

1        **Effects of Kisspeptin-10 on in vitro proliferation and Kisspeptin receptor**  
2        **(Kiss-1R) expression in primary epithelial cell cultures isolated from bovine**  
3        **placental cotyledons of embryos/fetuses at the first trimester of pregnancy**

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6                NA Martino<sup>1\*</sup>, A Rizzo<sup>1</sup>, F Pizzi<sup>2</sup>, ME Dell'Aquila<sup>3#</sup>, RL Sciorsci<sup>1#</sup>

7                **Running title: Bovine placental cotyledon cell cultures and Kisspeptin-10**

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10        <sup>1</sup>Veterinary Clinics and Animal Production Unit – Dipartimento dell’Emergenza e Trapianti  
11        D’Organo (DETO), Università degli Studi di Bari Aldo Moro, Bari, Italy.

12        <sup>2</sup>Istituto di Biologia e Biotecnologia Agraria (IBBA) Consiglio Nazionale delle Ricerche (CNR),  
13        Milan, Italy

14        <sup>3</sup>Dipartimento di Bioscienze, Biotecnologie e Biofarmaceutica (DBBB), Università degli Studi di  
15        Bari Aldo Moro, Bari, Italy.

16  
17  
18  
19        \*Correspondence to: nicmartino@libero.it

20        #These two authors contributed equally

27 **ABSTRACT**

28 Kisspeptins (Kps) and Kiss-1 receptor (Kiss-1R) expression have been reported in the placenta and  
29 a possible involvement of the Kiss-1R/Kps system in regulating trophoblast invasion and  
30 proliferation has been hypothesized. The aim of the present study was to investigate whether Kiss-  
31 1R activation by Kisspeptin 10 (Kp-10) could modulate *in vitro* proliferation and progesterone (P<sub>4</sub>)  
32 secretion of bovine primary placental cell lines isolated from cotyledons of foetuses in the first  
33 pregnancy trimester. The involvement of Kiss-1R in the cell responses observed was also analyzed.  
34 Uteri from cows at the first trimester of pregnancy were obtained from local abattoirs. Fetal  
35 cotyledon fragments were digested with collagenase in Low Glucose Dulbecco's Modified Eagle's  
36 Medium (LG-DMEM) and cell lines were isolated. After being characterized for epithelial  
37 polygonal morphology, presence of binucleate cells, male gender and expression of cytokeratin and  
38 zona occludens 2, cell lines were cultured in a LG-DMEM based expansion medium in the presence  
39 of 0.01, 0.1, 1 and 10 µM Kp-10. Control cells were cultured in the absence of Kp-10. Cell  
40 population doubling time was evaluated for each culture passage (P) from P1 to P10. Cells were  
41 tested for Kiss-1R mRNA expression analysis by Real Time RT-PCR and culture media were  
42 analyzed for P<sub>4</sub> concentration by radioimmunoassay. Kp-10 modulated *in vitro* proliferation of  
43 epithelial cell lines isolated from cotyledons recovered from bovine fetuses in the first trimester of  
44 pregnancy. Inhibitory (line A) or stimulatory (line B) effects of Kp-10 on cell proliferation were  
45 found in different cell lines and observed cell responses were found to be related to Kiss-1R mRNA  
46 levels. Inhibition of cell proliferation matched with not significant variation of Kiss-1R expression,  
47 whereas stimulation of cell proliferation was found to be related to Kiss-1R up-regulation. In both  
48 cell lines, no effect of Kp-10 on P<sub>4</sub> secretion was found at any tested concentration. These results  
49 lead to the conclusion that the Kiss-1R/Kps system is involved in the regulation of cell proliferation  
50 of bovine placental cotyledon cell lines isolated at the first trimester of pregnancy but, at this  
51 gestational stage, it may be not involved in modulating placental P<sub>4</sub> secretion.

52

53 **Key Words** Bovine placental cotyledon cell line, *in vitro* proliferation, Kiss-1R, Kisspeptin-10,  
54 progesterone secretion.

55

## 56 **1. Introduction**

57 Kisspeptins (Kps) are a family of peptides derived from the primary translation product of the Kiss-  
58 1 gene, which was identified and designated in 1996 [1]. The main Kiss-1 gene product is a 145  
59 amino acid polypeptide which, by proteolytic processing, results in shorter Kps, with 54 (Kp-54;  
60 metastin), 14 (Kp-14), 13 (Kp-13) or 10 (Kp-10) amino acids, respectively. All small peptides share  
61 a common carboxyl-terminal amidated 10-aa sequence which is enough to efficiently bind the Kp  
62 receptor (Kiss-1R) and induce its biological activity [2]. Kiss-1R is a G protein-coupled receptor  
63 also known as GPR54 and transduces the kps signal through Gq/11- $\alpha$  subunits of heterotrimeric G  
64 proteins. The activation of phospholipase C (PLC), upon Kiss-1R stimulation, induces the  
65 hydrolysis of phosphatidylinositol bisphosphate (PIP<sub>2</sub>) into inositol 1,4,5-trisphosphate (IP<sub>3</sub>) and  
66 subsequent mobilization of calcium (Ca<sup>2+</sup>) ions from intracellular stores. In addition, the rise of PIP<sub>2</sub>  
67 hydrolysis following Kp stimulation, leads to diacylglycerol (DAG) formation and, thereby, protein  
68 kinase C (PKC) activation [3,4]. In turn, activated PKC is thought to cause phosphorylation of  
69 mitogen-activated protein kinases (MAPKs), such as ERK1/2 and p38, which are involved in  
70 several biological processes, such as pituitary [5,6,7,8] and luteal hormone secretion [9],  
71 neuroendocrine function and the regulation of cell proliferation and migration [4,10].

72

73 The expression of Kiss-1R and Kp has been demonstrated in a variety of human tissues. They are  
74 mainly expressed in the placenta, with much lower expression levels in ovary, testis, liver, pancreas,  
75 small intestine and brain e.g., hypothalamus and pituitary gland [4,6,7,11,12,13]. The Kiss-1 gene  
76 was originally identified as a human metastasis suppressor gene; however, Kps have emerged as  
77 important gatekeepers of key aspects of reproductive maturation and function, from sexual  
78 differentiation of the brain and puberty onset to adult regulation of gonadotropin secretion and the

79 metabolic control of fertility [4,14]. Kisspeptins and Kiss-1R play fundamental roles in pregnancy  
80 establishment. Saadeldin et al. [15], in fact reported that porcine oocyte maturation and oocyte  
81 maternal gene expression (MOS, GDF9 and BMP15), blastocyst formation rate, blastocyst hatching  
82 and blastocyst total cell count increased significantly when oocytes were cultured in medium  
83 containing Kp [15]. Moreover, the highest expression of Kps and Kiss-1R in trophoblastic cells  
84 during the first trimester in humans and, at embryonic day 12.5 in rats, coincides with the peak of  
85 trophoblastic invasion which is critical for pregnancy establishment. In addition, the fact that Kp-10  
86 inhibits the migration and invasion of trophoblastic cells provides further support to the notion that  
87 Kps may play a role in controlling trophoblastic invasion and migration during human [3,16, 17]  
88 and rat placentation [18]. The invasion of extravillous trophoblasts into the uterine wall is of crucial  
89 importance for placental and fetal development, and its deregulation has been implicated in a wide  
90 spectrum of abnormal pregnancies [3]. Reduced expression of Kiss-1 and increased expression of  
91 Kiss-1R at both mRNA and protein levels have been involved in the pathogenesis of pre-eclampsia  
92 [14]. In humans, decreased Kp expression in trophoblasts is associated with recurrent pregnancy  
93 loss and Kp has been hypothesized as engaging the regulation of decidual natural killer cell  
94 infiltration [19]. To date, no studies have been performed on the role of Kps on placental cell  
95 proliferation and progesterone (P<sub>4</sub>) secretion.

96

97 In the present study, we evaluated the effects of Kp-10 addition in culture media on proliferation  
98 and P<sub>4</sub> secretion of bovine placenta cotyledon cell lines isolated from fetuses in the first trimester of  
99 pregnancy. The involvement of Kiss-1R in observed cell responses was investigated by analyzing  
100 the relative abundance variations of Kiss-1R mRNA by quantitative Real Time RT-PCR.

101

## 102 **2. Materials and methods**

103

### 104 *2.1. Chemicals and Drugs*

105 All chemicals were purchased from Sigma-Aldrich (Milan Italy) unless otherwise indicated.

106

## 107 *2.2. Tissue collection*

108 Uteri from pregnant cows in the 2<sup>nd</sup> or 3<sup>rd</sup> month of gestation were retrieved at a local  
109 slaughterhouse within 20 min after slaughter, kept at 4°C in phosphate-buffered saline (PBS) with  
110 penicillin (100 U/mL), streptomycin (100 mg/mL), and amphotericin (0.25 mg/mL) solution during  
111 transport to the cell culture laboratory and processed immediately. Prior to incision of the uterine  
112 wall, the site was cleaned with PBS supplemented with antibiotics and antimycotic. Placentomes  
113 were put in a sterile beaker and immediately placed under the laminar flow hood for subsequent  
114 processing. They were manually separated into fetal cotyledon and maternal caruncle. Depending  
115 on the size of each tissue, a 1–2 cm<sup>3</sup> piece of fetal cotyledon was dissected into small pieces (2x2  
116 mm) before enzymatic digestion.

117

## 118 *2.3. Isolation of cells from cotyledons*

119 Fetal cotyledon fragments were collected in falcon tubes and washed two times in PBS  
120 supplemented with antibiotics and antimycotic at 300 g for 5 min, and then digested with 0.75  
121 mg/ml collagenase in Low Glucose Dulbecco's Modified Eagle's Medium (LG-DMEM) at 37 °C  
122 for 4h, until each fragment was completely digested. After incubation, collagenase activity was  
123 stopped with LG-DMEM supplemented with 10% Fetal Calf Serum (FCS), the suspension was  
124 filtered through 80µm filters (Millipore, Milan, Italy), centrifuged at 300 g for 10 min and cells  
125 were collected from the pellet.

126

## 127 *2.4. Cell culture and expansion*

128 Cells were cultured in expansion medium consisting of LG-DMEM supplemented with 10% FCS,  
129 100 U/ml penicillin, 100 mg/ml streptomycin, 0.25 mg/ml amphoterycin solution, 2 mM L-  
130 glutamine and 10 ng/ml Epidermal Growth Factor (EGF), as reported in previous studies carried out

131 by our research unit of extra fetal tissues [20,21]. The number of viable cells was counted by the  
132 trypan blue dye exclusion test, using a Bürker chamber. For culture maintenance, cells were plated  
133 at a density up to  $1 \times 10^5$  cells/cm<sup>2</sup> and incubated at 38.5 °C in a humidified atmosphere (95%) with  
134 5% CO<sub>2</sub>. The medium was replaced for the first time after 72 h to remove non-adherent cells, and  
135 then changed either twice per week or according to the experimental design.

136

## 137 *2.5. Cell line characterization*

### 138 *2.5a Cell morphology*

139 The epithelial origin of primary placental cotyledonary cell lines was assessed following the criteria  
140 by Hambruch et al., 2010 [22]. Morphologically, these cells form monolayers composed of  
141 approximately polygonal cells, clearly distinguishable from maternal caruncular epithelial cells  
142 which displayed curl-like morphology within the colonies (Hambruch et al., 2010). For assessing  
143 the presence of binucleate cells,  $1 \times 10^4$  cells were seeded onto cover slips and cultured, in the  
144 conditions described above, until 50% confluence was reached. Cells were fixed with 2%  
145 paraformaldehyde in PBS for 10 min, stained with 2.5 µg/ml Hoechst 33258 in 3/1 glycerol/PBS  
146 solution and mounted onto slides in the same solution. Evaluation of cell morphology, by phase  
147 contrast microscopy, and nuclear chromatin, by epifluorescence microscopy, were performed under  
148 a Nikon Eclipse 600 (x 400 magnification) epifluorescence microscope. Stainings were carried out  
149 in triplicates.

150

### 151 *2.5b XY-chromosome specific PCR*

152 After thawing, cell lines underwent genomic DNA extraction using the Genelute genomic DNA kit  
153 (Sigma-Aldrich, Milan, Italy) following the protocol recommended by the manufacturers. The  
154 tissue was disrupted by liquid nitrogen and, before going through the extraction procedure,

155 incubated with proteinase K overnight at 54 °C, whereas cell samples were centrifuged and then  
156 incubated with lysis buffer supplemented with proteinase K for 10 min at 72 °C. As positive  
157 control, for the presence of Y-chromosome, genomic DNA extracted from a skin sample collected  
158 at a local slaughterhouse from a male calf was used. As Y-chromosome negative control, bovine  
159 genomic DNA isolated from ovarian surface epithelial cells was used. The sequence length  
160 polymorphisms between bovine amelogenin X and amelogenin Y genes (AMLX/Y) were used as  
161 markers for sexing the bovine fetal cells, as previously described in other bovine cell systems [23].  
162 The oligonucleotide sequences of used primers (Primm S.R.L., Milano, Italy) were (forward) 5'-  
163 CAGCCAAACCTCCCTCTGC-3' and (reverse) 5'-CCCGCTTGGTCT TGTCTGTTGC-3'. This  
164 primer set was designed to amplify a single fragment of 280 bp on the X-chromosome (female  
165 cells) and two fragments of 280 and 217 bp on X and Y chromosomes, respectively (male cells).  
166 Therefore, our sexing method was used to confirm the presence of male fetus cells in our samples.  
167 The PCR reaction mixture (50 µL/tube) consisted of 1.0 U Taq DNA polymerase, 5x Taq buffer,  
168 0.2 mM of each dNTPs, 1 µM specific primers, and 100 ng template DNA. The PCR conditions  
169 consisted of an initial denaturing cycle at 94 °C for 5 min followed by 37 cycles of 94 °C for 45 s.,  
170 64.7 °C for 45 s., 72 °C for 30 s. and, after the last cycle, samples were kept at 72 °C for 5 min for  
171 the final extension, and then the PCR procedure was completed. The PCR products were analyzed  
172 by electrophoresis in 2 % (w/v) agarose gel and visualized under UV light after ethidium bromide  
173 staining. The analysis was carried out in triplicates.

174

## 175 *2.5c Gene expression study*

### 176 *2.5.c1 Total RNA extraction*

177 The RNeasy kit (QIAGEN, Milan, Italy) was used to extract total RNA. After thawing, cells were  
178 washed in PBS and then disrupted in Lysis Buffer RLT and homogenized at room temperature. A  
179 70% ethanol solution was then added to the cell lysate, thus creating conditions that promoted

180 selective binding of RNA to the RNeasy membrane. The lysate was transferred to an RNeasy spin  
181 column and centrifuged for 15 s at 10000 rpm. The column was washed twice with washing buffer  
182 and finally, the total RNA was eluted in 30  $\mu$ l of RNase-free water and stored at  $-80^{\circ}\text{C}$ . The total  
183 amount of RNA of each sample was measured with the BioPhotometer plus (Eppendorf, Milan,  
184 Italy).

185

### 186 *2.5.c2 Reverse Transcription PCR*

187 The High Capacity cDNA Reverse Transcription kit (LifeTechnologies, Monza, Italy) was used to  
188 convert RNA to cDNA. After thawing, 2  $\mu$ g of total RNA of each samples were added to 2  $\mu$ L 10X  
189 RT Buffer, 0.8  $\mu$ L 25X dNTP Mix, 2  $\mu$ L RT Random Primers, 1  $\mu$ L M-MLV Reverse  
190 Transcriptase, 1  $\mu$ L RNase Inhibitor, nuclease-free H<sub>2</sub>O for a total volume of 20  $\mu$ l and then mixed  
191 gently and centrifuged briefly. Reaction tubes were incubated at  $10^{\circ}\text{C}$  for 10 min, then at  $37^{\circ}\text{C}$  for  
192 120 min and finally at  $85^{\circ}\text{C}$  for 5 min. Specific bovine cDNAs of Cytokeratin (CK), Tight junction  
193 protein 2 (zona occludens 2, ZO-2) and Glyceraldehyde 3-phosphate dehydrogenase (GAPDH)  
194 were amplified by PCR, using the primers as shown in Table 1. PCR conditions were as follows: 95  
195  $^{\circ}\text{C}$  for 2 min, 35 cycles of denaturation at  $94^{\circ}\text{C}$  (45 s), annealing at  $60^{\circ}\text{C}$  for GAPDH and CK and  
196  $56^{\circ}\text{C}$  for ZO-2 (45 s) and extension at  $72^{\circ}\text{C}$  (45 s), final extension at  $72^{\circ}\text{C}$  for 5 min. PCR  
197 products were separated using a 2% agarose gel and visualized by ethidium bromide staining. The  
198 analysis was carried out in triplicates.

199

### 200 *2.6. Proliferation study*

201 The proliferation study was carried out on placenta cotyledon cell lines from fetuses in the first  
202 trimester of pregnancy. Gestational age was determined on the basis of fetal body length [24]. In  
203 order to calculate doubling time (DT, indicative of the proliferation rate), cells were seeded at a  
204 density of 1000 cell/cm<sup>2</sup> in expansion medium containing 0.01, 0.1, 1 or 10  $\mu$ M of human Kp-10  
205 (Sigma M2816). Cell cultures performed in the absence of Kp-10 were used as controls. Culture



206 passages were performed every 3 days and Kp-10 was added at each passage (P). Cells of each well  
207 were detached using 0.05% trypsin/0.02% EDTA in PBS and were counted, by dilution (1:1) in  
208 Trypan blue, with Bürker's chamber. The DT data were calculated by using the following formula:  
209  $CD = \ln(N_f / N_i) / \ln 2$  and  $DT = CT / CD$ , where DT is the cell-doubling time, CD is the cell-doubling  
210 number, and CT is the cell culture time. The proliferation rate was calculated from each passage,  
211 where  $N_f$  is the final number of cells and  $N_i$  the initial number of cells. The total number of live  
212 cells was obtained for each passage and DT values were calculated from P1 to P10.

213

### 214 *2.7. Placenta cells and culture media cryopreservation*

215 At the end of each passage, cells were divided into two parts, one of which was used for the  
216 proliferation study, as described above, and the other was suspended in a cryopreservation medium  
217 consisting of 40% LG-DMEM, 50% FCS and 10% Dimethyl sulfoxide (DMSO) and stored at -80  
218 °C, whereas the culture media for each passage and culture condition were removed and stored in  
219 cryotubes at -80 °C for P<sub>4</sub> radioimmunoassay. After thawing, cells were used for molecular  
220 analysis.

221

### 222 *2.8. Progesterone measurements*

223 Frozen culture media were lyophilized (freeze-dried) and then the P<sub>4</sub> levels were measured by the  
224 use of a chemiluminescent immunoassay kit (LKPG1, MEDICAL SYSTEM, Genoa, Italy). The  
225 assay detection range was 0.2–100 ng/mL.

226

### 227 *2.9. Real Time PCR*

228 Real Time PCR was performed using Real Time TaqMan® technology and analyzed on the  
229 automated StepOne System (Applied Biosystems, Monza, Italy). All TaqMan bovine primers and  
230 probes were purchased from Life Technologies. The Glyceraldehyde 3-phosphate dehydrogenase  
231 (GAPDH; Bt03210913\_g1) gene assay was inventoried by Life Technologies. For the specific

232 bovine Kiss-1R, primers and probes were properly designed, across exon-exon junction, using the  
233 Primer Express 3.0 software (Applied Biosystem) based on NCBI Bos Taurus available sequences  
234 or on mammal multi-aligned sequences. Kiss-1R Real Time PCR primers: forward primer: 5'-  
235 GCGGTCACGGACTTAACGTT -3'; reverse primer: 5'- CACCGAGACCTGCTGGATGT -3';  
236 Kiss-1R hybridization probe: -5'TGTGCTGCGTGCCCT-3'. The TaqMan® MGB (minor groove-  
237 binder) hybridization probe was labeled with a reporter dye (6-carboxy-fluorescein, FAM) on the 5'  
238 nucleotide and a quenching dye with NFQ (non fluorescent quencher) on the 3' nucleotide where  
239 MGB hyper-stabilized duplexes with complementary DNA. Samples were run in duplicates on  
240 Microamp fast optical 48-well reaction plate (Life Technologies) where twenty-microliter reactions  
241 for each well contained: 10 µL TaqMan gene expression Master Mix 2X (Life Technologies), 1 µL  
242 800 nM Primers and 250 nM Probe, 1-4 µL cDNA and RNase Free H<sub>2</sub>O. Cycle parameters were: 2  
243 min at 50°C, 10 min at 95°C followed by 40 cycles of 15 s at 95°C and 1 min at 60°C. Data were  
244 collected using the StepOne Software and relative quantification was performed using a  
245 comparative method after determining the Ct (threshold cycle) values for the reference endogenous  
246 control (GADPH) and the target gene in each sample set, according to the  $2^{-\Delta\Delta Ct}$  method, as  
247 described by the manufacturer. Changes in mRNA expression levels were calculated after  
248 normalization to GAPDH. The program calculates the  $\Delta Ct$  and the  $\Delta\Delta Ct$  with the formulas below:  
249  $\Delta Ct = Ct\_Mean(GAPDH) - Ct\_Mean(Kiss-1R)$ ;  $\Delta\Delta Ct = \Delta Ct - \Delta Ct\_Mean$ , so that the gene  
250 expression level =  $2^{-\Delta\Delta Ct}$ . Changes in gene expression were reported as percentage changes relative  
251 to controls.

252

## 253 2.10. Statistical analysis

254 Cell proliferation (doubling time) data were analyzed by using the General Linear Model procedure  
255 (SAS™ package v. 9.1, SAS Institute, Cary, NC) considering line, passage and Kp-10  
256 concentration as fixed effects. Least-Square means (LS-means) ± standard error (SE) are shown.  
257 Within-passage Kp-10 concentration effects on DT data were evaluated by one-way ANOVA

258 (GraphPad Prism Software 5.0) followed by Dunnett's multiple comparison post-hoc test and  
259 means  $\pm$  standard deviations (SD) are shown. Relative abundance of Kiss1R mRNA and P<sub>4</sub> levels  
260 were analyzed by the GLM procedure and LS means  $\pm$  SE are shown. Values with P<0.05 were  
261 considered significantly different.

262

### 263 **3. Results**

#### 264 *3.1. Bovine primary placenta cotyledon cell cultures*

265 The proliferation study was carried out on two placenta cell lines, namely line A, which was  
266 isolated from a fetus in the 2<sup>nd</sup> month of gestation (thus having a body length of 8 cm), and line B,  
267 which was isolated from a fetus in the 3<sup>rd</sup> month of pregnancy (with a body length of 12 cm). Upon  
268 *in vitro* culture, primary cotyledon cell lines formed monolayers composed of polygonal epithelial  
269 cells (Fig. 1 A, B). Morphologically, they were clearly distinguishable from maternal caruncular  
270 epithelial cells which display curl-like morphology as shown in previous studies [22].  
271 Characterisation of both cell lines via epifluorescence revealed the presence of larger binucleated  
272 cells (Fig. 1 C, D). The presence of a Y-chromosome specific sequence in both cell lines confirmed  
273 their fetal origin, as reported in previous studies [23, literature cited in ref. 23]. By gel  
274 electrophoresis of PCR products amplified from the two cell lines, two fragments of 280 and 217  
275 bp, indicating the presence of X- and Y-chromosome specific sequences, were observed (male  
276 pattern; Fig. 1, E). The expression study demonstrated that both cell lines express CK and ZO-2  
277 (Fig. 1, F), as revealed by gel electrophoresis of PCR products amplified from mRNA extracted  
278 from the two cell lines, showing specific bands for CK (152 bp) and ZO-2 (224 bp).

279

#### 280 *3.2. Effects of Kp-10 on in vitro proliferation of bovine placental cotyledon cell lines*

281 Significant "cell line" effect was observed, as line A showed significantly higher DT values than  
282 line B (LS means  $\pm$  SE: 1.69 $\pm$ 0.12 versus 1.32 $\pm$ 0.12 for lines A and B, respectively; GLM  
283 procedure: P<0.05). Significant "cell passage" effect was detected in both cell lines and the two cell

284 lines behaved differently. Line A showed a slower DT at P1 which then settled into a faster rate  
285 from P2 to P10 (GLM procedure: significance level ranging from  $P < 0.05$  to  $P < 0.001$ ; Fig. 2, panel  
286 A); whereas line B had a steady DT until P9 which then increased at P10 (GLM procedure:  
287 significance level ranging from  $P < 0.05$  to  $P < 0.001$ ; Fig. 2, panel B). As for the “within-line Kp-10  
288 concentration effect”, no effect of Kp-10 addition was found in either cell line at any tested  
289 concentration (Table 2, GLM procedure: not significant). Instead, by analyzing the “within-passage  
290 Kp-10 concentration effect”, it was found that, within each specific cell passage, the two cell lines  
291 responded differently to Kp-10 addition (One-way ANOVA followed by Dunnett’s post-hoc test).  
292 Specifically, in line A, a stimulatory effect of Kp-10 was found during early cell passages (P1 and  
293 P2 with significance level ranging from  $P < 0.05$  to  $P < 0.01$ ), followed by a switch to an inhibitory  
294 effect at P3 and from P5 to P10 (significance level ranging from  $P < 0.05$  to  $P < 0.001$ ; Fig. 2 panel  
295 A’). In line B, a stimulatory effect was observed during intermediate passages (P3, P4, P6, P7;  
296 significance level up to  $P < 0.01$ ; Fig. 2, panel B’) whereas no effects were observed in other  
297 passages. In both cell lines, cell viability did not change with passage number and was not affected  
298 by Kp-10 treatment (no significant differences, data not shown).

299

300

### 301 3.3. *Effects of Kp-10 on Kiss-1R mRNA relative abundance of bovine placental cotyledon cell lines*

302 Based on the results of the proliferation study, the involvement of Kiss-1R on observed cell  
303 response was analyzed at mRNA level by quantitative Real Time RT-PCR. In each cell line,  
304 relevant passages representative of the intermediate phase (P3 to P9) were selected for molecular  
305 analysis, due to their higher and more stable proliferation rate. Among them, those showing  
306 inhibition (P7 and P9 for cell line A), switching (P4 for cell line A) or stimulation (P3, P4 and P6  
307 for cell line B) of cell proliferation upon Kp-10 addition were used. In cell line A, no significant  
308 differences were found in Kiss1R mRNA relative abundance, upon Kp-10 addition, at any  
309 examined passage and concentration (Fig. 3, panel A; GLM procedure: not significant). In cell line  
310 B, the Kiss-1R transcript level significantly increased upon Kp-10 stimulation at P3 and P6 (Fig. 3,

311 panel B, GLM procedure). At P3, the relative abundance was doubled in cells treated with 0.01  $\mu\text{M}$   
312 Kp-10 and increased approximately 2.5-fold in cells treated with 10  $\mu\text{M}$  Kp-10 ( $P < 0.05$ ) and at P6,  
313 the Kiss-1R transcript level was approximately 6-fold higher in cells treated with 0.01  $\mu\text{M}$  Kp-10  
314 ( $P < 0.001$ ) and 3-fold higher in cells treated with 0.1 and 10  $\mu\text{M}$  Kp-10 ( $P < 0.01$ ; Fig. 3, panel B).

315

### 316 3.4. *Effect of Kp-10 on P<sub>4</sub> secretion in bovine placental cells*

317 The effects of kp-10 on P<sub>4</sub> secretion in *in vitro* cultured placenta cotyledon cell lines were analyzed  
318 in order to elucidate the role of Kp-10 on the secretion of this pregnancy supporting hormone. In  
319 preliminary tests performed in media of cells cultured in control conditions, P<sub>4</sub> levels were found to  
320 be below minimal detectable values (less than 0.20 ng/ml). For this reason, media were lyophilized  
321 and used for single determinations. Data from culture media collected at a representative early (P2),  
322 intermediate (P4) and late cell passage (P10) were averaged. In both cell lines, P<sub>4</sub> production was  
323 not affected by Kp-10 addition at any tested concentration and LS mean values ranged from 0 to 2.9  
324 ng/ml (GLM procedure: not significant; Fig. 4).

325

## 326 4. Discussion

327 The prime purpose of this study was to establish primary cultures of cell lines from bovine placental  
328 cotyledons, which may provide a valuable tool for the study of bovine feto-maternal interactions.  
329 These cell lines may be useful as *in vitro* models to elucidate the roles of factors involved in the  
330 regulation of bovine placenta development and function, such as the Kiss-1R/Kps system, and to  
331 establish new pregnancy maintenance strategies for practical application in bovine reproductive  
332 management. Moreover, the bovine placenta stands as an ideal model for the study of factors  
333 involved in human placenta development and function due to the similar nine-month pregnancy  
334 duration, comparable to that in humans, despite the known differences in placenta distribution  
335 (cotyledonary versus discoid) and structure (syndesmochorial versus hemochorial). In the present  
336 study, the isolation of bovine placental cotyledonary cell lines was confirmed by the epithelioid

337 polygonal morphology, presence of binucleate cells, presence of the Y-chromosome, and the  
338 simultaneous expression of cytokeratin and zonula occludens 2 as trophoblast-specific markers (Fig.  
339 1) [22]. To our knowledge, this is the first study reporting a proliferation study through 10  
340 consecutive passages of bovine placental cotyledon cell lines, which provides useful indications of  
341 their behavior in *in-vitro* cultures.

342  
343 After having successfully isolated and expanded epithelioid cotyledon cell lines, we decided to  
344 determine whether - in these cell lines - Kiss-1R activation, by means of its endogenous ligand Kp-  
345 10, could modulate their *in vitro* proliferative activity. Kiss-1R expression in human placenta  
346 [16,17,25] as well as in animal models [18,26] has been reported and a possible involvement of the  
347 Kiss-1R/Kps system in regulating trophoblast invasion and proliferation has been hypothesized  
348 [3,16,17,27]. In the present study, cell line effect and cell passage effect were observed, due to the  
349 use of primary cell lines isolated from bovine placenta of fetuses in the first trimester of pregnancy.  
350 In detail, line A showed high DT value at P1, followed by increased cell proliferation rate, maybe  
351 due to gradual adaptation of the cell line to *in vitro* culture conditions (Fig. 2, panel A), whereas cell  
352 line B showed a prolonged phase of high proliferation rate and inhibition of cell proliferation in late  
353 passages, perhaps due to loss of cell viability (Fig. 2, panel B). Taking into account these effects,  
354 the effects of Kp-10 concentration were first evaluated by grouping data from all passages, which  
355 showed no effects of Kp-10 addition at any tested concentration (Table 2). However, this analysis  
356 turned out to be very weak, as very low R-square values were found in both cell lines (R-square  
357 value = 0.004035 and 0.000971 for cell line A and B, respectively), indicating that the effects of  
358 Kp-10 are masked by greater effects related to cell line and cell passages. After excluding cell line  
359 and cell passage effects, we demonstrated using within-cell passage analysis that Kp-10 modulates  
360 *in vitro* proliferation in these cells. The two analyzed cell lines behaved differently upon Kp-10  
361 addition, with inhibition of cell proliferation prevailing in line A, isolated from embryos/fetuses  
362 during the 2<sup>nd</sup> month of gestation (Fig. 2, panel A') and stimulation of cell proliferation prevailing

363 in line B, isolated from embryos/fetuses at the 3<sup>rd</sup> month of pregnancy (Fig. 2, panel B'). To our  
364 knowledge, this is the first study in which Kp-10 has been reported as having a stimulatory effect on  
365 cell proliferation, as we observed in line B. In previous studies, analogously with its originally  
366 identified antimetastatic role [1], Kp was reported as having inhibitory effects in cell proliferation  
367 [28-32], although the biological mechanisms remains to be further elucidated. It has already been  
368 clarified that the activation of Kiss-1R leads to phosphorylation of different MAPKs, which might  
369 contribute to the antimetastatic and/or antiproliferative effects of kisspeptins (Kps, [28-30]).  
370 However, the subset of intracellular kinases activated upon kisspeptin stimulation appears to be, at  
371 least partially, dependent on the cellular context [4], and maybe on the cell type isolated from  
372 different tissues.

373

374 The Kp-10 effects observed on bovine primary placenta cotyledon cell lines were not dose-  
375 dependent. Previous studies refer to the variable effects of Kp-10 addition, depending on cell type,  
376 whether primary or established or embryonic or cancer cell line, and the nature of Kiss-1R  
377 expression, whether endogenous or artificially-induced. Kotani et al., 2001 [29] reported inhibition  
378 of cell proliferation in CHO-K1 cells transfected and expressing rat or human GPR54 (Kiss-1R) and  
379 treated with 1  $\mu$ M Kp-10. Cho et al., 2009 [30] reported no dose-dependent effect of Kp-10 tested at  
380 concentrations ranging from 1.0 to 100  $\mu$ M on cell proliferation of human umbilical vein  
381 endothelial cells (HUVEC) and prostate cancer cells (PC-3). Huma et al., 2013 [31] reported a dose-  
382 dependent inhibitory effect of Kp-10 on cell proliferation of Rhesus Monkey derived embryonic  
383 stem cells (R366.4) at concentrations ranging from 0.1 to 100 nM. Ziegler et al., 2013 [32] reported  
384 no effects in breast cancer cell lines (such as MDA-MB-231; MDA-MB-435s; HCC 1806 and  
385 MCF-7) expressing the receptor endogenously but significant inhibition of cell proliferation in  
386 transfected neuronal cells overexpressing Kiss-1R. To the best of our knowledge, no data have been  
387 reported to date on the effects of Kp-10 addition on cell proliferation of primary placenta cell lines.  
388 Primary cell lines isolated from placenta cotyledons recovered in the first trimester of pregnancy in

389 the bovine model can be considered as a physiological model for studying the role of the Kiss1-  
390 R/Kp-10 system in trophoblastic growth and invasion at the early stages of pregnancy other than  
391 immortalized trophoblastic cell lines, as also reported in a recent study by Francis et al., 2014 on  
392 human trophoblastic cells isolated in the first trimester of pregnancy [33].

393

394 In order to identify the reasons for different *in vitro* effects of Kp-10 on the proliferative activity of  
395 the two cell lines, we considered it important to assess whether Kp-10 stimulation could differently  
396 modify Kiss-1R expression. Thus, we investigated the effects of Kp-10 on the relative abundance of  
397 mRNA in the two cell lines. Specifically, in cell line A, in which a prevailing inhibition of cell  
398 proliferation was found upon Kp-10 stimulation, the Kiss-1R transcript level did not change in Kp-  
399 10 treated cells compared with control conditions. In cell line B, in which a stimulatory effect on  
400 cell proliferation was observed upon Kp-10 addition, the relative abundance of Kiss-1R  
401 significantly increased in Kp-10 treated cells compared with controls. At the present time, these  
402 findings can only be interpreted in the light of results from previous studies. In human [3,16,17,30]  
403 and rat placenta [18], it has been reported that Kiss-1R mRNA expression increases significantly  
404 during pregnancy, with distinct spatial and temporal variations. Indeed, the two cell lines analyzed  
405 came from embryos/fetuses at the 2<sup>nd</sup> and the 3<sup>rd</sup> month of gestation, respectively. The recent study  
406 by Ziegler et al., 2013 [33], evidencing a connection between the anti-proliferative effect of Kp-10  
407 and Kiss-1R overexpression, suggests possible correlations between Kiss-1R expression status in  
408 the bovine primary placenta cotyledon cell lines used in the present study and observed proliferative  
409 and molecular cell responses. Further studies are needed to verify this hypothesis. The modulation  
410 of the Kiss-1R mRNA transcription in bovine cotyledons raises the hypothesis that the Kiss-1R/Kps  
411 system plays a pivotal role in embryo/placenta development and reinforces its suggested role in the  
412 regulation of trophoblast proliferation. Although Kp is already known to play a key role in the first  
413 trimester of pregnancy in humans [3], to our knowledge, detailed information about cell  
414 proliferation and related underlying mechanisms during the first trimester of bovine placenta



415 formation have not been reported to date. Our data lead us to hypothesize that, in line B, kp-10  
416 addition significantly up-regulated kiss-1R expression and, perhaps consequently, increased cell  
417 proliferation whereas, in line A, kp-10 had no effect on the relative abundance of Kiss-1R mRNA.  
418 Possible explanations for Kiss-1 R involvement in the observed inhibitory effect on cell  
419 proliferation in line A could only be hypothesized as being related to: 1) receptor desensitization,  
420 even not associated with its mRNA transcription down-regulation; 2) different signaling pathways  
421 elicited by the ligand, such as transactivation of other receptors, as reported by Roseweir et al., 2012  
422 [17]; 3) different functional status of Kiss-1R in placental cells recovered at early gestational age.  
423 The qRT-PCR analysis performed in this study, using bovine primary placental cotyledonary cell  
424 lines, is of particular interest because the results may show more precisely what would happen in  
425 native tissue, compared with the use of immortalized cells which have been shown to have a  
426 differential gene expression to primary cells brought about by the immortalization process [17].

427

428 As a further approach to elucidate the role of the Kiss-1R/Kps system on bovine cotyledon placenta  
429 cell lines, we investigated the effects of Kp-10 on the P<sub>4</sub> secretory activity of these cells. Although  
430 considerable efforts have been addressed to unravelling the major role of Kps as regulatory factors  
431 for the neurons producing gonadotropin-releasing hormone, which stimulates pituitary gonadotropin  
432 secretion [4,6,10], the isolation and identification by mass spectrometry of Kps from human  
433 placenta [4] opens the way for investigating their effects on P<sub>4</sub> secretion. Previous studies were  
434 reported on the effects of kp-10 on P<sub>4</sub> secretion in ovarian cell systems involved in P<sub>4</sub> production. In  
435 chicken *in vitro* cultured granulosa cells, Kp-10 was found to stimulate P<sub>4</sub> secretion in a dose-  
436 dependent manner and depending on follicle developmental stage [34]. In rat *in vitro* cultured luteal  
437 cell lines, kp-10 was found to stimulate P<sub>4</sub> secretion via the ERK1/2 mitogen-activated protein  
438 kinase signalling pathway and by increasing the synthesis of three key steroidogenic enzymes, such  
439 as StAR, CYP11A and 3-beta-HSD [9]. To our knowledge, no studies have been reported to date on  
440 the effects of Kp-10 on P<sub>4</sub> secretion of placenta cell lines in any species and at any stage of

441 pregnancy and this is the first study reporting the effects of Kp-10 addition on P<sub>4</sub> secretion in  
442 bovine primary placenta cotyledon cell lines. The ability of fetal cotyledon to secrete P<sub>4</sub> is  
443 documented in early as well as in late pregnancy [35] even though P<sub>4</sub> concentration in cotyledons  
444 increased sharply in later stages from the 7<sup>th</sup> to the 9<sup>th</sup> month [36]. In the present study, in  
445 supernatants of culture media of both cell lines, P<sub>4</sub> was found at basal levels, ranging from 0 to 2.9  
446 ng/ml, but its secretion was not influenced by Kp-10 addition. The P<sub>4</sub> values found in the present  
447 study can be compared with those reported in previous studies in other P<sub>4</sub> cell systems after Kp-10  
448 stimulation. Xiao et al., 2011 [34] in chicken granulosa cell lines found values within 0.8 ng/ml or  
449 within 8.5 ng/ml, depending on follicle size after stimulation with Kp-10 at concentrations ranging  
450 from 10 nM to 1 μM. On investigating rat luteal cells, Peng et al., 2013 [9] found a higher range of  
451 P<sub>4</sub> values (ranging from 45 to 80 ng/ml) after stimulating with Kp-10 at concentrations comprised  
452 between 40 to 160 nM. Our data lead us to conclude that the Kiss1R/Kps system may not be  
453 involved in modulating P<sub>4</sub> synthesis/secretion in bovine placenta cotyledon cell lines isolated in  
454 early gestational age. Rather, at this stage, it could be involved in initiating the signalling pathways  
455 leading to P<sub>4</sub> synthesis which could become more consistent/relevant in later stages of pregnancy.  
456 Further studies are needed to elucidate this hypothesis.

457  
458 Taken together, these results lead to the conclusion that the Kiss-1R/Kps system could play a role in  
459 the fine developmental regulation of bovine placental cotyledonal cell proliferation in the first  
460 trimester of pregnancy, but it may not yet be competent enough to activate P<sub>4</sub> secretion. In the cell  
461 line isolated from an embryo/fetus at the 2<sup>nd</sup> month of pregnancy, inhibition of cell proliferation  
462 matching with non inducible Kiss-1R gene expression, as revealed by not significant variation of  
463 Kiss-1R expression was observed; whereas in the cell line isolated from an embryo/fetus at the 3<sup>rd</sup>  
464 month of pregnancy, stimulation of cell proliferation associated with inducible gene expression, as  
465 revealed by Kiss-1R up-regulation, was found. This study contributes to elucidate the role exerted  
466 by the Kiss-1R/Kps system in bovine placenta development and function. Primary cultures of

467 epithelial cell lines from bovine placental cotyledons provide a valuable tool for the study of the  
468 role of the Kiss-1R/Kps system in feto-maternal interactions in bovines and could be used in  
469 translational research programs aimed at human reproductive medicine.

470

#### 471 **AUTHOR'S ROLES**

472 RLS, NAM, AR and MED designed the study. NAM conducted most of the experimental  
473 procedures used in the study, i.e. the kinetic/proliferation, the expression study and data analysis  
474 and drafted the manuscript. AR performed tissue selection and setting up and P<sub>4</sub> determinations.  
475 NAM and FP performed data analysis and statistics. MED, FP and RLS supervised experiments and  
476 critically revised the manuscript. All authors have read and approved the manuscript.

477

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483

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485

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619  
620 **Figure legends**

621  
622 **Figure 1.** Characterization of primary epithelial cell lines isolated from bovine placental cotyledons  
623 of embryos/fetuses at the first trimester of pregnancy. Photomicrographs of cell lines exhibiting



624 polygonal epithelial morphology by phase contrast microscopy (A, B for cell line A and B,  
625 respectively; Scale bars: 10 $\mu$ m) and presence of larger binucleated cells (white arrows) by  
626 epifluorescence microscopy (C, D for cell line A and B, respectively; Scale bars: 20 $\mu$ m). Presence  
627 of a Y-chromosome specific sequence confirming the fetal origin of cell lines: gel electrophoresis of  
628 PCR products amplified from genomic DNA isolated from the two cell lines, A and B (E). In lanes  
629 1 and 2, corresponding to cell lines A and B, respectively, two fragments of 280 and 217 bp are  
630 shown (male pattern). A single fragment of 280bp is evident in the Y-chromosome negative control  
631 (female pattern, lane 3) whereas two fragments are shown in the Y-chromosome positive control  
632 (lane 4). Lane M: 100bp markers. CK and ZO-2 expression study: gel electrophoresis of PCR  
633 products amplified from mRNA extracted from the two cell lines, A (F, lane 1, 2 and 3) and B (F,  
634 lane 4, 5 and 6). Bovine cotyledonary cells show specific bands for CK (152 bp, lane 2 and lane 5)  
635 and ZO-2 (224 bp, lane 3 and lane 6). GAPDH (197 bp, lane 1 and 4) was used as a housekeeping  
636 gene. Lane 7 shows the negative PCR control. Lane M: 100bp markers.

637

638 **Figure 2.** Effects of Kp-10 on cell proliferation (Doubling Time, DT) of primary epithelial cell  
639 lines isolated from bovine placental cotyledons. Cell passage effect (panel A and panel B for cell  
640 line A and B, respectively). Results are Least Square means  $\pm$  SE of 20 data points (4 data points  
641 for 5 conditions; GLM procedure). Line A (panel A) shows a higher DT at P1 compared with all  
642 other passages and then a faster proliferation rate from P2 to P10; line B (panel B) has a steady DT  
643 until P9, which then significantly increases at P10. In panels A and B, each passage is compared  
644 with all other passages; each symbol indicates a specific passage and crescent symbol numbers  
645 indicate increasing levels of significance (i.e. P1: \*, \* P<0.05; \*\*P<0.01;\*\*\*P<0.001). Within-  
646 passage Kp-10 concentration effect (panel A' and panel B' for cell line A and B, respectively):  
647 results are means  $\pm$  SD of 4 data points analyzed by one-way ANOVA followed by Dunnett's post-  
648 hoc test. In line A (panel A'), stimulatory effect of Kp-10 can be seen at P1 and P2, followed by a  
649 switch to inhibitory effect at P3 and from P5 to P10; in Line B (panel B'), stimulatory effect is

650 visible at P3, P4, P6 and P7 whereas no effects can be seen in other passages. In panel A' and B',  
651 bars with different letters indicate statistically significant differences: a,b P<0.05; a,c P<0.01; a,d  
652 P<0.001.

653

654 **Figure 3.** Effects of Kp-10 on Kiss-1R mRNA relative abundance of primary epithelial cell culture  
655 isolated from bovine placental cotyledons analyzed by quantitative Real Time RT-PCR. Data were  
656 analyzed using the General Linear Model procedure (see Materials and Methods). In cell line A  
657 (panel A), no statistical significance on Kiss-1R expression levels, upon Kp-10 addition, was found  
658 in cells at any examined passage. In cell line B (panel B), Kiss-1R transcription was significantly  
659 up-regulated upon Kp-10 stimulation in P3 and P6. For each sample, data (Least Square means  $\pm$   
660 SE of two independent experiments in duplicate, Kiss-1R average Ct) were normalized relatively to  
661 the abundance of GAPDH mRNA (endogenous control) and normalized values were compared  
662 among groups. Bars with different letters indicate statistically significant differences: a,b P<0.05;  
663 a,c P<0.01; a,d P<0.001.

664

665 **Figure 4.** Within-line Kp-10 concentration effects on progesterone secretion of primary epithelial  
666 cell culture isolated from bovine placental cotyledons. Results are Least Square means  $\pm$  SE of 3  
667 data points (GLM procedure). Values from an early (P2), an intermediate (P4) and a late passage  
668 (P10) were averaged. In both cell lines, Kp-10 had no effect on P<sub>4</sub> secretion at any tested  
669 concentration.

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677 Table 1. Sequence of primers used for PCR

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Name	Accession number		Sequence	Product length
CK	NM_001033610.1	Forward: Reverse:	5'TGGAAGGGCTGACTGATGAG 3' 5'CTTGACCTCAGCGATGATGC 3'	152 pb
ZO-2	NM_001102482.1	Forward: Reverse:	5'GGATATTTGTGGCCGGCATT 3' 5'ATGAATCCCCTCTGCCACAA 3'	224 pb
GAPDH	NM_001034034	Forward: Reverse:	5'GTCTTCACTACCATGGAGAAGG- 3' 5'TCATGGATGACCTTGGCCAG 3'	197 pb

679

680 Table 2. Within-line Kp-10 concentration effect on cell proliferation of primary epithelial  
681 cell lines from bovine placental cotyledons, expressed as DT values.

682

Cell line	Kp-10 concentration ( $\mu$ M)	Least Square (LS) means $\pm$ standard error (SE)
line A	0 (CTRL)	1.37 $\pm$ 0.12
	0.01	1.30 $\pm$ 0.11
	0.1	1.30 $\pm$ 0.12
	1	1.33 $\pm$ 0.11
	10	1.23 $\pm$ 0.12
line B	0 (CTRL)	1.31 $\pm$ 0.11
	0.01	1.29 $\pm$ 0.11
	0.1	1.34 $\pm$ 0.11
	1	1.34 $\pm$ 0.11
	10	1.35 $\pm$ 0.11

683

684 For each cell line and Kp-10 concentration, data are LS means  $\pm$  SE of DT values from 40 data points (4 data  
685 points for 10 passages). The overall F test of the model is not significant ( $F = 0.19$ ;  $p = 0.9427$ ), indicating that  
686 the model containing only within-line Kp-10 concentration effect does not accounts for a significant amount of  
687 the variation in DT.

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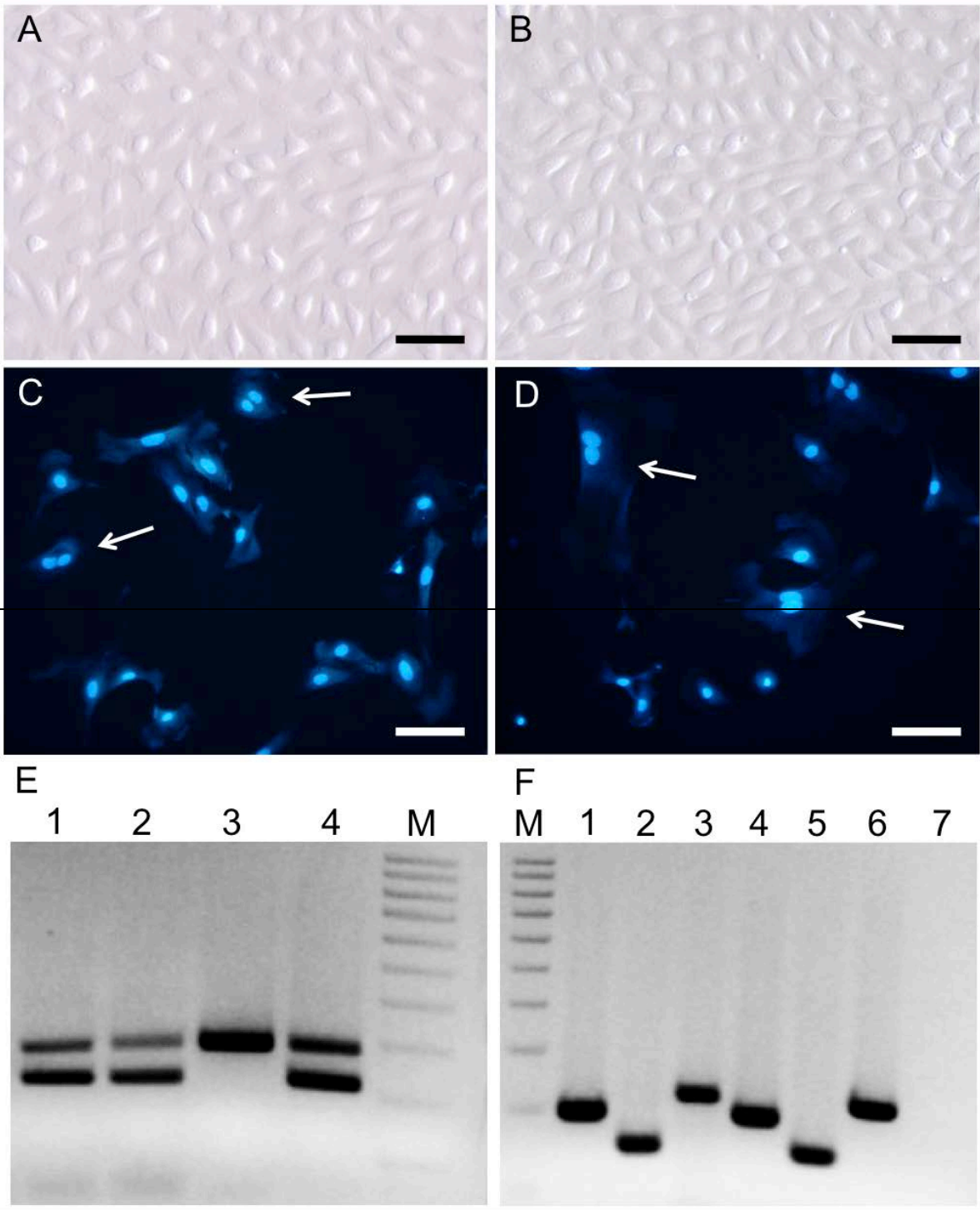
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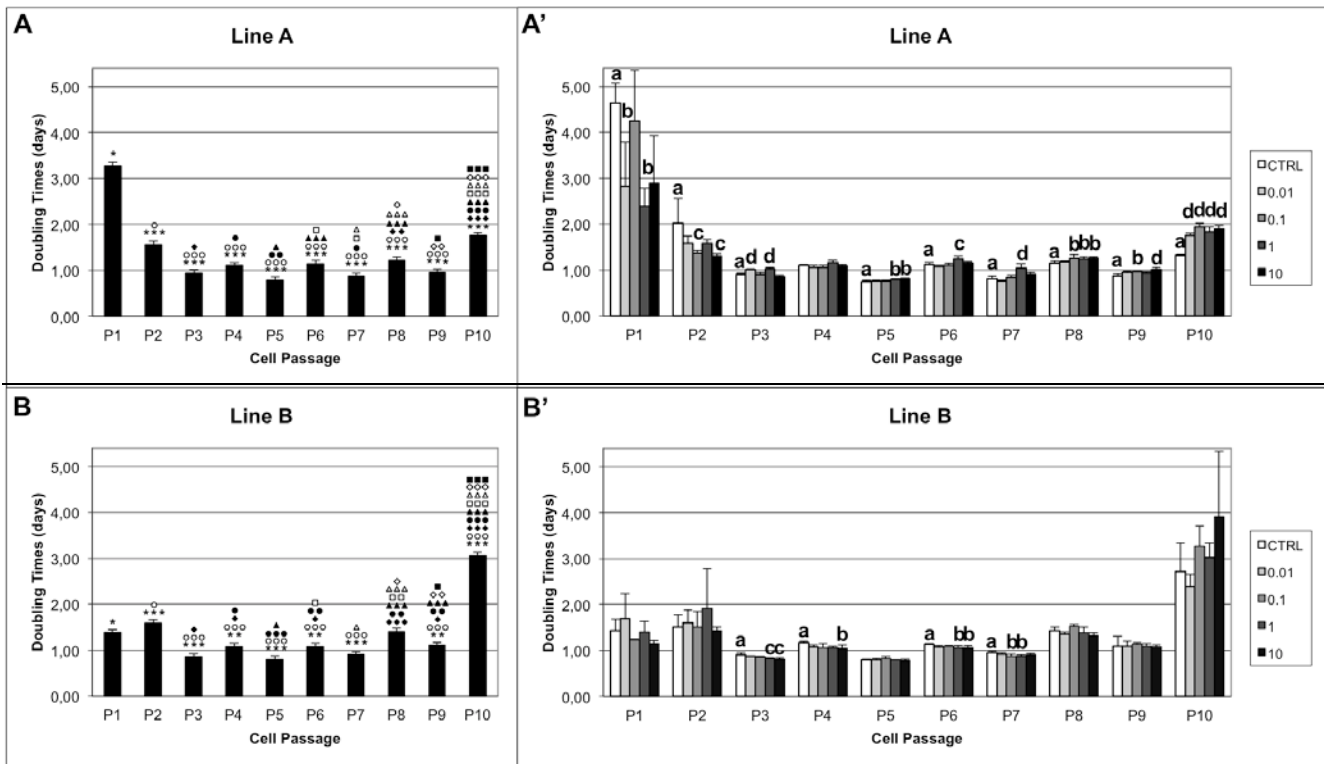
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715 Figure 2



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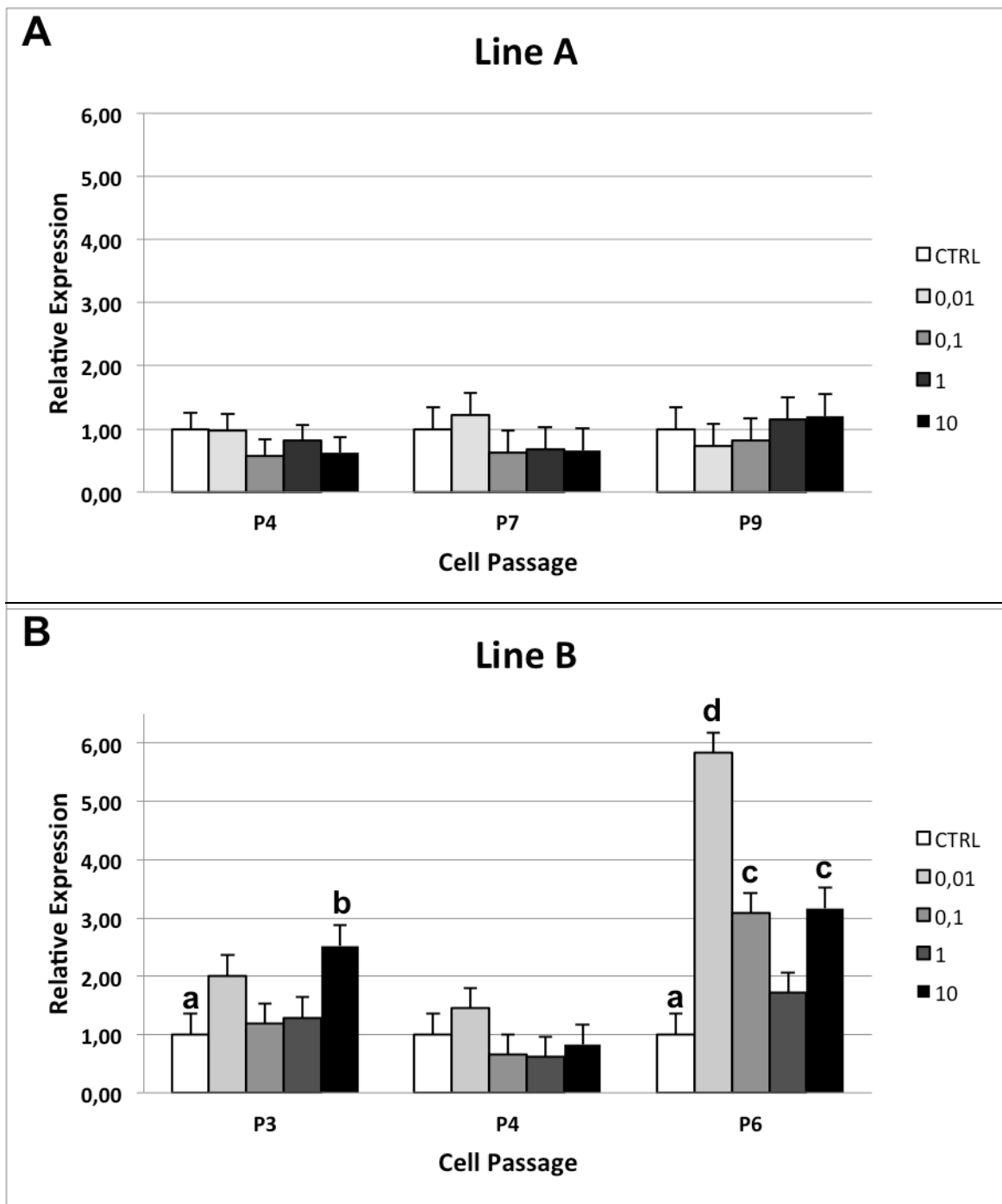
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731 Figure 3



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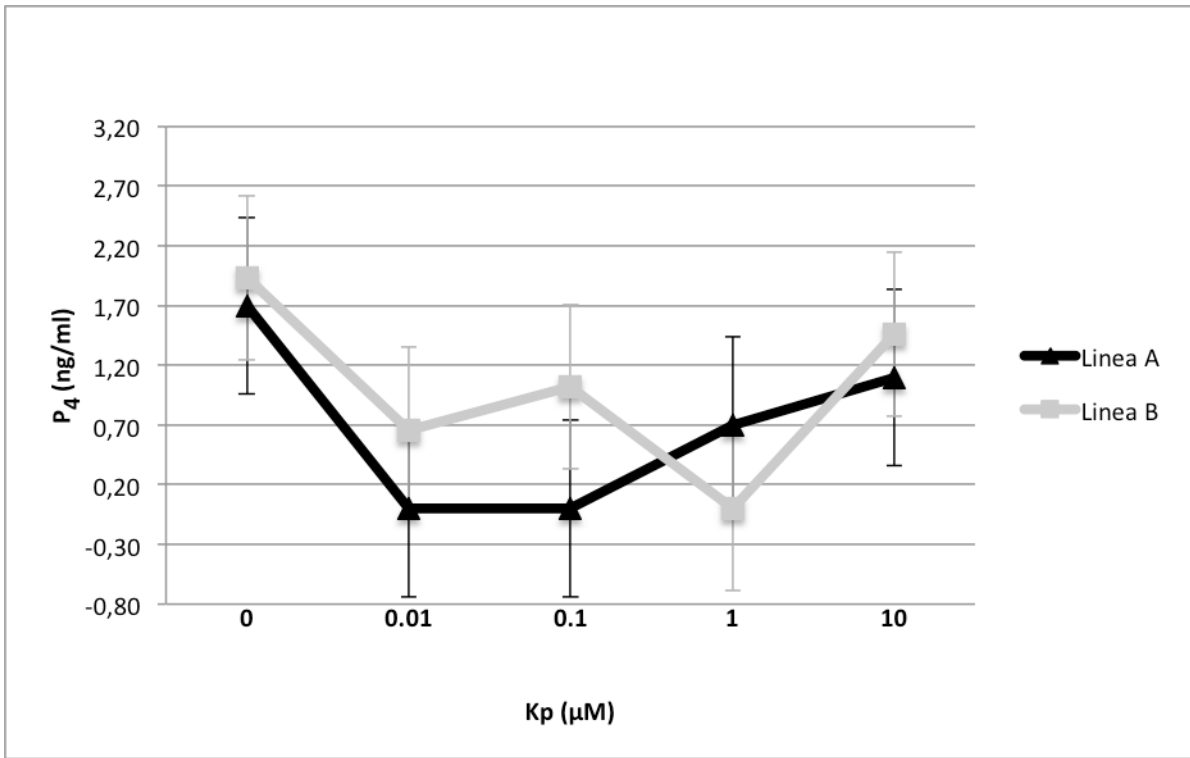
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738 Figure 4



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740 Supplementary file 1:  
 741 Within-passage Kp-10 concentration effect on cell proliferation of primary epithelial cell line from bovine placental cotyledons.

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 743 Line A

Cell passage	Kp-10 concentration ( $\mu\text{M}$ )				
	0 (CTRL)	0.01	0.1	1	10
P1	4.64 $\pm$ 0.43 <b>a</b>	2.82 $\pm$ 0.97 <b>b</b>	4.25 $\pm$ 1.11	2.39 $\pm$ 0.39 <b>b</b>	2.89 $\pm$ 1.03
P2	2.02 $\pm$ 0.55 <b>a</b>	1.59 $\pm$ 0.15	1.38 $\pm$ 0.05 <b>c</b>	1.59 $\pm$ 0.07	1.31 $\pm$ 0.04 <b>c</b>
P3	0.91 $\pm$ 0.02 <b>a</b>	1.01 $\pm$ 0.02 <b>d</b>	0.91 $\pm$ 0.05	1.03 $\pm$ 0.02 <b>d</b>	0.86 $\pm$ 0.02
P4	1.11 $\pm$ 0.02	1.06 $\pm$ 0.04	1.06 $\pm$ 0.03	1.16 $\pm$ 0.05	1.09 $\pm$ 0.03
P5	0.75 $\pm$ 0.03 <b>a</b>	0.77 $\pm$ 0.01	0.77 $\pm$ 0.02	0.80 $\pm$ 0.01 <b>b</b>	0.80 $\pm$ 0.03 <b>b</b>
P6	1.12 $\pm$ 0.05 <b>a</b>	1.08 $\pm$ 0.02	1.11 $\pm$ 0.04	1.24 $\pm$ 0.06 <b>c</b>	1.15 $\pm$ 0.03
P7	0.82 $\pm$ 0.06 <b>a</b>	0.77 $\pm$ 0.01	0.83 $\pm$ 0.05	1.04 $\pm$ 0.10 <b>d</b>	0.90 $\pm$ 0.04
P8	1.15 $\pm$ 0.05 <b>a</b>	1.18 $\pm$ 0.03	1.26 $\pm$ 0.07 <b>b</b>	1.25 $\pm$ 0.05 <b>b</b>	1.26 $\pm$ 0.03 <b>b</b>
P9	0.88 $\pm$ 0.03 <b>a</b>	0.95 $\pm$ 0.04	0.97 $\pm$ 0.01 <b>b</b>	0.94 $\pm$ 0.03	1.00 $\pm$ 0.05 <b>d</b>
P10	1.33 $\pm$ 0.02 <b>a</b>	1.76 $\pm$ 0.04 <b>d</b>	1.94 $\pm$ 0.07 <b>d</b>	1.83 $\pm$ 0.11 <b>d</b>	1.89 $\pm$ 0.08 <b>d</b>

744  
 745 Line B

Cell passage	Kp-10 concentration ( $\mu\text{M}$ )				
	0 (CTRL)	0.01	0.1	1	10
P1	1.43 $\pm$ 0.26	1.70 $\pm$ 0.54	1.22 $\pm$ 0.03	1.40 $\pm$ 0.23	1.15 $\pm$ 0.07
P2	1.51 $\pm$ 0.27	1.60 $\pm$ 0.27	1.51 $\pm$ 0.34	1.91 $\pm$ 0.87	1.43 $\pm$ 0.08
P3	0.90 $\pm$ 0.04 <b>a</b>	0.87 $\pm$ 0.00	0.85 $\pm$ 0.02	0.82 $\pm$ 0.02 <b>c</b>	0.81 $\pm$ 0.03 <b>c</b>
P4	1.16 $\pm$ 0.05 <b>a</b>	1.08 $\pm$ 0.04	1.07 $\pm$ 0.08	1.06 $\pm$ 0.04	1.04 $\pm$ 0.08 <b>b</b>
P5	0.80 $\pm$ 0.01	0.80 $\pm$ 0.04	0.82 $\pm$ 0.04	0.79 $\pm$ 0.01	0.78 $\pm$ 0.02
P6	1.14 $\pm$ 0.02 <b>a</b>	1.08 $\pm$ 0.03	1.08 $\pm$ 0.02	1.06 $\pm$ 0.04 <b>b</b>	1.06 $\pm$ 0.05 <b>b</b>
P7	0.95 $\pm$ 0.03 <b>a</b>	0.92 $\pm$ 0.03	0.87 $\pm$ 0.05 <b>b</b>	0.88 $\pm$ 0.03 <b>b</b>	0.90 $\pm$ 0.04
P8	1.43 $\pm$ 0.07	1.36 $\pm$ 0.04	1.52 $\pm$ 0.05	1.39 $\pm$ 0.12	1.32 $\pm$ 0.07
P9	1.09 $\pm$ 0.22	1.10 $\pm$ 0.11	1.14 $\pm$ 0.03	1.09 $\pm$ 0.07	1.08 $\pm$ 0.04
P10	2.72 $\pm$ 0.62	2.39 $\pm$ 0.26	3.27 $\pm$ 0.44	3.03 $\pm$ 0.31	3.92 $\pm$ 1.42

746 Data are means  $\pm$  standard deviations. One-way ANOVA followed by Dunnett's post hoc test.  
 747 For Kp-10 concentration effect: comparisons treated versus CTRL (between columns); a,b P<0.05; a,c P<0.01; a,d P<0.001.  
 748 No passage effect (between lines comparisons) is represented in this table, as it is shown in Figure 2.