

Seasonal expression of androgen receptor, aromatase, and estrogen receptor alpha and beta in the testis of the wild ground squirrel (*Citellus dauricus* Brandt)

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Abstract

The aim of this study was to investigate the seasonal expression of androgen receptor (AR), estrogen receptors α and β (ER α and ER β) and aromatase cytochrome P450 (P450arom) mRNA and protein by real-time PCR and immunohistochemistry in the wild ground squirrel (WGS) testes. Histologically, all types of spermatogenic cells including mature spermatozoa were identified in the breeding season (April), while spermatogonia and primary spermatocytes were observed in the nonbreeding season (June), and spermatogonia, primary spermatocytes and secondary spermatocytes were found in pre-hibernation (September). AR was present in Leydig cells, peritubular myoid cells and Sertoli cells in the breeding season and pre-hibernation with more intense staining in the breeding season, whereas AR was only found in Leydig cells in the nonbreeding season; P450arom was expressed in Leydig cells, Sertoli cells and germ cells during the breeding season, whereas P450arom was found in Leydig cells and Sertoli cells during pre-hibernation, but P450arom was not present in the nonbreeding season; Stronger immunohistochemical signal for ER α was present in Sertoli cells and Leydig cells during the breeding season; ER β was only expressed in Leydig cells of the breeding season. Consistent with the immunohistochemical results, the mean mRNA level of AR, P450arom, ER α and ER β were higher in the testes of the breeding season when compared to pre-hibernation and the nonbreeding season. These results suggested that the seasonal changes in spermatogenesis and testicular recrudescence and regression process in WGSs might be correlated with expression levels of AR, P450arom and ERs, and that estrogen and androgen may play an important autocrine/paracrine role to regulate seasonal testicular function.

Introduction

In mammalian testes, testosterone appears to be responsible for maintaining adequate blood-testis barrier function,¹ inducing meiosis and postmeiotic development of germ cells^{2,3} and inhibition of germ cell apoptosis.⁴ It has been clearly established that the AR is expressed in Sertoli, Leydig and peritubular cells. However, immunodetection of the AR in testicular germ cells is controversial, with reports indicating its detection and absence, although functional AR in germ cells is not essential for spermatogenesis and male fertility.⁵ Testosterone is converted in target cells to estradiol by the enzyme aromatase. Aromatase is a member of the cytochrome P450 superfamily that catalyzes the conversion of androgens (C19), namely testosterone and androstenedione, into estrogens (C18), estradiol and estrone, respectively. The presence of P450arom in the testes has been reported in numerous species, including American black bears,⁶ raccoon dogs,⁷ Shiba goats,⁸ brown bears,⁹ Göttingen miniature pigs,¹⁰ bank voles,¹¹ rats,¹² mice,¹³ ground squirrels¹⁴ and muskrats.¹⁵ These evidences indicated that estrogens act on the initiation and maintenance of spermatogenesis and on germinal stem cell division and survival in a paracrine/autocrine manner(s).

Estrogen action is displayed by means of two different estrogen receptors (ERs), estrogen receptor-alpha (ER α) and estrogen receptor-beta (ER β), localized in the different testicular cells types. The localization of ERs in testicular cells is not only species-specific but also varies depending on the type of receptor and the developmental stage of the germ cell.¹⁶ In most species analyzed, such as human, rat, cat and dog, ER α and ER β co-localize in spermatogonia, spermatocytes and spermatids as well as in Sertoli, Leydig, and peritubular myoid cells and these locations appeared to change with age in some reports.^{17,18} In other species, such as the boar, ER α and ER β localize separately to spermatogonia/primary spermatocytes and Sertoli cells, respectively.¹⁹ Thus, ER α and ER β , and their localization and function in testicular cells vary depending on the species, cellular developmental stage and type of receptor.

The wild ground squirrel (*Citellus dauricus* Brandt, WGS) is a typical seasonal breeder which has a strict and extremely compressed breeding period from April to May, a long period of sexual dormancy from June to the following March, and a 6-month hibernation (from October to March).^{20,21} The testis and epididymis of this species exhibits a distinct seasonal morphology changes from the breeding season to the nonbreeding season.²⁰⁻²³ Our published results have indicated that

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immunoreactivity of P450c17 and P450arom in testicular tissues was accompanied by changes of testicular activity during the breeding and nonbreeding seasons.¹⁴ In this study, we investigated the immunohistochemical localization of AR, P450arom, ER α and ER β as well as their mRNA expression in the breeding, nonbreeding seasons and pre-hibernation, to gain insight of the relation between steroid hormones and testicular function throughout the reproductive cycle of WGS.

Materials and Methods

Animals

The wild male ground squirrels were captured by box traps in September 24 (n=12) of 2012 and in April 13 (n=15) and June 27 (n=10) of 2013 in Hebei Province, China. WGSs were thought to be adult based on their body weights (242-412 g). All procedures involving animals were carried out in accordance with the Policy on the Care and Use of

Animals by the Ethical Committee, Beijing Forestry University and approved by the Department of Agriculture of Hebei Province, China (JNZF11/2007). An overdose of pentobarbital (BioDee Co., Beijing, China) was applied afterwards for euthanasia. Testicular tissues were excised quickly after necropsy. On necropsy, the testes were excised, weighed, measured and tissues fixed in 4% paraformaldehyde in 0.05 M PBS (pH 7.4) for histological and immunohistochemical observations, and the others were immediately frozen in liquid nitrogen and stored at -80°C until used for RNA isolation.

Histology

Tissue samples were dehydrated in ethanol series and embedded in paraffin wax. Serial sections (4 μm) were mounted on slides coated with poly-L-lysine (Sigma Chemical Co., St. Louis, MO, USA) and stained with hematoxylin-eosin (HE) for general histology observation. The sections were screened using an Olympus photomicroscope with a $\times 20$ objective lens and imaged with software Image-Pro Plus 4.5 (Media Cybernetics, Bethesda, MD, USA).

Evaluation of spermatogenesis

The score count for determining the stage of spermatogenesis¹⁴ was modified and used to evaluate spermatogenesis. The modified criteria of the score count (Table 1), principally based on the most advanced spermatogenetic cells. Counting was usually performed while the whole sample was examined under a light microscope using a low (100 \times) and then high (400 \times) magnification.

Antibodies

The primary antibodies used in the present study included rabbit polyclonal anti-AR (sc-816, Santa Cruz Biotechnology, Santa Cruz, CA, USA), rabbit polyclonal anti-ER α (sc-542, Santa Cruz Biotechnology), rabbit polyclonal anti-ER β (sc-8974, Santa Cruz Biotechnology), rabbit polyclonal anti-P450arom (derived from human Aromatase C-terminus, sc-30086, Santa Cruz Biotechnology). The dilution ranges of AR, ER α , ER β and Aromatase antibodies for immunohistochemistry were all 1:500. The specificity of AR, ER α , ER β and P450arom antibodies have been described in our previous studies on WGS.^{14,24} The immunohistochemistry kits with the secondary antibody of goat anti-rabbit was applied corresponding with the primary antisera.

Immunohistochemistry

Testicular serial paraffin sections were incubated with 10% normal bovine serum to reduce background staining caused by the secondary antibody, the sections were then incubated with primary antibody for 12 h at 4°C . Subsequent incubation with the secondary antibody, goat

anti-rabbit IgG conjugated with biotin and peroxidase with avidin, using rabbit ExtrAvidinTM Peroxidase staining kit (Sigma Chemical Co., St. Louis, MO, USA) was performed, followed by visualizing with 30 mg 3,3-diaminobenzidine (Wako, Tokyo, Japan) solution in 150 mL of 0.05 mol Tris-HCl buffer, pH 7.6, plus 30 μL H_2O_2 . Only the testicular sections reacted with anti-P450arom were counterstained with haematoxylin solution (Merck, Tokyo, Japan). Control sections were treated with normal bovine serum (Sigma Chemical) instead of the primary antisera. The negative controls were divided into two groups, stained with the haematoxylin or not, to be as the controls for P450arom and controls for AR, ER α and ER β respectively. The immunostained slides were scanned using the software Image-Pro Plus 4.5 (Media Cybernetics, Rockville, MD, USA) at 40 \times magnification. The immunohistochemical staining was determined as positive (+), strong positive (++) , very strong positive (+++), and negative (-). Staining that was weak but higher than control was set as positive (+); the highest intensity staining was set as very strong positive (+++); staining intensity between + and +++ was set as strong positive (++) .

Total RNA isolation and cDNA synthesis

Total RNA was isolated from testicular tissues of wide ground squirrels using Trizol[®] Reagent (Invitrogen, Carlsbad, CA, USA). Approximately 0.1 g of testicular tissues were thawed and

immediately homogenized in 1 mL of TRIzolTM Reagent. The homogenate was incubated for 5 min at room temperature to allow the complete dissociation of nucleoprotein complexes. After the addition of 0.2 mL of chloroform, the mixture was vigorously shaken for 15 s at room temperature and centrifuged at 12,000g for 15 min at 4°C . The aqueous phase was then transferred to a fresh tube and an equal volume of isopropanol was added. Then the sample was kept for 10 min at room temperature. RNA was precipitated by centrifugation at 12,000g for 10 min at 4°C . The RNA pellet was washed twice with 70% ethanol and dissolved in 50 μL of diethylprocarbonate-treated water. The integrity of RNA was tested by gel electrophoresis and its concentration was measured with spectrophotometer.

Real-time PCR

The first-strand cDNA from total RNA was synthesized using Superscript II Reverse Transcriptase (Invitrogen) and Oligo (dT) according to the manufacturer's protocol. The 20 μL of reaction mixture contained 4 μg of total RNA, 0.5 μg of oligo (dT), 2.5 mM MgCl_2 , 0.5 mM dNTP, 10 mM dithiothreitol, 20 mM Tris-HCl (pH 8.4) and 200 U of Superscript II enzyme. Real-time PCR was utilised to quantify targeted cDNAs using an ABI PRISM 7900HT system (Applied Biosystems, Foster City, CA, USA). Oligonucleotide primers were designed using Web-based Primer3 software; the primers are listed in Table 2. Polymerase chain reaction (PCR) was carried out using ABI PRISM[®]

Table 1. The modified criteria of the score count.

Most mature germ cell type	Score
Spermatogonia	1
Primary spermatocytes	2
Secondary spermatocytes	3
Round spermatid	4
Mature-phase spermatozoa	5

Table 2. Oligonucleotide primers used for quantitative real time PCR.

Gene	Sequence of primer (5'-3')	Product size (bp)
AR	F: TGGGACCTTGGATGGAGAAC R: CTCCGTAGTGACAGCCAGAA	150
ERalpha	F: GCATGATGAAAAGCGGCATA R: AAGGACAAGGCAGGGCTATT	184
ERbeta	F: TCTGGGTGATTGCGAAGAGT R: CCCCAGATTGAGGACTTGT	215
CYP19	F: CAGAGGTCGAAGCAGCAATC R: AACCGGGTAGCCGTCAATTA	188
β -actin	F: TTGTGATGGACTCCGGAGAC R: TGATGTCACGCACGATTTC	186

7500 Fast Real-Time PCR System (Applied Biosystems) according to manufacturer's instruction, and levels of each target mRNA relative to β -actin mRNA were determined using the $2^{-\Delta\Delta CT}$ method. All quantitative reactions were subjected to 95°C for 30 sec, followed by 35 cycles of 95°C for 5 s, 55°C for 10 s, and 72°C for 10 s. Melting curve analysis was applied to all reactions to ensure homogeneity of the reaction product. In addition, the amplified size was checked by electrophoresis and then sequenced. The experiments were repeated four times. The purified PCR products were ligated into pCR 2.1-TOPO (Invitrogen) and the ligation products were used to transform the competent *Escherichia coli* using TOPO TA Cloning Kit (Invitrogen). Plasmids were extracted from the bacteria and positive clones containing the proper insert were sequenced in both directions using Thermo Sequenase II Dye Terminator Cycle Sequencing Premix Kit (Amersham Pharmacia Biotech, Amersham, UK) with an automatic sequencing system (ABI PRISM 377, Applied Biosystems Japan, Tokyo, Japan).

Statistical analysis

Statistical analysis was calculated and analyzed by Student's *t*-test using the SPSS computer package. Values of $P < 0.05$ were considered as significant.

Results

Morphology and histology

The morphological differences of testes could be observed in the breeding season (Figure 1a left), the nonbreeding season (Figure 1a middle) and pre-hibernation (Figure 1a right). There were seasonal changes with testicular weight, size and seminiferous tubules diameter in the WGS during the breeding, the nonbreeding seasons and pre-hibernation (Figure 2). The largest values of testicular weight (Figure 2A), size (Figure 2B) and seminiferous tubules diameter (Figure 2C) were found in the breeding season (April), and the smallest values in the nonbreeding season (June). There was significant

increase in testicular weight, size and seminiferous tubules diameter from the nonbreeding season to pre-hibernation. Meanwhile, HE staining showed that all stages of sperm cells could be identified in the seminiferous epithelium in the breeding season (Figure 1b), only spermatogonium and primary spermatocyte in the nonbreeding season (Figure 1c), but there was no secondary spermatocyte in the nonbreeding season, which could be found in the seminiferous epithelium of pre-hibernation (Figure 1d). The stages of sperm cells were evaluated and summarized in Figure 2D. This score count showed that sperm cells existed in pre-hibernation were in a more mature phase than those in the nonbreeding season.

Immunohistochemistry

Immunostaining for AR, ER α , ER β and P450arom was evaluated in testicular tissues during the breeding, the nonbreeding seasons and pre-hibernation (Figure 3). The staining of AR was observed in the Leydig cell, Sertoli cell and peritubular myoid cells during the breeding season (Figure 3a) and pre-hibernation (Figure 3c), but much stronger in the

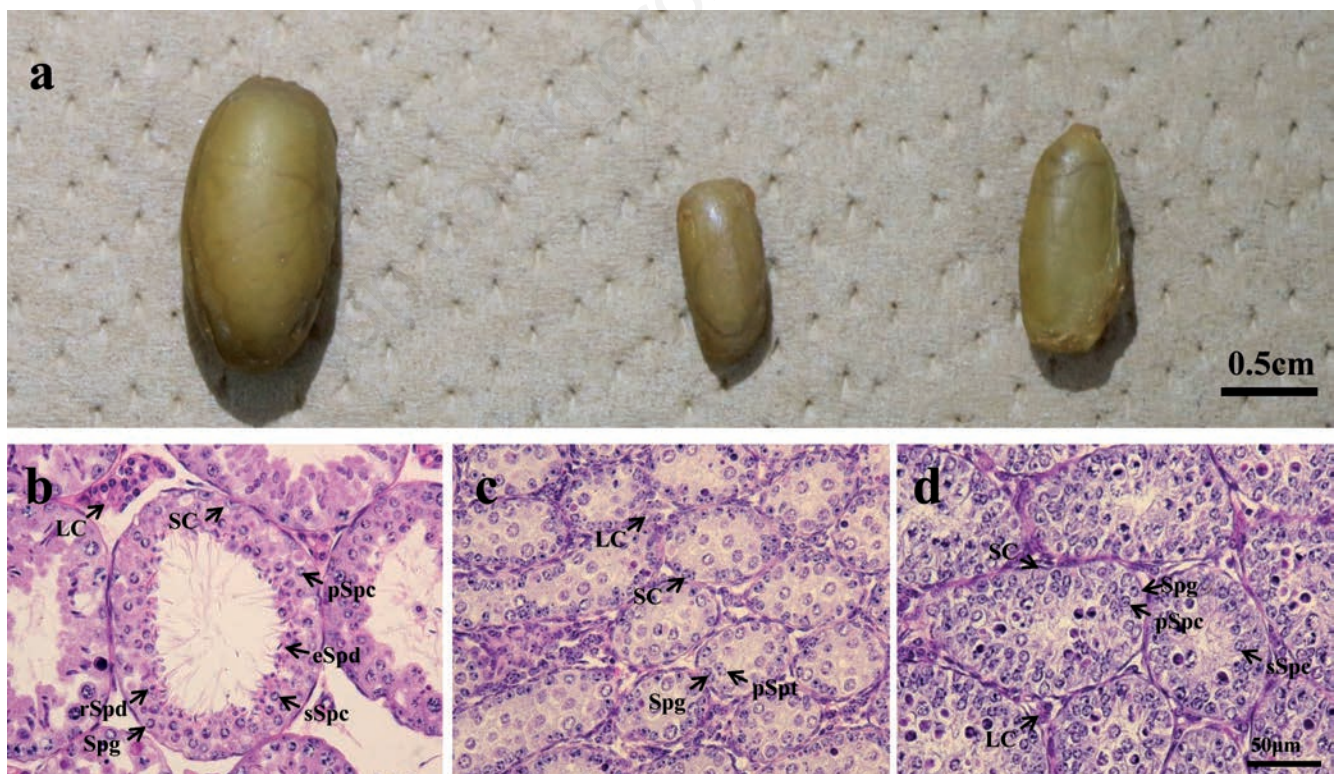


Figure 1. Seasonal changes of testicular morphology in the breeding season (a, left), nonbreeding season (a, middle) and pre-hibernation (a, right). The b, c and d were histological structure of testicular tissues of the breeding season, the nonbreeding season and pre-hibernation, respectively. SC, Sertoli cell; LC, Leydig cell; Spg, spermatogonia; pSpc, primary spermatocyte; sSpc, secondary spermatocyte; rSpd, round spermatid; eSpd, elongate spermatid.

breeding season. In the nonbreeding season, only weak immunostaining was shown in the Leydig cells (Figure 3b). Immunoreactivity of ER α was detected in the Leydig cells and Sertoli cells during the breeding season and in the Sertoli cells during pre-hibernation (Figure 3 d,f), whereas no staining was detected in the nonbreeding season (Figure 3e). With regard to ER β , positive immunostaining

was shown in the Leydig cells and weak signal was found in germ cells during breeding season (Figure 3g), yet no signal in the nonbreeding season and pre-hibernation (Figure 3 h,i). Stronger immunohistochemical signal for P450arom was present in Sertoli cells, Leydig cells and germ cells during the breeding season (Figure 3m); P450arom was also presented in Sertoli cells and Leydig cells during pre-

hibernation (Figure 3o), while there was no positive P450arom signal in the nonbreeding season (Figure 3n). No signal was seen in the negative controls without or with haematoxylin (Figure 3 j-l, p-r). The immunoreactivity of each staining was quantified and summarized in Table 3.

Table 3. Immunohistochemical localization of AR, ER α , ER β and P450arom, in testicular cells of the wild ground squirrel.

	AR			ER α			ER β			P450arom		
	B	NB	Pre-H	B	NB	Pre-H	B	NB	Pre-H	B	NB	Pre-H
Leydig cell	+++	+	++/+	++	-	-	+++	-	-	+++	-	++
Sertoli cell	+++	-	++	+++	-	++/+	-	-	-	+++	-	++

B, breeding season; NB, nonbreeding season; Pre-H, pre-hibernation; -, negative staining; +, positive staining; ++, strong positive staining; +++, very strong positive staining and/or no such cell type.

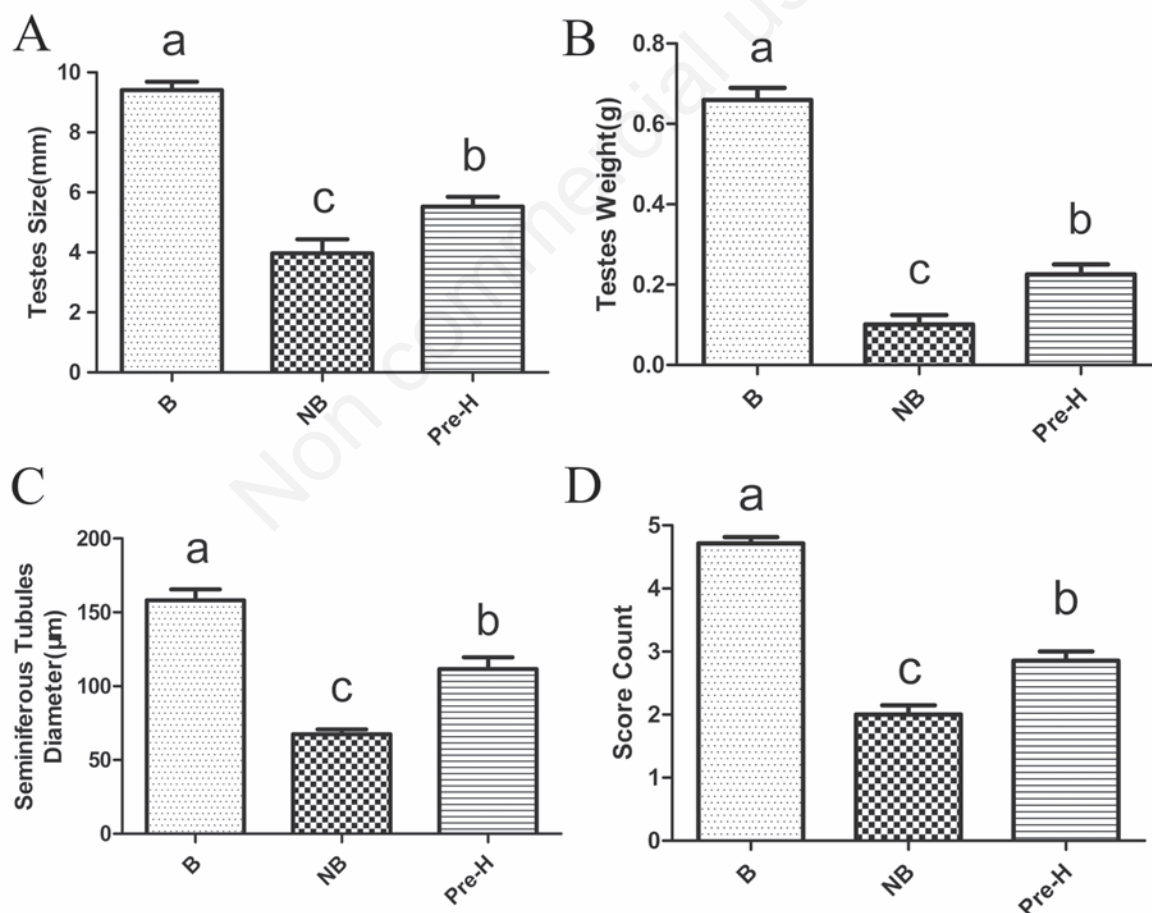


Figure 2. Changes of statistical data about testicular index. The seasonal profile of testicular weight (A), size (B), seminiferous tubule diameter (C) and spermatogenic score (D). B, the breeding season; NB, the nonbreeding season; Pre-H, pre-hibernation. Bars represent means + SD for five independent experiments. Means within the columns marked with different letters indicate significant difference ($P < 0.05$).

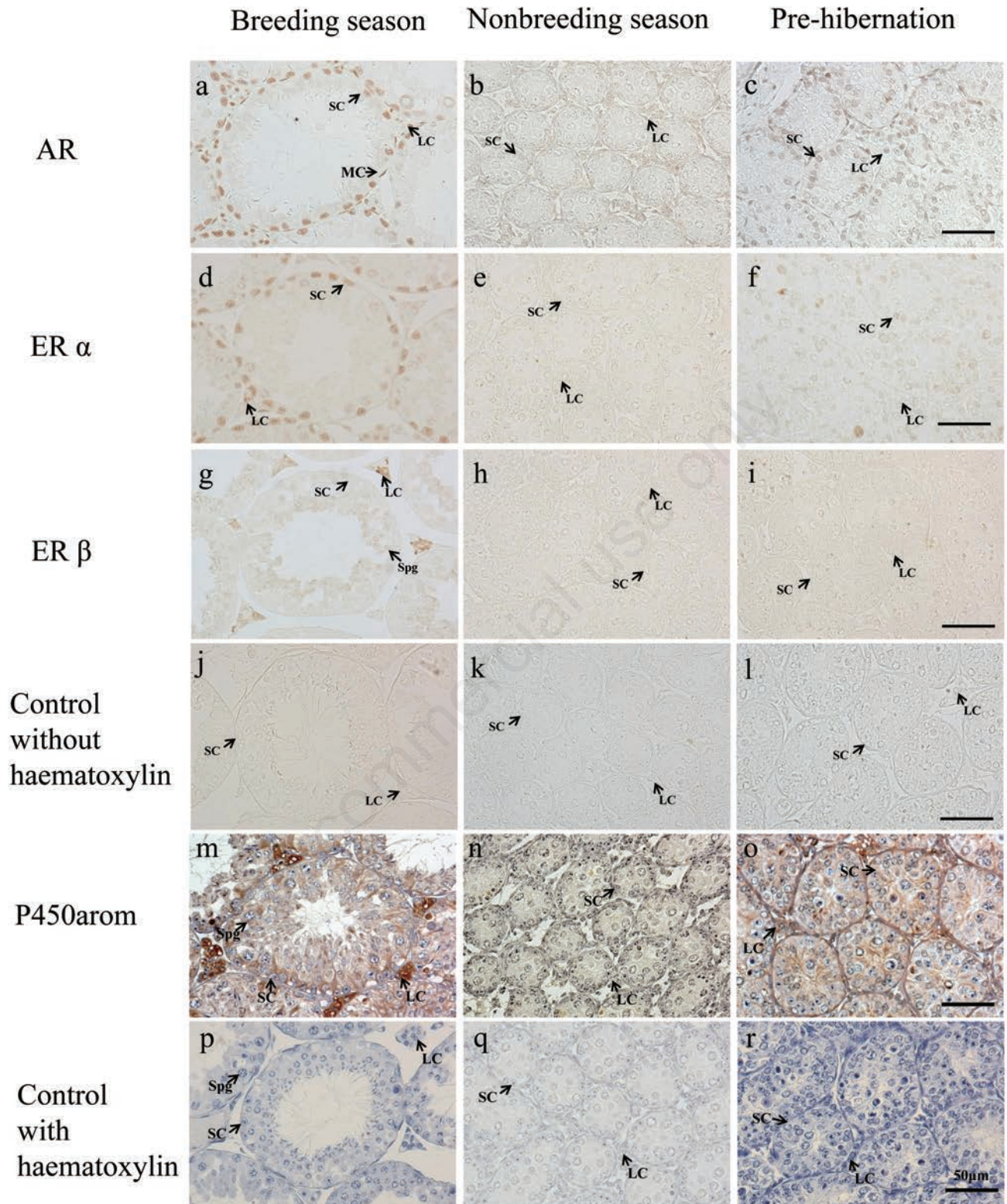


Figure 3. Immunolocalization of AR, ER α , ER β and P450arom in the testes of the wild ground squirrels. The first column (a, d, g, j, m, p) represents staining in the breeding season; the second column (b, e, h, k, n, q), and the third column (c, f, i, l, o, r) represent immunostaining in the nonbreeding season and pre-hibernation, respectively. Since AR (a-b), ER α (e-f) and ER β (g-i) were expressed in nuclear, the relative negative controls (j-l) were not stained with haematoxylin. P450arom (m-o) was shown in cytoplasm of the testes, so the relevant negative control (p-r) were stained with haematoxylin. SC, Sertoli cell; LC, Leydig cell; MC, myoid cell; Spg, spermatogonia.

Real-time PCR

The mRNA expression levels of *AR*, *P450arom*, *ER α* and *ER β* were detected and the results were shown in Figure 4. The *AR* gene was started at a high level during the breeding season, and there was no remarkable difference in expression during the rest of the reproductive cycle (Figure 4A). For *ER α* mRNA, the expression peaked in the breeding

season, rapidly reduced during the nonbreeding season, and returned to a relatively high level during pre-hibernation (Figure 4B). With regard to *ER β* gene, the level decreased drastically from the breeding to the nonbreeding seasons, and remained low in pre-hibernation (Figure 4C). The expression of *CYP19* gene was significantly higher in the breeding season. After the obvious reduction of the non-

breeding season, the level of *P450* mRNA showed a marked increase in pre-hibernation (Figure 4D). After obtaining the sequence of each PCR product, we blasted with the known mRNA sequences of mouse, rat, bovine and human, found the homologous sequence fragments in each species and compare for homology (Table 4).

Table 4. Nucleotide sequence identity in testis of wild ground squirrel in comparison with rat, mouse, human and bovine.

	Rat (%)	Mouse (%)	Human (%)	Bovine (%)
<i>AR</i>	94.67	92.76	85.33	84.67
<i>ER alpha</i>	91.85	97.28	79.89	82.07
<i>ER beta</i>	97.21	94.42	84.65	82.79
<i>CYP19</i>	94.34	87.23	80.32	81.91
<i>Beta-actin</i>	94.62	91.94	88.17	86.56

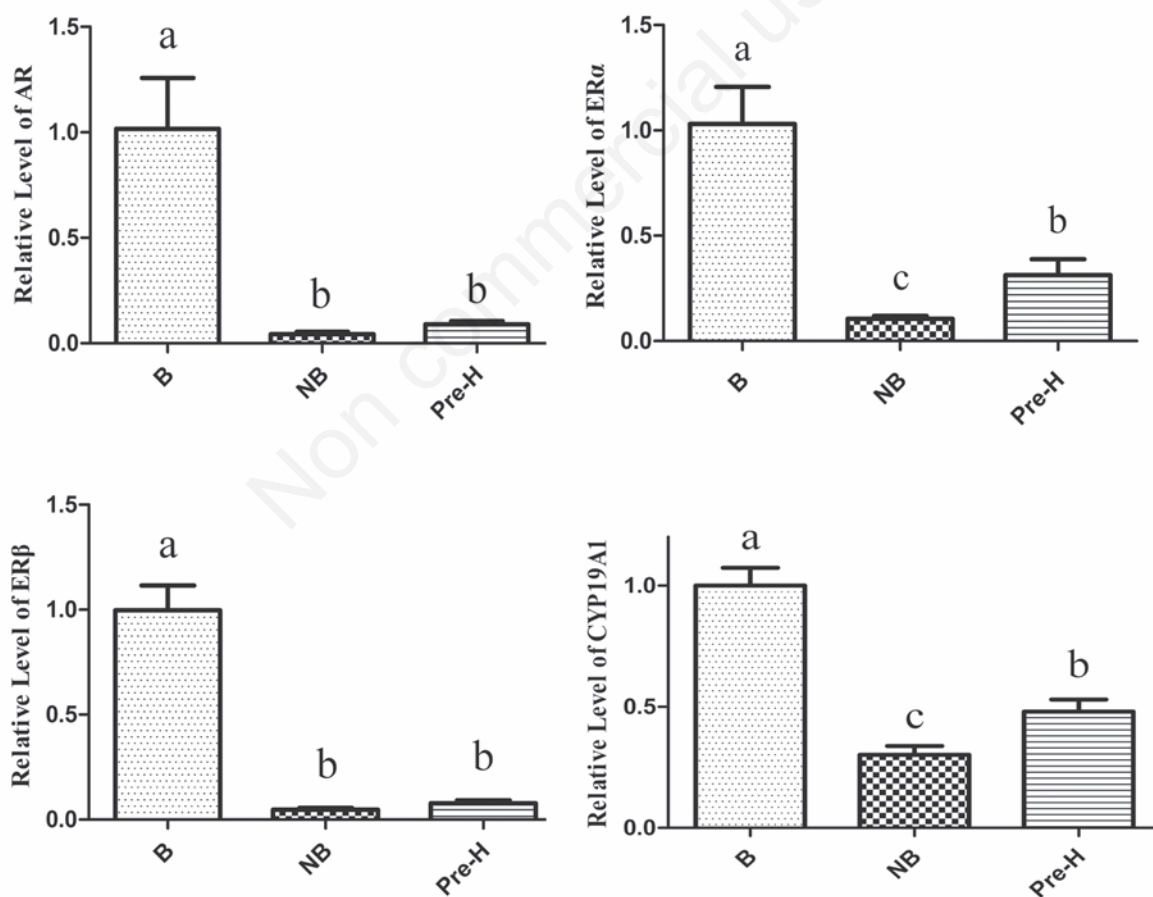


Figure 4. Real-time RT-qPCR analysis of the mRNA level of *AR*, *P450arom*, *ER α* and *ER β* during the annual cycle. The expressions of genes *AR* (A), *ER α* (B), *ER β* (C) and *CYP19* (D) showed the changes during the breeding season, the nonbreeding season and pre-hibernation. B, breeding season; NB, nonbreeding season; Pre-H, pre-hibernation. Bars represent means + SD for five independent experiments. Means within the columns marked with different letters indicate significant difference ($P < 0.05$).

Discussion

In this study, we revealed the immunolocalization and expression patterns of *AR*, *P450arom*, *ER α* and *ER β* in testes of a wild seasonal breeding rodent. We showed that the expression levels of *AR*, *P450arom*, *ER α* and *ER β* were correlated with the changes of testicular functions during the breeding, non-breeding seasons and pre-hibernation. These findings suggested that *AR*, *ER α* , *ER β* and *P450arom* may be involved in the regulation of seasonal changes in the testicular functions of the WGS.

Many of the biological actions of androgens are mediated by AR. Androgen and AR play important roles in male spermatogenesis and fertility by way of a paracrine/autocrine manner.^{25,26} AR is expressed in Sertoli cells, peritubular myoid cells, Leydig cells and perivascular smooth muscle cells of the testis depending on the species, but its presence in germ cells remains controversial. In the present study, AR was expressed in Leydig cells, Sertoli cells and peritubular myoid cells in the breeding season and pre-hibernation and Leydig cells in the nonbreeding season, but not in germ cells. These findings are consistent with previous observations in rat, horse and human testes,²⁷⁻³⁰ suggesting that androgens in these species are important for initiation and maintenance of spermatogenesis and steroidogenesis *via* the Leydig, Sertoli and peritubular myoid cells but not the germ cells. It is well known that androgen has been demonstrated to regulate the genes required for *de novo* cholesterol synthesis in Leydig cells.³¹ The present results showed that seasonal change in localization of AR in the breeding season, the nonbreeding season and pre-hibernation were different, and that the changes in testicular weight, size and histological appearance were in parallel with the expression levels of testicular AR. These findings suggest that androgen acts as a paracrine factor regulating germ cell development in seminiferous tubules and an autocrine factor regulating steroid hormones synthesis in the Leydig cells of the WGS.

Aromatase that irreversibly transforms androgens into estrogens is present in the smooth endoplasmic reticulum of nearly all cell types in the mammalian testis.³² In this study, there was a marked seasonal change in the immunolocalization and expression pattern of *P450arom*, and immunoreactivity specific to *P450arom* was observed in Leydig cells, Sertoli cells and germ cells in the breeding season. These findings were similar to some models of spermatogenesis arrest in the other seasonal breeders. Study in American and Japanese black bears have shown that *P450arom* was

present in Leydig cells, Sertoli cells, round and elongating spermatids during the mating season.^{6,33} In previous study of WGS, positive immunoreactivity for aromatase has also been found in Leydig and Sertoli cells as well as in spermatogenic cells during the breeding season.¹⁴ Similar observations have been reported in the wild raccoon dog, suggested that Leydig cells, Sertoli cells and germ cells were the sites of estrogen synthesis, which might be critical for male gonadal functions as well.³⁴ In addition, the present data showed that *P450arom* was not immunostained in testicular tissues of the nonbreeding season, and was immunolocalized in Sertoli cells at the beginning of testicular recrudescence. Meanwhile, the mRNA expression level of *P450arom* in the testes was significantly higher in pre-hibernation than those of the nonbreeding season. These findings were in general agreement with those observed in other mammals. In American black bear, the presence of aromatase has been reported at the beginning of testicular recrudescence in Sertoli cells.³³ In the Siberian hamster, estrogen was able to induce initiation of spermatogenesis, independently of FSH in photo-regressed adult male.³⁵ Besides, the stimulatory effects of estrogen on initiating spermatogenesis have also been reported in the bank vole,³⁶ showed that an *in vivo* treatment with low dose of estrogens induced a recrudescence of spermatogenesis. Taken together, the present results supported the proposal that the presence of *P450arom* in Sertoli cells at the beginning of testicular recrudescence may imply a substantial role of aromatase and estrogen in re-initiating spermatogenesis.⁶

Unlike AR, which is localized mainly in testicular somatic cells, ERs are present in both somatic and germ cells in the testis.³² The localization of ERs in testicular cells varies depending on the species, developmental stage of the cell and type of receptor.^{17,37,38} In the present study, *ER α* and *ER β* were strongly present in testicular tissues of the ground squirrel during the breeding season, suggesting that estrogen could function *via* both receptor subtypes. Both *ER α* and *ER β* regulate the expression of a variety of different genes. The present results showed that *ER α* was present in Sertoli and Leydig cells, and *ER β* was identified in Leydig cells and spermatids during the breeding season. These findings suggested that estrogen might regulate steroidogenesis by acting through *ER α* and *ER β* in the Leydig cells and promote gametogenesis by acting through *ER α* in the Sertoli cells and *ER β* in the germ cells. Study in the stallion showed that both ERs were immunodetected in Sertoli and Leydig cells before, during and after puberty but showed differential expression, with *ER β* being expressed until sexual maturity. In

the boar, *ER α* was localized in spermatogonia and primary spermatocytes, whereas *ER β* immunoreactivity was localized in Sertoli cells.^{39,40} In the South American Plains Vizcacha fetal and prepubertal testes,¹⁶ *ER α* and *ER β* were expressed in seminiferous tubules and the interstitium, respectively. Yet in adult testes, both *ER α* and *ER β* co-localized in Leydig and peritubular cells. Taken together, our data implicated that estrogen can act in a paracrine/autocrine manner(s) *via* *ER α* and *ER β* throughout the developing, mature and recrudescence testis.

In summary, the present results demonstrated that seasonal changes in testicular weight, size and seminiferous tubule diameters in WGSs are correlated with changes in spermatogenesis and testicular distribution of AR, *P450arom* and ERs during the breeding and nonbreeding seasons and pre-hibernation. The data presented here will greatly aid the dissection of steroid hormones endocrine pathways in WGSs, and population control of a fecund rodent species. Future studies will investigate seasonal changes in plasma concentrations of steroid hormones, and clarify the molecular mechanisms of steroid hormones regulating testicular recrudescence and regression.

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