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A specific role for PRND in goat foetal Leydig cells is suggested by prion family gene expression during gonad development in goats and mice

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The prion gene family is mainly composed of three genes; Prnp, encoding the prion protein (PrP), Prnd, encoding Doppel (Dpl), both located on the same genomic locus on mouse chromosome 2 and Sprn, encoding Shadoo (Sho), located on mouse chromosome 7. These proteins are glycophasphatidylinositol (GPI)-anchored glycoproteins and they shared some structural homology one with each other. The N-terminal regions of PrP and Sho are composed of basic repeat regions and of an hydrophobic domain, whereas the C-terminal regions of PrP and Dpl contain alpha helices [1]. PrP plays a pivotal role in transmissible spongiform encephalopathy (TSE), a fatal neurodegenerative disorder affecting animals and humans [2–4]. PrP is almost ubiquitously expressed with higher amount of expression occurring in the central nervous system (CNS). In mice and rams, PrP was found to be expressed in germ cells [5–7] but the genetic ablation of its gene in mice [8,9], cattle [10] and goats [11] does not induce a fertility-associated phenotype and/or major neuronal disorders [12]. Thus, the PrP biological function remains elusive even if various roles have been proposed [13,14]. These observations suggested a biological redundancy between PrP and another PrP-like protein in mammals.

Abbreviations
CNS, central nervous system; dpc, days post coitum; Dpl, Doppel; dpp, days post partum; ECL, enhanced chemiluminescence; GPI, glycophasphatidylinositol; IHC, immuno-histo-chemistry; mpp, months post partum; PBS, phosphate saline buffer; qRT-PCR, quantitative RT-PCR; Sho, Shadoo; TSE, transmissible spongiform encephalopathy.
Sho is expressed in the CNS and both Sho and PrP share neuro-protective properties [15]. Using Sprn reporter mice, a recent study describes expression of Sprn in the male and female gonads suggesting an involvement of Sho in reproduction [16]. The Sprn mRNA knockdown in Prnp\(^{0/0}\) embryos produces early embryonic lethality, suggesting that Sho and PrP could play a role in early developmental stages [17,18]. However, the Sprn ablation (Sprn\(^{0/0}\)) or the double-knockout of Sprn and Prnp (Sprn\(^{0/0}\)/Prnp\(^{0/0}\)) in mice resulted in no drastic developmental phenotype [19].

Dpl is mainly expressed in the testis of adult mammals. Its ectopic expression induces some neuro-degeneration in the CNS only in the absence of PrP [20–23], suggesting a biological link between Dpl and PrP. Prnd ablation in mice (Prnd\(^{0/0}\)) resulted in male infertility characterized by the sperm’s inability to perform the acrosome reaction and by an elevated level of oxidative DNA damage [24,25]. The double-knockout of Prnd and Prnp (Prnd\(^{0/0}\)/Prnp\(^{0/0}\)) only mimicked the effect of the Prnd single inactivation [25]. Immunohistochemical studies of Dpl were performed in gonads of various species, such as humans, rodents, boars and bovidae. The cellular localization of Dpl depends on the maturation stage of the gonads, on the studied species and the antibodies [12]. For instance, in rodents and sheep, Dpl was only detected in germinal and somatic cells in mature testis, whereas in humans, boars and bovine, DPL seems to be present during most of the developing stages of the germ cells and in the Sertoli cells of foetal and mature gonads [26–29]. In goats and bovine, DPL was detected both in immature testis and in young female follicles [28,30]. Nevertheless, these different observations suggested a role of Dpl in early and/or mature sex differentiation [12].

To get deeper into the potential role of the prion protein gene family during gonad development, we report the comparative expression profiles of the three members of the prion protein gene family and the comparative localizations of their encoded proteins during ovari and testis development in two different species: goats and mice. These data suggest that Prnd may exert a yet unknown specific role in goat foetal Leydig cells.

**Materials and methods**

**Animals and tissue samples**

Procedures for handling goats were conducted in compliance with the guidelines for Care and Use of Agricultural Animals in Agricultural Research and Teaching (authorization no. 78–34). All goat foetuses and young goats were obtained from pregnant females, following hormonal treatment as previously described [31].

For mice, animal experiments were carried out in strict accordance with the recommendations in the guidelines of the Code for methods and Welfare Considerations in Behavioral Research with Animals (Directive 86/609EC). Experiments were approved by the Local ethics committee of Jouy-en-Josas on the Ethics of Animals Experiments of the author’s institution, INRA ( Permit Number RTA06-091). All transgenic animal manipulations were performed according to the recommendations of the Haut Conseil des Biotechnologies (Permit number 6461). All mouse foetuses and pups were obtained from pregnant FVB/N, FVB/N Prnp\(^{0/0}\) [quoted in 17] and Prnd\(^{0/0}\) females [24].

Day 0 post coitum corresponds to the day of mating. The genetic sex of all foetuses was determined by PCR amplification of SRY and ZFY/ZFX genes, on liver genomic DNA [31]. For each goat sample, one gonad was frozen in liquid nitrogen for molecular analysis; the other was fixed for immuno-histological studies. For mice, samples of a same sex were pooled before molecular analysis. Two or 3 gonads were fixed for immuno-histological studies at each developmental stage. Table 1 summarized the number of individuals used at each developmental stage in mice and goats.

**PCR primers**

PCR primers were designed using primer express Software for Real-Time PCR 3.0 (Applied Biosystems, ThermoFisher Scientific, Courtaboeuf, France) analysis of Prnp, Prnd and Sprn expression in mice and goats (Table 2). Mice and goats gene sequences were obtained from GenBank. Primer efficiencies and specificities were evaluated on genomic DNA. The chosen sets of primers share similar efficiencies (not below 90%).

**Quantitative RT-PCR**

RNAs were extracted using the RNeasy Mini kit (Qiagen, Courtaboeuf, France). Super-Script II (Invitrogen, Thermo-Fisher Scientific) was used to synthesize cDNA for qRT-PCR from 1µg (mice) or 2 µg (goats) of gonad RNA (Table 1). To identify appropriate qRT-PCR normalizing genes for foetal and postnatal gonads in mice, expression stability of seven genes (Gapdh, Actb, B2 m, Mapk1, H2afz, Ywhaz and Hprt1) was tested at each time point and the GeNorm program [32] used to select a combination of the most stable genes. The three retained genes were Ywhaz, H2afz and Hprt1 (Table 2). For goats, the previously described ACTB, YWhAZ and H2AFZ genes were used [33] (Table 2). qRT-PCR was performed on all genes at all time points, in triplicates, using the Absolute Blue SYBR Green ROX mix (ThermoFisher Scientific) and the StepOnePlus Real-Time PCR System (Applied Biosystems). The results were analysed...
| Species | Stages | Sex | Tissues | Number of individual | Number of independent RT |
|----------|--------|-----|---------|----------------------|--------------------------|
| Mouse   | 12.5 dpc | Female | 2 Gonads + Mesonephros | 9 | 2 |
|         |        | Male | 2 Gonads + Mesonephros | 7 | 2 |
|         | 13.5 dpc | Female | 2 Gonads + Mesonephros | 6 | 2 |
|         |        | Male | 2 Gonads + Mesonephros | 12 | 2 |
|         | 14.5 dpc | Female | 2 Gonads | 4 | 2 |
|         |        | Male | 2 Gonads | 6 | 2 |
|         | 18.5 dpc | Female | 2 Gonads | 6 | 2 |
|         |        | Male | 2 Gonads | 9 | 2 |
|         | 5 dpp | Female | 2 Gonads | 11 | 2 |
|         |        | Male | 1 Gonad | 4 | 2 |
|         | 25 dpp | Female | 1 Gonad | 5 | 2 |
|         |        | Male | 1/2 Gonad | 4 | 2 |
|         | 50 dpp | Female | 1 Gonad | 3 | 3 |
|         |        | Male | 1/2 Gonad | 3 | 3 |
|         | 4 mpp | Female | 1 Gonad | 3 | 3 |
|         |        | Male | 1/2 Gonad | 3 | 3 |
|         | 6 mpp | Female | 1 Gonad | 2 | 2 |
|         |        | Male | 1/2 Gonad | 3 | 3 |
|         | 6–8 mpp | Female | 1 Gonad | 3 | 3 |
|         |        | Male | 1/2 Gonad | 3 | 3 |
|         | 10–12 mpp | Female | 1 Gonad | 3 | 3 |
|         |        | Male | 1/2 Gonad | 3 | 3 |
| Goat    | 30 dpc | Female | 1 Gonads + Mesonephros | 1 | 2 |
|         |        | Male | 1 Gonads + Mesonephros | 2 | 4 |
|         | 32 dpc | Female | 1 Gonads + Mesonephros | 1 | 2 |
|         |        | Male | 1 Gonads + Mesonephros | 2 | 4 |
|         | 34 dpc | Female | 1 Gonads + Mesonephros | 2 | 3 |
|         |        | Male | 1 Gonads + Mesonephros | 2 | 4 |
|         | 36 dpc | Female | 1 Gonads + Mesonephros | 3 | 3 |
|         |        | Male | 1 Gonads + Mesonephros | 2 | 4 |
|         | 41 dpc | Female | 1 Gonad | 4 | 2 |
|         |        | Male | 1 Gonad | 4 | 4 |
|         | 45 dpc | Female | 1 Gonad | 9 | 3 |
|         |        | Male | 1 Gonad | 4 | 4 |
|         | 50 dpc | Female | 1 Gonad | 4 | 3 |
|         |        | Male | 1 Gonad | 3 | 4 |
|         | 60 dpc | Female | 1 Gonad | 2 | 3 |
|         |        | Male | 1 Gonad | 2 | 4 |
|         | 70 dpc | Female | 1 Gonad | 2 | 3 |
|         |        | Male | 1 Gonad | 2 | 4 |
|         | 90 dpc | Female | 1 Gonad | 2 | 3 |
|         |        | Male | Piece of gonad | 2 | 4 |
|         | 130 dpc | Female | 1 Gonad | 1 | 3 |
|         |        | Male | Piece of gonad | 1 | 2 |
|         | 5 dpp | Female | Piece of gonad | 1 | 3 |
|         |        | Male | Piece of gonad | 1 | 3 |
|         | 1 mpp | Female | Piece of gonad | 1 | 3 |
|         |        | Male | Piece of gonad | 1 | 4 |
|         | 3 mpp | Female | Piece of gonad | 1 | 3 |
|         |        | Male | Piece of gonad | 1 | 4 |
|         | 7 mpp | Female | Piece of gonad | 1 | 3 |
|         |        | Male | Piece of gonad | 1 | 4 |
|         | 4 ypp | Male | Piece of gonad | 1 | 3 |

* Three different pieces of the same testis was used to realize three independent RT.
by the relative standard curve method with the qBASE Software [34]. Data points were plotted using Excel. Statistical analyses were performed using the INVIVOSTAT software [35] that combines an ANOVA approach followed by a Fisher’s Least Significant Difference (LSD)-test.

**Immunostaining**

Freshly dissected gonads were fixed in 4% paraformaldehyde in phosphate saline buffer (PBS) at 4 °C for 1 h or overnight (according to the size of the gonad). After washes in PBS with increasing concentrations of sucrose (0, 12%, 15% and 18%), tissue specimens were embedded in Jung Tissue Freezing Medium (Leica Biosystems, Nanterre, France) and frozen at −80 °C. Cryo-sections (7 μm thick) were obtained and stored at −80 °C until used. The sections were air-dried, rehydrated in PBS and permeabilized during 30 min in PBS with 0.5% triton and 1% BSA. The primary antibodies were then applied overnight at 4 °C. Table 3 describes the antibody references and concentrations [15,28,36–38]. After several washes, the sections were incubated with secondary antibodies for 1 h at room temperature. The slides were then rinsed in PBS, mounted in Vectashield mounting medium with DAPI (Vector) and observed as above.

**Western blot analysis**

Frozen tissues of adult mice and goats were homogenized in 50 mM Tris HCl, 150 mM NaCl, 0.5% sodium deoxycholate (w/v), 0.1% sodium deoxychylate (w/v), 1% of a nonionic non-denaturing detergent (NP-40), one complete EDTA free mini-protease inhibitor tablet per 10 mL (Roche Diagnostic, Saint-Egrève, France). Whole extracts (20 μg of total protein) were subject to 4–15% gradient SDS/PAGE and transferred to a poly(vinylidene fluoride) membrane (GE healthcare Life Sciences, Velizy-Villacoublay, France). The membrane was probed with anti-biotinylated-PrP antibody (bSha 31; Table 3). The secondary antibody used was horseradish peroxidase/streptavidin conjugated antirabbit (Table 3). Immunodetection using the enhanced chemiluminescence (ECL) method (PIERCE) was performed according to the manufacturer’s instruction and the images were recorded on an image analysis station (Luminescent Image analyse Las-1000plus Fujifilm).

**Results and discussion**

**By contrast with mice, PRND is highly expressed in goat foetal testis**

We have previously carried out an expressional study of the Prion gene family in the goat species [30] that
suggested an involvement of PRND in early gonadal differentiation. The aim of the present study was to complete this observation by (i) including the recently discovered SPRN gene and (ii) establishing a comparative view of the expression of the three members of the Prion gene family throughout all gonad developmental stages, from differentiation to adulthood (Fig. 1A), in (iii) two mammalian species, mice and goats. Gonadal expression profiles of PRNP, SPRN and PRND have been precisely determined using quantitative RT-PCR instead of classical RT-PCR as previously used [30], at 7 and 15 developmental stages in mice and goats respectively (Fig. 1B–G), and during ageing in mice where four additional stages were studied. A stable gonadal expression of these three genes in the mouse species was observed during ageing since 50 dpp until 10–12 months of age (Fig. 2). In both species, Prnp/PRNP gene expression slightly increases during development and appears to be more intense around birth (Fig. 1B,C). During gonadal development, SPRN was found to be expressed in male and female gonads of mice and goats at all tested stages (Fig. 1B–E) but only faintly when compared to PRNP and PRND (as for example in goat testis samples, the cycle thresholds, CT, values are for PRNP and PRND between 21 and 30, but only between 31 and 36 for SPRN). Sprn/SPRN appears to be more expressed in ovaries of both species during early follicles formation (i.e. from 18.5 dpc to 5 dpp in mice and 70–90 dpc in goats); and specifically in goat ovaries before the beginning of germ cell meiosis (Fig. 1D,E). Indeed, the highest SPRN expression level is found at this premeiotic stage (i.e. 41–50 dpc) only in the goat species. The duration of this premeiotic stage, which starts after gonad commitment in one sex or the other (i.e. 12.5 dpc in mice, 36 dpc in goats), appears quite different in mice and goats, 24 h in mice instead of 2 weeks in goats. Although no profound change could be noticed in mouse ovaries during this period, goat ovaries were organized into cortical and medullar compartments where germ cells were concentrated in the cortex and estrogens were produced by the medulla part under the control of the FOXL2 gene that has been shown to be a major ovarian-determining gene in goats, by contrast with mice [39–41].

Finally and as previously noticed [30], Prnd/PRND expression is found higher in testes than in ovaries at adulthood for both species, but this sex-dimorphism appears since early testicular differentiation specifically in the goat species (Fig. 1F,G).

**Table 3. Antibodies used in the present study.**

| Primary antibody | Reference | Source | IHC dilution | WB dilution |
|------------------|-----------|--------|--------------|-------------|
| DPL – purified boDpl67-81 | Gift of Dr Paltrinieri (Rondena et al. [28]) | Rabbit | 1 : 150 | |
| PRP – Sha31 | (Feraudet et al. [36]) | Mouse | 1 : 1000 | |
| Biotinylated PRP – b Sha31 | (Feraudet et al. [36]) | Mouse | 1 : 50 000 | |
| Murine SHO | Gift of Dr Westaway (Watts et al. [15]) | Rabbit | 1 : 500 | |
| Ovine SHO | Gift of Dr Peelman (Lampo et al. [37]) | Rabbit | 1 : 50 | |
| Cyp17A | Gift of Dr Hales (Hales et al.[38]) | Rabbit | 1 : 200 | |

| Secondary antibody | Reference | Conjugate | IHC dilution | WB dilution |
|-------------------|-----------|-----------|--------------|-------------|
| Anti-rabbit IgG-Dylight 488 | 072-03-15-06, KPL | Fluorescein | 1 : 200 | |
| Anti-rabbit IgG-Dylight 594 | 072-09-15-06, KPL | Fluorescein | 1 : 200 | |
| Anti-mouse IgG-Cy3 | AP160C, Millipore | Cyanine 3 | 1 : 200 | |
| Anti-rabbit IgG-B | BA-1000, Vector | Biotin | 1 : 200 | |
| Anti-mouse-IgG-HRP strep | Horseradish peroxidase | Streptavidin | 1/20 000 | |
| Fluorescein Streptavidin | SA-5001, Vector | | 1 : 200 | |
| Texas Red® Streptavidin | SA-5006, Vector | | 1 : 200 | |

| Mounting medium | Reference |
|-----------------|-----------|
| Vectashield for fluorescence with DAPI | H-1200, Vector |
Fig. 1. Expression of Prion family genes during gonad development. (A) Chronology of gonad differentiation in mice and goats. Expression of the Prnp/PRNP (B, C), Sprn/SPRN (D, E) and Prnd/PRND (F, G) genes was quantified using real-time RT-PCR in ovaries (pink histograms) and testes (blue histograms) in mice (B, D, F) and goats (C, E, G). From 12.5 dpc to 13.5 dpc (mice) and 30 dpc to 36 dpc (goats): gonad + mesonephros, others stages: gonad only. Prnp/PRNP, PrP protein gene; Sprn/SPRN, Sho gene; Prnd/PRND, Dpl gene; dpc, days post coitum; dpp, days post partum; mpp, months post partum. Values are expressed in percentage according to the highest one noted 100%. Means ± SD were plotted. Planned comparisons were made on the predicted means with ‘sex’ and ‘stage’ as treatment factors (two-way ANOVA approach, followed by a Fisher’s LSD-test). For each stage, significant differences between the two sexes are showed by stars (*P-value ≤ 0.05; **P-value ≤ 0.01; ***P-value < 0.001).
Sho and PrP cellular localizations remain difficult to ascertain

In order to gain more information on the 3 proteins encoded by the 3 Prion gene family members, we carried out IHC (Immuno-Histo-Chemistry) and western blot experiments using available antibodies (Table 3). Two antibodies have been tested against Sho. Each gives a specific staining by IHC that remains different (i) from each other, (ii) from one species to another and also (iii) from Sho gonadal expression already described by additive transgenesis of a Sprn-LacZ mini-gene in mice [16]. Thus, we are unable to ascer-
tain in what gonadal cell type Sho could be detected. The recent derivation of Sho-knockout mice [19] might help to decipher the real expression pattern of this protein by comparative IHC analysis, but these mice were not available to us at the time of this experiment.

In the same way, the PrP protein was detected by IHC and western blot at adulthood in both species and at 130 d_{pc} in goats (Fig. 3). By discarding the strong nonspecific staining in the interstitial area of adult mouse testes, identified by using Prnp^{0/0} testicular samples (Fig. 3A), the PrP protein appears to be present at all stages mainly inside the seminiferous tubules, most likely in the Sertoli cells (clearly visible at 130 d_{pc} in goats) and in the germ cells at the end of spermiogenesis (see the staining of elongated spermatids on Fig. 3B,C). Presence of PrP in the testis was confirmed by western blot performed on adult mice and goats testes and sperm cells (Fig. 3F). No protein was detected in Prnp^{0/0} brain mice used as negative control.

**By contrast with mice, DPL is detected in goat Leydig cells**

We have previously shown that the DPL protein could be detected in germ cells of both sexes and in foetal Leydig cells of early goats developing testes at 44 and 62 d_{pc} [30]. According to qRT-PCR results (Fig. 1F,G) showing a high expression of PRND in goat developing testes, we checked its cellular localization throughout testis development in immature and mature testes and on spermatozoa of both species (Fig. 4). First, the specificity of anti-DPL antibody boDpl67-81 [28] was confirmed as no signal was obtained on adult Prnd^{0/0} mouse testes (Fig. 4A,B). Using this antibody, DPL is detected in the cytoplasm of some germ cells in both species; germ cells that have passed the zygotene stage of meiosis (i.e. pachytene and spermatids)(Fig. 4C,F and K,N) and its presence persists on spermatozoa with a high staining of the acrosomal vesicle (Fig. 4G,H and O.
These results are in complete agreement with the role of DPL in spermiogenesis [24,25].

More interestingly, a strong staining is detected in some cells of the interstitial testicular space, specifically in the goat species and at immature stages, as the staining disappears in adult testes (compare Fig. 4I,L with M,N). In order to precisely define this DPL staining in the interstitial testicular compartment, we carried out double IHC with an anti-CYP17 antibody detecting a Leydig cell-specific marker corresponding to a key enzyme of steroid synthesis, the cytochrome P450 17alpha-hydroxylase/17,20-lyase [38]. DPL and CYP17 are found colocalized in the same interstitial cells (Fig. 5A–F). DPL staining in goat testes disap-
pears between the prepubertal 3-month and the pubertal 7-month stages (Fig. 4K–N), suggesting that PRND is specifically expressed in the foetal Leydig cell population. These cells disappear after birth and are replaced by adult Leydig cells at puberty [42]. Finally, the PRND expression profile appears hugely similar to that of 3βHSD, another Leydig cell-specific marker, until 1 month after birth indicating that during testis development the major part of PRND expression is Leydig specific (Fig. 5G). This observation could explain the high testicular levels of PRND transcripts detected from 41 dpc to 1 mpp specifically in goat testes [41] and not in the mouse testes of the corresponding stages, from 12.5 dpc to 5 dpp (Fig. 1F). 

During the prepubertal period, spermatogenesis starts (25 dpp in mice, 3 mpp in goats), the number of germ cells increases and more importantly differentiated meiotic germ cells expressing PRND/Prnd appear. From these prepubertal stages, the PRND/Prnd expression profiles are similar between mice and goats (Fig. 1F,G), but diverge from that of 3βHSD in goats (Fig. 5G). It indicates that around puberty PRND expression increases in postmeiotic germ cells and disappears from foetal Leydig cells because of their own disappearance.

In conclusion, we report the differential expression of the three members of the prion protein gene family in the developing gonads of mice and goats. Only relatively low levels of expression were detected for Sho, an observation that might relate with the lack of reproductive-associated phenotype in mouse Sprn-knockout mice [8,19]. By contrast and in addition to its conserved role in spermiogenesis, PRND seems to be a key marker of foetal Leydig cells in goats. This observation adds PRND to the list of genes having potentially different biological roles in mice and humans or ruminants (see [39,43] for recent examples). The present results highlight that PRND is a key candidate gene for functional studies in goats because its involvement in foetal Leydig cells cannot be studied in the widely used mouse mammalian model. Deciphering this role may have important implications for human reproduction. In order to determine if PRND could be a crucial actor of foetal Leydig cell develop-
ment, its targeting is currently under way in the goat species by using genome editing technologies, recently proven successful by us in this species [39].

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Author contributions

AAB, MP, KMG, JLV and EP participated in the conception of the study, interpretation of data and in the drafting of the article. BP and JC participated in the design and realization of animal experiments. AAB, ME, MA and FMT performed experiments.

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