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<b>Article Type</b>	Research article
<b>Article Title (within 20 words without abbreviations)</b>	Rapamycin Treatment During Prolonged <i>In Vitro</i> Maturation Enhances the Developmental Competence of Immature Porcine Oocytes
<b>Running Title (within 10 words)</b>	Rapamycin Treatment During Prolonged IVM Improves Development of Porcine Oocytes
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ACCEPTED

9 **Rapamycin Treatment During Prolonged *In Vitro* Maturation Enhances the Developmental Competence of**  
10 **Immature Porcine Oocytes**

11 **Running head:** Rapamycin Treatment During Prolonged IVM Improves Development of Porcine Oocytes

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24

25 **Abstract**

26 Porcine oocytes undergo *in vitro* maturation (IVM) for 42–44 h. During this period, most oocytes proceed to  
27 metaphase and then to pro-metaphase if the nucleus has sufficiently matured. Forty-four hours is sufficient for oocyte  
28 nuclear maturation but not for full maturation of the oocyte cytoplasm. This study investigated the influences of  
29 extension of the IVM duration with rapamycin treatment on molecular maturation factors. The phospho-p44/42  
30 mitogen-activated protein kinase (MAPK) level was enhanced in comparison with the total p44/42 MAPK level after  
31 52 h of IVM. Oocytes were treated with and without 10 µM rapamycin (10 R and 0 R, respectively) and examined  
32 after 52 h of IVM, whereas control oocytes were examined after 44 h of IVM. Phospho-p44/42 MAPK activity was  
33 upregulated the 10 R and 0 R oocytes than in control oocytes. The expression levels of maternal genes were highest  
34 in 10 R oocytes and were higher in 0 R oocytes than in control oocytes. Reactive oxygen species (ROS) activity was  
35 dramatically increased in 0 R oocytes but was similar in 10 R and control oocytes. The 10 R group exhibited an  
36 increased embryo development rate, a higher total cell number per blastocyst, and decreased DNA fragmentation. The  
37 mRNA level of development-related (*POU5F1* and *NANOG*) mRNA, oocyte-apoptotic (*BCL2L1*) genes were highest  
38 in 10 R blastocysts. These results suggest that prolonged IVM duration with rapamycin treatment represses ROS  
39 production and increases expression of molecular maturation factors. Therefore, this is a good strategy to enhance the  
40 developmental capacity in porcine oocytes.

41

42 Key words: IVM extension; Rapamycin; Molecular maturation factors; ROS; Porcine oocyte

43

## 44 Introduction

45 *In vitro* maturation (IVM) is essential for better understanding of process of oocyte development and maturation  
46 in various species, including pigs [1]. IVM induces meiotic maturation from prophase I to metaphase II (MII) *in vitro*.  
47 Oocytes generated by IVM are used to produce high-quality embryos upon *in vitro* fertilization (IVF) and somatic cell  
48 nuclear transfer (SCNT). Modulation of follicle size [2, 3], medium composition [4, 5], culture temperature [6, 7],  
49 antioxidants [8, 9] and cumulus features [10] can enhance the quality of porcine oocytes in IVM. Specifically,  
50 supplementation of antioxidant to IVM medium improved embryo quality, reduced early apoptosis levels, increased  
51 GSH content and reduced ROS accumulation, promoting embryonic development [11]. However, further research is  
52 needed to improve IVM of porcine oocytes.

53 Porcine oocytes, IVF embryos, and parthenotes are matured for 42–44 h and SCNT oocytes are matured for 36–  
54 38 h *in vitro*. During this period, most oocytes proceed to metaphase and then to pro-metaphase if the nucleus has  
55 sufficiently matured [12, 13]. While in meiotic arrest, the nuclear status and structural morphology of matured oocytes  
56 remain unchanged. This amount of time is sufficient for oocyte nuclear maturation but not for full maturation of the  
57 oocyte cytoplasm [14]. Many studies have been conducted on different maturation starts to acquire oocyte capacitation,  
58 which plays an important role in oocytes reaching MII [15-17]. Lin *et al.* extended the duration of IVM to 52 h in  
59 order to increase the poor-quality oocytes and performed treatment with melatonin to inhibit reactive oxygen species  
60 (ROS) production, apoptosis, and DNA damage [15]. However, oocyte maturation for an excessive amount of time  
61 causes cytoplasmic changes that negatively affect oocyte quality and increase the risk of spontaneous oocyte activation  
62 [18, 19] and subsequent aberrant cleavage characterized by unequally sized blastomeres [19, 20]. It is possible to  
63 prevent, delay, or reverse these cellular and molecular abnormalities [21].

64 Rapamycin has antifungal and immunosuppressant properties [22] and binds to FK506-binding protein 12 to  
65 form a complex that prevents the kinase activity and function of mTOR [23]. mTOR activity inhibition by rapamycin  
66 affects establishment of the cortical granule-free zone and actin cap, and disrupts alignment of the surrounding spindle  
67 and division during oocyte meiotic maturation [24]. Administration of rapamycin during IVM of porcine oocytes  
68 dose-dependently enhances cytoplasmic and nuclear maturation by inducing autophagy [25]. Aged porcine oocytes  
69 treated with 10  $\mu$ M rapamycin enhances blastocyst quality by regulating the mitochondrial distribution, autophagy,  
70 apoptotic cells, and mTOR signaling [26]. Supplementation of tissue culture medium (TCM)-199 with 0.5  $\mu$ M

71 rapamycin increases expression of matrix metalloproteinase in the trophoblast and inner cell mass (ICM), while it  
72 inhibits apoptosis [27]. Therefore, in our study, we extended the IVM duration to fully mature porcine oocytes and  
73 performed rapamycin treatment to reduce apoptosis, ROS production, and oocyte aging. This study investigated the  
74 impacts of rapamycin treatment during prolonged IVM on molecular maturation factors and the developmental  
75 capacity of porcine oocytes *in vitro*. Activated mitogen-activated protein kinase (MAPK) and ROS levels were  
76 modulated upon extension of the IVM duration with rapamycin treatment. The *in vitro* development rate, total cell  
77 numbers, and level of apoptosis were determined using blastocysts generated from these oocytes.

## 78 **Materials and Methods**

### 79 *Chemicals and reagents*

80 Unless otherwise specified, all chemicals and reagents utilized in this study were procured from Merck (St. Louis,  
81 MO, USA). The oocytes and embryos were maintained in CO<sub>2</sub> incubator at 38.8°C in a humidified atmosphere  
82 containing 5% CO<sub>2</sub> and 95% air.

### 84 *IVM of rapamycin-treated porcine oocytes*

85 Porcine ovaries from pre-pubertal sows were obtained from a provincial slaughterhouse. We used an 18-gauge  
86 needle attached to 10mL syringe to aspirate follicles measuring 2-8mm from the follicles, and selected cumulus-  
87 oocyte-complexes (COCs). After washing in TCM-199-HEPES supplemented with 0.1% (w/v) bovine serum albumin  
88 (BSA), a group of approximately 50 selected COCs were cultured in 500 µL of TCM-199 (M-199; Gibco, Grand  
89 Island, NY, USA) containing Earle's salts, 0.57 mM cysteine, 10% (v/v) porcine follicular fluid, 10 ng/mL epidermal  
90 growth (E-9644), 0.5 µg/mL follicle-stimulating hormone (F-2293), and 0.5 µg/mL luteinizing hormone (L-5269).  
91 The maturation process was conducted for 44 h, with the COCs placed beneath a layer of mineral oil. Thereafter, MII  
92 oocytes were transferred into TCM-199 supplemented with 0 or 10 µM rapamycin (R-8781) and incubated for 8 h  
93 (total IVM duration of 52 h). According to our previous studies, the concentration of rapamycin was set at the 10 µM  
94 [26].

### 95 *Parthenogenetic activation and embryo culture*

96 Subsequent IVM, adherent cumulus cells were eliminated using 0.1% (w/v) hyaluronidase. Collected oocytes  
97 were parthenogenetically activated by incubation in 5  $\mu\text{M}$   $\text{Ca}^{2+}$  ionomycin (Merck) for 5 min. Following culture in  
98 porcine zygote medium-5 (PZM-5) with 7.5  $\mu\text{g}/\text{mL}$  cytochalasin B (Merck) for 4 h, the oocytes were washed with  
99 PZM-5 containing 0.4% (w/v) BSA. Following 6 days of incubation in the same medium, the oocytes were washed  
100 with Dulbecco's phosphate-buffered saline (PBS). Finally, oocytes and embryos were fixed in 3.7% (w/v)  
101 paraformaldehyde for 20 min at 4°C or they were snap freezing using liquid nitrogen and subsequently stored at -70°C.

#### 102 *Detection of intracellular ROS activity*

103 Intracellular ROS activity in denuded oocytes were quantified using the 2,7-dichlorofluorescein protocol  
104 described previously [26]. Concisely, oocytes (25–30 oocytes per sample, four replicates) were cultured in 100  $\mu\text{M}$   
105 2',7'-dichlorodihydrofluorescein-diacetate (DCHF-DA) for 20 min, washed five times with PZM-5, and promptly  
106 observed under epifluorescence microscopy (Olympus, Tokyo, Japan) with an ultraviolet filter (450–490 nm and 515–  
107 565 nm). A microscope-mounted digital camera (Nikon, Tokyo, Japan) was used to capture grayscale images, and  
108 ImageJ software (NIH, Bethesda, MD, USA) was employed to acquire mean grayscale values. Image analysis was  
109 conducted with the Adobe Photoshop CS6 software package (version 13, Adobe Systems Inc.) by quantitating the  
110 average pixel intensities in various regions of the raw image. Before statistical analysis, background fluorescence  
111 values were calculated by subtracting them from the final values.

#### 112 *TUNEL assay*

113 Blastocysts on day 6 were washed with PBS (pH 7.4) with 1 mg/mL polyvinylpyrrolidone (PBS/PVP). After  
114 fixing with 3.7% formaldehyde prepared with PBS overnight at 4°C. Following washing with PBS/PVP, the oocytes  
115 were subsequently permeabilized by culture in the dark for 1 h with 0.3% Triton X-100 at room temperature. After  
116 washing with PBS/PVP, the blastocysts were cultured at 37°C for 1 h in fluorescein-conjugated dUTP and terminal  
117 deoxynucleotidyl transferase using the In Situ Cell Death Detection Kit (Roche, Mannheim, Germany). Following  
118 counterstaining with Hoechst 33342 in the dark for 30 min at 37°C to label the nuclei, the samples were washed with  
119 PBS/PVP and then carefully set on glass slides. The experiment was independently repeated three times.

#### 120 *Real-time quantitative PCR*

121 mRNA extraction from oocytes at the MII stage and *in vitro*-cultured embryos at day 6 (20 embryos per sample,  
122 three replicates) by a Dynabeads mRNA Direct Kit (DynaLAsa, Oslo, Norway). SuperScript™ III reverse transcriptase  
123 (Invitrogen, Grand Island, NY, USA) and oligo (dT)<sub>12-18</sub> primers were used to synthesize first-strand cDNA. Real-  
124 time quantitative PCR was conducted on a Step One Plus Real-time PCR system (Applied Biosystems, Warrington,  
125 UK) using the primers specified in Table 1. The total reaction volume for the final PCR consisted of 20 µL, including  
126 SYBR Green PCR Master Mix (Applied Biosystems). As follows were the amplification conditions: 10 min at 94°C,  
127 followed by 39 cycles of denaturation for 30 sec at 94°C, annealing for 30 sec at 55°C, and extension for 55 sec at  
128 72°C, and a final extension for 5 min at 72°C. Relative mRNA expression levels were determined according to the 2<sup>-</sup>  
129  $\Delta\Delta C_t$  protocol [28] by normalization to *GAPDH*.

### 130 *Western blot analysis*

131 The experimental assay followed a previously described protocol [24]. For protein extraction, 20 µL of 1× SDS  
132 sample buffer containing 5 mM Tris-HCl, pH 6.8 at 25°C, 2% (w/v) SDS, 10% (v/v) glycerol, 50 mM DTT, and 0.01%  
133 (w/v) bromophenol blue or phenol red was added to oocytes (20 oocytes per sample, three replicates), and the samples  
134 were boiled for 5 min at 95°C. Subsequently, proteins were separated by electrophoresis on a 5–12% Tris-SDS-PAGE  
135 gel for 1.5 h at 80–100 V. Thereafter, proteins were electrophoretically transferred to a nitrocellulose membrane  
136 (Hybond-ECL; Amersham, Buckinghamshire, UK) at 300 mA for 2 h using transfer buffer (25 mM Tris, 200 mM  
137 glycine, and 20% methanol, pH 8.5). After being blocked with 5% (w/v) skim milk prepared in PBS for 1 h, the  
138 membranes were cultured with an anti-Cdc2 p34, anti-p44/42 MAPK, or anti-phospho-p44/42 MAPK antibody (1:500;  
139 Cell Signaling Technology, Danvers, MA, USA) in blocking solution (1× Tris-buffered saline containing 0.1% (v/v)  
140 Tween® 20 and 5% (w/v) skim milk) for at least 2 h. After washing with TBST (20 mM Tris-HCl, pH 7.5, 250 mM  
141 NaCl, and 0.1% (v/v) Tween® 20), the membranes were cultured with anti-rabbit IgG-horseradish peroxidase (1:2000,  
142 Cell Signaling Technology) with blocking solution for 1 h. Following washing with TBST, binding of antibody was  
143 analyzed with chemiluminescence luminol reagent (Invitrogen).

### 144 *Statistical analysis*

145 The Statistical Analysis System (SAS User's Guide, 1985; Statistical Analysis System Inc., Cary, NC, USA)  
146 was employed to analyze data using the general linear model procedure. Significant differences were determined using

147 Tukey's multiple range test. Relative gene expression levels were compared by the Student's *t*-test. Differences were  
148 regarded as biologically significant at  $p < 0.05$  and  $p < 0.01$ .

## 149 **Results**

### 150 *Rapamycin treatment increases the levels of maturation factors in porcine oocytes*

151 To explore the influence of extension of the IVM duration on molecular maturation of porcine oocytes, we  
152 examined expression of maturation/M phase-promoting factor (MPF), activated MAPK, and maternal genes (Fig. 1  
153 and 2). We monitored total and phospho-p44/42 MAPK levels at 2 h intervals after 42–52 h of IVM (Fig. 1A). The  
154 phospho-p44/42 MAPK level was normalized against the total MAPK level at 44 h (set to 1). The phospho-p44/42  
155 MAPK level was upregulated in comparison with the total MAPK level at 52 h (42 h,  $1.02 \pm 0.09$ ; 46 h,  $0.94 \pm 0.07$ ;  
156 48 h,  $0.82 \pm 0.07$ ; 50 h,  $0.90 \pm 0.11$ ; and 52 h,  $1.36 \pm 0.20$ ;  $p < 0.05$  and  $p < 0.01$ ; Fig. 1B). The phospho-p44/42  
157 MAPK level was higher at 52 h than at the other time points (Fig. 1A and B).

158 Oocytes were supplemented with and without 10  $\mu$ M rapamycin and examined after 52 h of IVM (10 R and 0  
159 R, respectively), whereas control oocytes were examined subsequent 44 h of IVM. The mRNA level of the maternal  
160 genes *BMP15*, *GDF9*, and *MOS* was analyzed using real-time quantitative PCR (Fig. 2A). The mRNA level of these  
161 genes was upregulated in 10 R oocytes compared to control and 0 R oocytes ( $p < 0.05$ ). There was an increase ( $p <$   
162  $0.01$ ) in the levels of *BMP15* and *GDF9* in 10 R oocytes compared with control oocytes. We monitored the levels of  
163 Cdc2 p34 and phospho-p44/42 MAPK (Fig. 2B). Expression of Cdc p34 and phospho-p44/42 MAPK was normalized  
164 to the total MAPK in each group (set to 1, Fig. 2C and D). The levels of Cdc2 p34 and phospho-p44/42 MAPK were  
165 higher in 10 R oocytes ( $p < 0.01$ ) than in control and 0 R oocytes (0 R,  $1.05 \pm 0.11$  and  $1.14 \pm 0.01$ , respectively; and  
166 10 R,  $1.21 \pm 0.02$  and  $1.29 \pm 0.06$ , respectively; Fig. 2C and D).

### 167 *Rapamycin treatment reduces ROS activity in porcine oocytes*

168 ROS activity was analyzed in control, 0 R, and 10 R groups using DCHFDA. ROS activity was lower in 10 R  
169 group ( $p < 0.01$ ) compared to 0 R oocytes and was similar to that in control oocytes (control,  $32.3 \pm 3.6$ ; 0 R,  $65.1 \pm 5.9$ ;  
170 and 10 R,  $28.9 \pm 7.6$ ; Fig. 3A and B). Treatment with 10  $\mu$ M rapamycin enhanced the developmental capacity of oocytes  
171 aged for 8 h by inhibiting ROS activity.

172 *Rapamycin treatment enhances the developmental capacity of porcine oocytes*

173 Control (n=400) and 0 R (n=345) and 10 R (n=310) groups were matured for 44 and 52 h, respectively. The  
174 percentage of matured oocytes was no differences among the control, 0 R, and 10 R groups ( $86.40\% \pm 1.92\%$ ,  $90.30\%$   
175  $\pm 2.03\%$ , and  $91.77\% \pm 2.86\%$ , respectively; Fig. 4B). After IVM, control (n=345), 0 R (n=312), and 10 R (n=284)  
176 groups were parthenogenetically activated. The morphology and percentage of embryos reaching the 2–4-cell stage  
177 showed no significant difference among the control, 0 R, and 10 R groups ( $68.73\% \pm 5.57\%$ ,  $56.58\% \pm 5.10\%$ , and  
178  $57.61\% \pm 4.60\%$ , respectively; Fig. 4A and B). The percentage of blastocyst at day 6 was highest in the 10 R group  
179 (n=75,  $48.36\% \pm 7.06\%$ ) and lowest in the 0 R group (n=38,  $22.81\% \pm 4.12\%$ ;  $p < 0.05$  or  $p < 0.01$ ; Fig. 4A and B).

180 Confirming the influence of rapamycin treatment on blastocyst quality, blastocysts at day 6 in the various groups  
181 were stained (Fig. 4C). The cell number per blastocyst at day 6 was highest in the 10 R group (control,  $50 \pm 6.92$ ; 0  
182 R,  $54 \pm 9.29$ ; and 10 R,  $76 \pm 12.49$ ;  $p < 0.05$ ; Fig. 4C). Reprogramming-related transcription factor genes *POU5F1*,  
183 *SOX2*, *NANOG*, and *CDX2* were examined for their expression levels by real-time RT-PCR (Fig. 4D). The 10 R group  
184 was upregulated ( $p < 0.05$ ) in *POU5F1* and *NANOG* expression, whereas it did not significant difference among the  
185 control and 0 R groups. *SOX2* expression tended to show a similar pattern, although there were not significantly  
186 differences among the groups. Expression of *CDX2* showed no significant difference among the groups.

187 *Rapamycin treatment of porcine oocytes decreases apoptosis in resultant blastocysts*

188 Using the TUNEL assay, individual embryos were assessed for genomic DNA fragmentation (an indicator of  
189 apoptosis). The percentage of fragmented DNA in the 10 R group ( $7.08\% \pm 0.50\%$ ) was significantly downregulated  
190 ( $p < 0.05$ ) compared with the 0 R group ( $13.06\% \pm 2.24\%$ ) and was similar to the control group ( $7.07\% \pm 1.37\%$ , Fig.  
191 5B). The 10 R group showed significantly higher levels of *BCL2L1*, the anti-apoptotic gene, compared to the 0 R  
192 group ( $p < 0.05$ ), while the control group exhibited the highest expression ( $p < 0.05$ ) (Fig. 5C). The mRNA level of  
193 the *FAS* and *CASP3*, pro-apoptotic genes, tended to be downregulated in the 10 R group, although these differences  
194 were not significant.

195

196 **Discussion**

197 Oocyte maturation is a multifaceted process involving both nucleus and cytoplasmic changes. An IVM  
198 duration of 42 to 44 h is sufficient for nuclear maturation through a mechanism such as oocyte capacity, but not

199 sufficient for complete maturation of the cytoplasm [14]. However, excessive maturation duration exposes oocytes  
200 to increased oxidative stress [29, 30]. ROS-induced oocyte damage reduces the ability and quality of subsequent  
201 embryonic development [31]. This study investigated the influences of extension of the IVM duration with  
202 rapamycin treatment on molecular maturation factors and embryonic development of porcine oocytes.

203 After 52 h of IVM, the phospho-p44/42 MAPK level was high in comparison with the total MAPK level (Fig.  
204 1A and B). Our previous study demonstrated that 10  $\mu$ M rapamycin increases blastocyst quality by affecting  
205 developmental rate and total cell number and reducing mitochondrial distribution, apoptosis, autophagy and ROS  
206 activity that regulates mTOR signaling [26]. The maternal genes expression levels of *BMP15*, *GDF9*, and *MOS*  
207 was higher in 0 R and 10 R oocytes than in control oocytes (Fig. 2A). Confirming the effect of rapamycin, levels  
208 of Cdc2 p34 and phospho-p44/42 MAPK were higher in 10 R oocytes than in control and 0 R oocytes (Fig. 2B–  
209 D). MAPK and MPF are critical for meiotic molecular maturation of oocytes. Protein phosphorylation and  
210 dephosphorylation are essential for the meiotic cell cycle of oocytes. MPF and MAPK, which are integral  
211 components of the key regulatory pathways involved in activation of extracellular signal-regulated kinase 1 and 2,  
212 are serine/threonine kinases that are phosphorylated and subsequent activated by MAPK kinase [32, 33]. MPF  
213 consists of the catalytic subunit p34/Cdc2, which possesses serine/threonine kinase activity, and the regulatory  
214 subunit cyclin B. Activation of MPF is regulated by various mechanisms, including binding of cyclin B to Cdc2,  
215 phosphorylation of threonine 161, and dephosphorylation of tyrosine 15 and threonine 14 [34]. MAPK activity  
216 usually peaks in porcine oocytes during IVM for 42–44 h. Many researchers have extended the IVM duration to  
217 improve oocyte maturity due to supply of insufficient maturation factors. The developmental rate reportedly  
218 increases when the IVM duration is extended (48–72 h). Specifically, the developmental rate is increased at 56 h  
219 due to maintenance of high MPF expression, but cytoplasmic senescence at 72 h decreases the developmental rate  
220 and suppresses MPF expression [35]. Another study confirmed that the developmental rate is improved by  
221 extending the IVM duration from 44 h, which is normally used. In the extended culture duration of 24 to 52 h, the  
222 cleavage rate of porcine oocytes was highest at 48 h as determined through the number of cumulus cell layers. [36].  
223 Although that study did not reveal whether this approach affects a molecular factor, it showed that long-term culture  
224 is more effective for oocyte maturation [36]. To induce resumption of meiosis, porcine COCs at the pre-IVM stage  
225 were preincubated for 12 h [17]. Extension of the total IVM duration to 52 h increases the development rate of IVF  
226 embryos at the blastocyst stage [17]. Excessive prolongation of oocyte maturation leads to aging, and reduced  
227 MAPK activity decreases the quality of oocytes [37, 38]. The IVM duration of poor-quality oocytes was previously

228 extended to 52 h, and treatment with the antioxidant melatonin improved the IVM rate and expression levels of  
229 maturation factors [15]. We previously showed that treatment of aged oocytes with rapamycin increases  
230 phosphorylated p44/42 MAPK activity and mRNA levels of maternal genes compared with untreated aged oocytes  
231 [26]. Rapamycin activates MPF and MAPK, which reduces oocyte activation susceptibility via inhibition of protein  
232 kinase A [39]. Therefore, we suggest that rapamycin treatment facilitates molecular maturation by preserving the  
233 ooplasm of MII oocytes, leading to enhanced transcription of maternal genes. This study indicates that extension  
234 of the IVM duration with rapamycin treatment, which maintains expression of molecular maturation factors, does  
235 not negatively affect porcine oocytes.

236 ROS function are crucial signaling molecules in diverse physiological processes, including resumption of the  
237 meiotic cell cycle, and contribute to pathological processes such as apoptosis and senescence [40, 41]. ROS were  
238 proposed to participate in oocyte meiotic arrest [42-44]. Oxidative stress perturbs bovine embryonic development  
239 after fertilization [45]. Furthermore, a change in the redox status of human oocytes during *in vitro* culture is related  
240 to an increased occurrence of apoptosis in gametes [46]. Porcine gametes can incur DNA damage and undergo  
241 apoptosis during IVM [47-49]. Treatment with rapamycin effectively reduces intracellular ROS levels and  
242 improves mitochondrial localization [26]. Extension of the IVM duration with rapamycin treatment demonstrated  
243 a significant reduction of levels of ROS (Fig. 3A and B). This finding suggests that this approach improves *in vitro*  
244 oocyte culture and maintains oocyte health.

245 We investigated the influence of extension of the IVM duration with rapamycin treatment on porcine  
246 embryonic development. The percentage of blastocyst formation at day 6 and total cell number per blastocyst were  
247 increased when the IVM duration was extended with rapamycin treatment (Fig. 4A–C). Gene expression is a major  
248 contributor to embryonic development, and any disruption in gene expression during culture of embryos *in vitro*  
249 can potentially hinder embryo production [50, 51]. We examined mRNA levels of *POU5F1*, *SOX2*, *NANOG*, and  
250 *CDX2* in the 0 R, 10 R, and control groups. Transcription factor genes such as these play essential roles in early  
251 development and are indispensable for proliferation of undifferentiated embryonic stem (ES) cells in culture.  
252 mRNA and protein expression of *POU5F1* has been detected in various cellular components, including blastomeres  
253 of preimplantation embryos, the ICM of blastocysts, epiblasts, primordial germ cells, and the majority of germ cells  
254 [52-54]. *SOX2* and *NANOG* form interactions with *POU5F1* to regulate the transcriptional hierarchy that specifies  
255 ES cell identity [55-57]. The transcription factors *POU5F1* and *SOX2* are expressed in both the ICM and

256 trophoctoderm (TE) of porcine blastocysts. *CDX2* expression is essential for TE formation [58]. *POU5F1* and  
257 *NANOG* exhibited significant upregulated ( $p < 0.05$ ) in 10 R blastocysts (Fig. 4D). However, *CDX2* expression  
258 was not significantly affected. These data show that extension of the IVM duration with rapamycin treatment  
259 upregulates specific transcription factors related to the ICM, including *POU5F1* and *NANOG*, but does not affect  
260 expression of the TE-related factor *CDX2* in blastocysts. Taken together, these observations demonstrate that  
261 extension of the IVM duration with rapamycin treatment significantly influences porcine embryonic development  
262 *in vitro*.

263 We explored the effect of extension of the IVM duration with rapamycin treatment on apoptotic cell death  
264 in porcine embryos. DNA fragmentation was significantly decreased in 10 R blastocysts extended duration of *in*  
265 *vitro* maturation (IVM) by rapamycin treatment (Fig. 5B). We also examined the apoptosis expression levels of  
266 *BCL2L1*, *FAS*, and *CASP3* (Fig. 5C). Although the pro-apoptotic expression levels of *FAS* and *CASP3* was not  
267 different in all groups, the anti-apoptotic expression level of *BCL2L1* in blastocysts was increased in the 10 R group.  
268 *BCL2L1* encodes the protein BCL-xL, an anti-apoptotic protein [59]. BCL-xL prevents cell death by inhibiting  
269 cytochrome c release from mitochondria, a key step in the cell death pathway [60, 61]. These observations suggest  
270 that extension of the IVM duration with rapamycin treatment facilitates embryonic development by suppressing  
271 apoptosis at the molecular level during preimplantation stages.

272 Our results indicate that extension of the IVM duration with rapamycin treatment enhances molecular  
273 maturation of porcine oocytes by repressing ROS production and improves porcine embryonic development. This  
274 study demonstrates the combination of extension of the IVM duration and treatment with rapamycin enhances  
275 maturation of porcine oocytes. These results should be further applied to assisted reproductive technology to  
276 produce high-quality embryos.

277

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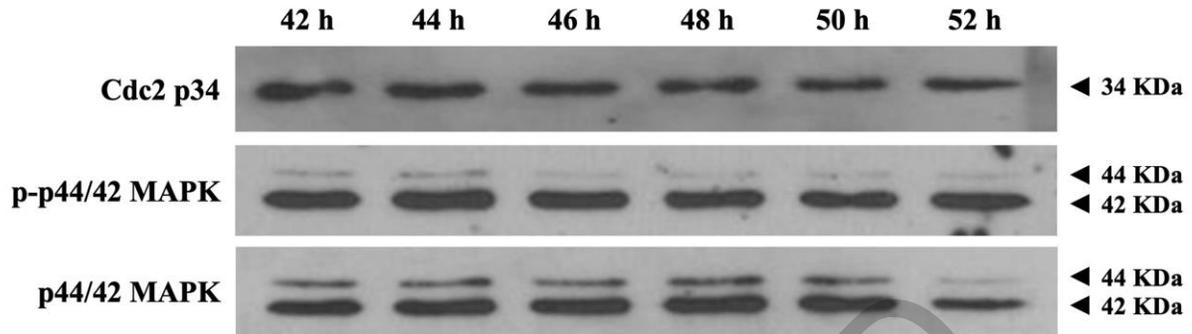
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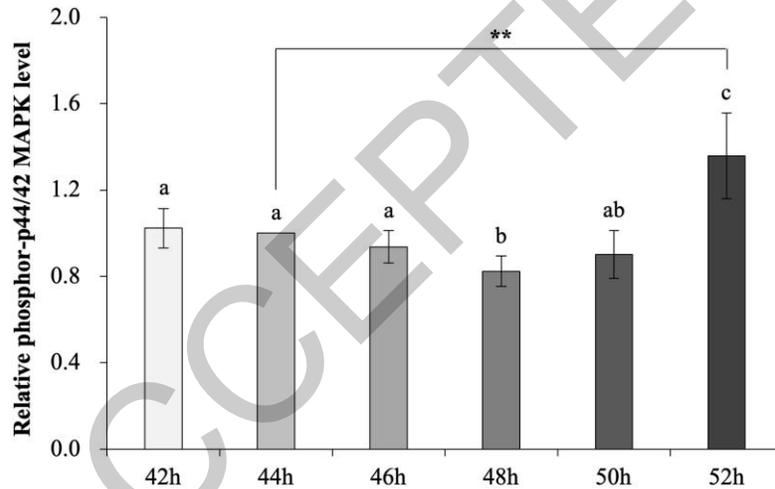
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A



B



425

426 **Figure 1.** Levels of maturation factors during the extended *in vitro* maturation (IVM) of porcine oocytes.

427 Western blot analysis of Cdc2 p34, p44/42 mitogen-activated protein kinase (MAPK), and phospho-p44/42

428 MAPK (A) and band intensities (B) of Cdc2 p34, p44/42 MAPK, and phospho-p44/42 MAPK in porcine

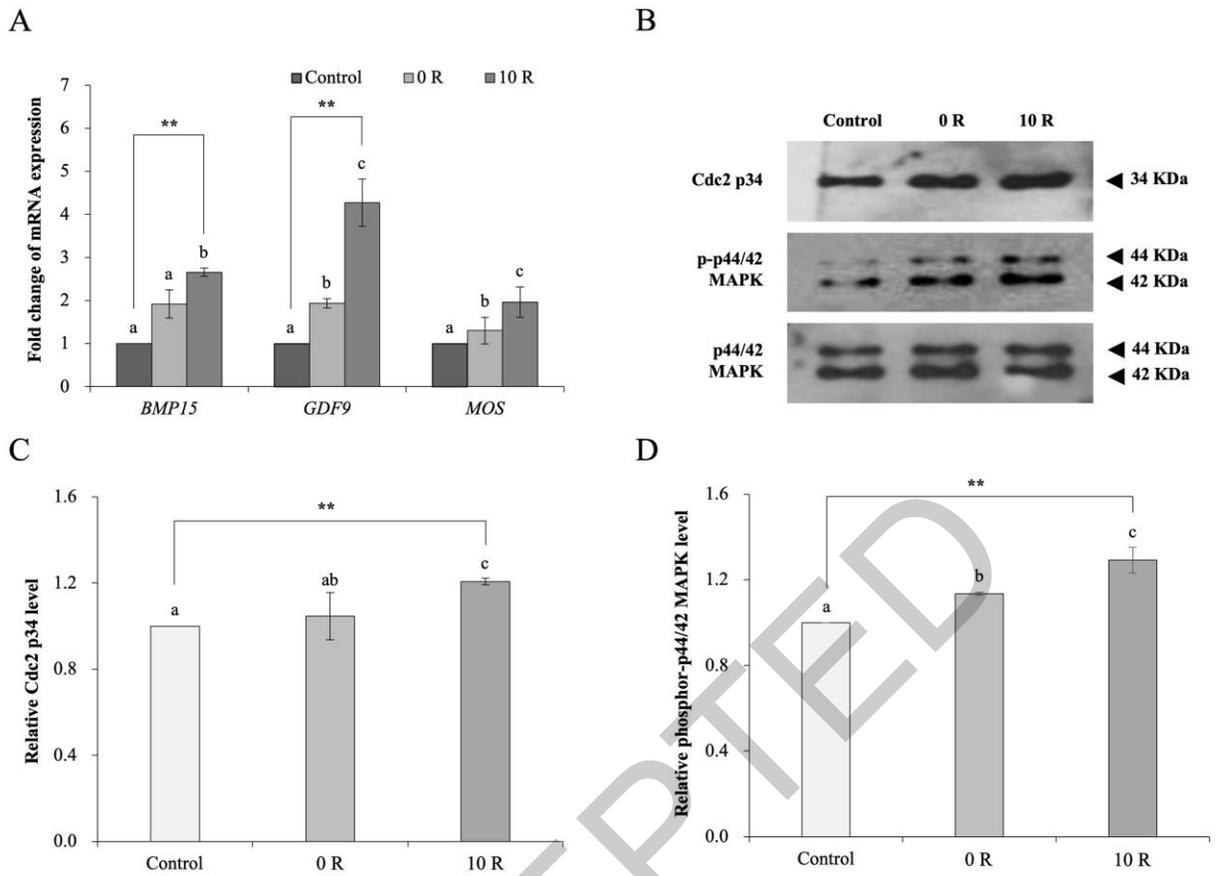
429 oocytes were examined every 2 h after 42–52 h of IVM. The experiment was independently repeated three

430 times. Significant differences compared with control oocytes are indicated (<sup>a-c</sup>*p* < 0.05 and \*\**p* < 0.01).

431 Values represent means ± standard error of the mean of independent experiments.

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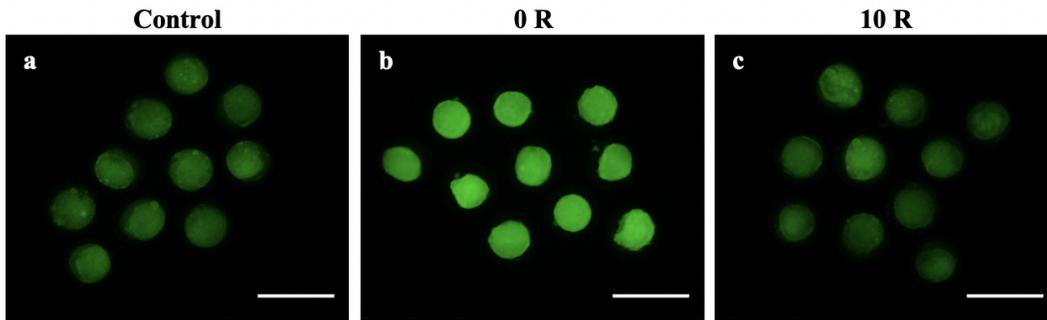
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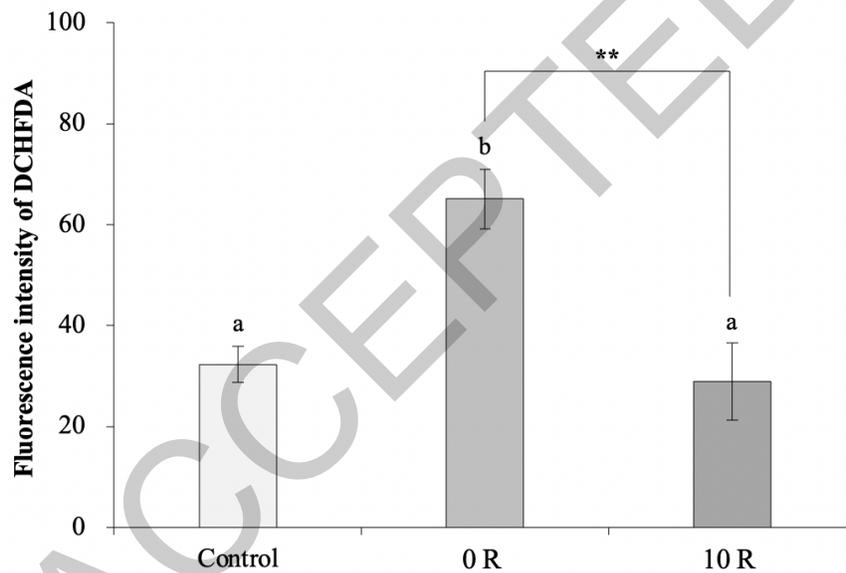
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**Figure 2.** Effect of rapamycin treatment on molecular maturation factors in porcine oocytes. Maternal gene expression (A) and levels of Cdc2 p34, p44/42 MAPK, and phospho-p44/42 MAPK (B) were examined in control, 0 R, and 10 R oocytes. Relative levels of Cdc2 p34 (C) and phospho-p44/42 MAPK (D) were determined. *GAPDH* was used as an internal standard. The experiment was independently repeated three times. Significant differences compared with control oocytes are indicated (<sup>a-b</sup>*p* < 0.05 and <sup>\*\*</sup>*p* < 0.01). Values represent means ± standard error of the mean of independent experiments.

A



B



443

444 **Figure 3.** Antioxidant effect of rapamycin treatment during IVM. Images of oocytes stained with DCHFDA

445 in rapamycin-untreated 44 h IVM (control), rapamycin-untreated 52 h IVM (0 R), and 10  $\mu$ M-rapamycin

446 treated 52 h IVM (10 R) (A) and the fluorescence intensity of DCHFDA (B) were evaluated in metaphase

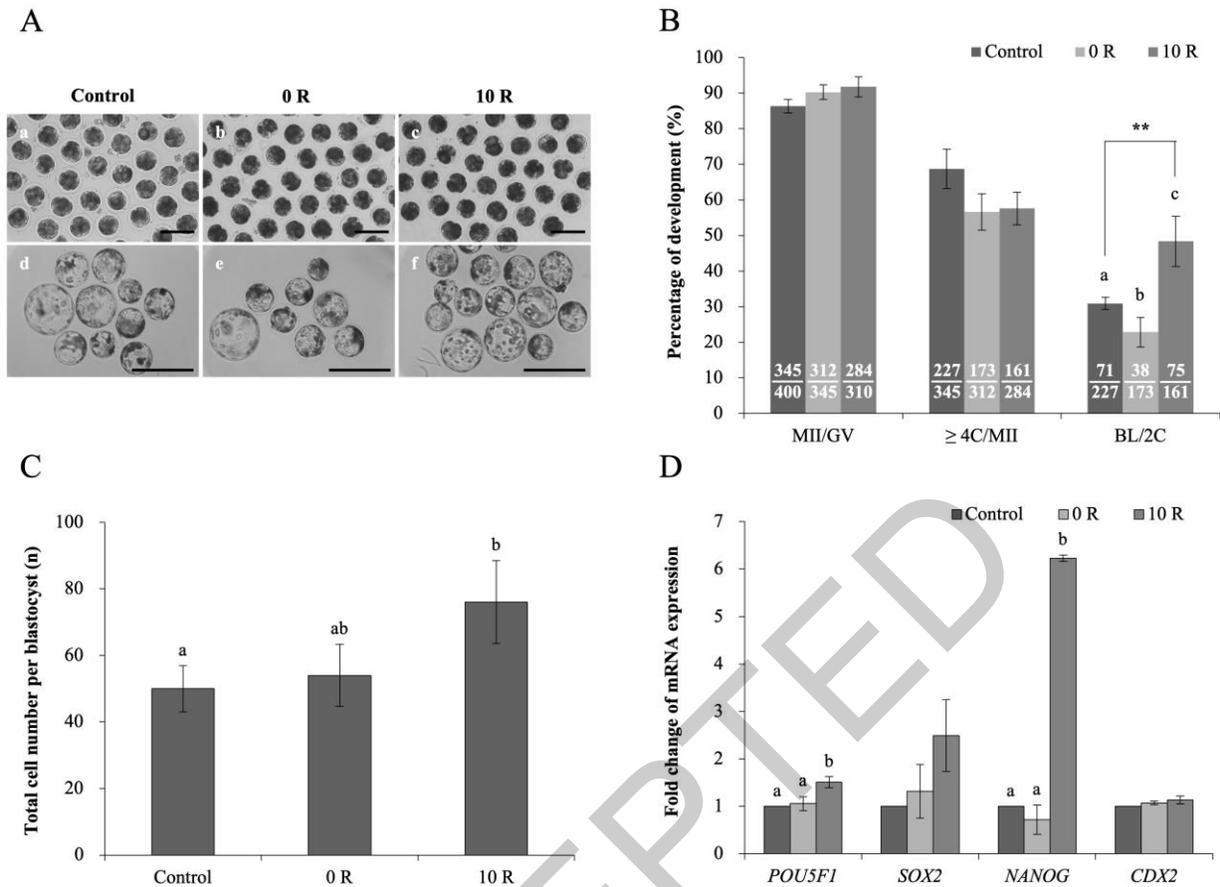
447 II oocytes in the control, 0 R, and 10 R groups. The experiment was independently repeated four times.

448 Significant differences compared with control group are indicated (<sup>a-b</sup> $p < 0.05$  and <sup>\*\*</sup> $p < 0.01$ ). Values

449 represent means  $\pm$  standard error of the mean of independent experiments. Bar = 200  $\mu$ m.

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452

453 **Figure 4.** Effect of rapamycin treatment on the developmental capacity of porcine oocytes. (A) Morphology

454 of embryos at day 2 (a–c) and blastocysts at day 6 (d–f) were examined in the control (a and d), 0 R (b and

455 e), and 10 R (c and f) groups. The *in vitro* development rate (B), total cell number per blastocyst at day 6

456 (C), and relative mRNA expression of the development-related genes *POU5F1*, *SOX2*, *NANOG*, and *CDX2*

457 (D) were examined in blastocysts in the control, 0 R, and 10 R groups. MII, Metaphase II; GV, Germinal

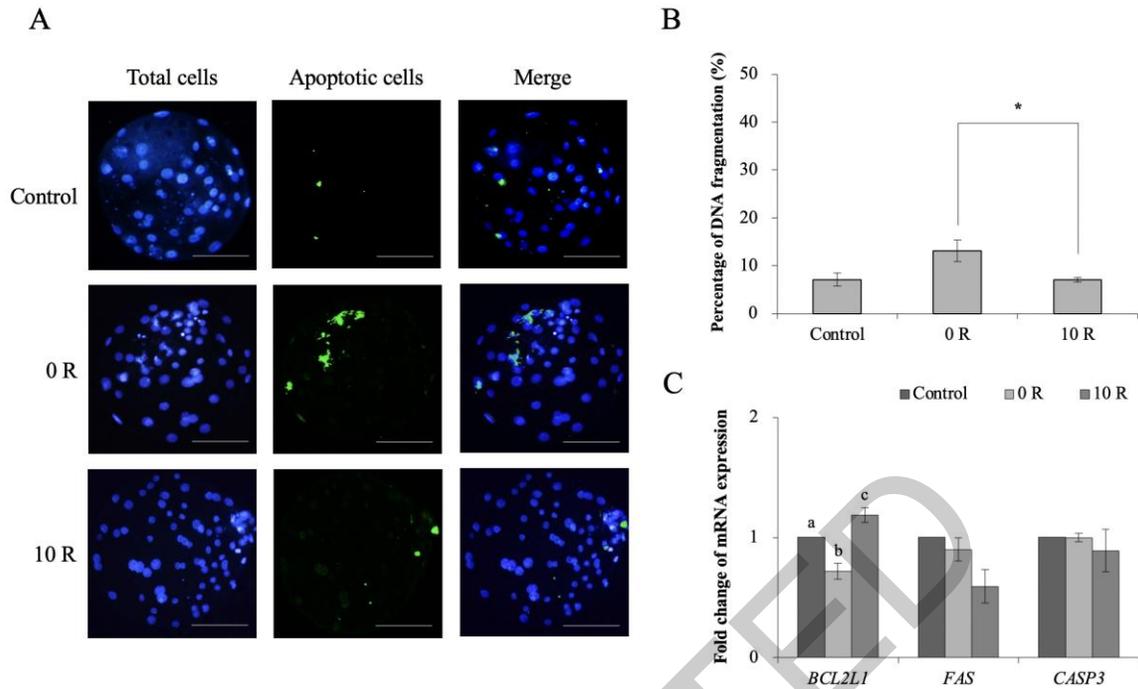
458 vesicle; BL, Blastocyst. The experiment was independently repeated five times (A and B) and three times

459 (C and D). Significant differences compared with the control group are indicated (<sup>a-c</sup>*p* < 0.05 and <sup>\*\*</sup>*p* <

460 0.01). Values represent the means ± standard error of the mean of independent experiments. Bar = 200 μm.

461

462



463  
 464 **Figure 5.** Effect of treatment of porcine oocytes with rapamycin on the level of apoptosis in blastocysts.  
 465 Morphology of blastocyst total cell and apoptotic cells (A), DNA fragmentation (B) and relative mRNA  
 466 expression of the apoptosis-related genes *BCL2L1*, *FAS*, and *CASP3* (C) were examined in blastocysts in  
 467 the control, 0 R, and 10 R groups. The experiment was independently repeated three times. Significant  
 468 differences compared with the control group are indicated (\* $p < 0.05$ ). Values represent means  $\pm$  standard  
 469 error of the mean of independent experiments. Bar = 50  $\mu\text{m}$ .

470  
 471

472 **Table 1.** Primers used for real-time PCR

Gene	GenBank accession no.	Primer sequence*	Annealing temperature (°C)	Product size (bp)
<i>GAPDH</i>	AF017079	F: GGGCATGAACCATGAGAAGT R: AAGCAGGGATGATGTTCTGG	60	230
<i>BMP15</i>	NM_001005155	F: CCCTCGGGTACTACACTATG R: GGCTGGGCAATCATATCC	60	192
<i>GDF9</i>	AY_626786	F: GAGCTCAGGACACTCTAAGCT R: CTTCTCGTGGATGATGTTCTG	60	272
<i>MOS</i>	NM_001113219	F: TGGGAAGAACTGGAGGACA R: TTCGGGTCAGCCCAGGTTCA	60	121
<i>POU5F1</i>	NM_001113060	F: AGTGAGAGGCAACCTGGAGA R: TCGTTGCGAATAGTCACTGC	60	166
<i>NANOG</i>	DQ447201.1	F: GAACTTTCCAACATCCTGAA R: TTTCTGCCACCTCTTACATT	55	87
<i>SOX2</i>	EU503117	F: GCCCTGCAGTACAACCTCCAT R: GCTGATCATGTCCCGTAGGT	60	216
<i>CDX2</i>	AM778830	F: AGCCAAGTGAAAACCAGGAC R: TGCGGTTCTGAAACCAGATT	60	178
<i>BCL2L1</i>	AF216205	F: ACTGAATCAGAAGCGGAAAC R: AAAGCTCTGATACGCTGTCC	60	249
<i>FAS</i>	AJ001202	F: AAGTTCCCAAGCAAGGGATT R: AATTTCCCATGTGGAGCAG	60	207
<i>CASP3</i>	NM_214131.1	F: GAGGCAGACTTCTTGTATGC R: ACAAAGTGACTGGATGAACC	55	93

473 \*F, forward; R, reverse.