (Xie *et al.*, 2007).

Reagentes químicos como a tunicamicina (TM) - que age inibindo a Nglicosilação (importante para o dobramento correto da proteína) - são geralmente usados para induzir o estresse do RE na PIVE. A TM foi negativamente relacionada com a maturação de oócitos e no desenvolvimento embrionário, promovendo o incorreto dobramento proteico no RE e ativando as vias de apoptose (Zhang *et al.*, 2011).

Em geral, os estudos descritos acima evidenciam os principais estressores encontrados no ambiente *in vitro* de produção embrionária. Eles ativam o estresse do RE e as vias de sinalização UPR em oócitos e embriões de mamíferos.

O uso de ácido tauroursodesoxicólico para aliviar o estresse do retículo endoplasmático

Os ácidos biliares são produtos naturais e componentes fundamentais da bile (Wang *et al.*, 2014). O ácido ursodesoxicólico (UDCA) e seu conjugado de taurina, o TUDCA, são ácidos biliares hidrofílicos endógenos aprovados pelo FDA (US Food and Drug Administration) para tratamentos clínicos de certas doenças hepáticas colestáticas (Vang *et al.*, 2014). Existem fortes evidências que os efeitos citoprotetores dessa molécula resultam, em parte, por sua capacidade de reduzir o limiar apoptótico em vários tipos de células através da modulação das vias mitocondriais clássicas (Rodrigues *et al.*, 1999; Rodrigues *et al.*, 2000). Além disso, foi observado um efeito positivo do uso do UDCA/TUDCA como um agente para tratar doenças não-hepáticas associadas ao aumento dos níveis de apoptose, como distúrbios neurodegenerativos (McMillin *et al.*, 2016).

O TUDCA foi definido funcionalmente como uma chaperona química, molécula que mimetiza as funções das chaperonas moleculares (Arakawa *et al.*, 2006). Além de induzir um efeito benéfico sobre o estresse mitocondrial, modulando a perturbação da membrana mitocondrial, a formação dos poros, a translocação de BAX, a liberação do citocromo c, a ativação da via caspase e, subsequentemente, a clivagem do substrato (Rodrigues *et al.*, 1999; Rodrigues *et al.*, 2000), recentemente o TUDCA se mostrou eficaz em regular a via mediada pelo estresse do RE (Lee *et al.*, 2010). Entretanto, até o momento, o mecanismo de ação do TUDCA sobre o estresse do RE não foi totalmente esclarecido.

Um estudo, em tipos celulares diferentes, demonstrou os efeitos de TUDCA modulando as vias da UPR e minimizando o estresse do RE. Os principais efeitos de TUDCA foram observados na redução do efluxo de cálcio e da ativação da caspase 12 (Xie *et al.*, 2002). Além disso, o TUDCA inibiu a fosforilação do eIF2a (Lee *et al.*, 2010) e regulou negativamente a via de sinalização IRE1/XBP1 (Luo *et al.*, 2019).

Embora os detalhes mecanicistas da ação do TUDCA sobre a via de estresse do RE não estejam totalmente elucidados, é sabido até o momento que essa chaperona química apresenta um efeito citoprotetor e com grande potencial terapêutico para tratamentos de doenças relacionadas com o estresse do RE, como a diabetes, a obesidade, as doenças cardiovasculares e o câncer.

Efeito da adição de ácido tauroursodesoxicólico na produção in vitro de embriões

Em várias espécies, o uso de inibidores de estresse do RE durante a maturação oocitária e o desenvolvimento inicial de embriões mostrou ser benéfico, diminuindo a apoptose mediada por esse estresse. Por sua função de chaperona, o TUDCA tem sido utilizado para aliviar o estresse do RE durante a maturação oocitária e/ou o desenvolvimento embrionário *in vitro*.

Foi sugerido que o estresse do RE e a sinalização UPR são intrínsecos em oócitos de mamíferos e que sua função adequada pode ser essencial para uma adequada maturação e qualidade dos oócitos (Zhang *et al.*, 2012). Em suínos, o uso de TUDCA na MIV não só aumentou a taxa de oócitos em metáfase II em 44 h, como diminuiu a produção de ERO no mesmo período. Além disso, o TUDCA mostrou um efeito reparador da maturação nuclear em oócitos suínos após as primeiras 22 horas de maturação sujeitas ao efeito de *knockdown* mediado pelo siRNA (*small interfering RNA*) #693 de BiP/GRP78 ou tunicamicina (Park *et al.*, 2017; Park *et al.*, 2018). Após a indução do estresse com tunicamicina, foi observada uma menor abundância de transcritos para ATF4 e CHOP, genes altamente expressos no estresse do RE e na apoptose. Da mesma forma, em camundongos, a adição de TUDCA demonstrou melhorar a viabilidade e o subsequente potencial de desenvolvimento embrionário de oócitos vitrificados, reduzindo o estresse do RE induzido pela criopreservação (Zhao *et al.*, 2015).

Contudo, até o momento, na espécie bovina não há relatos sobre os efeitos da suplementação de TUDCA na maturação *in vitro* oocitária, com ou sem a adição de estressores químicos. Entretanto, a suplementação de TUDCA durante o CIV mostrou ser benéfico para o desenvolvimento do embrião bovino. Além de aliviar o estresse do RE em embriões clonados por transferência nuclear de célula somática, aumentando a

taxa de blastocistos produzidos (Song *et al.*, 2011), também reduziu os níveis de ERO e a abundância de mRNA de marcadores de estresse do RE em sistema de cultivo com 20% de O_2 (Yoon *et al.*, 2014). Ainda em bovinos, um recente estudo relatou uma melhora na crio tolerância dos embriões com a suplementação de TUDCA no cultivo, o que aumentou a taxa de eclosão e reduziu o número de células apoptóticas no embrião após serem submetidos ao estresse de vitrificação (Khatun *et al.*, 2020).

De forma semelhante, em suínos e camundongos, a adição de TUDCA no CIV foi benéfica para o alívio do estresse do RE em embriões. Em suínos, o número total de células aumentou e a taxa de apoptose reduziu em embriões cultivados com TUDCA (Kim *et al.*, 2015). Além disso, foi relatado um aumento da expressão do gene antiapoptótico BCL2 e uma diminuição da expressão dos genes pro-apoptóticos, BCL2L1 (Bcl-xl) e TP53 em embriões tratados com TUDCA (Kim *et al.*, 2015). Embriões suínos clonados por transferência nuclear de célula somática e tratados com TUDCA apresentaram uma redução do estresse oxidativo pela regulação positiva do antioxidante GSH e uma redução na abundância de transcritos para sXBP1 (Lin *et al.*, 2016), ressaltando a ação do TUDCA em mitigar o estresse do RE.

Em camundongos, foi observado uma melhora significativa no desenvolvimento embrionário e na taxa de produção de filhotes após embriões derivados de oócitos livres de células do cumulus serem tratados com TUDCA (Mochizuki *et al.*, 2018).

Coletivamente, esses estudos sugerem que o estresse do RE é um evento comum na MIV de oócitos e/ou CIV de embriões de mamíferos. Além disso, o alívio do estresse do RE pelo tratamento com TUDCA pode melhorar o potencial de desenvolvimento dos oócitos e embriões.

Vitrificação de embriões produzidos *in vitro* e seu efeito no metabolismo embrionário

A criopreservação de embriões é extremamente importante para a conservação de material genético de valor e para aumentar a eficiência da produção de embriões produzidos *in vitro* em sistemas comerciais de aprimoramento genético na criação de gado (A-Na *et al.*, 2014). A vitrificação, uma das técnicas de criopreservação, pode ser alcançada utilizando altas temperaturas de resfriamento e alta concentração de soluções crioprotetoras (Sudano *et al.*, 2016a).

O primeiro relato sobre a criopreservação foi na utilização em embriões de camundongos, por Whittingham *et al.* (1971). Posteriormente, outros estudos surgiram e em diversas espécies (Arav *et al.*, 1987; Vajta *et al.*, 1998; Kuwayama *et al.*, 2005), sendo uma delas especialmente estudada, a bovina. Devido ao impacto econômico da criação de bovinos, a produção de embriões *in vitro* tem crescido a cada ano. Os embriões obtidos pela produção *in vitro* (PIV) possuem baixa resistência à criopreservação (isto é, baixa crio tolerância) a qual, aparentemente, está associada com os lipídios (sua composição na membrana celular, conteúdo e o tamanho das gotículas citoplasmáticas), bem como a composição e a morfologia das organelas presentes nas células embrionárias (Leibo *et al.*, 1995; Abe *et al.*, 2002; Camargo *et al.*, 2011; Paschoal *et al.*, 2017; López-Damián *et al.*, 2012; 2018).

Apesar de os lipídios serem um obstáculo na PIV de embriões, eles possuem extrema importância nos processos metabólicos e auxiliam a coordenar eventos fundamentais durante o desenvolvimento embrionário inicial (Sudano *et al.*, 2016b). Uma das possíveis explicações para esse elevado depósito lipídico está relacionada com a suplementação de soro fetal bovino nos meios de cultivo *in vitro* (Paschoal *et al.*, 2017).

Lipídios como os triglicerídeos (TAG), a principal classe lipídica encontrada no citoplasma das células de mamíferos, são armazenados como gotículas lipídicas (Sudano et al., 2016b) e, nos processos de criopreservação, a sua presença é um fator possivelmente comprometedor devido às injúrias celulares que causa. Em estudo recente foi demostrado que embriões das duas subespécies (Bos taurus e Bos indicus) produzidos in vivo sofreram maior peroxidação lipídica, número de células apoptóticas e ROS, comparados a embriões que não passaram pela criopreservação (López-Damián et al., 2019). Uma das consequências da peroxidação lipídica é aumento do conteúdo lipídico intracitoplasmático, impactando diretamente na sobrevivência após aquecimento Assim, após a criopreservação, os embriões de PIV ficam mais vulneráveis aos danos oxidativos, por motivo de sua estrutura ser delicada e não carecer de mecanismos de defesa desenvolvidos (Nedambale et al., 2006).

Em situações de alta demanda energética, pode ocorrer a hidrólise dos TAG no organismo, gerando ácidos graxos e glicerol em um processo denominado lipólise (Ravnskjaer *et al.*, 2015; Razza *et al.*, 2018). Sendo assim, todos os efeitos gerados pela técnica e manipulação dos oócitos e embriões podem interferir no metabolismo celular.

Dessa forma, a utilização de antioxidantes no meio de cultivo e nos meios pósaquecimento pode viabilizar o potencial de desenvolvimento dos embriões (Nedambale *et al.*, 2006).

A inclusão de inibidores do estresse do RE, como o ácido tauroursodesoxicólico (TUDCA), ácido fenilbutírico (PBA) e salubrinal (Song *et al.*, 2012; Zhang *et al.*, 2012; Sutton-McDowall *et al.*, 2015) nos meios de produção *in vitro* de embriões, foi benéfica e melhoraram a competência e o desenvolvimento embrionário podendo, eventualmente, inclusive impactar na sobrevivência pós-implantação subsequentes ao processo de criopreservação.

HIPÓTESES

1. HIPÓTESE GERAL

A adição de TUDCA em etapas distintas da PIVE alivia o estresse do RE em oócitos e embriões, melhorando a competência oocitária e aumentando a taxa de produção e a criotolerância embrionária.

2. HIPÓTESES ESPECÍFICAS

- 2.1. A suplementação de TUDCA durante a MIV melhora a competência oocitária, afetando positivamente a maturação nuclear, a atividade mitocondrial, a produção de ERO e o perfil de transcritos do oócito, além de aumentar a taxa de oócitos fertilizados e, consequentemente, a produção embrionária (Capítulo 1).
- 2.2. A suplementação de TUDCA durante o CIV melhora o desenvolvimento embrionário, afetando positivamente o perfil de transcritos e a criotolerância dos blastocistos produzidos (Capítulo 2).

OBJETIVOS

1. OBJETIVO GERAL

Investigar os efeitos da suplementação de TUDCA em oócitos e embriões bovinos, durante as etapas distintas da MIV e da CIV.

2. OBJETIVOS ESPECÍFICOS

2.1. Investigar o efeito da adição de TUDCA durante a MIV sobre a competência oocitária (maturação nuclear, atividade mitocondrial, produção de ERO e abundância de transcritos).

2.2. Investigar o efeito da adição de TUDCA durante a MIV sobre a fertilização (taxa de formação de pronúcleo) e o desenvolvimento embrionário (taxa de formação de blastocistos e abundância de transcritos no embrião).

2.3. Investigar o efeito da adição de TUDCA durante o CIV sobre a competência embrionária e a criotolerância do embrião (cinética de eclosão e abundância de transcritos de embriões eclodidos antes e depois da vitrificação).

CAPÍTULO 1

MODULATING THE ENDOPLASMIC RETICULUM STRESS IN *IN VITRO* MATURATION OF BOVINE OOCYTES: COULD TAUROURSODEOXYCHOLIC ACID IMPROVE DEVELOPMENTAL COMPETENCE?

| 1 2 3 4 | Modulating the endoplasmic reticulum stress in <i>in vitro</i> maturation of bovine oocytes: Could tauroursodeoxycholic acid improve developmental competence? |
|------------------|--|
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28 ABSTRACT

Tauroursodeoxycholic acid (TUDCA), which is a bile acid that acts as a potent chemical 29 chaperone to inhibit endoplasmic reticulum (ER) stress in vitro, has been used 30 during IVM of oocyte and/or embryo development. This study aimed to evaluate the 31 effect of three concentrations of TUDCA (50 µM, 200 µM and 1,000 µM) during IVM 32 33 of bovine *cumulus*-oocyte complexes for 24 hours. For this, after maturation, oocytes were analyzed for nuclear maturation, reactive oxygen species (ROS) production, 34 mitochondrial activity, and abundance of target-transcripts in oocytes. Besides, after 35 36 IVM treatment, oocytes were fertilized and cultured in vitro to assess their developmental competence. After 18-20 hours of insemination, oocytes were evaluated 37 38 according to the number of pronuclei, and after 7 and 8/9 days the rates of blastocysts 39 and hatched blastocysts were evaluated, respectively. We also evaluated the transcript abundance of embryonic quality markers. As a main finding, the addition of 200 µM 40 41 TUDCA in IVM seemed to modulate ER stress by decreasing oocyte ROS levels and increasing the abundance of transcripts reported in antioxidant defense in oocytes and 42 embryos. On the other hand, treatment with 1,000 µM TUDCA appeared to be toxic to 43 the oocyte, decreasing oocytes that reached the metaphase II stage, reducing 44 mitochondrial activity, and upregulating the expression of pro-apoptotic genes in 45 oocyte. These results suggest that treatment with 200 µM of TUDCA can relieve 46 oxidative stress during in vitro maturation of bovine oocytes. 47

48

49 INTRODUCTION

50 *In vitro* maturation (IVM) is one of the main restrictive steps for the 51 optimization of *in vitro* production (IVP) since during this period the oocyte acquires

the intrinsic capacity for gradual development until activation of the embryonic genomeafter fertilization [1, 2].

However, studies have indicated that in vitro conditions in which oocytes are 54 55 exposed cause a variety of cellular stresses that contribute to the loss of competence in embryo development [3, 4]. The endoplasmic reticulum (ER) is an important organelle 56 responsible for protein folding, transport and synthesis, trafficking and metabolism of 57 lipids, and cellular Ca²⁺ storage [5, 6]. In vitro however, the ER microenvironment can 58 be disturbed by Ca²⁺ depletion, hypoxia, and N-terminal glycosylation dysfunction, 59 causing ER stress [6]. This stress is triggered when misfolded or unfolded proteins 60 61 accumulate in the lumen of the ER. As a result, this stress can surpass its threshold inducing cellular damage such as apoptosis, degeneration, and carcinogenesis [7, 8]. As 62 a pro-survival response, unfolded protein response (UPR) works to alleviate the 63 64 accumulation of unfolded proteins and restore ER function [9]. The UPR regulation is coordinated by three main sensors: PKR-like ER kinase (PERK), activating 65 transcription factor 6 (ATF6), and inositol-requiring enzyme 1 (IRE1) [10]. However, if 66 misfolded or unfolded protein accumulation is persistent and the stress cannot be 67 relieved, apoptosis is induced by activation of the CHOP, Jun N-terminal kinase (JNK), 68 69 and caspase 12 pathways [11, 12].

An alternative to relieve ER stress is the addition of ER stress inhibitors to the culture media. Tauroursodeoxycholic acid (TUDCA), which is a bile acid that acts as a potent chemical chaperone [13], has been used to alleviate ER stress during *in vitro* oocyte maturation and/or embryo development [14, 15, 16, 17]. The beneficial role of TUDCA is typically attributed to suppression of the UPR [13, 18, 19]. In swine, studies described that ER stress on *in vitro* development of oocytes can be prejudicial [14]. In addition, TUDCA supplementation of maturation media improved oocyte quality,
increasing the oocyte rate that reached metaphase II and decreasing the production ofROS [14, 15].

However, little is known about the mechanism of action of TUDCA and itseffects on *in vitro* matured bovine oocyte.

We hypothesized that *in vitro* maturation of bovine oocytes supplemented with an optimal TUDCA concentration (*i.e.* the maximum concentration without causing toxicity) can alleviate the ER stress caused by the *in vitro* environment. In this way, the proposed model could improve oocyte maturation and, consequently, fertilization and *in vitro* embryonic development.

Therefore, the aim of this study was to evaluate the effect of different TUDCA concentrations during *in vitro* maturation on 1) oocyte (50, 200 or 1,000 μ M), and 2) on fertilization and *in vitro* blastocyst development in cattle (50 or 200 μ M). The effects were evaluated by nuclear maturation, ROS production, mitochondrial activity, pronucleus formation, and expression of ER stress markers (oocyte and embryo).

91

92 MATERIALS AND METHODS

93

94 Experimental Design

Tauroursodeoxycholic acid sodium salt (TUDCA; Selleckchem, Houston, TX, USA) was dissolved in sterile, distilled water to make a 100 mM stock solution (stored at -80° C). This stock solution was diluted into IVM media to make 50 μ M (T50), 200 μ M (T200), and 1,000 μ M (T1000) solutions of TUDCA [14, 15, 20] (Figure 1). The objectives of experiments 1 to 4 were to evaluate the influence of TUDCA during IVM of bovine oocytes (T50, T200 and T1000). The other experiments, 5, 6 and 7, aimed to evaluate the influence of TUDCA supplemented during IVM on fertilization andembryonic development (T50 and T200).

103

104 Experiment 1: The effect of TUDCA during IVM on oocyte nuclear maturation

105 After 24 hours of the IVM with the treatments, denuded oocytes were stained 106 with the Hoechst assay to evaluate nuclear maturation. This experiment was replicated

107 five times using 87 to 110 *cumulus*-oocyte complexes (COCs) per treatment.

108

109 *Experiment 2: The effect of TUDCA during IVM on oocyte ROS production*

110 After 24 hours of the IVM with the treatments, denuded oocytes were stained 111 with the Cell-ROX assay to measure reactive oxygen species. This experiment was 112 replicated five times using 87 to 95 COCs/treatment.

113

114 *Experiment 3: The effect of TUDCA during IVM on oocyte mitochondrial activity*

115 After 24 hours of the IVM with the treatments, denuded oocytes were stained 116 with the Mito-Tracker assay to evaluate mitochondrial activity. This experiment was 117 replicated five times using 87 to 105 COCs/treatment.

118

119 *Experiment 4: The effect of TUDCA during IVM on the abundance of target-transcripts*120 *in oocytes*

After 24 hours of the IVM with the treatments, denuded oocytes were stored at -80° C until samples were processed for gene expression analysis. After RNA extraction and cDNA reverse transcription, the Biomark HD platform was used to relatively quantify the mRNA of interest. This experiment was replicated five times using 100 COCs/treatment. 126

127 Experiment 5: The effect of TUDCA during IVM on pronucleus formation in
128 presumptive zygotes

After 24 hours of the IVM with the treatments, COCs were inseminated for 18-20 hours and presumptive zygotes were separated from *cumulus* cells and stained with the Hoechst assay to evaluate for spermatozoon penetration (number of pronuclei). This experiment was replicated five times using 74 to 84 COCs/treatment.

133

134 Experiment 6: The effect of TUDCA during IVM on developmental competence of135 embryos

After 24 hours of the IVM with the treatments, COCs were inseminated and presumptive zygotes were cultured for 9 days. Blastocyst and hatched blastocyst rates were assessed on day 7 and days 8-9, respectively. This experiment was replicated five times using 218 to 224 COCs/treatment.

140

141 *Experiment 7: The effect of TUDCA during IVM on transcript abundance of embryonic*142 *quality markers*

After 24 hours of the IVM with the treatments, COCs were inseminated and presumptive zygotes were cultured for 9 days. The hatched blastocysts were collected on days 8-9 and stored at -80°C until samples were processed for gene expression analysis. After RNA extraction and cDNA reverse transcription, the Biomark HD platform was used to relatively quantify the mRNA of interest. This experiment was replicated five times using 218 to 224 COCs/treatment.

149

151 In vitro Maturation

152 Bovine ovaries (mainly Bos t. indicus and its crossbreeds) were obtained from a 153 commercial abattoir located at Assis (São Paulo, Brazil). They were transported to the laboratory in sterile saline (0.9% NaCl) at 37°C for 30 minutes at maximum. COCs 154 were collected by aspiration of follicles with 3-8 mm in diameter [21]. After 155 sedimentation, COCs were recovered and selected using a stereomicroscope. Only 156 157 COCs with a homogeneous cytoplasm and a compact multilayer of *cumulus* cells were 158 used (grades 1 and 2) [22]. COCs were washed and transferred to 500 µL drops of maturation medium (10 µL/COCs) in a four-well plate, which consisted of TCM199 159 160 containing Earle salts supplemented with 0.1 IU/mL rhFSH (Gonal-f, Merck Serono, Rockland, MA, USA), 0.22 mg/mL sodium pyruvate, 75 µg/mL amikacin, 4 mg/mL 161 BSA and concentrations of TUDCA according to the experimental design previously 162 163 described. All experiments had a Control group, *i.e.* maturation medium without TUDCA. Drops were incubated at 38.5°C in humidified air with 5% CO₂ for 24 h. 164

165

166 In vitro Fertilization and Culture

In experiments that were performed in vitro fertilization (IVF) and culture 167 (IVC), groups of 25 COCs were transferred to 90 µL drops of Tyrode Albumin Lactate 168 169 Pyruvate (TALP) supplemented with fatty-acid-free BSA (6 mg/mL), pyruvate (0.22 mg/mL), amikacin (75 µg/mL), heparin (30 µg/mL) and PHE (20 µM penicillamine, 10 170 μ M hypotaurine and 1 μ M epinephrine). Oocytes were subjected to insemination step 171 172 with frozen-thawed semen from a single sample of a Nelore breed bull (Ópio). Spermatozoa were selected by the Select SPERM (Botupharma Animal Biotechnology, 173 174 Botucatu, São Paulo, Brazil) method, and the concentration was adjusted to 1×10^6 spermatozoa/mL. Oocytes and spermatozoa were co-incubated under the same 175

conditions as during IVM, and the day of insemination was designated as Day 0. At 18-176 20 hours post-insemination, presumptive zygotes were denuded from *cumulus* cells and 177 transferred to 500 µL drops of SOF medium (synthetic oviduct fluid; 10 µL/zygotes) in 178 179 a four-well plate, supplemented with pyruvate (0.22 µg/mL), amikacin (75 µg/mL), 2.5% v/v of fetal calf serum and BSA (5 mg/mL). Subsequently, they were cultivated in 180 physiological oxygen tension (5%) in small sealed plastic bags with a gas mixture of 181 182 5% O₂, 5% CO₂ and 90% N₂ [23], and high humidity in an incubator at 38.5° C. To all 183 experiments, the culture was maintained for 9 days after insemination to reach the hatching stage of the embryos. Blastocyst and hatched blastocyst rates were assessed as 184 185 the percentage by Day 7 and Days 8/9, respectively, of observed structures and based on the number of COCs used on IVM. 186

- 187
- 188 Nuclear staining: Hoechst 33342

189 The nuclear staining was used in two studies. COCs (Experiment 1) and 190 presumptive zygotes (Experiment 5) were collected at 24 h of IVM and 20-18 h of IVF, 191 respectively, and vortexed in wash medium for 2 min. Then, denuded oocytes and presumptive zygotes were fixed in 4% (v/v) paraformaldehyde for 30 min in humidifier 192 chamber at room temperature (RT), incubated with 5 µg/µL Hoechst 33342 (Sigma-193 194 Aldrich, St. Louis, MO, USA) for 30 min at RT, and transferred to poly-L-lysine-coated 195 slides mounted with a coverslip. The oocytes and presumptive zygotes were analyzed by epifluorescence inverted microscope (Eclipse Ti-E, Nikon, Japan) with A4 filter 196 197 (emission 420 nm and excitation 330-385 nm). Nuclear maturation was graded into 2 categories: immature oocytes (germinal vesicle stage and metaphase I without first polar 198 body), and mature oocytes (metaphase II with primary polar body) [24]. Pronucleus 199 200 formation was graded into 3 categories: (1) unfertilized oocytes: there was a single

pronucleus in the ooplasm, (2) fertilized oocytes: there were two pronuclei in the
ooplasm, and (3) polyspermic oocytes: there were more than two pronuclei in the
ooplasm [25].

204

205 Detection of reactive oxygen species and mitochondrial activity: Cell Rox Green/Mito

206 Track Red CMX ROS

207 To Experiments 2 and 3, COCs were collected at 24 h of IVM and vortexed in 208 wash medium for 2 min. Separately, denuded oocytes were incubated in 5 µM Cell Rox Green (Life Technologies, Foster City, CA, USA), or 0.5 µM Mito Track Red CMX 209 210 ROS (Invitrogen, Ltd), for 30 min in humidifier chamber at 38.5 °C at RT, according to the purpose of the study. Then, the oocytes were fixed in 4% (v/v) paraformaldehyde for 211 212 15 min in humidifier chamber at RT and transferred to poly-L-lysine-coated slides 213 mounted with a coverslip. Among the staining, fixation and slides mounted processes, 214 oocytes were washed 3 times in PBS containing 1 mg/mL PVP. Oocytes were analyzed 215 by epifluorescence microscopy (Eclipse Ti-E, Nikon) equipped with L5 filter (emission 216 519 nm and excitation 495 nm), for ROS detection, and with N21 filter (emission 615 nm and excitation 587 nm), for mitochondria evaluation. In both experiments, a digital 217 218 camera attached to the microscope was used to acquire images, and the fluorescent pixel intensity values of the total area of each oocyte were measured using a freehand tool to 219 220 delimitate the cytoplasm of each oocyte (ImageJ software, https://imagej.nih.gov/ij/index.html). Background fluorescence was subtracted from 221 222 each image before fluorescence measurement and quantification [26, 27]

223

224 Target-Transcripts Relative Quantitation: RT-qPCR

225 RNA isolation and reverse transcription

Total RNA from oocytes and blastocysts were extracted with the PicoPure RNA Isolation kit (Life Technologies, Foster City, CA, USA) following the manufacturer's protocol. Extracted RNA was stored at -80°C until further analysis by qPCR. RNA concentration was quantified by a spectrophotometer (Nanodrop, ThermoFischer Scientific, MA, USA).

We used for each sample a pool of 20 oocytes and a pool of 3 blastocysts to reverse transcription. The cDNA synthesis was performed using High Capacity Reverse Transcription kit (Applied Biosystems, Foster City, CA, USA), following manufacturer's instructions. DNAse treatment was performed in all samples before reverse transcription as manufactures' instructions.

236

237 *Pre-amplification and quantitative polymerase chain reaction*

238 Gene expression analysis of bovine oocyte and blastocysts was performed independently, using Applied BiosystemsTM TaqMan[®] Assays, specific for Bos t. 239 240 taurus. A total of 86 target genes was analyzed (Table 1). Prior to qPCR thermal 241 cycling, each sample was submitted to sequence-specific preamplification process as follows: 1.25 µL assay mix (TaqMan[®] Assay was pooled to a final concentration of 242 0.2X for each of the 96 assays), 2.5 µL TaqMan PreAmp Master Mix (Applied 243 244 Biosystems, #4391128) and 1.25 µL cDNA (5 ng/µL). The reactions were activated at 245 95°C for 10 min followed by denaturing at 95°C for 15 s, annealing and amplification at 60°C for 4 min for 14 cycles. These preamplified products were diluted 5-fold (oocyte 246 247 and embryos) prior to RT-qPCR analysis.

Assays and preamplified samples were transferred to an integrated fluidic circuits (IFC) plate. For gene expression analysis, the sample solution preparation consisted of 2.25 µL cDNA (preamplified products), 2.5 µL of TaqMan Universal PCR Master Mix (2X, Applied Biosystems) and 0.25 μ L of 20X GE Sample Loading Reagent (Fluidigm, South San Francisco, CA, USA); and the assay solution: 2.5 μ L of 20X TaqMan Gene Expression Assay (Applied Biosystems) and 2.5 μ L of 2X Assay Loading Reagent (Fluidigm). The 96.96 Dynamic ArrayTM Integrated Fluidic Circuits (Fluidigm) chip was used to data collection. After priming, the chip was loaded with 5 μ L of each assay solution and 5 μ L of each sample solution and loaded into an automated controller that prepares the nanoliter scale reactions.

The qPCR thermal cycling was performed in the Biomark HD System (Fluidigm) using the protocol TaqMan GE 96x96 Standard, that consisted of one stage of Thermal Mix (50°C for 2 min, 70°C for 20 min and 25°C for 10 min) followed by a Hot Start stage (50°C for 2 min and 95°C for 10 min), followed by 40 cycles of denaturation (95°C for 15 s), primer annealing and extension (60°C for 60 s).

263

264 Statistical analysis

265 The fluorescence intensity data from ROS detection and mitochondrial activity 266 experiments were compared using the non-parametric Kruskal-Wallis test and a post hoc test of Dunn. Data of nuclear maturation rate along with sperm penetration rate as 267 268 well as the rate of blastocysts and hatched blastocysts were arcsines transformed and 269 subjected to analysis of variance (ANOVA), and the means were compared using the post hoc test of Tukey. The normality was assessed with the tests of Shapiro-Wilk and 270 Bartlett. The results are presented as the mean \pm standard error of the mean (SEM) or 271 median and 1st and 3rd interquartile interval according data normality. For quantitative 272 PCR data, it was calculated the ΔC_q values relative to the geometric mean of the best 273 274 reference genes - *i.e.* HPRT1, PPIA and HPRT1 (experiment 4) and GAPDH, HPRT1 275 and PPIA (experiment 7) - among the 96-gene set. Fold-changes were calculated as 2⁻

- Δ^{Cq} . All analyses were performed using JMP software (SAS Institute, Cary, NC, USA).
- 277 Moderate statistical significance was determined based on a 0.01 < P-value ≤ 0.06 while
- a strong significance was considered when P-value ≤ 0.01 .
- 279

280 **RESULTS**

281

282 The effect of TUDCA during IVM on oocyte nuclear maturation

Considering the number of matured oocytes after 24 h of IVM, *i.e.* that reached metaphase II (MII, the analysis of meiosis progression showed a decreasing in the T1000 group (P=0.0002) in comparison with Control, T50, and T200 groups. Moreover, the percentages of oocytes in MII were similar between Control, T50 and T200 groups (Figure 2).

288

289 The effect of TUDCA during IVM on oocyte ROS production

290 Considering that the fluorescence intensity evaluated in this experiment reflects 291 the presence of ROS in the oocytes, after 24 hours of maturation it was observed a 292 decreasing of ROS production (P=0.001) in the T200 group when compared to Control, 293 T50 and T1000. Furthermore, immature oocytes were not significantly different from 294 mature oocytes (with or without TUDCA treatment; Figure 3).

295

296 The effect of TUDCA during IVM on oocyte mitochondrial activity

297 Considering that the fluorescence intensity evaluated in this experiment reflects 298 the oocyte mitochondrial activity, after 24 hours of maturation it was observed a 299 decreasing of mitochondrial activity (P=0.001) in the T1000 treatment when compared 300 to Control, T50 and T200. In addition, mitochondrial activity was significantly higher in immature oocytes (P=0.001) than matured groups with 200 and 1,000 μM of TUDCA
(Figure 4).

303

304 The effect of TUDCA during IVM on the abundance of target-transcripts in oocytes

Transcript abundance from 6 genes was significantly affected in oocyte after TUDCA treatment (Figure 5). When compared to Control group, upregulation was observed in T50 (2 genes), T200 (3 genes) and T1000 groups (3 genes). Of the differentially expressed genes there were transcripts related to endoplasmic reticulum stress (HSPA5), oxidative stress and response to cellular stress (CAT, GPX1 and HMOX1), and apoptosis (CASP3 and CD40; Figure 5).

311

312 The effect of TUDCA during IVM on pronucleus formation in presumptive zygotes

After 24 hours of maturation, oocytes were inseminated and analyzed after 18-20 hours. The rate of pronuclear formation of the oocytes with one, two or more than two pronuclei, were not statistically different between Control, T50 and T200 groups (Table 2).

317

318 The effect of TUDCA during IVM on developmental competence of embryos

After 24 hours of maturation, oocytes were inseminated and cultured for 9 days. On day 7, the percentage of blastocyst production was evaluated and there was no statistical difference between groups. On days 8 and 9, the percentage of hatched blastocyst was evaluated and there was no statistical difference between groups (Table 323 3).

325 The effect of TUDCA during IVM on transcript abundance of embryonic quality 326 markers

Transcript abundance from 4 genes was significantly affected in blastocyst after TUDCA treatment during IVM (Figure 6). When compared to Control group, upregulation was observed in one gene in T50 group and 4 genes in T200 group. Of the differentially expressed genes there were transcripts related to oxidative stress and response to cellular stress (GPX1 and PRDX3), metabolism (AGPAT9), and pluripotency and cell differentiation (OCT4; Figure 6).

333

334 **DISCUSSION**

335 This study demonstrated the effects of TUDCA supplementation on IVM of 336 bovine oocytes. The major results reported here suggest that supplementation of the MIV medium with 200 µM of TUDCA is associated with the greater acquisition of 337 338 resistance to oxidative stress in the oocyte, decreasing the oocyte ROS production and increasing the abundance of transcripts reported to be related with antioxidant activity 339 both in oocytes and embryos. Furthermore, there was observed that 1,000 µM 340 341 concentration of TUDCA seems to be toxic to the oocyte, decreasing nuclear maturation 342 rate, mitochondrial activity and upregulating the expression of pro-apoptotic genes.

In the present study, there was no increase in nuclear maturation rates (*i.e.*, oocytes that reached the MII stage), when COCs were submitted to TUDCA treatment with 50 and 200 μ M for 24 hours. However, the higher concentration (1,000 μ M) decreased the number of metaphase II oocytes among the groups. Differently, in mice, the concentration of 1,000 μ M of TUDCA in the maturation of *cumulus*-free oocytes was not significantly different from the control [20]. In addition, in swine, 200 μ M TUDCA inducted a restorative effect on nuclear maturation after the first 22 hours of

maturation subjected to the effects mediated by knockdown of siRNA #693 of 350 BiP/GRP78 or tunicamycin [28, 29]. Also, in pigs, the 50 µM TUDCA concentration 351 increased the number of oocytes that reached the MII stage when compared to control, 352 353 and this positive modulation in nuclear maturation was associated with a decrease in oocyte ROS production [14]. Collectively, the data suggest that the action mechanisms 354 of TUDCA in the oocyte during IVM can be highly influenced by the species, the tested 355 concentration and by the *in vitro* conditions that the oocytes are submitted to, *e.g.* the 356 357 use of ER stress inducers in the environment, justifying the different results found between those reports and our results. 358

359 Overproduction and the accumulation of ROS overloading the antioxidant defense mechanism is a form of cellular damage called oxidative stress [30]. There is an 360 intimate connection between ER stress and oxidative stress [31, 32]. Oxidative stress 361 362 can cause a reduction-oxidation (redox) imbalance and trigger ER stress [7, 33], which can increase the excessive production of ROS causing oxidative stress [7, 33]. Thereby, 363 364 in this study, we investigated the TUDCA supplementation effects during IVM on 365 oocyte ROS production. The results showed a significant reduction in ROS levels in the T200 group compared with the other groups, unlike the reported in pigs, which 366 observed a reduction in the production of ROS in the oocyte with only 50 µM of 367 368 TUDCA in the IVM [14]. Besides, in our study, the T200 group did not differ from the immature oocyte group, demonstrating to be potentially an alternative to reduce 369 oxidative stress in oocytes during in IVM when necessary. 370

In addition to increasing the production of mitochondrial ROS, ER stress can contribute to the decoupling of oxidative phosphorylation, reduction of mitochondrial membrane potential, matrix swelling and subsequent release of various apoptotic factors, including cytochrome C and effector caspases that lead to apoptosis cells [34].

These effects on mitochondria demonstrate the importance of balancing the flow of Ca²⁺ 375 376 between these organelles [35, 36], where the imbalance of one of them, such as ER stress, can seriously affect the other. In mice, it was observed that mitochondrial 377 378 dysfunction induced in the oocyte can be improved by the therapeutic direction of ER stress, such as the use of salubrinal (ER stress inhibitor) in IVM [36]. When we assessed 379 mitochondrial activity, as measured by fluorescence intensity, the T1000 group 380 381 decreased mitochondrial activity compared to Control and immature oocytes, while the 382 T200 group reduced mitochondrial activity only when compared to the immature oocyte group, but similar to the control. Reports in the literature indicate that increased 383 384 mitochondrial activity is apparently necessary to allow nuclear maturation and completion of meiosis II [37, 38]. Contrary to what was expected in oocyte maturation, 385 the T1000 group decreased mitochondrial activity and reduced nuclear maturation rate, 386 387 suggesting that it was a harmful concentration for in vitro matured bovine oocytes. We 388 have no knowledge about previous description of this information (i.e. toxic 389 concentration of TUDCA for bovine oocytes under IVM).

390 Complementing the previous findings, we verified the abundance of oocyte transcripts previously related to oocyte competence markers. Treatment of the oocyte 391 with TUDCA affected the mRNA abundance of CASP3, CAT, CD40, GPX1, HMOX1 392 393 and HSPA5. A significant increase in the expression of apoptosis markers (CASP3, 394 CD40), and ER stress marker (HSPA5) was observed in oocytes of the T1000 group when compared to the Control. HSPA5 is an important molecular chaperone responsible 395 396 for activating UPR pathways in response to ER stress [6, 8, 9]. In oocyte maturation, its high expression is associated with low oocyte competence [29, 36]. In addition, the 397 increase in the mRNA abundance of HSPA5 is associated with cell apoptosis in cases of 398 severe ER stress, which can positively regulate the expression of pro-apoptotic genes 399

[11, 13, 39], such as the case of Caspase-3 in our experiment [13]. Like CASP3,
positive regulation of CD40 in the oocyte is associated with low competence and death
of the oocyte [40, 41]. Again, our findings reinforce the detrimental effect of T1000 on
bovine oocyte IVM.

In addition, a significant increase in the mRNA abundance of CAT and GPX1 404 was observed in the T50 and T200 groups, and of HMOX1 in the T200 group when 405 compared to the control. The positive regulation of CAT, GPX1 and HMOX1 are 406 407 related to the antioxidant defense mechanism, important for maintaining cellular homeostasis in cases of high ROS levels [42, 43]. The balance between ROS production 408 409 and antioxidant cell defense mechanisms is essential for IVM, otherwise oxidative 410 stress can result in cell damage and apoptosis [44, 45, 46]. Although the T50 treatment has not shown an effect on reducing oocyte ROS, as seen in T200, our gene expression 411 412 data suggest that both treatments, 50 and 200 µM TUDCA, can be used as an alternative 413 to minimize oxidative stress.

Based on all the negative data retrieved from T1000 group, we choose not to
include the 1,000 μM TUDCA concentration in subsequent evaluations.

As already mentioned, balancing calcium oscillations are of great importance in 416 417 maintaining cellular communication [36, 47]. In mammals, this oscillation is a feature 418 of fertilization and plays a central role in activating development. The calcium required for these oscillations is derived mainly from ER, which accumulates in clumps in the 419 microvillar subcortex during oocyte maturation [47]. ER migration to the cortex plays 420 421 an important role in making the ER competent to generate calcium oscillations during oocyte maturation [48]. However, in stressful situations the ER redistribution is altered 422 and calcium oscillations impaired, making sperm penetration difficult [49]. There was 423 evaluated the oocyte fertilization according the pronuclei number, and no statistical 424

difference was found between the groups. Likewise, when we evaluate embryonic 425 development, the blastocyst and hatched blastocyst rates, respectively on days 7 and 8/9 426 of culture, were not statistically different among the groups. As a similar way, in mice, a 427 428 study reported that the 10, 100 and 1,000 µM concentrations of TUDCA on IVM did not increase fertilization and blastocyst rate from *cumulus* free oocytes when compared 429 to the control group [20]. It is not clear why the apparent decoupling of TUDCA effect 430 on the fertilization and blastocyst rates was observed even when beneficial markers (ER 431 432 and oxidative stress mitigation) were observed. The complex process of release of oscillatory calcium waves by spermatozoa phospholipase during fertilization [47, 48] 433 434 could only partially rely on a non stressed ER. The ability to supply calcium by the ER could be retained in early stages of ER stress and an alternative calcium source could 435 theoretically explain that decoupling. Moreover, the absence of effect (at least 436 437 statistically) on the blastocyst production could be attributed to missing the beneficial 438 TUDCA effect at the end of maturation after 7 to 9 days of embryo culture without 439 TUDCA. A possible way to assess that possibility is to conjugate a TUDCA treatment 440 both on IVM and IVC.

Although we did not find a significant difference in the rate of blastocyst with 441 442 the TUDCA treatment, when we evaluated in the embryo the expression of transcripts 443 previously related to markers of embryonic competence, a beneficial modulation was 444 observed in the mRNA abundance. A significant increase in the mRNA abundance for AGPAT9 was observed in the T50 and T200 groups, and GPX1, PRDX3 and OCT4 in 445 446 the T200 group when compared to the control. In the T200 group, was observed a positive regulation of genes evolved in the antioxidant mechanism and gene involved 447 with cell pluripotency. Like GPX1, the transcription factor PRDX3 is involved in 448 antioxidant protection and cell signalling [50, 51], being related to the potential role in 449

oocyte maturation and embryo development [45, 50, 51]. OCT4, encoded by the 450 POU5F1 gene, is a key component of the pluripotency regulatory network [52]. OCT4 451 is postulated to play a critical role in defining totipotency and inducing pluripotency 452 453 during embryonic development [52, 53], being essential to retainment of pluripotency of the inner cell mass and epiblast. We also found T50 and T200 groups upregulated to the 454 isoform AGPAT9, which has been identified as a key regulator of lipid accumulation in 455 456 adipocytes [54], suggesting that it can influence with the highest lipid droplet content in 457 the embryo. From the reports found in the literature and with our result of evaluating the abundance of transcripts in the embryo, we can infer that the treatment of TUDCA in 458 459 IVM is beneficial for the embryo. However, the T200 group appeared to be more effective than the T50 group. However, those beneficial markers need a definitive 460 461 functional proof as the pregnancy rate after fresh embryo transfer.

462 Although the TUDCA treatment has had no effect, both on the oocyte and the 463 embryo, on the gene expression reported in ER stress, we cannot rule out its possible 464 effect in relieving ER stress. Yoon et al. (2014) demonstrated that embryo development 465 in vitro depended on the orchestration between ROS and ER stress [7]. Similarly, our results demonstrate that the oocyte treatment with 200 µM of TUDCA, a supposed ER 466 stress inhibitor, was able to reduce the production of ROS and positively regulate the 467 468 action of antioxidants. Moreover, the *in vitro* conditions used in this study (*i.e.* low O₂ tension without any ER stress inductor as tunicamicin) could not be robust enough to 469 470 highlight TUDCA effect. A functional proof (*i.e.* pregnancy rate) and/or increasing the 471 sampling (to increase the sensitivity of the statistical test) could answer if TUDCA has effectively the potential to improve the *in vitro* embryo production. 472

473 In summary, the use of 200 μM of TUDCA during IVM of bovine oocytes
474 seems to be beneficial in relieving oxidative in the oocyte, reducing ROS production

and increasing the abundance of transcripts reported in antioxidant defense. In addition, 475 when we evaluated the embryonic gene expression, the T200 group seems to be 476 beneficial in positively modulating genes involved in antioxidant defense and cell 477 478 differentiation. In contrast, supplementation of 1,000 µM of TUDCA in IVM was detrimental to the development of oocytes, reducing nuclear maturation and 479 480 mitochondrial activity, and increasing the expression of pro-apoptotic genes in the 481 oocyte. Our study is the first to report the action of TUDCA during IVM of bovine oocytes, however, further studies are needed to confirm the ideal concentration of 482 TUDCA and its possible effect on pregnancy rate from produced embryo. 483

484

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| Gene Symbol | Function | Assay ID* | |
|--------------|--|---------------|--|
| PPIA | Reference gene | PA5-16914 | |
| GAPDH | Reference gene | Bt03210912_g1 | |
| ACTB | Reference gene | Bt03224617_g1 | |
| RPL30 | Reference gene | Bt03226330_g1 | |
| B2M | Reference gene | Bt03251628_m1 | |
| HMBS | Reference gene | Bt03234763_m1 | |
| HPRT1 | Reference gene | Bt03225311_g1 | |
| RLP15 | Reference gene | Bt03288449_g1 | |
| MAPK1 | Oocyte maturation | Bt03216718_g1 | |
| BMP15 | Oocyte maturation | Bt03286494_u1 | |
| H1FOO | Oocyte maturation | Bt03228652_g1 | |
| HS2 | Oocyte maturation | Bt03212695_g1 | |
| PTX3 | Oocyte maturation | Bt03249011_m1 | |
| VCAN | Oocyte maturation | Bt03217333_m1 | |
| NFE2L2 | Oxidative stress and response to cellular stress | Bt03817661_m1 | |
| KEAP1 | Oxidative stress and response to cellular stress | Bt03228713_m1 | |
| CAT | Oxidative stress and response to cellular stress | Bt03215423_g1 | |
| SOD1 | Oxidative stress and response to cellular stress | Bt03215423_g1 | |
| SOD2 | Oxidative stress and response to cellular stress | Bt03244551_m1 | |
| GPX1 | Oxidative stress and response to cellular stress | Bt03259217_g1 | |
| GPX4 | Oxidative stress and response to cellular stress | Bt03259611_m1 | |
| PRDX1 | Oxidative stress and response to cellular stress | Bt03223684_m1 | |
| PRDX3 | Oxidative stress and response to cellular stress | Bt03214402_m1 | |
| ARO(CYP19A1) | Oxidative stress and response to cellular stress | Bt03213774_m1 | |
| GFPT2 | Oxidative stress and response to cellular stress | Bt03250351_m1 | |
| GLRX2 | Oxidative stress and response to cellular stress | Bt03229700_m1 | |
| FOX3 | Oxidative stress and response to cellular stress | Bt03649334_s1 | |
| TXNRD1 | Oxidative stress and response to cellular stress | Bt03215471_m1 | |
| VNN1 | Oxidative stress and response to cellular stress | Bt03220248_m1 | |
| H1F1A | Oxidative stress and response to cellular stress | Bt03259341_m1 | |
| HMOX1 | Oxidative stress and response to cellular stress | Bt03218624_m1 | |
| EIF2A | Endoplasmic reticulum stress | Bt03274460_m1 | |
| HSPA5 | Endoplasmic reticulum stress | Bt03244880_m1 | |
| HSPD1 | Endoplasmic reticulum stress | Bt04301470_g1 | |
| HSPA1A | Endoplasmic reticulum stress | Bt03292670_g1 | |
| ATF4 | Endoplasmic reticulum stress | Bt03221057_m1 | |
| ATF6 | Endoplasmic reticulum stress | Bt03287802_s1 | |
| XBP1 | Endoplasmic reticulum stress | Bt03227621_g1 | |
| DDIT3 | Endoplasmic reticulum stress | Bt03251320_g1 | |
| DNMT1 | DNA methylation | Bt03224737_m1 | |
| DNMT3A | DNA methylation | Bt01027164_m1 | |
| DNMT3B | DNA methylation | Bt03259810_m1 | |
| HP1 | DNA methylation | Bt03246076_m1 | |
| PAF1 | DNA methylation | Bt03239371_g1 | |

Table 1. Gene symbol, functions and primers assay ID used for microfluidicexpression analyses (Biomark HD System - Fluidigm).

| NANOG | Related to pluripotency and cell differentiation | Bt03220541_m1 | | |
|--------------|--|---------------|--|--|
| POU5F1(OCT4) | Related to pluripotency and cell differentiation | Bt03223846_g1 | | |
| HAND1 | Related to pluripotency and cell differentiation | Bt04318733_g1 | | |
| REST | Related to pluripotency and cell differentiation | Bt03278318_s1 | | |
| IGFBP2 | Related to embryo development and cell proliferation | Bt01040719_m1 | | |
| IGFBP4 | Related to embryo development and cell proliferation | Bt03259500_m1 | | |
| SOX2 | Related to embryo development and cell proliferation | Bt03278318_s1 | | |
| IFITM3 | Related to embryo development and cell proliferation | Bt03292973_g1 | | |
| GSK3A | Related to embryo development and cell proliferation | Bt03273698_g1 | | |
| KRT8 | Related to embryo development and cell proliferation | Bt03225178_g1 | | |
| LUM | Related to embryo development and cell proliferation Bt03211920_m1 | | | |
| MTIF3 | Related to embryo development and cell proliferation | Bt03231844_m1 | | |
| TNF | Related to embryo development and cell proliferation | Bt03259156_m1 | | |
| S100A10 | Related to embryo development and cell proliferation | Bt03215645_m1 | | |
| S100A14 | Related to embryo development and cell proliferation | Bt03230771_g1 | | |
| GATM | Related to embryo development and cell proliferation | Bt03237896_m1 | | |
| BAX | Related to apoptosis | Bt03211777_g1 | | |
| CASP9 | Related to apoptosis | Bt04282453_m1 | | |
| CASP3 | Related to apoptosis | Bt03250954_g1 | | |
| BID | Related to apoptosis | Bt03241255_m1 | | |
| CD40 | Related to apoptosis | Bt03817804_g1 | | |
| 1L-1b | Related to apoptosis | Bt03212740_m1 | | |
| NFKB2 | Related to apoptosis | Bt03272789_g1 | | |
| SREBF1 | Related to metabolism | Bt03276370_m1 | | |
| SREBF2 | Related to metabolism | Bt04283467_m1 | | |
| ACACA | Related to metabolism | Bt03213360_m1 | | |
| ACSL3 | Related to metabolism | Bt04282138_m1 | | |
| ELOVL3 | Related to metabolism | Bt00907566_m1 | | |
| ELOVL5 | Related to metabolism | Bt03235956_m1 | | |
| ELOVL6 | Related to metabolism | Bt00907566_m1 | | |
| FADS2 | Related to metabolism | Bt03256255_g1 | | |
| FASN | Related to metabolism | Bt03210471_g1 | | |
| AQP3 | Related to metabolism | Bt03253663_m1 | | |
| PGK1 | Related to metabolism | Bt03225854_mH | | |
| SLC2A3 | Related to metabolism | Bt03259513_g1 | | |
| SLC2A5 | Related to metabolism | Bt03258299_g1 | | |
| AKR1B1 | Related to metabolism | Bt03218049_g1 | | |
| G6PD | Related to metabolism | Bt03649181_m1 | | |
| AGPAT9 | Related to metabolism | Bt04292093_m1 | | |
| PPARA | Related to metabolism | Bt03220821_m1 | | |
| PLIN2 | Related to metabolism | Bt03212182_m1 | | |
| PLIN3 | Related to metabolism | Bt03230537_m1 | | |

* ThermoFischer Scientific

| Number of replicates: 5 | Control | T50 | T200 | Р |
|--|------------------|------------------|------------------|------|
| Total number of oocytes | 79 | 74 | 84 | |
| Unfertilized oocytes rate (%; mean \pm SEM) ^a | 19.43 ± 1.42 | 20.36 ± 4.10 | 17.07 ± 2.97 | 0.79 |
| Fertilized oocytes rate (%; mean \pm SEM) ^a | 74.68 ± 2.12 | 75.61 ± 3.34 | 77.19 ± 1.04 | 0.76 |
| Polyspermic oocytes rate (%; mean \pm SEM) ^a | 5.89 ± 2.49 | 4.03 ± 2.80 | 5.74 ± 2.44 | 0.86 |

Table 2. Pronucleus formation data of Control, T50 and T200 groups.

^a Total pronuclei/ presumptive zygotes

| Number of replicates: 5 | Control | Т50 | T200 | Р |
|--|------------------|------------------|----------------|------|
| Total number of oocytes | 223 | 224 | 218 | |
| Blastocyst rate (%; mean \pm SEM) ^a | 37.79 ± 0.85 | 43.10 ± 2.00 | 42.04 ± 2.32 | 0.14 |
| Hatched blastocyst rate (%; mean \pm SEM) ^b | 37.49 ± 6.01 | 38.07 ± 6.61 | 35.60 ± 3.62 | 0.96 |
| | | | | |

Table 3. Embryo production data of the Control, T50 and T200 groups.

^aTotal blastocysts/total oocyte - from D7.

^b Total hatched blastocysts/total blastocysts - from D8 and D9.

Figure 1. Illustrative experimental design. $IVM = in \ vitro$ maturation; $IVF = in \ vitro$ fertilization; $IVC = in \ vitro$ culture; $IO = in \ vitro$ mature occyte; $MO = mature \ occyte$; $PZ = presumptive \ zygote$; BL = blastocyst and hatched blastocyst.



Figure 2. Effect of TUDCA concentrations during IVM on the percentage of oocytes that reached the MII stage. Results are least-squares means \pm SEM of five replicates using 87 to 110 COCs/treatment. The bar marked with asterisk differ significantly (P \leq 0.06). Control = 0 µM of TUDCA; T50 = 50 µM of TUDCA; T200 = 200 µM of TUDCA; T1000 = 1000 µM of TUDCA.



Figure 3. Effect of TUDCA concentrations during IVM on fluorescence intensity in the presence of ROS in oocyte. A) Illustrative images of fluorescence intensity representing experimental groups in the presence of Cell Rox Green. Bar = 50 μ m. B) Results are presented as the median and 1st and 3rd interquartile interval of five replicates using 87 to 95 COCs/treatment. Different letters in each box represents significant difference (P≤0.06). IO = Immature oocyte; Control = 0 μ M of TUDCA; T50 = 50 μ M of TUDCA; T200 = 200 μ M of TUDCA; T1000 = 1000 μ M of TUDCA.



Figure 4. Effect of TUDCA concentrations during IVM on fluorescence intensity in the presence of mitochondrial activity in oocyte. A) Illustrative images of fluorescence intensity representing experimental groups in the presence of Mito Track Red CMX ROS. Bar = 50 μ m. B) Results are presented as the median and 1st and 3rd interquartile interval of five replicates using 87 to 105 COCs/treatment. Different letters in each box represents significant difference (P≤0.06). IO = Immature oocyte; Control = 0 μ M of TUDCA; T50 = 50 μ M of TUDCA; T200 = 200 μ M of TUDCA; T1000 = 1000 μ M of TUDCA.



Figure 5. Effect of TUDCA concentrations during IVM on differential gene expression in oocyte. Data represent the foldchange of level of expression relative to Reference gene. Results are least-squares means + SEM. Different letters in each bar represent a significant difference ($P \le 0.06$).



Figure 6. Effect of TUDCA concentrations during IVM on differential gene expression in embryo. Data represent the foldchange of level of expression relative to Reference gene. Results are least-squares means + SEM. Different letters in each bar represent a significant difference ($P \le 0.06$).



CAPÍTULO 2

MOLECULAR AND CELLULAR EFFECTS OF THE INCLUSION OF TAUROURSODEOXYCHOLIC ACID IN *IN VITRO* CULTURE BEFORE TO CRYOPRESERVE BOVINE BLASTOCYSTS 1 <u>Title:</u> Molecular and cellular effects of the inclusion of tauroursodeoxycholic acid in *in*

2 *vitro* culture before to cryopreserve bovine blastocysts

3

4 **<u>Running title:</u>** Effect of tauroursodeoxycholic acid in bovine blastocysts

- 5 **<u>Summary sentence</u>**: Addition of tauroursodeoxicolic acid in *in vitro* culture before the
- 6 cryopreservation of bovine blastocysts has been shown to be beneficial in reducing the
- 7 stress of the endoplasmic reticulum

8 Keywords: ER stress, *in vitro* culture, TUDCA, vitrification.

9 **<u>Grant support:</u>** $FAPESP^1$; $CAPES^2$

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35 ABSTRACT

36

During embryo development, the endoplasmic reticulum (ER) acts as an important site 37 for protein biosynthesis, however, in vitro culture (IVC) can negatively affect ER 38 homeostasis. Thereby, the aim of our study was to evaluate the effects of 39 40 supplementation of tauroursodeoxycholic acid (TUDCA), an ER stress inhibitor, in the IVC of bovine embryos. Two experiments were carried out: Exp. 1: evaluation of 41 42 blastocyst rate, hatching kinetics and gene expression of hatched embryos after being treated with different concentrations of TUDCA (0, 50, 200 or 1,000 μ M) in the IVC; 43 Exp. 2: evaluation of the re-expansion and hatching rate (12-, 24- and 48-hours post-44 warming), and gene expression of hatched embryos previously treated with 200 µM of 45 TUDCA at IVC and submitted to vitrification. There was no increase in the blastocyst 46 and hatched blastocyst rates treated with TUDCA in the IVC. Then 1,000 µM 47 48 concentration appeared to be toxic for embryonic development, as it reduced the rate of hatched embryo and upregulated the expression of reported genes in ER stress, 49 50 oxidative stress, lipid accumulation and cell survival. However, embryos submitted to vitrification after treatment with 200 µM of TUDCA, increased hatching rate post-51 warming, together with a downregulation in the expression of ER stress-related genes 52 53 and the accumulation of lipids. We concluded that the treatment with TUDCA (200 μ M) in conditions of low tension of O₂ in the IVC did not improve the competence of the 54 55 development of the embryo unless it be subjected to vitrification.

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57

58 **INTRODUCTION**

59 Disturbances of the endoplasmatic reticulum (ER) homeostasis cause protein 60 folding or misfolding in the ER lumen, a condition called ER stress, which triggers 61 unfolded protein response (UPR) [1]. A paradox of the UPR pathway is that it leads to a

response with simultaneous activation of cell survival and pro-apoptotic pathways. 62 63 Under ER stress conditions, activation of the UPR reduces unfolded protein load through several pro-survival mechanisms, including the expansion of the ER membrane, 64 65 the selective synthesis of key components of the protein folding and quality control machinery and the attenuation of the influx of proteins into the ER [2]. When ER stress 66 is not mitigated and homeostasis is not restored, the UPR triggers apoptosis. Among the 67 68 UPR signaling pathways, there are three predominant and unique signaling transduction mechanisms: inositol-requiring enzyme 1(IRE1), protein kinase RNA (PKR)-like kinase 69 (PERK) and activating transcription factor 6 (ATF6) [3,4]. The IRE1 endoribonuclease 70 71 is activated through dimerization and transphosphorylation. This leads to the removal of 72 a 26-nucleotide intron from the premature unspliced form of XBP1 (XBP1-u) gene to produce the spliced XBP1 (XBP1-s) form [5]. XBP1-s moves to the nucleus and 73 74 induces UPR-responsive genes. XBP1-s is usually regarded as a reliable marker for the 75 induction of the IRE1 pathway of the UPR, because XBP1 is spliced exclusively under 76 ER stress conditions [6]. However, if ER stress is excessive or prolonged, the UPR fails, 77 and cellular apoptosis is induced by activation of CCAAT-enhancer-binding-protein homologous protein (CHOP), Jun N-terminal kinase, and cleaved caspase 3 [7,8]. 78

79 Early embryonic stages are one of the most critical periods of the mammalian development [9]. These early stages involve various morphological and biochemical 80 changes related to genomic activity and a complex set of physiological processes, many 81 82 of which are still unknown [10]. These processes are controlled by several molecular 83 mechanisms and pathways that have a fundamental role in the coordination of 84 homeostatic and metabolic processes [11,12]. Within in vitro systems, disturbances in the embryo's culture environment after fertilization can have detrimental effects on 85 embryonic gene expression [13] which, in turn, can have serious implications for the 86

normality of the blastocyst's physiology. However, the exact influence of *in vitro*culture conditions during each of these critical events/steps is still unknown.

Recent studies in several species have shown that ER stress in the embryo impairs embryo developmental competence [14,15] and that stress relief can improve embryo quality [15-18]. In the bovine species, supplementation of the *in vitro* culture (IVC) medium with tauroursodeoxycholic acid (TUDCA) - a bile acid that acts as a potential chemical chaperone against ER stress *in vitro* [19] - has been an alternative to relieve ER stress and improve the developmental competence in embryos cloned by somatic cell nuclear transfer [16] and embryos subjected to high O₂ tension [20].

96 However, little is known about the effects of ER stress on cryogenic tolerance of embryos generated in vitro [21]. It is known that the in vitro produced embryo (IVPE) 97 98 has distinct characteristics when compared to its counterpart produced in vivo. These 99 differences between IVPE and in vivo derived embryos involve morphological [22] and 100 molecular aspects that affect embryo quality and development [23] decreasing cryo-101 tolerance and pregnancy rates [24]. Therefore, maintaining cell viability after warming 102 is a prerequisite to achieving high outcomes within cryopreservation protocols. To help bridge this gap between IVPE and in vivo derived embryos to improve post-103 cryopreservation results, the ER stress could be an alternative target for 104 105 pharmacological approach.

We hypothesized that supplementation of TUDCA during IVC decreases the endoplasmic reticulum stress of bovine embryos and improves the cryogenic competence of the embryo related to re-expansion and post-warming hatching rate. Therefore, the present study aimed: 1) to evaluate the developmental competency of bovine embryos after being treated with TUDCA in IVC; 2) to investigate the effects of
111 TUDCA treatment in IVC on subsequent developmental competency post-warming of112 the vitrified blastocysts.

113

114 MATERIALS AND METHODS

115

116 In vitro Production

117 Bovine ovaries (mainly Bos t. indicus and its crossbreeds) were obtained from a 118 commercial abattoir located at Assis (São Paulo, Brazil) and during the months of August and September of 2019. They were transported to the laboratory in sterile saline 119 120 (0.9% NaCl) at 37°C for 30 minutes at maximum. *Cumulus* oocyte complexes (COCs) were collected by aspiration of follicles with 3-8 mm in diameter [25]. After 121 sedimentation, COCs were recovered and selected using a stereomicroscope. Only 122 123 COCs with a homogeneous cytoplasm and a compact multilayer of *cumulus* cells were 124 used (grades 1 and 2) [26]. COCs were washed and transferred to 500 µL drops of 125 maturation medium (10 µL/COCs) in a 4-well dishes (Nunc, Roskilde, Denmark), which consisted of TCM199 containing Earle salts supplemented with 0.1 IU/mL 126 rhFSH (Gonal-f, Merck Serono, Rockland, MA, USA), 0,22 mg/mL sodium pyruvate, 127 75 µg/mL amikacin, 4 mg/mL BSA. Drops were incubated at 38.5°C in humidified air 128 129 with 5% CO_2 for 24 h.

Following *in vitro* maturation (IVM), groups of 25 COCs were transferred to 90 μ L drops of Tyrode Albumin Lactate Pyruvate (TALP) supplemented with fatty-acidfree BSA (6 mg/mL), pyruvate (0,22 mg/mL), amikacin (75 μ g/mL), heparin (30 μ g/mL) and PHE (20 μ M penicillamine, 10 μ M hypotaurine and 1 μ M epinephrine). Oocytes were subjected to *in vitro* fertilization (IVF) step with frozen-thawed semen from a single sample of a Nelore breed bull (named "Ópio"). Spermatozoa were

selected by the Select SPERM (Botupharma Animal Biotechnology, Botucatu, São 136 Paulo, Brazil) method, and the concentration was adjusted to 1 x 10⁶ spermatozoa/mL. 137 Oocytes and spermatozoa were co-incubated under the same conditions as during IVM, 138 and the day of insemination was designated as Day 0. At 18-20 hours post-insemination, 139 presumptive zygotes were denuded from *cumulus* cells and transferred to 500 µL drops 140 of SOF medium (synthetic oviduct fluid; 10 µL/zygotes) in a 4-well dishes, 141 supplemented with pyruvate (0,22 µg/mL), amikacin (75 µg/mL), 2.5% v/v of FCS and 142 143 BSA (5 mg/mL), and concentrations of TUDCA according to the experimental design described below. All experiments had a Control group (i.e., culture medium without the 144 addition of TUDCA). Subsequently, they were cultivated in physiological oxygen 145 tension (5%) in small sealed plastic bags with a gas mixture of 5% O₂, 5% CO₂ and 90% 146 N₂ [27], and high humidity in an incubator at 38.5° C. To all experiments, the culture 147 148 was maintained for 9 days after insemination to reach the hatching stage of the embryos.

149

150 Chemical Treatment

Tauroursodeoxycholic acid sodium salt (TUDCA; Selleckchem, Houston, TX, USA) was dissolved in sterile, distilled water to make a 100 mM stock solution (stored at -80° C). This stock solution was diluted into culture media to make 50 μ M, 200 μ M, and 1,000 μ M solutions of TUDCA at experiment 1 (respectively, groups T50, T200 and T1000) [28, 29, 30]. The experiment 2 (vitrification) was performed only with the T200 group.

157

158 Target-Transcripts Relative Quantitation: RT-qPCR

159 RNA isolation and reverse transcription

Total RNA from hatched blastocysts was extracted with the PicoPure RNA
Isolation kit (Life Technologies, Foster City, CA, USA) following the manufacturer's
protocol. Extracted RNA was stored at -80°C until further analysis by qPCR. RNA
concentration was quantified by a spectrophotometer (Nanodrop, ThermoFischer
Scientific, MA, USA).

We used for each sample a pool of 3 hatched blastocysts to reverse transcription.
The cDNA synthesis was performed using High Capacity Reverse Transcription kit
(Applied Biosystems, Foster City, CA, USA), following manufacturer's instructions.
DNAse treatment was performed in all samples before reverse transcription as
manufactures' instructions.

170

171 *Pre-amplification and quantitative polymerase chain reaction*

172 Gene expression analysis of blastocysts was performed independently, using Applied BiosystemsTM TaqMan[®] Assays, specific for *Bos t. taurus*. A total of 80 173 174 candidate genes was analyzed (Supplementary Material-Table S1). Prior to qPCR 175 thermal cycling, each sample was submitted to sequence-specific preamplification process as follows: 1.25 µL assay mix (TaqMan[®] Assay was pooled to a final 176 concentration of 0.2X for each of the 96 assays), 2.5 µL TaqMan PreAmp Master Mix 177 178 (Applied Biosystems, #4391128) and 1.25 µL cDNA (5 ng/µL). The reactions were 179 activated at 95°C for 10 min followed by denaturing at 95°C for 15 s, annealing and amplification at 60°C for 4 min for 14 cycles. These preamplified products were diluted 180 5-fold prior to RT-qPCR analysis. 181

Assays and preamplified samples were transferred to an integrated fluidic
circuits (IFC) plate. For gene expression analysis, the sample solution prepared
consisted of 2.25 μL cDNA (preamplified products), 2.5 μL of TaqMan Universal PCR

Master Mix (2X, Applied Biosystems) and 0.25 μ L of 20X GE Sample Loading Reagent (Fluidigm, South San Francisco, CA, USA); and the assay solution: 2.5 μ L of 20X TaqMan Gene Expression Assay (Applied Biosystems) and 2.5 μ L of 2X Assay Loading Reagent (Fluidigm). The 96.96 Dynamic ArrayTM Integrated Fluidic Circuits (Fluidigm) chip was used to data collection. After priming, the chip was loaded with 5 μ L of each assay solution and 5 μ L of each sample solution and loaded into an automated controller that prepares the nanoliter reactions.

The qPCR thermal cycling was performed in the Biomark HD System (Fluidigm) using the protocol TaqMan GE 96x96 Standard, that consisted of one stage of Thermal Mix (50°C for 2 min, 70°C for 20 min and 25°C for 10 min) followed by a Hot Start stage (50°C for 2 min and 95°C for 10 min), followed by 40 cycles of denaturation (95°C for 15 s), primer annealing and extension (60°C for 60 s).

197

198 Vitrification of embryos

All media used for vitrification and for warming were from Vitrogen Ltd.
(Cravinhos, São Paulo, Brazil). The Vitrific[®] device were purchased from WTA Ltd.
(Cravinhos, São Paulo, Brazil).

202 Quality grades 1 and 2 [26] expanded blastocysts were collected on days 7 and 8 203 of culture (D7 and D8; n=205). Blastocysts were removed from the culture medium 204 (Control group and T200) and washed three times in holding medium (washing medium at 37°C). After this, a vitrification solution 1 (V1) was used at room temperature, in 205 206 which the embryos were submitted to two sequential baths for 8 minutes each. 207 Subsequently, the embryos were transferred to a second solution (vitrification solution 2; V2). In this solution, they remained by just 40 seconds and 3 to 5 structures were 208 allocated to the Vitrific[®] device with the aid of a glass micropipette (approximate inner 209

diameter of 150 µm) to ensure the loading with the minimum possible of the medium. 210 Immediately after, the structures were deposited on the device, excess solution was 211 removed, leaving only a thin layer of medium on the structures. After then, the Vitrific[®] 212 device was plunged directly into the liquid nitrogen (N_2) in a vertical position. The 213 214 protective cap was placed with the device still submerged in N_2 . The blastocysts remained vitrified for an average of 12 to 24 hours. 215

- 216
- 217

Warming and culture of cryopreserved embryos

For warming, the Vitrific $^{\ensuremath{\mathbb{R}}}$ device was removed from the N_2 with the aid of 218 tweezers. The device's cap was removed and immediately after, its tip containing the 219 220 structures was submerged in the warmed solution 1 (D1) at 37° C for 1 minute. Subsequently, the embryos were washed in two drops of warmed solution 2 (D2) and 221 222 two drops of warmed solution 3 (D3) at room temperature, with an average time of 6 223 minutes for all steps. After then, the embryos were washed in 5 drops of culture 224 medium. After this, the embryos were transferred to 4-well dishes with 500 µL of 225 culture medium (maximum of 50 structures per well) and they were incubated in 38°C and 5% O2 tension in humidified air (IVC conditions). Embryos were evaluated for re-226 expansion and hatching rate at 12-, 24- and 48-hours post-warming. 227

228

Experimental Design 229

230

231 Experiment 1: Effects of TUDCA during IVC on developmental competence and gene expression of embryos 232

To investigate the effect of TUDCA on embryo development and quality, 233 234 blastocyst and hatched blastocyst rates were analyzed, respectively, on day 7 and days 8-9 (Day 0 being the day of insemination). Hatched blastocysts were collected on days
8-9 for gene expression analysis. After RNA extraction and cDNA reverse transcription,
the Biomark HD platform was used to relatively quantify the mRNA abundance of
markers of interest. This experiment was replicated five times using 250 presumptive
zygotes/treatment.

240

241 Experiment 2: Effects of TUDCA during IVC on developmental competence and gene 242 expression of post-warmed vitrified blastocysts

After treatment with TUDCA in the culture, expanded blastocysts were collected 243 244 on days 8-9 and subjected to the technique of vitrification. After 12-24 hours the cryopreserved blastocysts were warmed and cultured at low tension of O₂. Embryo re-245 246 expansion and hatching rates were analyzed at 12-, 24- and 48-hours post-warming. 247 Hatched embryos were collected 24- and 48-hours post-warming for the analysis of 248 gene expression. After RNA extraction and cDNA reverse transcription, the Biomark 249 HD platform was used to relatively quantify the mRNA abundance of markers of 250 interest. This experiment was replicated five times using 88 to 117 vitrified embryos/treatment. 251

252

253 Statistical analysis

To the embryonic development (experiments 1 and 2), the blastocysts and hatched blastocysts rates were arcsines transformed and subjected to analysis of variance (ANOVA), and the means were compared using the *post hoc* Tukey test. The normality was assessed with the Shapiro–Wilk test and Bartlett's test. The results are presented as the mean \pm standard error of the mean (SEM). For quantitative PCR data, we calculated the ΔC_q values relative to the geometric mean of the best reference genes

| 260 | - <i>i.e.</i> B2M, HPRT1 and PPIA (experiment 1) and ACTB, HPRT1 and PPIA (experiment |
|-----|--|
| 261 | 2) - among the 96-gene set. Fold-changes were calculated as $2^{-\Delta Cq}$. All analyses were |
| 262 | performed using JMP software (SAS Institute, Cary, NC, USA). Statistical significance |
| 263 | was determined based on a P-value ≤ 0.1 , the significance being considered weak |
| 264 | (between 0.051 and 0.1), moderate (between 0.010 and 0.050) or strong (less than 0.01). |
| 265 | |

266 <u>RESULTS</u>

267

268 The effects of TUDCA during IVC of bovine embryos

269 Developmental competence

270 After 7 days of culture, in the absence and presence of TUDCA, there was no 271 difference in the rate of blastocysts among the Control group and the treatments with TUDCA. However, group T200 presented a higher rate of blastocysts when compared 272 to group T1000 (P=0.0268; Figure 1). Evaluating the embryo hatching kinetics, there 273 was no difference between treatments on day 7. However, on days 8 and 9, the T1000 274 group (7.3% and 6.95%, respectively) was statistically different from day 8 Control 275 276 (26.4%) and all treatments by day 9 (23.8%, 24.2% and 27.9%; respectively to Control 277 group, T50 and T200; Figure 2). The results suggest a toxic effect in the group treated 278 with 1,000 µM of TUDCA altering the embryonic development kinetics and, 279 subsequently, decreasing its ability to hatch.

280 *Gene expression*

Transcript abundance from 10 genes was significantly affected in hatched blastocyst after TUDCA treatment, with all genes up regulated in the T1000 group in comparison to the other groups (Figure 3). Of the differentially expressed genes there

were transcripts related to endoplasmic reticulum stress (ATF6, EIF2A and HSPD1), 284 oxidative stress and response to cellular stress (GFPT2, HMOX1 and TXNRD1), 285 metabolism (ACACA, ELOVL5 and SREBF2), and embryo development and cell 286 287 proliferation (GSK3A; Figure 3). Corroborating with the analysis of the developmental competence, the T1000 group appeared to be toxic to the embryo, increasing the 288 abundance of transcripts related to ER stress and oxidative stress. In addition, was 289 290 observed an increase in the mRNA abundance for ACACA, ELOVL5 and SREBF2, 291 lipid metabolism markers involved with ER stress (Figure 4).

292

293 The effects of TUDCA during IVC on post-warmed vitrified blastocysts

294 Developmental competence

There was no difference in the re-expansion rate evaluated after 12, 24 and 48 hours of warming up (P>0.1; Table 1). However, a larger number of hatched blastocysts were observed from TUDCA treatment in the assessment post-warming (Table 1). After 24 and 48 hours post warming, there was a higher significantly number of hatched blastocysts in embryos cultivated with TUDCA compared to control (P=0.09 and 0.0423, respectively; Table 1). The results suggest a beneficial effect of treatment with TUDCA by increasing the hatching rates of embryos subjected to vitrification.

302 *Gene expression*

Transcript abundance from 11 genes was affected in hatched blastocyst after vitrification and post-warming. When compared to Control group, in T200 group 5 genes were downregulated, and 6 genes were up-regulated (Figure 5). Downregulation was observed for genes related to endoplasmic reticulum stress (HSPA5 and XBP1), oxidative stress and response to cellular stress (GLRX2), and some markers for

metabolism (PLIN3 and SREBF2; Figure 5). Up-regulation was observed for genes 308 related to oxidative stress and response to cellular stress (CAT, GPX1, NFE2L2 and 309 310 PRDX1) and metabolism (G6PD and SLC2A3; Figure 5). The treatment with TUDCA 311 could have relieved the ER stress and modulated the lipid metabolism of the hatched 312 embryo, but direct evaluations of these pathways were not performed. String analysis of those genes statistically different reveals a correlation between genes related to 313 314 endoplasmic reticulum stress and genes related to oxidative stress and response to 315 cellular stress (Figure 6).

316

317 **DISCUSSION**

Developing embryos may be subjected to several sources of exogenous stress in 318 in vitro culture system [15, 31, 32, 33]. These include oscillating temperature, DNA 319 320 damage or DNA damaging agents, osmotic stress and availability of organic osmolytes, 321 oxygen and oxidative stress, hyperglycemia and carbon substrate availability, 322 hyperlipidemia and oxidized lipids, calcium ionophores, cytokines, amino acid 323 deprivation, insulin signaling, and serum components [15, 31, 32, 33]. Cold stress associated with cryopreservation affects embryo development and gene regulation [22, 324 24, 34]. These adverse factors negatively impact ER functions and protein synthesis and 325 326 folding, resulting in the activation of ER stress and the UPR signaling pathways in in 327 vitro produced embryo [14, 15, 16, 17]. In the present study, we demonstrated that the addition of TUDCA during *in vitro* culture can improve the cryo-tolerance of the bovine 328 329 blastocyst through the putative modulation of ER and oxidative stress. However, and in the culture conditions, there was no observed effect on embryo development with the 330 TUDCA. 331

TUDCA supplementation in IVC was associated with improved embryonic 332 developmental rates in mice [17, 30], pigs [18, 28], and cows [20, 21]. In our 333 experiment, we did not find a significant increase in the rates of blastocyst formation 334 with TUDCA treatments. Similarly, in conditions of low O_2 tension (5%), 335 supplementation of 50 µM of TUDCA in IVC of bovine embryos did not modulate 336 embryo competence [20]. However, embryos submitted to high tension O_2 (20%) in the 337 IVC showed an increased blastocyst rate in cattle and pigs with the supplementation of, 338 respectively, 50 and 200 µM of TUDCA [18, 20]. In this way, the potential beneficial 339 effect of TUDCA supplementation seems to be linked with the culture conditions. When 340 341 stringent condition of embryo culture is used (e.g., high oxygen tension as source to generate an increasing of the reactive oxygen species) the effect of TUDCA to alleviate 342 343 ER stress was observed [18, 20] that was not the case of our study. Corroborating with 344 our result, recent studies have shown that the increase in reactive oxygen species (ROS), 345 due to O₂ tension, is closely related to the increase in ER stress in embryos [20]. In 346 addition, unlike the purpose of our study, TUDCA proved to be beneficial for the 347 development of in vitro produced embryos in conditions where ER stress was chemically or physically actively induced (e.g., using tunicamycin or heat stress, 348 respectively) [18, 20, 21, 35, 36, 37]. 349

Complementing these results, when we evaluated the hatching kinetics of embryos treated with TUDCA, it was observed a significant reduction in the hatched embryos rate with the T1000 group when compared to the other groups (with or without the addition of TUDCA). Contrary to the reported in mice - where a positive effect on embryonic development and newborn rate was described with the addition of 1,000 μ M of TUDCA in the culture [30] - the higher concentration of TUDCA used in our study proved to be toxic to the bovine embryo and impaired its development. In addition to the factor already mentioned (O_2 tension linked to the TUDCA effect) it seems that the species (mouse or cattle) also plays a role on the upper threshold of beneficial effect of TUDCA (*i.e.*, when the threshold is exceeded and the toxic effect is observed).

In the analysis of the mRNA abundance involved in ER stress, oxidative stress,
metabolism and embryonic quality, the negative effect on hatching rate of the T1000
group was reinforced.

363 In case of misfolded proteins in the ER lumen, molecular chaperones (HSPD1 364 and HSPA5) are activated with the function of correcting this misfolding and maintaining homeostasis in the ER [38; 39]. For instance, in cases of ER stress, HSPA5 365 366 dissociates from PERK, ATF6, and IRE1 receptors, activating the UPR pathway [3, 4, 38]. Activated PERK can recognize and phosphorylate eIF2α, which in turn regulates 367 positively the translation of ATF4, an important inducer of CHOP, GADD34, ATF-3, 368 369 and genes involved in apoptosis [40]. In the T1000 group, there was an increase on the 370 transcript abundance for HSPD1, ATF6, and EIF2A in hatched embryos. This suggests 371 that the higher concentration of TUDCA of this study paradoxically induced ER stress 372 in the blastocysts. Also, an increase in the gene's expression involved with oxidative stress (e.g., GFPT2, HMOX1, and TXNRD1) was observed and reinforces the close 373 374 relationship between ER stress and oxidative stress, where ROS functions as a mediator 375 of these two events [41; 42]. Oxidative stress in embryos can lead to DNA damage [31] 376 and to inhibit preimplantation development [43]. For transcripts involved in lipid 377 metabolism (SREBF2, ELOVL5 and ACACA), there were an up regulation in embryos 378 of the T1000 group. Crosstalk between ER stress and lipid metabolism is well established [44, 45, 46, 47]. Several reports indicate that the pathways that regulate 379 UPR also induce the lipid accumulation in the cell. For instance, the ATF6α pathway 380 plays a role in lipid accumulation interacting with the nuclear form of SREBP-2 [46, 381

382 47]. In the literature, the lipid accumulation in embryos is associated with lower rates of 383 embryonic survival after cryopreservation and deviations in the relative abundance of transcripts of important genes for embryonic development [23, 48]. Additionally, and 384 385 corroborating the results, the T1000 group has been shown to increase the transcript abundance for GSK3A (related to embryo development and cell proliferation). GSK3A 386 387 is a negative regulator in the hormonal control of glucose homeostasis, cell division, 388 proliferation, motility, and survival. In the literature, the high expression of GSK3A in 389 the embryo is associated with low embryo competence. [49].

Unlike the T1000 group, the T50 and T200 groups did not significantly affect gene expression in hatched blastocysts when compared to Control, corroborating the results of embryonic development. Although without molecular and cellular evidence of any beneficial effect of TUDCA (T50 and T200 groups), there was no assessment on the pregnancy rate of those embryos (fresh transfer) or the effect of high oxygen culture system.

396 There is evidence that IVPEs are more sensitive to cryopreservation than in vivo 397 derived embryos [48, 50] and that this reduced cryotolerance may be associated with the high lipid content present in the cytoplasm as well as the lipid profile of the cell 398 membrane of these embryos [23, 24, 48]. Although TUDCA has been demonstrated to 399 400 exert efficient cytoprotective activity in relieving ER stress [17, 18, 20, 21], recent studies reported on its new potential and molecular modes of action as a weight-401 reducing agent, modulating lipid metabolism through or independently of ER 402 403 modulation [51, 52].

404 In the present study, we demonstrated that 200 μ M of TUDCA during IVC 405 enhanced the cryotolerance of bovine embryo. At 24- and 48-hours post-warming,

406 embryos treated with TUDCA during IVC had an increase in hatching rate when407 compared to Control group.

408 Unlike our results, in cattle was reported that 10 μ M TUDCA was able to 409 improve cryotolerance of embryos after vitrification, increasing hatching rates and 410 decreasing the number of apoptotic cells in the embryo 48-hours post-warming [21]. 411 Once again, the action of TUDCA is highly dependent on the complex combination of 412 variables such the species, breed, the used concentration and the *in vitro* culture 413 conditions.

When we evaluated the 96 markers of transcript abundance in the hatched 414 415 embryos post-warming, TUDCA treatment induced the decreasing of the mRNA 416 abundance related to ER stress and lipid metabolism pathways. An increase in the mRNA abundance related to antioxidant activity was also observed in embryos in the 417 418 T200 group. After warming post-vitrification, hatched blastocysts treated with TUDCA 419 showed less mRNA abundance for HSPA5 and XBP1. In several species, the increasing 420 of XBP1-s expression is widely used as a molecular marker of ER stress in vivo and in 421 vitro [53, 54]. Increasing mRNA abundance for XBP1-s and HSPA5 was associated with the low competence of embryonic development in several species [18, 20, 50]. In 422 423 addition, after warming vitrified embryos treated with TUDCA showed a change in the 424 expression of antioxidants, with high mRNA abundance for CAT, GPX1, NFE2L2 and 425 PRDX1, and less abundance for GLRX2. NFE2L2 is a promising target against oxidative stress, responsible for inducing the expression of several endogenous 426 427 cytoprotective enzymes [55, 56]. In vitro study with the human neuroblastoma SH-SY5Y cell line, observed that TUDCA prevented oxidative stress through the high 428 expression of NRF2, DJ-1, and antioxidant enzymes heme oxygenase-1 (HO-1) and 429 glutathione peroxidase (GPx) [57], corroborating with our results. Previous study has 430

431 shown that embryos produced in vivo and cryopreserved undergo greater oxidative 432 stress when compared to embryos that have not been subjected to cryopreservation [34]. 433 In the conditions of this study, the treatment with TUDCA could has prevented high 434 levels of oxidative stress in vitrified embryos cultured after warming. Collectively, 435 those data could partially explain the higher cell competence to hatch observed with 436 TUDCA treatment, since a mitigated stress (ER and oxidative) improves the cellular 437 activity.

The hatched embryos also modulated markers related to metabolism. While the
mRNA abundance for PLIN3 and SREBF2 decreased, the abundance for G6PD and
SLC2A3 were increased with TUDCA treatment.

Reduced post-warming cell viability is associated with the abnormal amount 441 and/or the type of lipids in the blastomeres that contribute to the occurrence of 442 cryogenic fractures during the freezing process [23, 48]. The cell membrane fluidity is 443 444 related to the lipid profile and the capacity to support cryo-injuries during 445 cryopreservation process [48]. IVPEs possessing different lipid profile than their in vivo 446 derived counterparts [48, 50] could have cryotolerance enhanced with changes in the lipid metabolism (e.g., TUDCA treatment). Although the evaluation of the lipid content 447 was not assessed in this work, we cannot rule out that TUDCA treatment may have 448 449 modulated the lipid content of the hatched embryo (downregulated mRNA abundance 450 of PLIN3 and SREBF2 in T200 group).

In addition, G6PD (Glucose-6-phosphate dehydrogenase) and SLC2A3 (Solute carrier family 2, facilitated glucose transporter member 3) mRNAs were up-regulated in the embryo treated with TUDCA. It was reported that those genes were essential for preimplantation embryonic development [58, 59, 60]. Partially, this finding add data to explain how the TUDCA treatment could improve the post-warming hatching rate of

vitrified embryos. Since the mammalian hatching process involves a coordinated
trophectoderm activity and is concomitant with the beginning of hypoblast appearance
in bovine species [61], the up regulation of G6PD and SLC2A3 genes suggests a
positive marker after TUDCA treatment.

In conclusion, this work showed that the addition of TUDCA during *in vitro* culture can improve the cryotolerance of the bovine blastocyst through the putative modulation of ER and oxidative stress. However, and in the culture conditions of this study, there was no observed effect on embryo development with the treatment of 50, 200 and 1,000 μ M of TUDCA. In addition, the highest concentration (1,000 μ M), proved to be detrimental to the development and kinetics of the embryo.

466

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Supplementary Material

| Gene Symbol | Function | Assay ID* | | |
|--------------|---|---------------|--|--|
| PPIA | Reference gene | PA5-16914 | | |
| GAPDH | Reference gene | Bt03210912_g1 | | |
| ACTB | Reference gene | Bt03224617_g1 | | |
| RPL30 | Reference gene | Bt03226330_g1 | | |
| B2M | Reference gene | Bt03251628_m1 | | |
| HMBS | Reference gene | Bt03234763_m1 | | |
| HPRT1 | Reference gene | Bt03225311_g1 | | |
| RLP15 | Reference gene | Bt03288449_g1 | | |
| NFE2L2 | Oxidative stress and response to cellular stress | Bt03817661_m1 | | |
| KEAP1 | Oxidative stress and response to cellular stress | Bt03228713_m1 | | |
| CAT | Oxidative stress and response to cellular stress | Bt03215423_g1 | | |
| SOD1 | Oxidative stress and response to cellular stress | Bt03215423_g1 | | |
| SOD2 | Oxidative stress and response to cellular stress | Bt03244551_m1 | | |
| GPX1 | Oxidative stress and response to cellular stress | Bt03259217_g1 | | |
| GPX4 | Oxidative stress and response to cellular stress | Bt03259611_m1 | | |
| PRDX1 | Oxidative stress and response to cellular stress | Bt03223684_m1 | | |
| PRDX3 | Oxidative stress and response to cellular stress | Bt03214402_m1 | | |
| ARO(CYP19A1) | Oxidative stress and response to cellular stress | Bt03213774_m1 | | |
| GFPT2 | Oxidative stress and response to cellular stress | Bt03250351_m1 | | |
| GLRX2 | Oxidative stress and response to cellular stress | Bt03229700_m1 | | |
| FOX3 | Oxidative stress and response to cellular stress | Bt03649334_s1 | | |
| TXNRD1 | Oxidative stress and response to cellular stress Bt02 | | | |
| VNN1 | Oxidative stress and response to cellular stress | Bt03220248_m1 | | |
| H1F1A | Oxidative stress and response to cellular stress | Bt03259341_m1 | | |
| HMOX1 | Oxidative stress and response to cellular stress | Bt03218624_m1 | | |
| EIF2A | Endoplasmic reticulum stress | Bt03274460_m1 | | |
| HSPA5 | Endoplasmic reticulum stress | Bt03244880_m1 | | |
| HSPD1 | Endoplasmic reticulum stress | Bt04301470_g1 | | |
| HSPA1A | Endoplasmic reticulum stress | Bt03292670_g1 | | |
| ATF4 | Endoplasmic reticulum stress | Bt03221057_m1 | | |
| ATF6 | Endoplasmic reticulum stress | Bt03287802_s1 | | |
| XBP1 | Endoplasmic reticulum stress | Bt03227621_g1 | | |
| DDIT3 | Endoplasmic reticulum stress | Bt03251320_g1 | | |
| DNMT1 | DNA methylation | Bt03224737_m1 | | |
| DNMT3A | DNA methylation | Bt01027164_m1 | | |
| DNMT3B | DNA methylation | Bt03259810_m1 | | |
| HP1 | DNA methylation | Bt03246076_m1 | | |
| PAF1 | DNA methylation | Bt03239371_g1 | | |
| NANOG | Related to pluripotency and cell differentiation | Bt03220541_m1 | | |
| POU5F1(OCT4) | Related to pluripotency and cell differentiation | Bt03223846_g1 | | |
| HAND1 | Related to pluripotency and cell differentiation | Bt04318733_g1 | | |
| REST | Related to pluripotency and cell differentiation | Bt03278318_s1 | | |

Table S1: Gene symbol, functions and primers assay ID used for microfluidic expression analyses (Biomark HD System - Fluidigm).

| IGFBP2 | Related to embryo development and cell proliferation | Bt01040719_m1 |
|---------|--|---------------|
| IGFBP4 | Related to embryo development and cell proliferation | Bt03259500_m1 |
| SOX2 | Related to embryo development and cell proliferation | Bt03278318_s1 |
| IFITM3 | Related to embryo development and cell proliferation | Bt03292973_g1 |
| GSK3A | Related to embryo development and cell proliferation | Bt03273698_g1 |
| KRT8 | Related to embryo development and cell proliferation | Bt03225178_g1 |
| LUM | Related to embryo development and cell proliferation | Bt03211920_m1 |
| MTIF3 | Related to embryo development and cell proliferation | Bt03231844_m1 |
| TNF | Related to embryo development and cell proliferation | Bt03259156_m1 |
| S100A10 | Related to embryo development and cell proliferation | Bt03215645_m1 |
| S100A14 | Related to embryo development and cell proliferation | Bt03230771_g1 |
| GATM | Related to embryo development and cell proliferation | Bt03237896_m1 |
| BAX | Related to apoptosis | Bt03211777_g1 |
| CASP9 | Related to apoptosis | Bt04282453_m1 |
| CASP3 | Related to apoptosis | Bt03250954_g1 |
| BID | Related to apoptosis | Bt03241255_m1 |
| CD40 | Related to apoptosis | Bt03817804_g1 |
| 1L-1b | Related to apoptosis | Bt03212740_m1 |
| NFKB2 | Related to apoptosis | Bt03272789_g1 |
| SREBF1 | Related to metabolism | Bt03276370_m1 |
| SREBF2 | Related to metabolism | Bt04283467_m1 |
| ACACA | Related to metabolism | Bt03213360_m1 |
| ACSL3 | Related to metabolism | Bt04282138_m1 |
| ELOVL3 | Related to metabolism | Bt00907566_m1 |
| ELOVL5 | Related to metabolism | Bt03235956_m1 |
| ELOVL6 | Related to metabolism | Bt00907566_m1 |
| FADS2 | Related to metabolism | Bt03256255_g1 |
| FASN | Related to metabolism | Bt03210471_g1 |
| AQP3 | Related to metabolism | Bt03253663_m1 |
| PGK1 | Related to metabolism | Bt03225854_mH |
| SLC2A3 | Related to metabolism | Bt03259513_g1 |
| SLC2A5 | Related to metabolism | Bt03258299_g1 |
| AKR1B1 | Related to metabolism | Bt03218049_g1 |
| G6PD | Related to metabolism | Bt03649181_m1 |
| AGPAT9 | Related to metabolism | Bt04292093_m1 |
| PPARA | Related to metabolism | Bt03220821_m1 |
| PLIN2 | Related to metabolism | Bt03212182_m1 |
| PLIN3 | Related to metabolism | Bt03230537_m1 |

* ThermoFischer Scientific

Figure 1: Effect of TUDCA concentrations on the percentage blastocyst. Results are least-squares means \pm SEM. Different letters in each bar represents significant difference (P \leq 0.1). Control = 0 µM of TUDCA; T50 = 50 µM of TUDCA; T200 = 200 µM of TUDCA; T1000 = 1000 µM of TUDCA.



Figure 2: Effect of TUDCA concentrations on the blastocyst hatching kinetics. Results are least-squares means \pm SEM. *P \leq 0.1. Control = 0 µM of TUDCA; T50 = 50 µM of TUDCA; T200 = 200 µM of TUDCA; T1000 = 1000 µM of TUDCA.



Figure 3: Effect of TUDCA concentrations on mRNA abundance of differentially expressed genes in embryo according to the following functional categories: endoplasmic reticulum stress (A), oxidative stress and response to cellular stress (B), related to metabolism (C), and related to embryo development and cell proliferation (D). Results are least-squares means \pm SEM. Different letters in each bar represent a significant difference (P \leq 0.1). Control = 0 μ M of TUDCA; T50 = 50 μ M of TUDCA; T200 = 200 μ M of TUDCA; T1000 = 1000 μ M of TUDCA.



Figure 4: STRING network of protein-protein interactions between genes with significant difference ($P \le 0.1$) of mRNA abundance among groups. Different line colors representes the relantionship between the proteins translated from analysed genes, as described on the legend.



Table 1: Re-expansion and hatching rate of embryos after vitrification on Days 7 and 8 (only expanded blastocysts) from groups Control or T200 (200 μ M of TUDCA). Total of five replicates (mean ± standard error mean).

| Treatment | Vitrified Embryos (D7/D8) | Re-exp. rate 12 h (%) | Re-exp. rate 24 h (%) | Re-exp. rate 48 h (%) | Hatch. rate 12 h (%) | Hatch. rate 24 h (%) | Hatch. rate 48 h (%) |
|-----------|------------------------------|--------------------------|--------------------------|--------------------------|-------------------------|-----------------------------|-------------------------------|
| Control | 88 | 55.54 ± 5.62 | 67.19 ± 4.81 | 67.19 ± 4.81 | 3.23 ± 1.33 | $11.71\pm3.28^{\mathrm{b}}$ | 26.67 ± 4.72^{b} |
| T200 | 117 | 61.88 ± 3.63 | 68.22 ± 5.16 | 79.00 ± 4.51 | 7.61 ± 2.44 | 20.50 ± 3.15^{a} | $45.87 \pm 4.62^{\mathbf{a}}$ |
| P-value | - | 0.4254 | 0.8885 | 0.1112 | 0.1552 | 0.09 | 0.0423 |

Abbreviations: Re-exp.: re-expansion; Hatch.: hatching.

Different letters represent a significant difference ($P \le 0.1$).

Figure 5: Effect of TUDCA concentrations on mRNA abundance of differentially expressed genes in embryo according to the following functional categories: endoplasmic reticulum stress (A), oxidative stress and response to cellular stress (B), and related to metabolism (C). Results are least-squares means \pm SEM. The effect of treatment was P \leq 0.1 for all genes in the figure. Control = 0 μ M; T200 = 200 μ M.



Figure 6: STRING network of protein-protein interactions between genes with significant difference ($P \le 0.1$) of mRNA abundance among groups. Different line colors representes the relationship between the proteins translated from analysed genes, as described on the legend.



CONSIDERAÇÕES FINAIS

Conclusões Gerais

A partir dos resultados apresentados nos Capítulos 1 e 2, todas as hipóteses propostas nesta Tese foram rejeitadas. Os principais resultados observados com este trabalho foram:

- A adição de TUDCA (50, 200 e 1.000 μM) durante a maturação *in vitro* (MIV) não melhorou a competência oocitária e o desenvolvimento do embrião.
- A adição de 200 µM de TUDCA, durante a MIV, pareceu modular o estresse do RE e o estresse oxidativo, diminuindo a produção de espécies reativas de oxigênio no oócito e aumentando a abundância de transcritos relacionados com a defesa antioxidante no oócito e no embrião.
- A adição de 1.000 µM de TUDCA, durante a MIV, se mostrou tóxica para o oócito e diminuiu as taxa de maturação nuclear e a atividade mitocondrial. Além disso, foi observado um aumento na abundância de transcritos pró-apoptóticos no oócito.
- A adição de TUDCA (50, 200 ou 1.000 μM) durante o cultivo *in vitro* (CIV) não melhorou a competência do desenvolvimento embrionário.
- A adição de 1.000 µM de TUDCA, durante a CIV, se mostrou tóxica para o embrião e diminuiu a taxa de eclosão. Além disso, houve um aumento da abundância de transcritos relacionados negativamente com o estresse do RE, o estresse oxidativo, o metabolismo lipídico e a sobrevivência celular.
- A adição de 200 µM de TUDCA, durante a CIV, pareceu modular o estresse do RE e o estresse oxidativo em embriões após a vitrificação. Além de aumentar a taxa de eclosão após o aquecimento, houve uma diminuição na abundância de transcritos relacionados negativamente com o estresse do RE e metabolismo lipídico, concomitante com o aumento na abundância de transcritos relacionados positivamente com a defesa antioxidante.

Resultados Suplementares

Por motivos técnicos, uma das análises que compõem o Capítulo 1 ainda não foi finalizada. Optamos por incluir na Tese a descrição desses experimentos, não concluídos, com o objetivo de aproveitar a análise crítica da banca examinadora.

Experiment: The effect of TUDCA during IVM on oocyte endoplasmic reticulum distribution

COCs were collected at 24h of IVM and vortexed in wash medium for 2 min. Separately, denuded oocytes were incubated in 100 nM ER-TrackerTM Red (glibenclamide BODIPY[®] TR) for 30 min in humidifier chamber at 38.5 °C at RT. For endoplasmic reticulum detection, oocytes were analyzed by Laser Scanning Confocal Microscope (TCS SP5 Leica) equipped with HeNe 543 laser (emission 564-663 nm and excitation 543 nm) and images were acquired by LAS AF Software. Each oocyte was visually classified according to three patterns of endoplasmic reticulum distribution previously described: ER I – presence of small ER clusters uniformly distributed throughout the cytoplasm with a few clusters concentrated at cortical region; ER II – predominance of ER clusters distributed on the cortical region of the oocyte; ER III – large ER clusters formation with disorganized distribution throughout the cytoplasm and presence of some open spaces.

Previous Results: This experiment was replicated four times using 30 COCs/treatment.



Illustrative images of fluorescence intensity representing endoplasmic reticulum distribution in the presence of ER-Tracker. Bar = 50 μ m. ER I = uniform distribution; ER II = cortical distribution; ER III = disorganized distribution.

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