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**Disturbed spermatogenic signaling in pituitary adenylate cyclase activating polypeptide deficient mice**

Running title: Spermatogenesis in PACAP deficient mice

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46

47 **Abstract**

48 PACAP is a neuropeptide with diverse functions in various organs, including reproductive  
49 system. It is present in the testis in high concentrations, and in addition to the stage-specific  
50 expression within the seminiferous tubules, PACAP affects spermatogenesis and the  
51 functions of Leydig and Sertoli cells. Mice lacking endogenous PACAP show reduced  
52 fertility, but the possibility of abnormalities in spermatogenic signaling has not yet been  
53 investigated. Therefore, we performed a detailed morphological analysis of spermatozoa,  
54 sperm motility, and investigated signaling pathways that play a role during spermatogenesis  
55 in knockout mice. No significant alterations were found in testicular morphology or motility  
56 of sperm in homo- and heterozygous PACAP deficient mice in spite of the moderately  
57 increased number of severely damaged sperms. However, we found robust changes in mRNA  
58 and/or protein expression of several factors that play an important role in spermatogenesis.  
59 Protein kinaseA expression was markedly reduced, while downstream phospho-ERK and p38  
60 were elevated in knockout animals. Expression of major transcription factors, such as Sox9  
61 and phospho-Sox9 was decreased, while that of Sox10, as a redundant factor, was increased  
62 in PACAP deficient mice. The reduced phospho-Sox9 expression was partly due to increased  
63 expression and activity of phosphatase PP2A in knockout mice. Targets of Sox transcription  
64 factors, such as Collagen type IV was reduced in knockout mice. In summary, our results  
65 show that lack of PACAP leads to disturbed signaling in spermatogenesis, which could be a  
66 factor responsible for reduced fertility in PACAP knockout mice, and further support the role  
67 of PACAP in reproduction.

68

69 **Keywords:** spermatogenesis, fertility, neuropeptide, neurotrophic factor, Sox transcription  
70 factors

71

72 **Introduction**

73

74 Pituitary adenylate cyclase activating polypeptide (PACAP) was originally isolated as a  
75 hypothalamic neuropeptide that stimulates adenylate cyclase activity, and thus, the release of  
76 several hormones in the pituitary (Miyata et al. 1989). PACAP plays diverse roles in the  
77 endocrine system, including the gonadal axis (Bardosi et al. 2016, Counis et al. 2007, Egri et  
78 al. 2016, Vaudry et al. 2009, Koves et al. 2014, 2016). The modulatory effects on

79 gonadotropin secretion suggest a role for PACAP in reproduction (Apostolakis et al. 2005,  
80 Kanasaki et al. 2016, Sherwood et al. 2000). An important discovery regarding male  
81 reproductive system came soon after the isolation of PACAP, showing that the peptide can be  
82 found in the testis in high concentrations, similar in range to those in the CNS (Arimura et al.  
83 1991). The high levels of PACAP in the testis suggest that it plays an important role in  
84 spermatogenesis and/or sperm functions (Daniel et al. 2000, Li and Arimura 2003, Li et al.  
85 2000). Indeed, PACAP has been shown to influence the development and functioning of  
86 spermatozoa (Gozes et al. 1998, Li et al. 2004, Koh et al. 2003, 2006). PACAP expression  
87 was found mainly in immature and, to a smaller extent, in mature sperms (Hannibal and  
88 Fahrenkrug 1995). PACAP regulates the activity of the supporting Sertoli cells, and acts at  
89 the hormonal level influencing the synthesis of testosterone by Leydig cells. Epididymis-  
90 derived PACAP may also influence the final stages of spermiogenesis (Csaba et al. 1997, El-  
91 Gehani et al. 2000, Heindel et al. 1992, Lacombe et al. 2006, Li et al. 2004, Matsumoto et al.  
92 2008, Rossato et al. 1997, Shioda et al. 1994, Tanii et al. 2011, Yanaihara et al. 1998, West et  
93 al. 1995).

94 In, mature sperms of golden hamster, it has been found that the addition of PACAP<sub>7-27</sub>  
95 hybrid antagonist results in a reduction in motility , implying the stimulatory effect of  
96 PACAP on sperm motility (Gozes et al. 1998). Indeed, PACAP has been reported to increase  
97 motility and penetration of ovum to promote fertilization in mice (Tanii et al. 2011). In an  
98 earlier study, we had confirmed this hypothesis in human sperms: we found that PACAP  
99 stimulated the slowly moving population, while it did not influence the cells with normal  
100 motility (Brubel et al. 2012). Furthermore, PACAP has been shown to regulate the synthesis  
101 of both secreted and intracellular proteins of spermatids and spermatocytes in vitro (West et  
102 al. 1995).

103 Mice deficient in PACAP are known to have several abnormalities during  
104 development (Nemeth et al. 2014, Sandor et al. 2016, Reglodi et al. 2012) and they also  
105 display reduced fertility (Reglodi et al. 2012, Sherwood et al. 2007). This is not limited to just  
106 one factor being abnormally altered but it seems that the reproduction of these animals is  
107 affected at several levels. PACAP deficient mice have been described to have impaired  
108 implantation (Isaac and Sherwood 2008), which, together with the increased rate of early  
109 postnatal death also leads to a reduced number of offspring (Gray et al. 2001, Wilson and  
110 Cumming, 2008). As PACAP is involved in several other processes, such as gametogenesis  
111 (Apa et al. 2002, Barberi et al. 2007, Canipari et al. 2016, Koppan et al. 2012, Li et al. 2004),  
112 placental development (Horvath et al. 2014, 2016) and gonadotropin regulation (Kanasaki et

113 al. 2016), it can be hypothesized that lack of PACAP may also influence reproduction in  
114 germ cells (Lacombe et al. 2006). In a previous study we found that the diameter of sperm  
115 heads in PACAP deficient mice was smaller, (Brubel et al. 2012), but no data was obtained  
116 as to whether this resulted in abnormal motility. Therefore, the aim of the present study was  
117 to perform detailed morphological analysis of spermatozoa, sperm motility, and to investigate  
118 signaling pathways playing a role during spermatogenesis in order to elucidate the role of  
119 endogenous PACAP at the molecular level.

120

121

## 122 **Materials and Methods**

123

### 124 **Animals and genotyping protocol**

125 Generation of male PACAP-deficient mice on CD-1 background was described earlier and it  
126 was demonstrated that heterozygous mice show a ~70% reduction of PACAP level  
127 (Hashimoto et al. 2001). Three-months-old wild-type (WT), heterozygous (HZ) and  
128 homozygous knockout (KO) mice were kept under light/dark cycles of 12/12 hours with free  
129 access to food and water. The study was carried out in accordance with ethical guidelines  
130 (ethical permission number for this study: BA02/2000-15024/2011, University of Pecs,  
131 Hungary). Genotyping was performed using Phire Animal Tissue Direct PCR Kit (Thermo  
132 Fischer Scientific, Waltham, MA, USA) according to the manufacturer's instructions. Primer  
133 sequences used for detection of wild-type and KO DNA signatures of PACAP were identical  
134 with those used earlier (Farkas et al. 2017, Hashimoto et al. 2001).

135

### 136 **Light and electron microscopical morphology**

137 Testes of 3-month-old wild-type (WT) (n=11), heterozygous (HZ) (n=10) and KO mice (n=8)  
138 were fixed in Bouin fixative containing (per 100 ml) 75 ml picric acid in saturated aqueous  
139 solution, 25 ml formalin (40% aqueous solution) and 5 ml glacial acetic acid, then embedded  
140 in paraffin. Serial sections were made and HE staining was performed (HE, Sigma-Aldrich,  
141 MO, USA). Photomicrographs were taken using an Olympus DP72 camera on a Nikon  
142 Eclipse E800 microscope. Sperms from a separate group of WT and KO animals were also  
143 investigated with electron microscopy (n=5-5 WT and KO). For electron microscopic  
144 evaluation, epididymis-derived sperms were fixed in 2% of formaldehyde and 2.5% of  
145 glutaraldehyde fixative overnight in 4°C after dissection. Samples were washed and  
146 dehydrated. One drop of the supernatant was dropped on a glass surface and it was sputter

147 coated with gold. Samples were examined and photographed in a JEOL 1200EX electron  
148 microscope.

149

### 150 **Evaluation of sperm quality**

151 Sperms were collected from the caudal part of the epididymis after cervical dislocation of 3-  
152 month-old adult male mice (n=18 WT, n=9 HZ and n=16 KO). The dissected caudal  
153 epididymis was placed in phosphate buffered saline (PBS) supplemented with 10% BSA and  
154 torn apart into small pieces followed by 5-minutes of incubation on a warming plate at 37.5  
155 °C. During incubation the spermatozoa had time to swim out of the epididymal duct.  
156 Obtained sperms were placed in PBS. Following 5 minutes of incubation on a warming plate  
157 at 37.5°C, sperm motility was measured with CASA System (Medealab™, Erlangen,  
158 Germany). For every sample, a total of 6 fields were examined giving an evaluation of  
159 approximately 900 cells per sample. Each field was recorded for 8 sec (total of 48 sec per  
160 sample). Sperms were divided into 4 groups based on motility: group A (rapid progressive),  
161 B (medium progressive), C (non-progressive), D (immotile). Sperm morphology (categories:  
162 normal, proximal or distal plasma droplets, detached head, bent tail, microcephaly,  
163 macrocephaly) and acrosome integrity (presence or absence of acrosome) were evaluated  
164 with Spermac™ staining (Beernem, Belgium). Minimum 200 cells were examined for the  
165 morphological analysis. After the measurement of untreated sperms, 5 µl of 100 nmol  
166 PACAP1-38 synthesized as previously described (Jozsa et al. 2005), dissolved in 495 µl PBS,  
167 was added to the samples which were incubated at 37°C for 5 minutes in order to investigate  
168 the effects of the exogenous peptide on motility (Brubel et al. 2012).

169

### 170 **Immunohistochemistry**

171 Immunohistochemistry was performed on WT (n=5), HZ (n=5) and KO (n=5) testis samples  
172 to visualize the localization of P-Sox9, Sox10 and collagen type IV (Col. IV). Testes were  
173 fixed in Saint-Marie's fixative and washed in 70% ethanol. After embedding serial sections  
174 were made, deparaffination was then followed by rinsing in PBS (pH 7.4). Non-specific  
175 binding sites were blocked with PBS supplemented with 1% bovine serum albumin (Amresco  
176 LLC, Solon, OH, USA), following which the samples were incubated with polyclonal P-  
177 Sox9 (Sigma-Aldrich, MO, USA), Sox10 (Abcam, Cambridge, UK), or Col. IV (Abcam,  
178 Cambridge, UK) antibodies at a dilution of 1:600 at 4°C overnight. For visualization of the  
179 primary antibodies, anti-rabbit Alexa fluor 555 secondary antibody (Life Technologies  
180 Corporation, Carlsbad, CA, USA) was used at a dilution of 1:1000. Samples were mounted in

181 Vectashield mounting medium (Vector Laboratories, Peterborough, England) containing  
182 DAPI for nuclear DNA staining. For negative controls anti-rabbit Alexa fluor 555 was used  
183 without the primary antibodies, which did not result in any positive signal (not shown).  
184 Photomicrographs of the tissues were taken using an Olympus DP72 camera on a Nikon  
185 Eclipse E800 microscope (Nikon Corporation, Tokyo, Japan). Images were acquired using  
186 cellSens Entry 1.5 software (Olympus, Shinjuku, Tokyo, Japan) with constant camera  
187 settings to allow comparison of fluorescent signal intensities. Images of Alexa555 and DAPI  
188 were overlaid using Adobe Photoshop version 10.0 software. The contrast of images was  
189 increased equally without changing constant settings.

190

### 191 **RT-PCR analysis**

192 Testes of WT, HZ and KO mice (n=5) were mechanically ground and were dissolved in  
193 Trizol (Applied Biosystems, Foster City, CA, USA), after 30 min incubation at 4°C total  
194 RNA was isolated. RNA was harvested in RNase-free water and stored at -70°C. Reverse  
195 transcription was performed by using High Capacity RT kit (Applied Biosystems, Foster  
196 City, CA, USA). For the sequences of primer pairs and details of polymerase chain reactions,  
197 see Table 1. Amplifications were performed in a thermal cycler (Labnet MultiGene™ 96-well  
198 Gradient Thermal Cycler; Labnet International, Edison, NJ, USA) as follows: at 95°C for 2  
199 min, followed by 35 cycles (denaturation at 94°C for 30 sec; annealing for 45 sec at  
200 optimized temperatures as given in Table 1; extension, 72°C, 90 sec) and then 72°C, 7 min.  
201 Glyceraldehyde 3-phosphate dehydrogenase (Gapdh) was used as internal control. PCR  
202 products were analyzed using a 1.2% agarose gel containing ethidium bromide. Optical  
203 densities of PCR product signals were determined by using ImageJ 1.40g freeware.

204

### 205 **Western blot analysis**

206 Testes of WT (n=5), HZ (n=5) and KO mice (n=5) were washed in physiological saline and  
207 stored at -70°C. Samples were mechanically disintegrated with a tissue grinder in liquid  
208 nitrogen. Then they were collected in 100 µL of homogenization RIPA (Radio Immuno  
209 Precipitation Assay)-buffer (150 mM sodium chloride; 1.0% NP40, 0.5% sodium  
210 deoxycholate; 50 mM Tris, pH 8.0) containing protease inhibitors (Aprotinin (10 µg/mL), 5  
211 mM Benzamidine, Leupeptin (10 µg/mL), Trypsine inhibitor (10 µg/mL), 1 mM PMSF, 5  
212 mM EDTA, 1 mM EGTA, 8 mM Na-Fluoride, 1 mM Na-orthovanadate). The suspensions  
213 were sonicated by pulsing burst for 30 sec at 40 A (Cole-Parmer, Illinois, USA). Total cell

214 lysates for Western blot analyses were prepared. Twenty  $\mu\text{g}$  protein was separated in 7.5%  
215 SDS–polyacrylamide gels for the detection of PKA, ERK 1/2, P-ERK 1/2, p38, P-p38, Sox9,  
216 P-Sox9, Sox10, PP2A, Collagen type IV (Col. IV), Collagen type IX (Col. IX), Testatin and  
217 Actin. Proteins were transferred by electrophoresis to nitrocellulose membranes and exposed  
218 to the primary antibodies overnight at 4°C in the dilution as given in Table 2. After washing  
219 for 30 minutes with PBST, membranes were incubated with the peroxidase-conjugated  
220 secondary antibody anti-rabbit IgG in a 1:1500 (Bio-Rad Laboratories, CA, USA) or anti-  
221 mouse IgG in 1:1500 (Bio-Rad Laboratories, CA, USA) dilution. Signals were detected with  
222 enhanced chemiluminescence (Advansta Inc., Menlo Park, CA, USA) according to the  
223 instructions provided by the manufacturer. Actin was used as an internal control. Signals  
224 were developed with gel documentary system (Fluorchem E, ProteinSimple, CA, USA).  
225 Optical densities of signals were measured by using ImageJ 1.40g freeware.

226

### 227 **In vitro protein phosphatase 2A (PP2A) activity assay**

228 PP2A activity was assayed with  $^{32}\text{P}$ -labelled myosin light-chain (5  $\mu\text{M}$ ) in testes of WT  
229 (n=5), HZ (n=5) and KO mice (n=5) as described previously (Erdodi et al. 1995). Briefly,  
230 pellets were suspended in 10 mM Tris/HCl (pH 7.4), 0.1 mM EGTA, 0.25 mM dithiothreitol,  
231 0.1 mM PMSF, 0.1 mM DFP, 0.1  $\text{mg mL}^{-1}$  leupeptin and 1 mM benzamidine (buffer A) and  
232 further diluted in buffer A supplemented with 1  $\text{mg mL}^{-1}$  BSA. One unit of the protein  
233 phosphatase activity releases 1  $\mu\text{mol}$  of  $\text{P}_i$  from the phosphosubstrate per min at 30°C.

234

### 235 **Statistical Analysis**

236 All data presented are representative of at least three independent experiments. Statistical  
237 analysis was performed by ANOVA and unpaired Student's *t*-test. The threshold for  
238 statistically significant differences as compared to controls was set at  $*P < 0.05$ .

239

240

## 241 **Results**

242

### 243 **Morphology of seminiferous tubules, sperm motility and morphology**

244 The general characteristics of seminiferous tubules were visualized with HE staining and  
245 morphological comparisons were made in WT and PACAP KO mice. No alterations were  
246 seen in tubular structures, and the general location of the differentiating cells was similar in

247 all experimental groups (Fig.1A). Regarding sperm morphology, there was no difference in  
248 the electron microscopical structure (Fig.1B). No difference was seen in the structure of HZ  
249 mice either (not shown).

250 Results of groups A and B sperms (rapid and medium progressive) were combined as  
251 well as those from groups C and D (non-progressive and immotile). The ratio of the sperms  
252 with good motility (A+B) and weak or no motility (C+D) was 65.6/34.4% in the wild-type  
253 animals, 60.6/39.8 in HZ mice and 68.7/31.3 in KO sperms (Fig.1D). There was no  
254 significant difference between any of the parameters. Addition of PACAP did not  
255 significantly change motility in any of the groups (data not shown). These results show that  
256 neither exogenous nor endogenous PACAP changed sperm motility in mice. Examining the  
257 morphology, we could not confirm statistical differences, in spite of the 10% less normal  
258 sperms in KO mice (percentage of normal sperms was  $69.6 \pm 3.1\%$ ,  $68.8 \pm 8.1$  and  $58.8 \pm 6.7$  in  
259 WT, HZ and KO mice, respectively). The standard deviation was high, but there were a few  
260 KO mice with no normal sperms at all, with the detached head being the most apparent  
261 abnormal sign (Fig.1C).

262

### 263 **PKA and MAPK pathways had modified activation in gene-deficient animals**

264 Expression of canonical targets of PACAP receptor activation such as PKA and MAPKs  
265 were examined by RT-PCR and Western blot. Although the mRNA expression of PKA did  
266 not show any alterations in HZ or KO mice, its protein expression was significantly  
267 decreased in KO animals, and it was expressed at higher levels in testis samples of HZ mice  
268 than in WT controls (Fig.2A and B). mRNA and protein expression of ERK1/2 were similar  
269 in the three experimental groups. The phosphorylated form of ERK1/2 represents a more  
270 active form of this MAPK, which may have an influence on cell proliferation. HZ and  
271 PACAP KO mice showed a significantly increased P-ERK1/2 level (Fig.2A and B). p38 is  
272 another possible MAPK which can be involved in PACAP-induced signaling pathways. Its  
273 mRNA expression did not show alterations, but significantly higher protein expression and P-  
274 p38 were detected in the HZ and KO mice than in WT controls (Fig.2A and B).

275

### 276 **Possible downstream targets of PKA in PACAP KO mice**

277 The activation of Sox9, a major transcription factor of testis development and  
278 spermatogenesis, can be regulated by PKA as a canonical downstream target of PAC1  
279 receptor activation. The mRNA expression of Sox9 was not altered in HZ or KO animals  
280 (Fig.3A). While Sox9 protein expression was unchanged in HZ mice, it was significantly



281 decreased in PACAP KO mice. The level of the more active phosphorylated form of Sox9  
282 also showed a reduction similar to the non-phosphorylated version in homogenous gene-  
283 deficient mice (Fig.3B), but again the protein expression was not significantly altered in HZ  
284 mice (Fig.3B). Next, we investigated the localization of P-Sox9 in seminiferous tubules and  
285 its immunopositivity was detected in the outer region of WT animals. In HZ animals  
286 expression of P-Sox9 was strongly localized close to the lumen of seminiferous tubules with  
287 a very weak peripheral presence (Fig.3D). Immunohistochemical results indicated lower  
288 immunopositivity of P-Sox9 detected in all parts of the seminiferous tubules of PACAP KO  
289 mice (Fig.3D). Sox10 is another important regulatory factor of testis  
290 development/spermatogenesis and it can substitute the function of Sox9. mRNA expression  
291 of this transcription factor was constant in the investigated samples (Fig.3A). Unexpectedly,  
292 Sox10 expression was significantly elevated in PACAP KO mice while it was not altered in  
293 HZ mice (Fig.3B). Immunopositivity of Sox10 in WT and HZ mice was very weak and  
294 randomly appeared in the cells of seminiferous tubules. On the contrary, strong Sox10 signals  
295 were detected in the middle portion of tubules in KO animals (Fig.3E).

296

### 297 **Phosphatase function**

298 The phosphorylation and, consequently, activation of Sox9 are regulated by the Ser/Thr  
299 phosphatase called PP2A, which can reversibly dephosphorylate the targets of PKA. As the  
300 decreased phosphorylated Sox9 expression can be the result of an increased PP2A activity we  
301 monitored the phosphatase function in the testis. Interestingly the mRNA and protein levels  
302 of PP2A showed a significant elevation in HZ and PACAP KO mice (Fig.3A and B). The  
303 increased expression of the phosphatase is not always followed by more intense activity;  
304 therefore, we measured the PP2A activity in the testis samples. In PACAP KO mice the  
305 phosphatase activity was dramatically increased. No differences were detected between the  
306 WT and HZ groups (Fig.3C).

307

### 308 **Target genes of Sox transcription factors in testis development**

309 Possible targets of Sox transcription factors can be collagen type IV (Col. IV), collagen type  
310 IX (Col. IX) or testatin, major components of the basal lamina of the tubules. The mRNA  
311 expression of testatin was not altered, while that of Col4a showed a significant decrease in  
312 HZ and KO mice (Fig.4A). However, the protein expression of this basement membrane  
313 component showed a significant decrease in HZ and PACAP KO mice (Fig.4B). The mRNA  
314 expression of Col. IX protein was moderately reduced in gene-deficient mice, while its

315 protein expression was elevated in both HZ and PACAP KO rodents (Fig.4A and B). The  
316 other testis-specific basement membrane component is testatin, the protein expression of  
317 which was dramatically increased in PACAP KO mice (Fig.4A and B). As these observations  
318 may reflect a possible alteration in the basement membrane integrity, immunohistochemistry  
319 was performed to detect structural differences of basal membrane Col. IV. In WT mice the  
320 integrity of the basement membrane was strongly detected around the seminiferous tubules.  
321 HZ animals showed a weaker Col. IV immunoreaction, which further decreased in PACAP  
322 KO animals (Fig.4C).

323

324

## 325 **Discussion**

326

327 In the present study we described that several factors playing an important role in  
328 spermatogenesis are significantly influenced by partial or complete lack of PACAP. In spite  
329 of the unaltered or only slightly altered morphology and motility of spermatozoa of PACAP  
330 deficient animals, the disturbed molecular mechanisms in spermatogenesis may account for  
331 some of the known perturbations in the fertility of PACAP KO mice. The possible effects of  
332 PACAP on signaling mechanisms are summarized in Figure 5 and discussed below.

333 Several studies have shown the presence of PACAP, its mRNA and receptors in the  
334 testis and epididymis of different vertebrate species and in the human testis (Agnese et al.  
335 2010, 2016, Konone et al. 1994, Rosati et al. 2014) after the pioneer study of Arimura and  
336 coworkers (1991), which showed that testis contained the highest levels of PACAP among  
337 the peripheral organs. Lv et al. (2011) described the gradual increase of PACAP mRNA in  
338 the rat testis from day 20 after birth, reaching the maximum level on day 60, mainly in  
339 spermatocytes and round spermatids. Similarly, the epididymis showed a gradual increase in  
340 PACAP during puberty (Lv et al. 2011). Other studies found stage-specific expression of  
341 PACAP: while high levels of expression could be found in most developing germ cells,  
342 PACAP expression is lacking in mature spermatids according to some reports (Yanaihara et  
343 al. 1998). Tanii and coworkers have described weak immunostaining in mature sperm heads  
344 (Tanii et al 2011). The expression of PACAP has been shown to be influenced by several  
345 factors, but its regulation in the testis is not yet entirely understood. Raising the temperature  
346 to body temperature by inducing experimental cryptorchidism can lead to a dramatic decrease  
347 of PACAP mRNA both in the testis and epididymis (Lv et al. 2011). It is possible that this  
348 and other, yet unknown, influencing factors are responsible for the stage-specific expression

349 of PACAP during the spermatogenic cycle and are responsible for the discrepancies found  
350 between descriptions (Rosati et al. 2014). A testis-specific splice variant of the PAC1  
351 receptor has also been reported (Daniel et al. 2001). Interestingly, unlike in other tissues,  
352 PACAP38 and 27 were found to have the same levels in human testis, shown by  
353 radioimmunoassay (Tamas et al. 2015). No differences were found in the level of PACAP  
354 peptides between normal tissues and seminomas (Tamas et al. 2015), but a different  
355 distribution pattern was observed in different tumor biopsies by Nakamura et al. (2014). In  
356 addition to the local production of PACAP, circulating PACAP can additionally contribute to  
357 testicular functions of the peptide, as it has been shown that PACAP can cross the blood-  
358 testis barrier (Banks et al. 1993, Mizushima et al. 2001). Testis functions in PACAP gene-  
359 deficient animals had only been studied by Lacombe et al. (2006). This study revealed  
360 interesting results in the testis: unlike in any other organ/system investigated so far, testicular  
361 aging was delayed in PACAP deficient mice with a decreased level of apoptosis and  
362 decreased oxidative stress markers (Lacombe et al. 2006). The authors argued that the  
363 stimulatory effect of PACAP on steroidogenesis may result in a higher level of its byproduct  
364 reactive oxygen species production, leading to testicular aging, but also supporting the role of  
365 PACAP played in normal testicular function at younger ages. Our present results of normal  
366 testicular histology in PACAP KO mice at a young age are in accordance with the findings of  
367 Lacombe and coworkers (Lacombe et al. 2006).

368         The high level of PACAP and its stage-specific expression pattern indicate that the  
369 peptide plays important roles in spermatogenesis. It has been described that PACAP  
370 influences hormonal secretion via regulating Leydig- and Sertoli cell growth and functions,  
371 as well as acrosome reaction during fertilization (Tanii et al. 2011). Influencing motility is  
372 also among the possible functions of PACAP, as our previous report showed that PACAP  
373 could increase the motility of abnormally slow-moving human sperms (Brubel et al. 2012).  
374 The results of Gozes et al. showing that addition of a PACAP antagonist peptide led to a  
375 decrease in motility provided a proof for the possible endogenous stimulatory effect of  
376 PACAP (Gozes et al. 1998). However, our present finding that mice deficient in PACAP  
377 show normal motility implies that even if present, the endogenous effects of PACAP on  
378 motility can be compensated by other factors. Furthermore, exogenous PACAP did not alter  
379 the sperm motility in the present study. This observation is in accordance with our earlier  
380 description in human sperms (Brubel et al. 2012), where we found that the motility of only  
381 slow-moving sperms could be increased by PACAP with normal moving sperms not  
382 influenced. Our previous study also described that sperm heads from PACAP deficient mice

383 were significantly smaller than those from wild types (Brubel et al. 2012). This observation  
384 further implies that PACAP is necessary for normal spermatogenesis. Our present findings  
385 also support this observation: the percentage of normal sperms was 10% lower in PACAP  
386 KO mice than in the other two groups, even if due to the high deviation this was not  
387 statistically significant. Among the abnormal sperms, more sperms had detached heads in the  
388 PACAP KO group. In spite of some differences, our present results show that reduced  
389 fertility in PACAP KO mice is most probably not due to altered morphology or motility,  
390 since differences were slight. However, we revealed marked differences in the molecular  
391 factors and targets of PACAP-induced signaling pathways.

392 PACAP, binding to its receptors, induces intracellular cAMP accumulation, and  
393 subsequently activates PKA, also known as the canonical downstream signaling pathway. As  
394 expected, the expression of PKA was reduced in the samples of KO mice. PKA has a crucial  
395 role in germ cell development and differentiation, therefore its lower activity can result in  
396 lower fertility, decreased sperm number, or reduced capacitation capability (Burton and  
397 McKnight, 2007). The inactivation of PKA by the lack of PACAP may influence  
398 spermatogenesis or spermiogenesis. However, PKA pathways have several cross-talks, thus  
399 the lack of PKA has compensation possibilities.

400 It is also known that PACAP receptor activation can communicate with other kinases  
401 such as ERK1/2 and p38 (Racz et al. 2007), which may influence cell survival, proliferation  
402 and differentiation. Therefore, we monitored the expression of these MAPKs. Protein  
403 expression of p38 and of the more active phosphorylated form of ERK1/2 and p38 increased  
404 in PACAP deficient testes. ERK1/2 functions have been described in the testis: it can regulate  
405 cell renewal in goat spermatogonia (Niu et al. 2015) and affects Sertoli cell junction, blood-  
406 testis barrier and division of spermatogonia (Siu et al. 2005, Wong and Cheng, 2005). Our  
407 data suggest that the elevated expression of P-ERK1/2 can be a compensatory effect of the  
408 lower PKA activation, ensuring the spermatogenesis with the activation of other downstream  
409 signalization in a PKA independent manner. p38 also plays role in the Sertoli cell adhesion  
410 stability and has some regulatory effect on elongated spermatids (Wong and Cheng, 2005)  
411 triggering normal motility. Thus, the effect of p38 in PACAP KO animals can be dual in  
412 testis, partly maintaining a functional blood-testis barrier and supporting the normal cell  
413 motility of spermatids. On the one hand, the increased phosphorylation of p38 alters the  
414 apoptotic processes in spermatogonia resulting in an abnormal differentiation with altered  
415 sperm morphology. On the other hand, PKA can phosphorylate several transcription factors,  
416 which play unique roles in testis development and spermatogenesis.

417 The members of SoxE transcription factors such as Sox8, Sox9 and Sox10 have  
418 fundamental functions in several developmental processes, including testis development  
419 (Barrinuevo and Scherer, 2010, Georg et al. 2012). Sox9 is required for maintenance of the  
420 integrity of Sertoli cells in adult testis (Barrinuevo et al. 2016). Loss of function of Sox10 had  
421 no effect on testis development (Barrinuevo and Scherer 2010) but a cross-regulation has  
422 been proven between Sox9 and Sox10 in other cells, such as melanocytes (Shakhova et al.  
423 2015). The reduced expression and phosphorylation of Sox9 in PACAP KO mice suggest a  
424 direct signalization with PKA as it has been shown in chondrogenic cells (Zakany et al.  
425 2005). Moreover, it has been demonstrated that PAC1 receptor activation increases the level  
426 of P-Sox9 in chondrogenic cell cultures (Juhasz et al. 2014a), which further supports the  
427 possible PKA-Sox9 signalization axis. The elevated Sox10 expression proves a possible  
428 substitution of Sox9 function, therefore, the increased level of Sox10 may compensate for the  
429 lower activity of Sox9, resulting in a normal blood-testis barrier. On the other hand,  
430 expression profiles in HZ animals were disturbed, also suggesting an interrupted Sox9-Sox10  
431 crosstalk. A recent study has demonstrated that some signs of neurobehavioural development  
432 are more disturbed in mice partially lacking PACAP than in mice with a complete lack of the  
433 peptide (Farkas et al. 2017), implying disturbed compensatory processes in some signaling  
434 pathways.

435 Transcription factors phosphorylated by PKA can be dephosphorylated by a Ser/Thr  
436 phosphatase, PP2A (Juhasz et al. 2014b; Zakany et al. 2002). PP2A was demonstrated  
437 influencing spermatid maturation (Hatano et al. 1993) and differentiation of Sertoli cells  
438 (Levallet et al. 2013). Furthermore, the inhibitory effect of PACAP on PP2A has been  
439 detected in chromaffin cells (Bobrovskaya et al. 2007) but no data is available for the testis.  
440 According to our results the absence of PACAP results in an increased PP2A activity, which  
441 can partly be responsible for the lower phosphorylation of Sox9 (Zakany et al. 2002).  
442 Therefore, the transcriptional inactivation can be a consequence of lower PKA activity, and a  
443 parallelly increased PP2A function. Sox9 can be translocated into the nuclei of Sertoli cells  
444 and can regulate certain gene expression. Col. IV is one of the major elements of the  
445 basement membrane in the seminiferous tubules, and regulates the formation of the blood-  
446 testis barrier (Harvey et al. 2006). Col. IX also has some functions in the maintenance of the  
447 basement membrane, consequently the blood-testis barrier (McClive and Sinclair, 2003).  
448 Furthermore, testatin has also been proven to have a role in the regulation of testis  
449 development (Georg et al. 2012). The genes of these proteins may be regulated by Sox9 as a  
450 termination of the PAC1-PKA-Sox9 axis. As we demonstrated, the protein level of Col. IV in

451 the basement membrane of the seminiferous tubules was decreased, therefore, the basement  
452 membrane was not well discernible in PACAP KO animals. On the other hand, the  
453 expression of testatin and Col. IX was elevated, suggesting a possible upregulation by other  
454 members of the SoxE family such as Sox10. On the contrary the mRNA expression of Col.  
455 IX and testatin was decreased indicating an altered transcriptional activation as it has been  
456 demonstrated in chondrogenic differentiation (Juhasz et al. 2014a). Our findings also support  
457 the idea that SoxE family members, especially Sox9, are important transcription factors in  
458 blood-testis barrier dynamics, which can be precisely regulated by PACAP. Indeed, we  
459 detected elevated expression of Sox 10 in testes of PACAP KO mice.

460 In summary, our findings suggest a role of PACAP in Sox9 expression and  
461 phosphorylation in mouse testis. In addition to the direct effects of Sox9 on testis  
462 determination and male germ cell production, PACAP seems to play a role in the  
463 maintenance of proper molecular composition of the basement membrane of seminiferous  
464 tubules. These results could be additional factors responsible for the reduced fertility in  
465 PACAP knockout mice as well as further supporting the role of endogenous PACAP in  
466 reproductive functions.

467

468

#### 469 **Acknowledgements**

470

471 NKFI K115874, PD109644, K119759, PTE AOK Research Grant, MTA-TKI14016, GINOP-  
472 2.3.2-15-2016-00050 “PEPSYS”, New National Excellence Program of the Ministry of  
473 Human Capacities (UNKP-16-4-IV.), KTIA\_13\_NAP-A-III/5, EFOP-3.6.1.-16-2016-00004  
474 Comprehensive Development for Implementing Smart Specialization Strategies at the  
475 University of Pecs, Centre for Neuroscience of the University of Pecs, University of  
476 Debrecen (RH/751/2015). The project is co-financed by the European Union and the  
477 European Social Fund. Bolyai Janos Research Scholarship (JT, AT). Szodoray Lajos and  
478 Magyary Zoltán Funds by Hungarian Academy of Science and the European Union and the  
479 State of Hungary, co-financed by the European Social Fund in the framework of TÁMOP  
480 4.2.4 (JT). A/2-11-1-2012-0001 “National Excellence Program”. The present scientific  
481 contribution is dedicated to the 650<sup>th</sup> anniversary of the foundation of the University of Pecs,  
482 Hungary.

483

#### 484 **Declaration of Interests**

485 The authors declare that there is no conflict of interest that could be perceived as prejudicing  
486 the impartiality of the research reported.

487

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## Figure Legends

### Fig. 1.

(A) Representative microphotograph of WT and KO seminiferous tubules stained with haematoxylin and eosin. Overview of several tubules and higher magnification view of the wall of one seminiferous tubule.

(B) Electronmicroscopical analysis of WT and KO sperms.

(C) Representative images of spermatozoa showing normal and most common abnormal morphologies. White arrow: plasma droplet; white arrowhead: hairpin tail; black arrowhead: acrosome deficiency; asterisk: detached head; n: normal morphology.

(D) Motility of sperms: A (rapid progressive), B (medium progressive), C (non-progressive), D (immotile). Data are represented as A+B (progressive) and C+D (non-progressive and immotile) together. Data are given as mean $\pm$ SEM.

### Fig. 2.

Optical density of signals was measured and results were normalized to the optical density of controls. For panels (A) and (B) numbers below the signals represent integrated densities of signals determined by ImageJ software. Asterisks indicate significant ( $*P < 0.05$ ) alteration of expression as compared to the respective control. Representative data of 3 independent experiments. WT (+/+), HZ (+/-), PACAP KO (-/-). For RT-PCR reactions GAPDH and for Western blot actin were used as controls.

### Fig. 3.

Optical density of signals was measured and results were normalised to the optical density of controls. For panels (A) and (B) numbers below the signals represent integrated densities of signals determined by ImageJ software. Asterisks indicate significant ( $*P < 0.05$ ) alteration of expression as compared to the respective control. Representative data of 3 independent experiments. (C) PP2A enzyme activity determined in a cell-free *in vitro* assay from lysates of WT, HZ and PACAP KO samples. Significant difference between PP2A enzyme activity in HZ and PACAP KO samples vs. control lysates is marked by asterisks ( $*P < 0.05$ ). (D) (E) Immunohistochemistry of P-Sox9 and

Sox10 in seminiferous tubules. Magnification was made with 20× objective. Scale bar: 50 μm. WT (+/+), HZ (+/-), PACAP KO (-/-). For RT-PCR Gapdh and for Western blot reactions actin were used as controls.

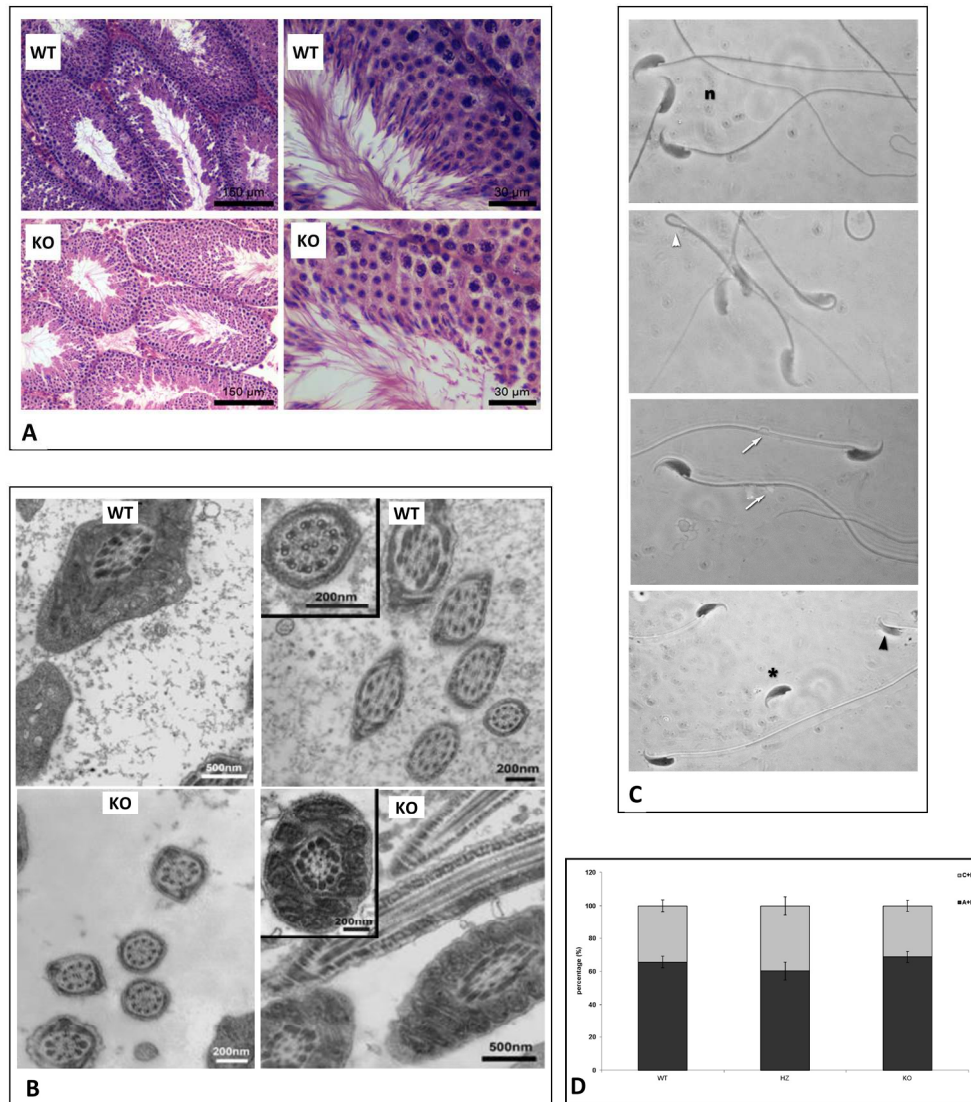
**Fig. 4.**

Downstream targets of Sox transcription factors altered in PACAP knockout mice. mRNA (A) and protein (B) expression of Col. IV, Col. IX and testatin. For RT-PCR Gapdh and for Western blot reactions actin were used as controls. Optical density of signals was measured and results were normalized to the optical density of controls. For panels (A) and (B) numbers below the signals represent integrated densities of signals determined by ImageJ software. Asterisks indicate significant (\*P<0.05) alteration of expression as compared to the respective control. Representative data of 3 independent experiments. (C) Immunohistochemistry of Col. IV in seminiferous tubules. Magnification was made with 20× objective. Scale bar: 50 μm. WT (+/+), HZ (+/-), PACAP KO (-/-).

**Fig. 5.**

Schematic drawing of the possible signaling pathways regulated by PACAP in spermatogenesis

PAC1 receptor activation leads to the increase of intracellular cAMP, which activates PKA. Downstream targets of PKA are partly regulated by Ser/Thr reversible phosphorylation. PP2A can regulate the dephosphorylation of the transcription factor Sox9, which can modify the expression or activation of Sox10. Additionally, PKA can influence ERK1/2 phosphorylation. Parallel with these events, p38 activation can also take place. Main targets of Sox9 transcription factor can be Collagen type IV, Collagen type IX or Testatin. These reversible phosphorylation events may regulate proliferation, motility, or formation of blood-testis barrier.

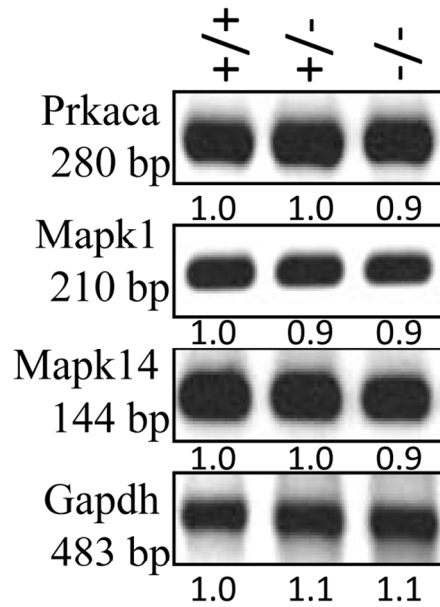
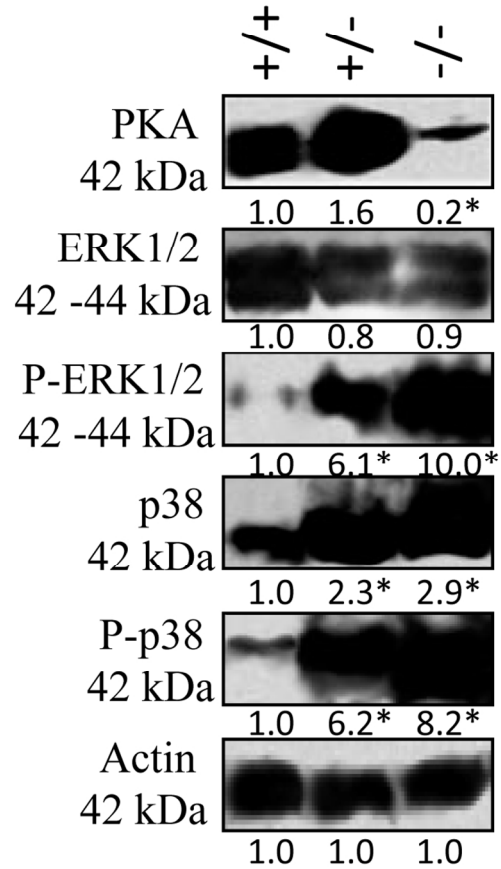


(A) Representative microphotograph of WT and KO seminiferous tubules stained with haematoxylin and eosin. Overview of several tubules and higher magnification view of the wall of one seminiferous tubule.

(B) Electronmicroscopical analysis of WT and KO sperms.

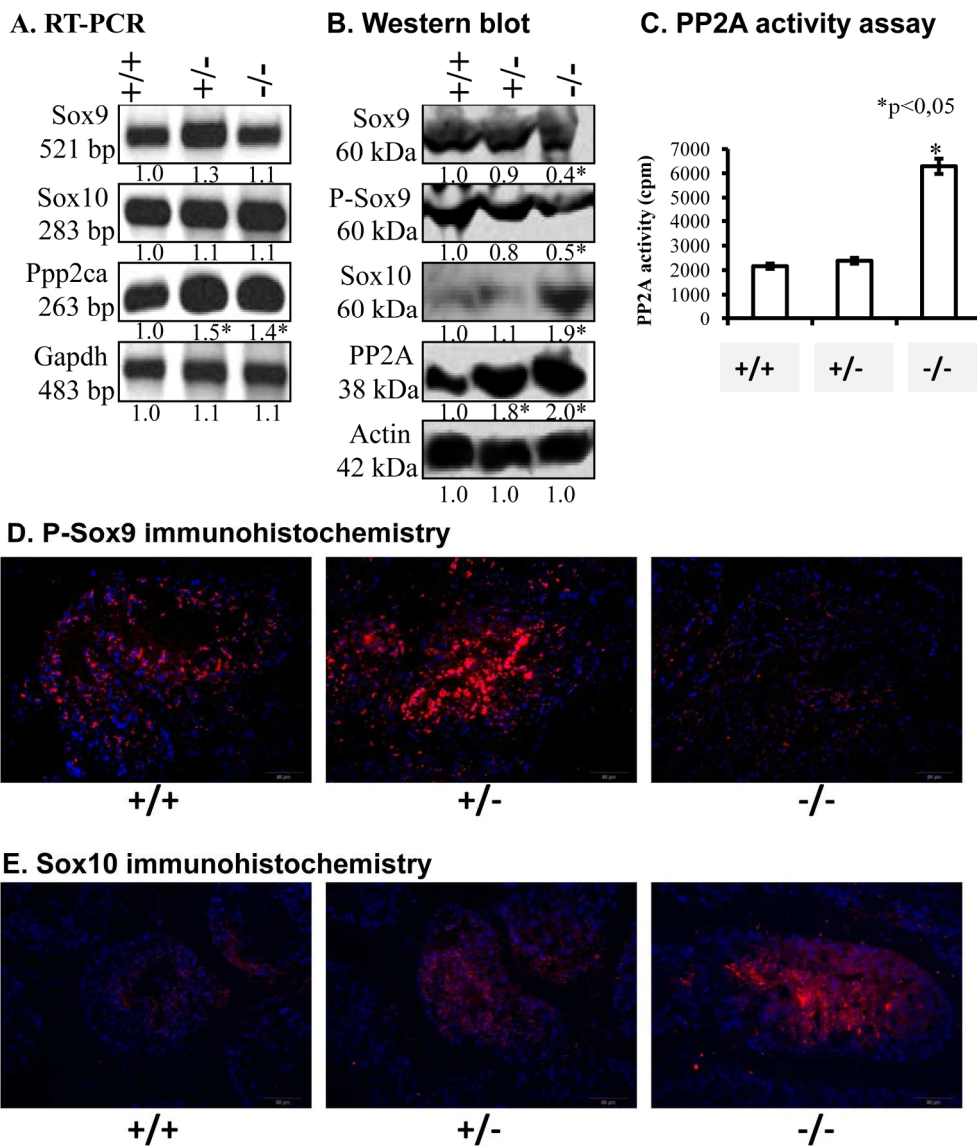
(C) Representative images of spermatozoa showing normal and most common abnormal morphologies. White arrow: plasma droplet; white arrowhead: hairpin tail; black arrowhead: acrosome deficiency; asterisk: detached head; n: normal morphology.

(D) Motility of sperms: A (rapid progressive), B (medium progressive), C (non-progressive), D (immotile). Data are represented as A+B (progressive) and C+D (non-progressive and immotile) together. Data are given as mean±SEM.

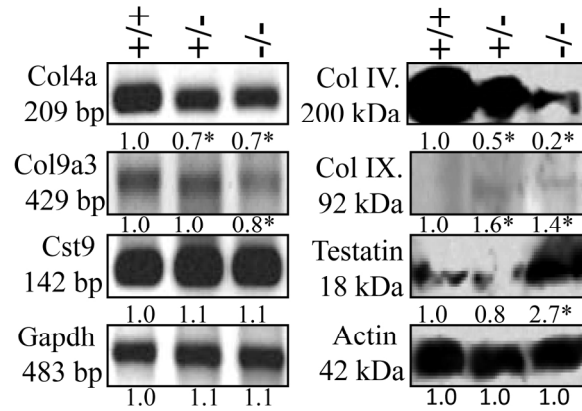
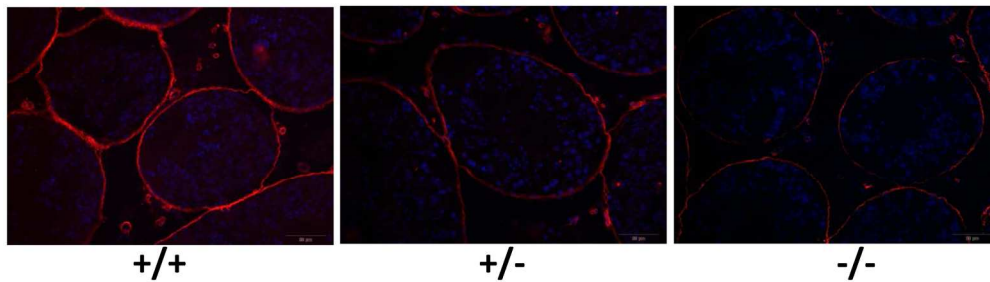
**A. RT-PCR****B. Western blot**

Optical density of signals was measured and results were normalized to the optical density of controls. For panels (A) and (B) numbers below the signals represent integrated densities of signals determined by ImageJ software. Asterisks indicate significant (\* $P < 0.05$ ) alteration of expression as compared to the respective control. Representative data of 3 independent experiments. WT (+/+), HZ (+/-), PACAP KO (-/-). For RT-PCR reactions Gapdh and for Western blot actin were used as controls.

112x107mm (300 x 300 DPI)



Optical density of signals was measured and results were normalised to the optical density of controls. For panels (A) and (B) numbers below the signals represent integrated densities of signals determined by ImageJ software. Asterisks indicate significant (\* $P < 0.05$ ) alteration of expression as compared to the respective control. Representative data of 3 independent experiments. (C) PP2A enzyme activity determined in a cell-free in vitro assay from lysates of WT, HZ and PACAP KO samples. Significant difference between PP2A enzyme activity in HZ and PACAP KO samples vs. control lysates is marked by asterisks (\* $P < 0.05$ ). (D) (E) Immunohistochemistry of P-Sox9 and Sox10 in seminiferous tubules. Magnification was made with 20 $\times$  objective. Scale bar: 50  $\mu$ m. WT (+/+), HZ (+/-), PACAP KO (-/-). For RT-PCR Gapdh and for Western blot reactions actin were used as controls.

**A. RT-PCR****B. Western blot****C. Coll type IV. immunohistochemistry**

Downstream targets of Sox transcription factors altered in PACAP knockout mice. mRNA (A) and protein (B) expression of Col. IV, Col. IX and testatin. For RT-PCR Gapdh and for Western blot reactions actin were used as controls. Optical density of signals was measured and results were normalized to the optical density of controls. For panels (A) and (B) numbers below the signals represent integrated densities of signals determined by ImageJ software. Asterisks indicate significant (\*P < 0.05) alteration of expression as compared to the respective control. Representative data of 3 independent experiments. (C) Immunohistochemistry of Col. IV in seminiferous tubules. Magnification was made with 20× objective. Scale bar: 50 μm. WT (+/+), HZ (+/-), PACAP KO (-/-).

168x142mm (300 x 300 DPI)

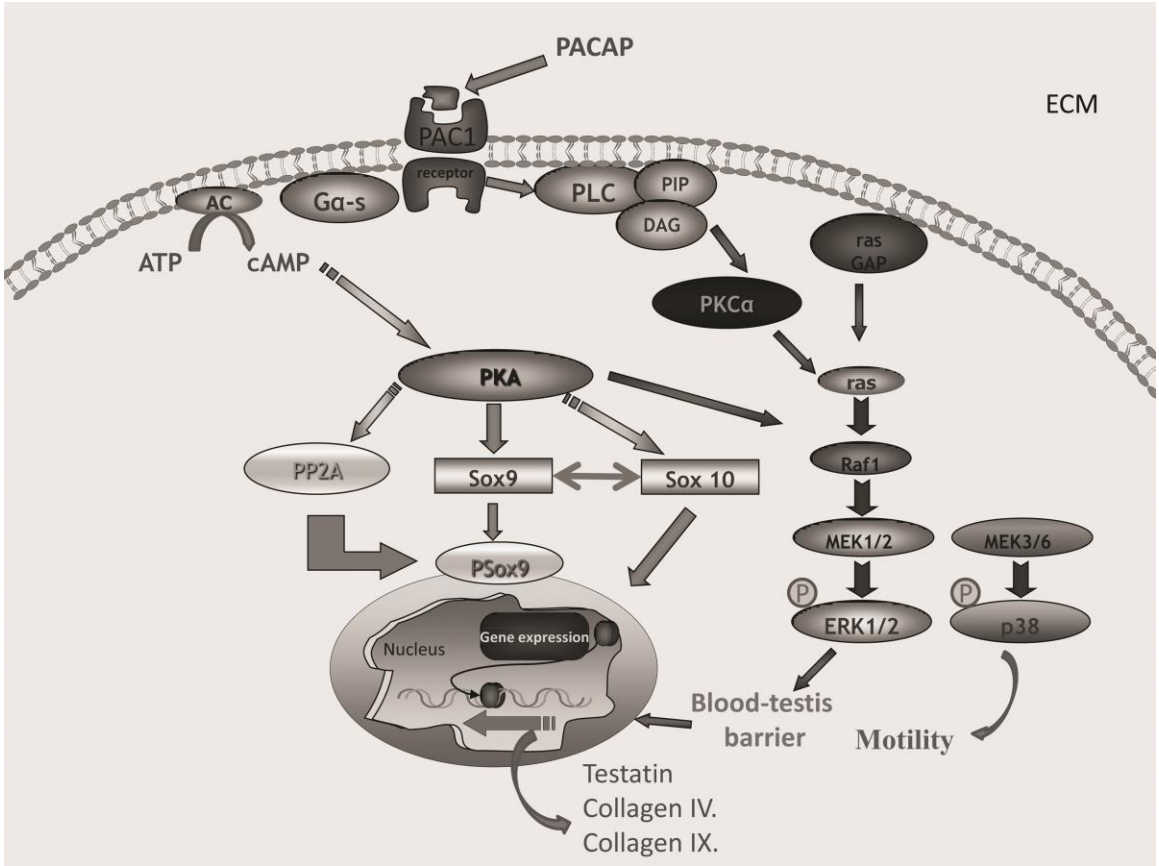


Table 1. Nucleotide sequences, amplification sites, GenBank accession numbers, amplicon sizes and PCR reaction conditions for each primer pair are shown.

| <i>Gene</i>                     | <i>Primer</i> | <i>Nucleotide sequence (5'→3')</i>             | <i>GenBank ID</i>     | <i>Annealing temperature</i> | <i>Amplicon size (bp)</i> |
|---------------------------------|---------------|--|-----------------------|------------------------------|---------------------------|
| <b>PKA<br/>(Prkaca)</b>         | sense         | GCA AAG GCT ACA ACA AGG C<br>(847–865)         | <b>NM_001277898.1</b> | 54°C                         | 280                       |
|                                 | antisense     | ATG GCA ATC CAG TCA GTC G<br>(1108–1126)       |                       |                              |                           |
| <b>ERK<br/>(Mapk1)</b>          | sense         | CCG TGA CCT CAA GCC TTC C<br>(1732–1750)       | <b>NM_011949.3</b>    | 58°C                         | 163                       |
|                                 | antisense     | ATG CAG CCC ACA GAC CAA A<br>(1877–1894)       |                       |                              |                           |
| <b>p38<br/>(Mapk14)</b>         | sense         | GAG GTG CCC GAA CGA TAC<br>(453–470)           | <b>NM_011951.3</b>    | 53°C                         | 144                       |
|                                 | antisense     | TTT GGC GTG AAT GAT GGA<br>(579–596)           |                       |                              |                           |
| <b>Sox9</b>                     | sense         | GTA CCC GCA TCT GCA CAA CG<br>(378–397)        | <b>NM_011448</b>      | 62°C                         | 521                       |
|                                 | antisense     | GTG GCA AGT ATT GGT CAA ACT CAT T<br>(874–898) |                       |                              |                           |
| <b>Sox10</b>                    | sense         | ACG ACT GGA CGC TGG TGC<br>(535–552)           | <b>NM_011437.1</b>    | 58°C                         | 283                       |
|                                 | antisense     | CGC CGA GGT TGG TAC TTG TAG<br>(797–817)       |                       |                              |                           |
| <b>PP2A<br/>(Ppp2ca)</b>        | sense         | CTC TGC GAG AAG GCT AAA<br>(288–305)           | <b>NM_017039.2</b>    | 54°C                         | 436                       |
|                                 | antisense     | TGA TTC CCT CGG AGT ATG<br>(706–723)           |                       |                              |                           |
| <b>Collagen IV<br/>(Col4a1)</b> | sense         | TCG GCT ATT CCT TCG TGA TG<br>(4963 – 4982)    | <b>NM_009931.2</b>    | 52°C                         | 209                       |
|                                 | antisense     | GGA TGG CGT GGG CTT CTT<br>(5154 –5171)        |                       |                              |                           |
| <b>Collagen IX<br/>(Col9a3)</b> | sense         | CAG GTT CCG ATG GTC TTC C<br>(1357-1375)       | <b>NM_009936.2</b>    | 55°C                         | 492                       |
|                                 | antisense     | CTG TTG CTC CCT TGT CCC<br>(1831-1848)         |                       |                              |                           |
| <b>Testatin<br/>(Cys9)</b>      | sense         | CTG GAG GGA GAA GGT AAA<br>(274-291)           | <b>NM_009979.1</b>    | 51°C                         | 142                       |
|                                 | antisense     | CAG GCA GGT GAA GGT ATT<br>(398-415)           |                       |                              |                           |
| <b>GAPDH</b>                    | sense         | TGG CAA AGT GGA GAT TGT TG<br>(238–258)        | <b>NM_008084.2</b>    | 59°C                         | 486                       |
|                                 | antisense     | GTC TTC TGG GTG GCA GTG AT<br>(775–793)        |                       |                              |                           |



Table 2. Tables of antibodies used in the experiments.

| <i>Antibody</i>       | <i>Host animal</i>  | <i>Dilution</i> | <i>Distributor</i>                        |
|-----------------------|---------------------|-----------------|---|
| <b>Anti-Col. IX.</b>  | rabbit, polyclonal, | 1:500           | Abcam, Cambridge, UK                      |
| <b>Anti-Col. IV.</b>  | rabbit, polyclonal, | 1:800           | Abcam, Cambridge, UK                      |
| <b>Anti-Testatin.</b> | rabbit, polyclonal, | 1:300           | Santa Cruz Biotechnology, Dallas, TX, USA |
| <b>Anti-PKA C</b>     | rabbit, polyclonal, | 1:800           | Cell Signaling, Danvers, MA, USA          |
| <b>Anti-ERK1/2</b>    | rabbit, polyclonal, | 1:1000          | Sigma-Aldrich, St. Louis, MO, USA         |
| <b>Anti-P-ERK/12</b>  | mouse, monoclonal,  | 1:800           | Sigma-Aldrich, St. Louis, MO, USA         |
| <b>Anti-p38</b>       | rabbit, polyclonal, | 1:1000          | Cell Signaling, Danvers, MA, USA          |
| <b>Anti-P-p38</b>     | rabbit, polyclonal, | 1:800           | Cell Signaling, Danvers, MA, USA          |
| <b>Anti-Sox9</b>      | rabbit, polyclonal, | 1:800           | Abcam, Cambridge, UK                      |
| <b>Anti-P-Sox9</b>    | rabbit, polyclonal, | 1:800           | Sigma-Aldrich, St. Louis, MO, USA         |
| <b>Anti-Sox10</b>     | rabbit, polyclonal, | 1:500           | Abcam, Cambridge, UK                      |
| <b>Anti-PP2A C</b>    | rabbit, polyclonal, | 1:600           | Cell Signaling, Danvers, MA, USA          |
| <b>Anti-Actin</b>     | mouse, monoclonal,  | 1:10000         | Sigma-Aldrich, St. Louis, MO, USA         |