



# The expression of androgen receptor messenger RNA is regulated by tri-iodothyronine in lizard testis

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## Abstract

The network of hormonal and non-hormonal signals required for testicular activity during the reproductive cycle of the seasonal breeding lizard, *Podarcis sicula*, are not yet well understood. Androgens are significantly involved in meiosis and spermiogenesis, and such an effect is mediated through their receptor (AR). Estrogens also affect the testicular activity down-regulating the expression of AR mRNA. Since over the last few years, extensive works have reported, in mammals, a clear influence of tri-iodothyronine (T<sub>3</sub>), the biologically active thyroid hormone, on Sertoli cell activities, we carried out a study to shed light on the effect/s exerted by T<sub>3</sub> in lizard testis. A thyroid hormone receptor mRNA (TR mRNA) has been found in the testis indicating that T<sub>3</sub> might be involved in the regulation of gonadal activity. In *in vivo* experiments, injection of T<sub>3</sub> to male lizards, captured during the recrudescence period (March) and maintained under experimental photothermal conditions (24°C and 15 h daylight), increased the expression of AR mRNA. The *in vitro* results confirmed the stimulatory effect of T<sub>3</sub> on AR mRNA levels. Thus, in testosterone (T) exposed cells, the highest values of AR mRNA were observed in T<sub>3</sub>-primed animals, indicating that T and T<sub>3</sub> increase AR gene transcription independently. The present data suggest that, in lizards, the combined action of androgens, estrogen and T<sub>3</sub> might regulate testicular activity, modulating AR mRNA levels. © 2000 Elsevier Science Ltd. All rights reserved.

**Keywords:** Steroid receptor; Lizard; Reproduction; mRNA expression

## 1. Introduction

In reptiles, many studies have been conducted to elucidate the dynamic relationships between the elements involved in the control of male gonadal activity, as well as the mechanisms responsible for the action of the regulating factors.

In the seasonal breeding lizard, *Podarcis sicula*, testicular activity is regulated by hormonal signals through mechanisms that are not yet well defined. Like most saurians inhabiting temperate zones, this species dis-

plays a reproductive cycle with full gonadal activity in spring and regression in summer. In July, the onset of an endogenous refractory phase blocks spermatogonial mitoses and causes the degeneration of the other germ cells [1–3]. As in other vertebrates, in lizards, specific changes in plasma sex hormone levels occur during the reproductive cycle [4,5] and it has been proposed that androgens are significantly involved in meiosis and spermiogenesis [1,2], and such an effect is mediated through their receptor (AR). Recently, *in vitro* studies carried out on primary cultures of lizard testis cells during full reproductive activity, have shown that both androgens (T) and estrogens (E) are involved in the regulation of AR mRNA expression [6]. T and E auto-

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induce the expression of androgen (AR) and estrogen receptor (ER) mRNAs, respectively. Conversely, a marked down-regulatory effect of E on the expression of AR mRNA has been observed [6]. Thus, in lizard testis cells, an interplay between estrogen and androgen receptors occurs.

The regulation of testicular activity, however, depends on several endogenous factors. Many *in vivo* studies have shown that, in mammals, the thyroid hormone affects testicular growth and differentiation [7–9]. Transient neonatal hypothyroidism in rats causes an increase in testis size [10], sperm production [11] and number of Sertoli cells [12] in adults. These effects *in vivo* seem to be due to the direct effect of thyroid hormone on the testis [13].

An interaction between thyroid hormone and reproductive activity has long been recognized in other vertebrates. In birds, it has been proposed that thyroxine may contribute to seasonal testicular regression by reducing testis sensitivity to LH [14]. Investigations upon agamids, lacertids and geckonids have shown that hypothyroidism, induced by thyroidectomy, causes a regression of the testis, involving spermatogenic arrest and reduction in androgen secretion [15,16]. Thyroid treatment stimulates spermatogenesis in *Rana pipiens* [17] and in the toad *Bufo regularis* [18].

Furthermore it has been found that the expression of thyroid hormone receptor (TR) is associated with the expression of other nuclear receptor genes. Thus in *Xenopus* liver the expression of TR potentiates estrogen activation of vitellogenin genes and autoinduction of ER in epatocytes [19]. Also in primary cultures of hamster Harderian gland cells, the expression of AR mRNA is increased when T<sub>3</sub> is present in the culture medium [20]. Since there are now studies regarding the cross-talk between different nuclear receptors in the view of multiple control of some physiological functions such as testicular activity, the aim of the present study was to determine if T<sub>3</sub> might be involved in the regulation of the AR mRNA levels in the testis of the lizard, *Podarcis sicula*. Therefore, we evaluated the expression of TR mRNA by Northern blot analysis and we investigated the role of T<sub>3</sub> on AR mRNA levels in the testis and testicular primary cultures.

## 2. Materials and methods

### 2.1. Animals

We utilized adult specimens of male lizards (*Podarcis sicula*) captured in the neighborhood of Naples (Italy) during March (pre-reproductive period), when the natural photoperiod is 11 h daylight with a temperature ranging from 12 to 14°C.

### 2.2. Chemicals

Glassware was sterilized and RNase free. Triiodothyronine (T<sub>3</sub>) and testosterone (T) were purchased from Sigma (Sigma-Aldrich, Italy). All other materials were of molecular biology grade.  $\alpha^{32}$ -P-dCTP (3000 Ci/mmol) was purchased from Amersham International (Amersham, Italy). Restriction enzymes were from New England Biolabs (Beverly, USA).

### 2.3. Animal treatments

Lizards were randomly divided into three groups ([a], [b], [c]) ( $n = 25$ /group). Group [a] was maintained in terraria under environmental photothermal regimes (control group). Groups [b], [c] were kept for 15 days in terraria under experimental photothermal conditions (24°C and 15 h daylight). Furthermore group [b] was treated every two days with intraperitoneal injection (100  $\mu$ l) of vehicle (saline) while group [c] was treated every two days with intraperitoneal injection (100  $\mu$ l) of T<sub>3</sub> ( $10^{-5}$  M). The experimental photothermal conditions (24°C and 15:9 L:D) utilized in our study induce a complete gonadal recovery many weeks earlier than the usual period [2]. The animals were fed on meal worms and fresh vegetables *ad libitum*.

At the end of the experiments, the lizards were killed under anesthesia, and the gonads were aseptically excised. Some testes were frozen in liquid nitrogen for RNA extraction, the others were washed in balanced saline solution for preparation of primary cell cultures.

### 2.4. Preparation of primary cultures of testis cells

Establishment of primary cultures of testis cells was carried out as described in Cardone et al. [6]. The analysis of cell types present in all preparations indicated that at least 80% of the total cells were primary and secondary spermatocytes and spermatids, whereas spermatogonia were about 8% and spermatozoa were absent. Interstitial cells and peritubular cells were about 2% and 10%, respectively, whereas Sertoli cells were rarely seen. The viability of the cells was evaluated by using the expression of cytoskeletal  $\beta$ -actin mRNA as control and was always >90%.

### 2.5. RNA extraction

Total cellular RNA was extracted following Chomczynski and Sacchi [21] with minor modifications. The yield and quality of RNA were assessed by the 260/280 nm optical density ratio ( $1.93 \pm 0.05$ ) and by electrophoresis under non-denaturing conditions on 1.2% agarose gels.

## 2.6. Northern blot analysis of TR expression

For Northern analysis, 30 µg of total RNA, isolated from whole lizard testes, was subject to electrophoresis on 1.2% agarose-formaldehyde gel. Single strand ( $\lambda$ -DNA/Hind III digest (23.130–0.564 kb, Stratagene, La Jolla, USA) and RNA molecular-weight markers (7.4–1.6 kb, Boehringer) were used to determine the sizes. Sample and markers were transferred overnight onto nylon membrane (Nytran, Schleicher & Schuell) using  $10 \times$  SSC (SSC: 0.015 M trisodium citrate, 0.15 M NaCl, pH 7.0). Filters were baked for 30 min/80°C in a vacuum oven and exposed to UV irradiation (254 nm/2 min). Prehybridization was performed at 42°C for 4 h in 50% deionized formamide,  $5 \times$  SSC, 0.1% SDS, 0.05 M phosphate buffer (pH 6.8), 0.005 M EDTA,  $5 \times$  Denhardt's solution [Denhardt's solution: 1% (w/v) ficoll, 1% (w/v) polyvinylpyrrolidone, 1% (w/v) BSA], 100 µg/ml yeast tRNA. Hybridization buffer was as above. A 840 bp fragment (Pst I) from *Xenopus* TR  $\alpha$  form cDNA (xTR $\alpha$ ) [22] and a 796 bp fragment (Hinf I) from *Xenopus* TR  $\beta$  form cDNA (xTR $\beta$ ) [22] were used as probe. Probes were labeled with  $\alpha$ -<sup>32</sup>P-dCTP by random priming to a specific activity of  $5 \times 10^8$  cpm/µg [23]. Hybridizations were performed at 42°C overnight. Filters were washed twice with  $2 \times$  SSC, 0.1% SDS at 68°C, twice with  $0.2 \times$  SSC, 0.1% SDS at 68°C and once with  $0.1 \times$  SSC, 0.1% SDS at 72°C. Dried filters were exposed to X-ray film (Fuji HR-H) for 48–96 h.

## 2.7. Effect of *in vivo* different experimental conditions on AR mRNA expression in lizard testis

Total cellular RNA, isolated from lizard testis of different experimental groups ([a], [b], [c]) was subjected to analysis of the relative amounts of AR mRNA expression. For this purpose, the analysis of RNA was performed by quantitative slot-blot technique as described elsewhere [24,25]. Thus, total RNA (15 µg/sample) was bound to nylon membrane (Nytran, Schleicher & Schuell) in a slot-blot apparatus (Schleicher & Schuell). Slot-blotting was performed in duplicate and hybridized as described by Cardone et al. [6]. Prehybridizations, hybridizations and washes were carried out as above.

The relative amounts of AR mRNA were determined by densitometric scanning of the autoradiograms and normalized to corresponding values of  $\beta$ -actin mRNA.

## 2.8. Effect of T on AR mRNA expression in primary cultures of testis cells within different experimental groups

Testicular cells from the different experimental

groups ([a], [b], [c]) were cultured in the presence or absence (control) of T ( $10^{-8}$  M). The incubations were performed for 8, 24, and 48 h, replacing the culture medium and drug every 24 h. At the end of each experimental time, the cells were washed and RNA was extracted as before.

Total RNA (15 µg/sample) was bound to nylon membrane in a slot-blot apparatus. Duplicate filters were hybridized to rAR and  $\beta$ -actin probes, respectively [6]. The experiment was performed three times using three different cell preparations. Prehybridizations, hybridizations and washes were carried out as above.

Two different types of analysis were performed by slot-blot method depending on the way of samples loading. The first one was the analysis of steady-state levels of AR and  $\beta$ -actin mRNAs in T-exposed and unexposed cells of each experimental group ([a], [b], [c]). For this purpose, RNA samples at each experimental time (i.e., in [a] group: T-exposed cells at 0, 8, 24, 48 h) were loaded consecutively and the densitometric values were obtained by scanning the slot-blot hybridization signals. Data were expressed as relative amounts of AR mRNA and  $\beta$ -actin mRNA with respect to total amount of signals (i.e., in [a] group T-exposed cells:  $0 + 8 + 24 + 48 = 100\%$ ). With the second way of analysis we studied the differences of AR mRNA levels between T-exposed and unexposed cells from time to time. For this purpose, samples having the same experimental time (i.e., in [c] group at 24 h: control cells and T-exposed cells) were loaded consecutively. The densitometric values were expressed as relative amounts of AR mRNA with respect to the total amount of signals (i.e., in [c] group at 24 h: control cells + T-exposed cells = 100%).

## 2.9. Analysis of AR mRNA expression in primary cultures of testis cells between the different experimental groups

We also examined the differences in AR mRNA levels between the T-exposed cells from the three experimental groups ([a] [b] [c]). For this purpose, RNA samples having the same experimental time (i.e., at 24 h: [a], [b], [c]) were loaded consecutively. The densitometric values, obtained by scanning the signals at each time, were expressed as relative amounts of AR mRNA with respect to the total signals (i.e., at 24 h: [a] + [b] + [c] = 100%).

## 2.10. Statistics

Significance of differences were evaluated by Duncan test for multigroup comparisons. Data were expressed as mean  $\pm$  SD.

### 3. Results

#### 3.1. Northern blot analysis of TR expression

Fig. 1 shows the Northern blot detection of TR mRNA in lizard testes from specimens captured during the pre-reproductive period (March). The probe used for TR $\alpha$  mRNA recognizes both TR $\alpha$ 1 and TR $\alpha$ 2 forms. The molecular sizes of these transcripts were 5.0 kb for TR $\alpha$ 1 and 2.6 kb for TR $\alpha$ 2 mRNA, a value similar to that reported for many mammals. A minor 6.0 kb band was also observed; it reportedly represents unprocessed heteronuclear RNA, encoding unspliced primary transcripts of RNA prior to their processing into the mature TR $\alpha$ 1 and TR $\alpha$ 2. As far as the occurrence of TR $\beta$  mRNA in lizard testis is concerned, no TR $\beta$  transcript was observed (data not shown).

It is worth to note that the probes used in this study contain both the steroid and the DNA binding domains and despite the high stringency conditions, it resulted evident that the testis of the lizard strongly expressed TR $\alpha$  mRNA.

#### 3.2. Effect of different *in vivo* experimental conditions on AR mRNA expression in lizard testis

Fig. 2 shows the levels of AR mRNA expression in lizard testis between the different experimental groups ([a], [b], [c]). Densitometric analysis of slot-blot hybridization, expressed as relative amounts of AR mRNA, indicated that in the lizard testis, experimental photothermal conditions (24°C and 15 h daylight) were able to stimulate the AR mRNA expression above the control levels. After *in vivo* T<sub>3</sub>-injection, the AR mRNA

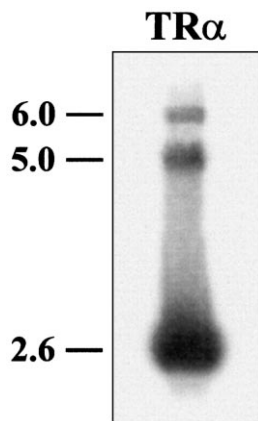


Fig. 1. Northern blot analysis of thyroid hormone receptor mRNA (TR mRNA) in lizard testis. Total RNA (30  $\mu$ g), obtained from the whole testis, was subjected to electrophoresis under denaturing conditions and transferred to nylon membrane. The blot was hybridized with a random-labeled 840 bp fragment (Pst I) from *Xenopus* TR $\alpha$  form (xTR $\alpha$ ) [22] or to 796 bp fragment (Hinf I) from *Xenopus* TR $\beta$  form cDNA (xTR $\beta$ ) [22] (data not shown). The sizes of the specific TR $\alpha$  transcripts are indicated on the left side of the figure.

levels were further enhanced, indicating that T<sub>3</sub> was in some way able to increase AR mRNA expression.

#### 3.3. Effect of T on AR mRNA expression in primary cultures of testis cells from different experimental groups

*In vitro* experiments confirmed that exposure to androgens affected AR mRNA levels in all experimental groups. Fig. 3 illustrates the steady-state levels of AR and  $\beta$ -actin mRNAs and the time course analysis of AR mRNA expression in primary cultures of testis cells obtained from animals of control group ([a]). The AR mRNA expression increased in T-exposed cells while there was a continuous decrease in control cells (Fig. 3A). In Fig. 3B, the autoregulatory effect of T is also evident.

The steady-state levels of AR and  $\beta$ -actin mRNAs and the time course analysis of AR mRNA expression in primary cultures of testis cells obtained from animals maintained under experimental photothermal conditions (24°C and 15 h daylight; [b]) are shown in Fig. 4. Also in this group, the exposure of the cells to T, increase the AR mRNA levels. In fact, the time course analysis (Fig. 4B) show that at the beginning of the experiment (time 0), no significant differences in AR mRNA levels were observed, both in control and T-exposed cells. AR mRNA was significantly increased in T-exposed cells at 8, 24, and 48 h after treatment. A similar trend was observed in primary cultures of testis cells obtained from animals maintained at 24°C and injected with T<sub>3</sub> ([c]) (Fig. 5). The steady-state levels (Fig. 5A) shows that T increased AR mRNA levels,

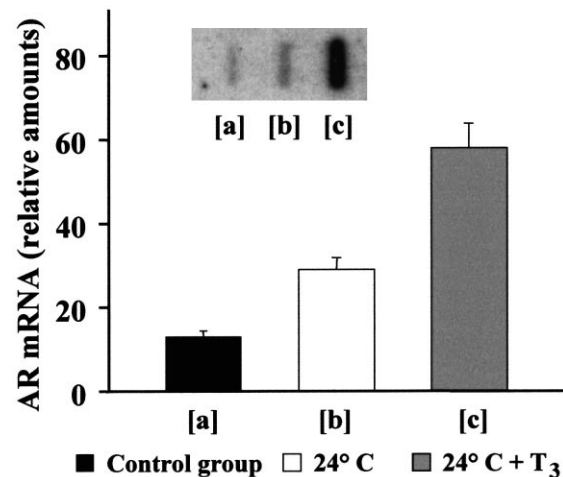


Fig. 2. Levels of AR mRNA expression in lizard testis between the different experimental groups ([a], [b], [c]). Insert: Representative autoradiogram of a slot-blot hybridization of the AR mRNA. The relative concentrations were determined by densitometric analysis of slot-blot hybridization and expressed as relative amounts of AR mRNA with respect to the total signals. Values are the means  $\pm$  SD.

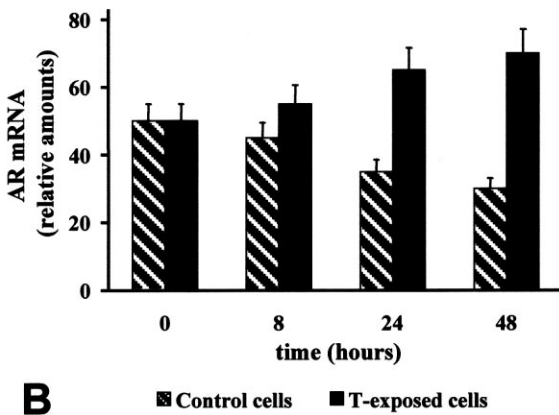
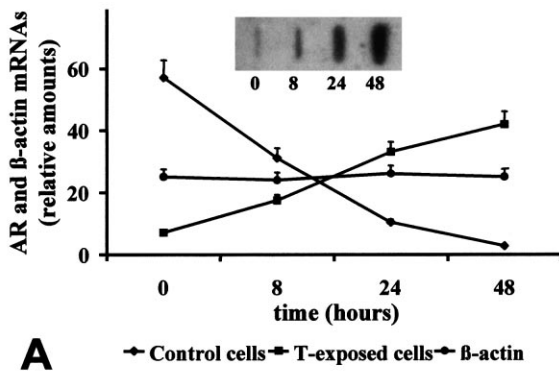


Fig. 3. Analysis of AR and  $\beta$ -actin mRNAs in primary cultures of testis cells of experimental group [a] (control group). The testicular cells were cultured in the presence or absence (control) of testosterone ( $T 10^{-8} M$ ). The relative values were determined by slot-blot hybridization. Values are the means  $\pm$  SD. A: Steady-state levels of AR and  $\beta$ -actin mRNAs as a function of the time. Values for  $\beta$ -actin are shown for T-exposed cells. Insert: Typical autoradiogram of a slot-blot of total RNA isolated from T-exposed cells, probed for AR. B: Time-point analysis of AR mRNA between the control and T-exposed cells. The values were obtained by scanning the hybridization signals at each time and were expressed as relative amounts with respect to the total signals.

while there was a continuous decrease in control cells. These results are also confirmed by time point analysis (Fig. 5B), although such a decrease was less evident than that seen in control cells from animals maintained at  $24^{\circ}C$  without  $T_3$  treatment (Fig. 4B). In all the experiments the levels of  $\beta$ -actin mRNA did not show variation between the control and T-exposed cells.

### 3.4. Analysis of AR mRNA expression in primary cultures of testis cells between the different experimental groups

The stimulatory effect of in vivo  $T_3$ -priming on AR mRNA levels was evident from the results of time point analysis of T-exposed testis cells from different experimental groups ([a], [b], [c]) (Fig. 6). Densito-

metric analysis of slot-blot hybridization, expressed as the relative amount of AR mRNA, indicated that the highest values of AR mRNA were found in the  $T_3$ -primed group. In fact, in T-exposed testis cells from animals injected in vivo with  $T_3$ , AR mRNA expression was higher than the other two groups at any time.

## 4. Discussion

It has been clearly demonstrated that hormones do not act separately, but they operate jointly in complex signalling networks [20,26–29]. An example of hormonal interplay is represented by the participation of

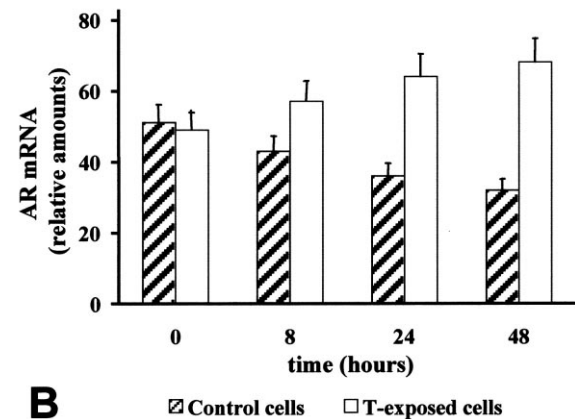
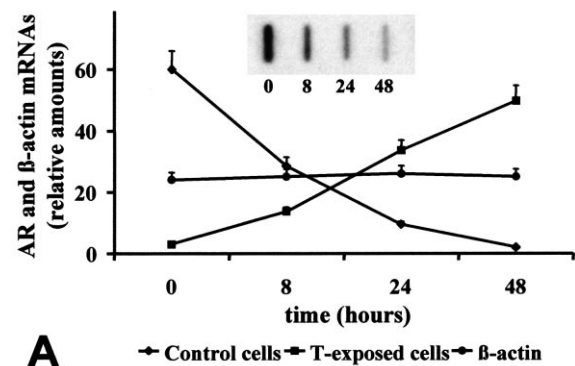


Fig. 4. Analysis of AR and  $\beta$ -actin mRNAs in primary cultures of testis cells of experimental group [b] ( $24^{\circ}C$  and 15 h daylight). The testicular cells were cultured in the presence or absence (control) of testosterone ( $T 10^{-8} M$ ). The relative values were determined by slot-blot hybridization. Values are the means  $\pm$  SD. A: Steady-state levels of AR and  $\beta$ -actin mRNAs as a function of the time. Values for  $\beta$ -actin are shown for T-exposed cells. Insert: Typical autoradiogram of a slot-blot of total RNA isolated from control cells, probed for AR. B: Time-point analysis of AR mRNA between the control and T-exposed cells. The values were obtained by scanning the hybridization signals at each time and were expressed as relative amounts with respect to the total signals.

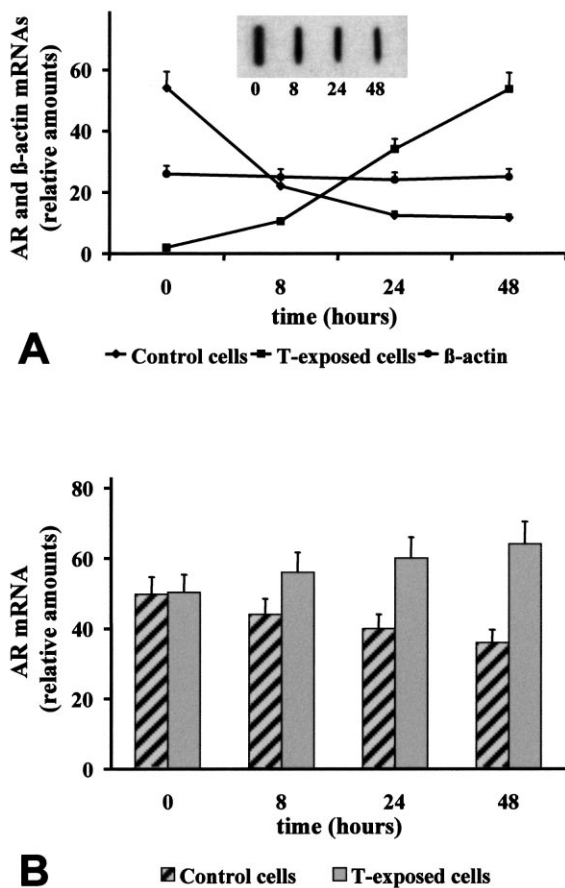


Fig. 5. Analysis of AR and  $\beta$ -actin mRNAs in primary cultures of testis cells of experimental group [c] ( $24^{\circ}\text{C}$  and 15 h daylight plus  $\text{T}_3$  injection). The testicular cells were cultured in the presence or absence (control) of testosterone ( $\text{T } 10^{-8} \text{ M}$ ). The relative values were determined by slot-blot hybridization. Values are the means  $\pm$  SD. A: Steady-state levels of AR and  $\beta$ -actin mRNAs as a function of the time. Values for  $\beta$ -actin are shown for control cells. Insert: Typical autoradiogram of a slot-blot of total RNA isolated from control cells, probed for AR. B: Time-point analysis of AR mRNA between the control and T-exposed cells. The values were obtained by scanning the hybridization signals at each time and were expressed as relative amounts with respect to the total signals.

thyroid hormones in multi-hormonal regulation of many physiological processes [19,20,26,30–33]. Recently, it has been established that  $\text{T}_3$  can increase Sertoli cell protein synthesis, providing further evidence that thyroid hormone has a fundamental role in the regulation of growth and differentiation of mammalian testis [34]. Binding studies and in situ hybridization indicate that Sertoli cells express TR and its mRNA during the neonatal period [35,36]. TR is encoded by two different genes,  $\alpha$  and  $\beta$ ; the alternative splicing of gene primary transcripts leads to the production of at least four distinct mRNA species generating separate TR isoforms ( $\text{TR}\alpha 1$ ,  $\text{TR}\alpha 2$ ,  $\text{TR}\beta 1$ ,  $\text{TR}\beta 2$ ). The  $\text{TR}\alpha 1$  and  $\text{TR}\beta$  isoforms bind  $\text{T}_3$  and appear to operate as transcriptional activators, while

the physiological significance of  $\text{TR}\alpha 2$ , a non- $\text{T}_3$ -binding variant, remains unknown [37,38].

The lizard testis expresses both  $\text{TR}\alpha 1$  and  $\text{TR}\alpha 2$  mRNA, having molecular sizes of 5.0 and 2.6 kb, respectively, similarly to what has been found in prepuberal rats [36,39]. The other major functional TR mRNA ( $\text{TR}\beta 1$ ), detected in rat Sertoli cells by PCR [36,40], has never been observed. However, the hybridization conditions and the molecular weight of the transcripts are consistent with the presence of  $\text{TR}\alpha 1$  and  $\text{TR}\alpha 2$  mRNAs in the testis of sexually mature lizards.

In mammals, adult testes are devoid of both  $\text{T}_3$  nuclear binding activity [35,41] and  $\text{TR}\alpha 1$  and  $\text{TR}\beta 1$  mRNA isoforms, while they exclusively express the non-hormone binding isoforms  $\text{TR}\alpha 2$  [36,42,43]. On the other hand, in rat testes,  $\text{TR}\alpha 1$  mRNA expression [36] and  $\text{TR}\beta 1$  transcript [40] have been detected only in immature Sertoli cells. As indicated by maximal binding capacity, the concentration of TR in whole testes is highest during the fetal and perinatal period and decreases significantly until puberty, virtually disappearing in the adult [35]. The presence of TR in fetal, perinatal and prepuberal rat testes suggests that  $\text{T}_3$  might regulate the growth of the seminiferous epithelium by modulating some different functions of Sertoli cells [44]. Although, our results are in contrast with those report in mammals, the occurrence of  $\text{TR}\alpha$  mRNA in adult lizard testes indicates that the thyroid hormone would have a direct effect on these cells. It has been proposed that the evolution of the thyroid function might have been paralleled by the endocrine control of reproduction and that the basic function for thyroid hormones might be associated primitively with

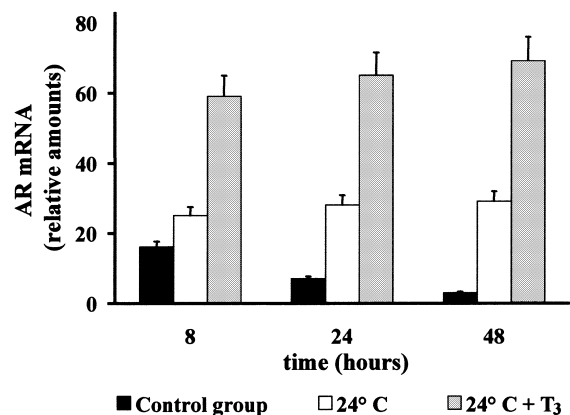


Fig. 6. Time-point analysis of AR mRNA in primary cultures of testis cells in all experimental group [a], [b], [c]. The testicular cells were cultured in the presence of testosterone ( $\text{T } 10^{-8} \text{ M}$ ). The relative values were determined by slot-blot hybridization. The values were obtained by scanning the hybridization signals between the different experimental group at each time and were expressed as relative amounts with respect to the total signals. Values are the means  $\pm$  SD.

gonadal maturation. Generally, the thyroid function in reptiles varies in proportion to changes in temperature, although the cause-effect relationships are still obscure [45]. Seasonal changes in thyroid function have been correlated with a number of reproductive events (spermatogenesis, ovulation, mating) in lizards, snakes, and turtles [46]. In the lizard, thyroidectomy causes gonadal regression, which can be overcome by thyroid hormone administration [47]. Moreover, hyperthyroidism, induced by thyroid hormone injection into intact lizards, blocks testis growth [15]. The effect of thyroid hormone, however, shows seasonal variations, which are greater when testes are fully active than during recrudescence [16]. The marked seasonality of thyroidal activities in reptiles further supports an involvement of this gland in the reproductive cycle [46]. Apart from these considerations, the expression of TR $\alpha$  mRNA in lizard testis also provides evidence that thyroid hormones can directly modulate AR mRNA expression in testicular cells. Recent works have indicated that T<sub>3</sub>-treatment regulates AR and ER content in cultured Sertoli cells from peripuberal rats [48] and results in increased AR mRNA levels in the cultures of the same cells from 5-day-old rats [39]. In this respect, our *in vivo* experiments showed that, in lizards, either temperature alone (24°C) or temperature and T<sub>3</sub> significantly stimulated AR mRNA levels above the control values. However, temperature and T<sub>3</sub> affected AR mRNA expression levels more markedly than the temperature alone, indicating that T<sub>3</sub> is able to up-regulate AR mRNA expression. It is clear that the increase in steady-state AR mRNA levels in lizard testes is due to a specific direct effect of T<sub>3</sub> on AR mRNA levels, rather than to a general stimulatory effect of T<sub>3</sub> on testis maturation. In fact, the analysis of germ cell percentage conducted on histological sections of both experimental groups did not reveal great differences (data not shown).

*In vitro* experiments confirmed previous results showing that exposure to androgens enhances AR mRNA levels [6]. It is worth noting that our previous results were gained using testicular cells from animals in full breeding season, while the present study was carried out during the pre-reproductive period (March). The results show the same pattern although the absolute values could be different, indicating that in this period the testis is able to respond to androgens up-regulating the AR mRNA. In fact, in T-exposed cells, AR mRNA significantly increased after 8, 24, and 48 h, with respect to the unexposed (control) cells in all experimental groups. The differences of AR mRNA levels, between unexposed and T-exposed cells from animals maintained at 24°C and T<sub>3</sub>-primed, however, were less evident with respect to the other two experimental groups. This result was probably due to the additional effect of the *in vivo* T<sub>3</sub>-priming and a

greater thyroid activity in the specimens of this experimental group. It is known that, in lizards from temperate regions, thyroid activity is highest during the warmer seasons. Artificial lowering of the environmental temperature during summer causes a reduction in thyroid activity. Lizards maintained in the laboratory at high temperatures (35°C) exhibit greater thyroid function than when maintained at 15°C [44]. The stimulatory effect of T<sub>3</sub>-priming on AR mRNA levels is evident from our results of time point analysis of T-exposed cells, obtained from the different experimental groups. These results indicate that the highest values of AR mRNA were found in the T<sub>3</sub>-primed group. In fact, time-course analysis showed that, in T-treated cells from animals exposed *in vivo* to T<sub>3</sub>, AR mRNA levels increased after 8, 24, and 48 h, although, the steady-state levels showed a rapid accumulation of AR mRNA in each group.

The effects of T<sub>3</sub> and T suggest that these hormones act in different pathways. Since receptors play a central role in the action of hormones, T<sub>3</sub>, binding its own receptor (TR), is able to stimulate AR mRNA levels and in turn influence the androgen responsiveness of testis cells during spermatogenesis. It is conceivable that thyroid hormone might act with other factors in the regulation of seasonal spermatogenetic cycle. In this respect it is worth to note that in a previous work carried out in our laboratory, we found that estrogens strongly down-regulated the expression of AR mRNA in primary cultures of lizard testis cells [6]. Increasing evidence in support of cross-talk between nuclear receptors and various types of signals has been observed [19,26–29,49,50]. Therefore, it appears that the cooperation of a number of hormonal (T, E, T<sub>3</sub>) and, probably, non-hormonal signals, is required for regulation of spermatogenetic cycle in the lizard. Taken together, our studies indicate the importance of interplay between different ligands of the steroid/thyroid hormone nuclear receptor gene family acting by modulating the expression of the gene encoding one of the members of this superfamily.

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