Thyroid hormone effects on androgen receptor messenger RNA expression in rat Sertoli and peritubular cells

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Abstract

Postnatal Sertoli cell maturation is characterized by a pronounced rise in androgen receptor (AR) expression, which increases several fold between birth and adulthood. Since both 3,3',5-triiodothyronine (T_3) and FSH regulate Sertoli cell proliferation and differentiation, we have determined the effects of T_3 and FSH on AR mRNA expression in cultured Sertoli cells from 5-day-old rats. These cultures contain 5–9% peritubular cells, which also express AR mRNA. To insure that the observed T_3 responses did not result from peritubular cells, we examined T_3 effects on AR mRNA expression in cultured 20-day-old Sertoli cells (which contain minimal peritubular contamination) and peritubular cells, and measured thyroid hormone receptor (TR) mRNA expression in both of these cell types. Sertoli cells from 5- and 20-day-old rat testes were grown in serum-free medium alone (controls) or with ovine FSH (100 ng/ml) and/or T_3 (100 nM) for 4 days. Peritubular cells purified from 20-day-old rat testes were grown in serum-containing medium for 8 days. These cells were split 1:4, and grown

an additional 8 days, the last 4 days in serum-free medium with or without T_3 . TR and AR mRNA levels in all cultures were determined by Northern blotting. AR mRNA levels in 5- and 20-day-old cultured Sertoli cells were significantly ($P < 0.05$) increased by both T_3 and FSH alone. Furthermore, AR mRNA levels in Sertoli cells treated with T_3 and FSH were greater than with either alone. TR mRNA expression was detected in cultured peritubular cells, but TR mRNA levels in these cells were only approximately 30% of that seen in 20-day-old cultured Sertoli cells. In contrast to Sertoli cells, T_3 did not affect peritubular AR mRNA expression. These results indicate that T_3 is an important regulator of the postnatal Sertoli cell AR mRNA increase. The additive effects of maximally stimulatory doses of FSH and T_3 suggest these hormones work through different mechanisms to increase AR mRNA. TR mRNA expression in peritubular cells indicates these cells may be direct T_3 targets, though the function of T_3 in these cells is unknown.

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Introduction

Postnatal Sertoli cell maturation is characterized by steadily increasing androgen receptor (AR) expression, which is barely detectable in these cells at 5 days postnatal but reaches maximal levels by 90 days of age (Bremner *et al.* 1994, Shan *et al.* 1995, 1997). Androgens are essential for the initiation of normal spermatogenesis and adult Sertoli cell function (Griswold 1995). This rise in Sertoli cell AR during neonatal and juvenile life may be necessary for the Sertoli cell to achieve full differentiation and androgen responsiveness, resulting in a cell competent to support complete spermatogenesis. The factors which regulate the increase in AR expression during Sertoli cell development are therefore of great importance. Follicle-stimulating hormone (FSH), the major tropic hormone for Sertoli cells (Griswold 1993), increases expression of AR and its mRNA in juvenile Sertoli cells (Sanborn *et al.* 1984, 1991, Verhoeven & Cailleau 1988), and androgens also appear to

regulate Sertoli cell AR (Shan *et al.* 1997). However, the overall hormonal mechanisms regulating Sertoli cell AR expression during development are not well understood.

Extensive work in the past few years has indicated that thyroid hormones are major regulators of Sertoli cell development. Transient neonatal hypothyroidism in rats produces unprecedented increases of 80 and 140% in adult testis weight and sperm production respectively (Cooke & Meisami 1991, Cooke *et al.* 1991). This increase in testis size results primarily from increased Sertoli cell proliferation during neonatal and juvenile life (van Haaster *et al.* 1992) and consequent larger populations of adult Sertoli and germ cells (van Haaster *et al.* 1992, Hess *et al.* 1993). Sertoli cells express thyroid hormone receptor (TR) and its mRNA during the neonatal period (Jannini *et al.* 1990, 1994) and 3,3',5-triiodothyronine (T_3), the biologically active thyroid hormone, directly inhibits Sertoli cell proliferation *in vitro* (Cooke *et al.* 1994). T_3 also appears to stimulate maturation of Sertoli cells (van Haaster *et al.*

1993), and has been shown to directly increase production of Sertoli cell secretory proteins (Cooke *et al.* 1994, Jannini *et al.* 1995) and decrease aromatase activity (Panno *et al.* 1994, Ullisse *et al.* 1994).

The aim of the present study was to use cultured neonatal Sertoli cells to determine if T_3 might be involved in the Sertoli cell AR increase during development. These experiments are complicated by the presence of peritubular cells in the cultures, which express high levels of AR neonatally and could make interpretation of the results difficult. Therefore, we also directly evaluated the expression of TR mRNA by cultured peritubular cells and the ability of T_3 to regulate AR mRNA in these cells *in vitro* to determine if these cells had the capacity to directly respond to T_3 and whether T_3 might also modulate AR mRNA expression in peritubular cells.

Materials and Methods

Animal purchase, care and breeding were as described previously (Cooke & Meisami 1991). All experiments described here involving animals were approved by the Laboratory Animal Care Advisory Committee of the University of Illinois, and were conducted in accordance with the Guiding Principles for the Care and Use of Research Animals.

Sertoli cell culture from 5-day-old rats

Sertoli cells from 5-day-old Sprague–Dawley rats (day of birth=day 0) were isolated using a sequential enzymatic procedure that we (Cooke *et al.* 1994) and others (Rong-Xi *et al.* 1987, Orth & Boehm 1990) have described previously. Briefly, for each culture, pools of Sertoli cells were obtained from eight to ten males from two litters. Sertoli cells were grown in 24-well plates coated with Matrigel (Collaborative Research, Waltham, MA, USA) diluted 1:5 with Hank's balanced salt solution. Cells were plated at a density of 4×10^5 cells/well. The nutritive media was Dulbecco's minimum essential medium (DMEM) supplemented with sodium pyruvate (1 mM), non-essential amino acids (0.1 mM) and an antimicrobial solution (Rong-Xi *et al.* 1987, Orth & Boehm 1990). Cells were grown for 4 days in a humidified atmosphere of 95% air/5% CO_2 at 34 °C. Media was changed every 24 h.

Some cultures were grown without hormonal supplementation, while others received FSH (ovine FSH-18, 100 ng/ml) and/or T_3 (Sigma, St Louis, MO, USA) at 100 nM. In all cases, T_3 was added at the initiation of the culture while FSH was added at the beginning of the third day, and fresh hormone was added when media were changed. After 4 days *in vitro*, some wells of cultured Sertoli cells were used to obtain RNA for Northern

analysis. The entire experiment was performed on 10 separate occasions.

Sertoli and peritubular cell culture from 20-day-old rats

Sertoli and peritubular cells from 20-day-old Sprague–Dawley rat testes were prepared using procedures that have been described previously (Tung & Fritz 1977, Buzek & Sanborn 1990). Briefly, testes were decapsulated in DMEM/Ham's F-12 medium and minced into 0.5 mm fragments. Testicular fragments were then subjected to enzymatic treatment, as described previously (Tung & Fritz 1977, Buzek & Sanborn 1990). The Sertoli cell-enriched fragment was then harvested by unit-gravity sedimentation, while the supernatant fraction was used to prepare peritubular cell cultures.

The digested Sertoli cell-enriched fragments were washed with DMEM/Ham's F-12, and the subsequent treatment of the cells was as described for the 5-day-old Sertoli cell isolation procedure (Cooke *et al.* 1994). Cells were plated at a density of 5×10^5 cells/well. These cells were then cultured under conditions identical to that described above for 5-day-old Sertoli cells, and the cells were similarly subjected to treatment with FSH and T_3 as above. The entire experiment was performed five times.

The peritubular cell-enriched preparation obtained from the collagenase supernatant fraction described above was used to obtain a cell pellet of peritubular cells, as described previously (Tung & Fritz 1977, Buzek & Sanborn 1990). This pellet was resuspended in DMEM/Ham's F-12 containing 1% BSA and washed. Peritubular cells were plated at a density of 4×10^5 cells/well and grown in a humidified atmosphere of 95% air/5% CO_2 at 34 °C on 24-well plates coated with Matrigel, as described for 5-day-old Sertoli cells. The nutritive medium was the same as used for Sertoli cells, but supplemented with 10% fetal calf serum (Sigma); medium was changed every 2 days. Cells were grown for 8 days, then removed from their substrate using trypsin (0.25%), split 1:4 and replated into 24-well plates as above. Cells were grown an additional 4 days in serum-containing medium, then cultured for 4 days without serum and either with or without T_3 (100 nM). The data shown are based on five separate isolation procedures.

Viability and purity of cultured cells

For all cell cultures described, the percentage of viable cells was determined by trypan blue exclusion before plating. To assess contamination with other cell types, cultures were examined daily using an inverted microscope. In addition, at the end of the culture, wells of all cell types were stained for alkaline phosphatase or 3 β -hydroxysteroid dehydrogenase (3 β -HSD), which are markers for peritubular and Leydig cells respectively (Chapin *et al.* 1987, Orth & Boehm. 1990, Cooke *et al.* 1994). Since Leydig

cells differentiate during early postnatal life and may not stain as intensely for 3 β -HSD during this period as fully mature adult Leydig cells, both intensely and weakly stained cells were scored as positive.

Northern analysis

Total RNA was prepared from all cell cultures using the RNeasy Mini Kit (Qiagen, Chatsworth, CA, USA). Total RNA was also isolated from 20-day-old testis and adult spleen, which served as positive and negative controls respectively, for the various probes. Purified total RNA was dissolved in diethylpyrocarbonate (DEPC)-treated water. Purity and concentration of the RNA were determined by UV (260/280 nm) absorbance in a spectrophotometer. Equal amounts of total RNA (8 μ g) from the various treatment groups were electrophoresed on a 1.5% agarose formaldehyde gel. The gel was blotted to nylon membrane and the RNA fixed to the membrane by UV cross linking.

The following probes were used in this study: 1) c-erbA α , a probe which detects both TR α 1 and α 2 mRNA (Murry *et al.* 1988), 2) rat antigen binding protein (ABP) (Joseph *et al.* 1987), 3) rat AR (Chang *et al.* 1988) and 4) human 28S rRNA (Erickson & Schimickel 1985). Initial Northern studies of AR mRNA expression were performed with the full-length 2.83 kb AR cDNA construct, while later experiments were performed using a 1.15 kb AR cDNA fragment (Chang *et al.* 1988). Both probes recognized the 9.5 kb AR mRNA transcript (Chang *et al.* 1988). Insert for each cDNA probe was isolated from the plasmid vector by restriction digestion and gel purification. Each insert was labeled with ³²P-dCTP using the Multiprime DNA labeling system (Amersham, Arlington Heights, IL, USA) and used to probe the membrane.

All hybridizations were carried out in QuikHyb (Stratagene, La Jolla, CA, USA) according to manufacturer's recommendations at 68 °C in a Robbins scientific hybridization oven (Sunnyvale, CA, USA). The hybridized membrane was washed, covered with plastic wrap and exposed to Kodak X-omat X-ray film with intensifying screens. After hybridization with the TR α cDNA probe, membranes were stripped of probe by incubation in 50% formamide at 65 °C for 1 h and then rehybridized with the AR cDNA probe. These Northern blots were also rehybridized with the ABP cDNA probe. For normalization of RNA load levels between lanes, the membrane was reprobed for a final time with labeled 28S rRNA cDNA probe.

The mRNA bands on the autoradiograms were scanned and quantitated using a computer-linked scanning laser densitometer and RFLPrint software (Pdi, Huntington Station, NY, USA). Relative levels of mRNA transcripts were adjusted to compensate for differences in total RNA loaded per gel lane as determined by densitometry of 28S

Figure 1 Northern blot detection of steady-state AR mRNA. A 9.5 kb transcript was detected in 20-day-old testis (lane 1) and Sertoli cells from 5-day-old rats cultured 4 days without hormonal supplementation (lane 2), with 100 nM T₃ (lane 3), with 100 ng/ml FSH (lane 4), or with FSH and T₃ (lane 5). A negative control, spleen, is also shown (lane 6).

rRNA hybridization signals (Bunick *et al.* 1994). All statistical analyses were performed using the SYSTAT statistical package (Wilkinson 1990). The optical densities (OD) of the bands in the various treatment groups were compared by two-way ANOVA, and differences between various treatment groups were compared using Tukey's HSD test. Differences were considered significant when $P < 0.05$.

Results

Characterization of Sertoli and peritubular cell cultures

Trypan blue exclusion experiments indicated that more than 95% of the isolated peritubular and Sertoli cells were viable. The Sertoli cell cultures from the 5-day-old rats had a morphology typical of Sertoli cells *in vitro* (Cooke *et al.* 1994). Approximately 5% of the cells were peritubular, as judged by their morphology and positive alkaline phosphatase staining, while Leydig cell contamination was minimal (<1%). The Sertoli cell cultures from the 20-day-old rats also showed a very low level of Leydig and peritubular cell contamination (<1%). Primary peritubular cell cultures consisted predominately of fibroblastic cells; small numbers of Sertoli cells could also be identified by their nuclear morphology and cell shape. However, after the peritubular cells were passaged, Sertoli cells were absent and cultures consisted exclusively of fibroblastic cells. Alkaline phosphatase staining was very strong in peritubular cell cultures, while less than 1% of the cells stained for 3 β -HSD, indicating minimal Leydig cell contamination.

Expression of AR and TR mRNA in Sertoli cell cultures from 5-day-old rats

In Northern blots for AR (Fig. 1), a 9.5 kb band which has previously been shown to correspond to AR mRNA (Chang *et al.* 1988, Sanborn *et al.* 1991, Shan *et al.* 1995) was detected in cultured Sertoli cells from 5-day-old rats. AR mRNA was also detected in 20-day-old rat testis, a positive control, but not in spleen, a negative control which lacks detectable AR mRNA (Shan *et al.* 1995). Densitometric analysis of hybridization results (Fig. 2)

Figure 2 Normalized densitometric data of AR mRNA expression by 5-day-old Sertoli cells cultured 4 days with or without hormonal supplementation, as in Fig. 1. Signals were normalized against a 28S rRNA cDNA probe to compensate for interlane differences in RNA loading. The values for each of the treatment groups were expressed as a fold increase relative to the control, and the data are presented as means $\pm$ S.E.M. of values from 10 separate experiments. The mean values for all treatment groups were different from each other ($P < 0.05$) with the exception of the groups treated with FSH or T₃ alone, which did not differ.

indicated that steady-state levels of AR mRNA expression in cells treated with T₃ or FSH alone were both slightly under 2-fold that of the untreated controls ($P < 0.05$ for both compared with controls). AR mRNA expression in cells treated with both of these hormones was approximately 2.5-fold that of control cultures, indicating that T₃ and FSH acted additively to increase AR mRNA expression to a level greater than with either hormone alone. Cultured 5-day-old Sertoli cells also expressed relatively high levels of TR α 1 and α 2 mRNA (data not shown).

Expression of AR, TR and ABP mRNA in Sertoli and peritubular cell cultures from 20-day-old rats

Hormonal regulation of AR mRNA expression in 20-day-old Sertoli cells (Fig. 3) was similar to that in 5-day-old Sertoli cells. Normalized densitometric data (Fig. 4) indicated that T₃, FSH, or T₃ and FSH significantly ($P < 0.05$) stimulated steady-state AR mRNA expression above control levels. The T₃ and FSH response was greater than that seen with either hormone alone, and the T₃ and FSH groups were not significantly different. In some cases, RNA from both 5- and 20-day-old Sertoli cell cultures were run on the same Northern blot, which allowed a direct comparison of mRNA levels of AR in the various

Figure 3 Northern blot detection of steady-state AR mRNA in testis, spleen and cultured testicular cells. A 9.5 kb transcript was detected in whole 20-day-old testis (lane 1). Peritubular cells from 20-day-old rats were grown for 16 days as described in the text; the last 4 days these cells were cultured either without (lane 2) or with (lane 3) 100 nM T₃. Sertoli cells from 20-day-old rats were cultured 4 days without hormonal supplementation (lane 4), with 100 nM T₃ (lane 5), with 100 ng/ml FSH (lane 6), or with FSH and T₃ (lane 7). Spleen (lane 8), which was used as a negative control, did not contain detectable levels of AR mRNA.

Figure 4 Normalized densitometric data of AR mRNA expression by 20-day-old Sertoli cells cultured 4 days with or without hormonal supplementation, as in Fig. 3. Signals were normalized against a 28S rRNA cDNA probe to compensate for interlane differences in RNA loading. The values for each of the treatment groups were expressed as a fold increase relative to the control, and the data are presented as means $\pm$ S.E.M. of values from five separate experiments. As in Fig. 2, the mean values for all treatment groups were different from each other ($P < 0.05$) with the exception of the groups treated with FSH or T₃ alone, which did not differ.

groups. These results indicated that 20-day-old Sertoli cells expressed approximately 2-fold more AR mRNA than 5-day-old cells from the same treatment group (not shown).

Although AR mRNA could be clearly detected in peritubular cell cultures, T₃ treatment did not have a significant effect on the steady-state AR mRNA level (Fig. 3), a conclusion confirmed by analysis of the normalized densitometric data (not shown). Thus, T₃ does not appear to stimulate AR mRNA in cultured peritubular cells as it does in Sertoli cells.

Northern blotting for TR mRNA in Sertoli cell cultures from 20-day-old rats indicated that they strongly

Figure 5 Northern blot detection of steady-state TR mRNA. The probe used recognizes both the TR α 1 (5.0 kb) and TR α 2 (2.6 kb) forms of TR α . A minor 6.0 kb band, thought to represent unprocessed heteronuclear RNA encoding unspliced primary transcripts of RNA prior to their processing into the mature α 1 and α 2 mRNAs, is also seen (Williams 1994). TR α 1 and TR α 2 mRNA were detected in 20-day-old testis (lane 1) and in peritubular cells from 20-day-old rats cultured the last 4 days either without (lane 2) or with (lane 3) 100 nM T $_3$, as in Fig. 3. Sertoli cells from 20-day-old rats cultured 4 days without hormonal supplementation (lane 4), with 100 nM T $_3$ (lane 5), with 100 ng/ml FSH (lane 6), or with FSH and T $_3$ (lane 7) all expressed TR α 1 and TR α 2 mRNA. Conversely, spleen (lane 8) did not. Analysis of normalized densitometric data indicated that TR mRNA expression by cultured peritubular cells was approximately 30% of that in Sertoli cells.

expressed TR α 1 and α 2 mRNA (Fig. 5). However, TR α 1 and α 2 mRNA expression was not significantly different in the 5- or 20-day-old Sertoli cells or amongst the different treatment groups at these ages. Therefore, in contrast to the results obtained with AR mRNA, neither FSH nor T $_3$ treatment nor age of the Sertoli cells used had a significant effect on the steady-state levels of TR α 1 and α 2 mRNA.

TR α mRNA was also detected in freshly isolated peritubular cells from 20-day-old rats (data not shown) and in these peritubular cells following 16 days of culture (Fig. 5). Both TR α 1 and α 2 mRNA were detected in cultured peritubular cells and the relative abundance of the TR α 2 mRNA was approximately 2-fold greater than that for TR α 1 in both the peritubular and Sertoli cells. TR α mRNA expression in peritubular cells was less than in Sertoli cells; in the control cultures, the relative abundance of TR α 1 and α 2 mRNA per μ g of RNA was about 3-fold greater in cultured 20-day-old Sertoli cells compared with 20-day-old peritubular cells (data not shown).

Expression of ABP mRNA in peritubular cell cultures was not detectable, in stark contrast to the 20-day-old Sertoli cell cultures, in which ABP mRNA was strongly expressed (Fig. 6). Therefore, Sertoli cell contamination is minimal in the peritubular cell cultures.

Discussion

Sertoli cell AR is barely detectable by immunohistochemistry in 5-day-old neonatal rats (Bremner *et al.* 1994). AR concentration per Sertoli cell then increases during juvenile life, reaching maximal levels during adulthood (Buzek *et al.* 1987, Buzek & Sanborn 1988, Shan *et al.*

Figure 6 Northern blot detection of steady-state ABP mRNA. Two ABP mRNA bands were detected in 20-day-old testis (lane 1). These bands, a major 1.7 kb ABP transcript and a minor 2.3 kb transcript, correspond to the sizes of the ABP transcripts reported previously (Joseph *et al.* 1987, Bunick *et al.* 1994). Peritubular cells from 20-day-old rats cultured the last 4 days either without (lane 2) or with (lane 3) 100 nM T $_3$ (as in Fig. 3) did not express detectable ABP mRNA, indicating that Sertoli cell contamination in these cultures was minimal. Sertoli cells from 20-day-old rats cultured 4 days without hormonal supplementation (lane 4), with 100 nM T $_3$ (lane 5), with 100 ng/ml FSH (lane 6), or with FSH and T $_3$ (lane 7) all expressed high levels of ABP mRNA. Spleen (lane 8) did not contain detectable ABP mRNA.

1997). The rise in Sertoli cell AR expression closely parallels that in Sertoli cell AR mRNA expression. For example, Sertoli cells show a 2-fold rise in both AR and AR mRNA between day 21 and adulthood (Shan *et al.* 1995, 1997).

The present results indicate that T $_3$ treatment results in increased AR mRNA expression in cultured Sertoli cells from 5-day-old rats and may be an important endocrine regulator of the developmental increase in Sertoli cell AR *in vivo*. Recent work has also indicated that T $_3$ increases androgen binding in cultured juvenile rat Sertoli cells (Panno *et al.* 1996). Our results suggest that the increased AR levels observed in this study in response to T $_3$ result from an increase in AR mRNA steady-state levels rather than changes in translation.

The mechanism by which T $_3$ treatment increases AR mRNA expression is not clear. T $_3$ has previously been shown to cause a number of changes which are associated with increasing maturation in the neonatal Sertoli cell, such as decreased proliferation, increased production of secretory proteins, and active proluminal secretion of fluid (Francavilla *et al.* 1991, van Haaster *et al.* 1993, Cooke *et al.* 1994). It therefore seems likely that the increases in steady-state AR mRNA levels may reflect general stimulatory effects of T $_3$ on Sertoli cell maturation rather than a specific, direct effect of T $_3$ on AR mRNA production, though the latter possibility can not be totally excluded.

The present results are the first demonstration that FSH stimulates AR mRNA expression in Sertoli cells of neonatal rats. In addition, although the stimulatory effects of FSH on AR mRNA in juvenile Sertoli cells have been shown (Sanborn *et al.* 1991), the present results are the first to show that the combined effects of T $_3$ and FSH on AR mRNA in neonatal and juvenile Sertoli cells are greater than with either alone. The additive effects of T $_3$ and FSH suggest that these hormones are not causing their effects through the same pathway, a conclusion consistent with

previous observations that T_3 and FSH effects on other Sertoli cell parameters, such as production of mRNA for an inhibin subunit, are also additive (Cooke *et al.* 1994).

Testosterone also affects AR expression in postnatal Sertoli cells (Verhoeven & Cailleau 1988, Sanborn *et al.* 1991). Therefore, it appears that a number of hormones, including T_3 , FSH and testosterone, and possibly other hormones as well, are necessary for the normal postnatal rise in Sertoli cell AR expression. Additional studies on the role of all of these hormones, alone and especially in combination, will be necessary to obtain a complete understanding of the factors that control Sertoli cell AR mRNA expression, the overall developmental sequence of postnatal Sertoli cell maturation, and the onset of functional activity in these cells.

One possible complication of using Sertoli cell cultures from early postnatal rats is that they contain significant peritubular cell contamination. Schlatt *et al.* (1996) showed that Sertoli cell cultures from 7-day-old rats contain approximately 6% peritubular cells after 3 days of culture, and 9% peritubular cells by the seventh day of culture. Peritubular cell AR expression exceeds that of Sertoli cells in 5-day-old rats (Bremner *et al.* 1994), indicating that even low levels of contaminating peritubular cells in these cultures could significantly affect total AR measurements. Therefore, data showing effects of T_3 on AR or its mRNA, such as our Northern results with Sertoli cells from 5-day-old rats and previous androgen binding data obtained from Sertoli cells derived from 2-week-old rats (Panno *et al.* 1996), must be interpreted cautiously.

Sertoli cell cultures from 20-day-old rats have been reported to contain less than 1% peritubular cell contamination (Raychoudhury *et al.* 1993), in contrast to the 5–9% peritubular contamination seen in neonatal Sertoli cell preparations (Cooke *et al.* 1994, Schlatt *et al.* 1996). The ability to obtain purer cultures with juvenile rat Sertoli cells allows the potentially confounding effects of peritubular cells to be minimized. Therefore, to obtain unequivocal information on regulation of AR mRNA in Sertoli cells, we also examined the effects of T_3 and FSH on AR mRNA expression in 20-day-old Sertoli cells.

The effects of T_3 and FSH, alone and in combination, on AR mRNA expression in the 20-day-old Sertoli cell cultures were similar to that seen in the 5-day-old cultures. These results, obtained in cultures where peritubular cell contamination is minimal, strongly suggest that the stimulatory effects of T_3 and T_3 and FSH seen in the 5-day-old Sertoli cell cultures were due to the effects of these hormones on the Sertoli cells themselves, rather than effects on the contaminating peritubular cells in these cultures. In addition, the 20-day-old Sertoli cells expressed approximately twice the AR mRNA of their 5-day-old counterparts, consistent with previous data suggesting a rise in AR between the neonatal and juvenile periods (Buzek *et al.* 1987, Bremner *et al.* 1994).

Previous work has indicated that TR α 1 and α 2 mRNA are produced at high levels in the testis and Sertoli cells neonatally. This expression declines with advancing age, reaching relatively low levels during adulthood (Jannini *et al.* 1990, 1994, Bunick *et al.* 1994). The major functional form of TR in the developing testis and Sertoli cells is TR α 1. The other major functional TR, TR β 1, can only be detected in Sertoli cells by PCR techniques (Jannini *et al.* 1994, Palmero *et al.* 1995), indicating very low abundance. TR α 2 (also called c-erbA α 2) is a splicing variant of the TR α locus, and does not bind T_3 despite its very high homology to TR α 1. Thus, TR α 2 is not a receptor in the strict sense. The function of TR α 2 is unknown, though it may have an inhibitory effect on thyroid hormone action (Williams 1994).

If peritubular cells were indeed contributory to the increase in AR mRNA in response to T_3 seen in Sertoli cell-enriched cultures from 5-day-old rats, peritubular cells would be expected to express TR mRNA and show increased AR mRNA expression when exposed to T_3 *in vitro*. We therefore directly evaluated TR mRNA expression and T_3 effects on AR mRNA production in 20-day-old peritubular cell cultures. Peritubular cells from 20-day-old rats were used because the small size of the 5-day-old rat testis makes it difficult to obtain sufficient peritubular cells to perform Northern blotting experiments. In addition, well-defined protocols for the purification, culture and assessment of 20-day-old peritubular cells are available (Tung & Fritz 1977, Sanborn *et al.* 1991).

Our results indicate that cultured 20-day-old peritubular cells expressed both TR α 1 and TR α 2 mRNA. However, overall levels of TR α 1 and α 2 mRNA per μ g of RNA in cultured peritubular cells were only about 30% of the levels in Sertoli cells cultured from 20-day-old rats. Although initial reports indicated that TR was present only in Sertoli cells of the testis and could not be detected in other cell types by either classical T_3 binding assays (Jannini *et al.* 1990) or *in situ* hybridization for TR mRNA (Jannini *et al.* 1994), Hardy *et al.* (1996) recently reported that TR α 1 and α 2 mRNA could be detected in immature and adult Leydig cells, as well as their juvenile mesenchymal precursors. These data and the present results with peritubular cells suggest that TR may be ubiquitous in the cell types of the developing testis, although levels in these other cell types during development are less than in Sertoli cells (Hardy *et al.* 1996). Thus, a complete determination of the role of thyroid hormones in the testis will require an understanding of the effects of thyroid hormone on all of these cell types both during development and adulthood.

One potential complication of measuring TR mRNA in peritubular cell cultures is that the strong expression of TR and its mRNA by young rat Sertoli cells (present results and Jannini *et al.* 1990, 1994) suggests that even low-level Sertoli cell contamination in peritubular cell

cultures would result in detectable TR mRNA expression. Sertoli cells produce large amounts of ABP, while Leydig and peritubular cells do not produce this protein (Pelliniemi *et al.* 1981). ABP therefore functions as a specific and sensitive marker for Sertoli cell contamination in peritubular cell cultures. The lack of detectable ABP mRNA in the cultured peritubular cells indicated that Sertoli cell contamination was minimal in the peritubular cell cultures. Therefore, peritubular cells themselves, rather than contaminating Sertoli cells, produce the TR mRNA detected in these cultures.

The expression of TR mRNA in peritubular cell cultures indicates that thyroid hormones could have direct effects on these cells, but T_3 did not increase AR mRNA in cultured peritubular cells. Thus, T_3 regulates AR mRNA in Sertoli, but not peritubular cells, a finding similar to what has been previously reported for the effects of FSH on AR mRNA expression in these two cell types (Sanborn *et al.* 1991). The role of T_3 in peritubular cells is presently unknown. Mendis-Handagama & Sharma (1994) reported an increase of approximately 50% in connective cells (which included peritubular myoid cells as well as other cells such as pericytes and fibroblasts) in adult testes of rats that had been made hypothyroid during neonatal life compared with age-matched untreated controls. These results suggest that T_3 may have an effect on the proliferation of peritubular cells, as it does in Sertoli cells (Cooke *et al.* 1994). Likewise, an effect of T_3 on peritubular cell differentiation is also plausible, based on similar effects in Sertoli cells. Peritubular cells communicate extensively with other testicular cell types such as Sertoli and Leydig cells (Skinner 1991) suggesting that in addition to possible direct T_3 effects, T_3 binding to TR in peritubular cells could also result in effects on other cell types in the testis through the production of paracrine factors by peritubular cells.

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