Variation in the End Products of Androgen Biosynthesis and Metabolism during Postnatal Differentiation of Rat Leydig Cells*

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ABSTRACT

The amount of testosterone (T) secreted by Leydig cells is determined by a balance between T biosynthetic and metabolizing enzyme activities. It has been established that 5α -androstan- 3α , 17β -diol (3α -DIOL) is the predominant androgen secreted by the testes of immature rats during days 20-40 postpartum, whereas T is the major androgen by day 56. However, the underlying changes in T biosynthetic and metabolizing enzymes during Leydig cell development and their magnitudes have remained unclear. The aim of the present study was to define the developmental trends for T biosynthetic and metabolizing enzymes in Leydig cells at three distinct stages of pubertal differentiation: mesenchymal-like progenitors on day 21, immature Leydig cells on day 35, and adult Leydig cells on day 90. Production rates for precursor androgen (androstenedione), T, and 5α -reduced and rogens [and rosterone (AO) and 3α -DIOL] were measured in progenitor, immature, and adult Leydig cells in spent medium after 3 h in vitro. Steady state messenger RNA (mRNA) levels and enzyme activities of biosynthetic and metabolizing enzymes were measured in fractions of freshly isolated cells at each of the three stages. Unexpectedly, progenitor cells produced significant amounts of androgen, with basal levels of total androgens (androstenedione, AO, T, and 3α -DIOL) 14 times higher than those of T alone. However, compared with immature and adult Leydig cells, the capacity for steroidogenesis was lower in progenitor cells, with a LH-stimulated production rate for total androgens of 84.33 ± 8.74 ng/10⁶ cells·3 h

NDROGEN stimulation is responsible for the maintenance of spermatogenesis and secondary sexual characteristics in the male. According to the literature, testosterone (T) produced by Leydig cells of the testis is the major androgen in the circulation of men and adult males of most mammalian species, including the rat. However, in rats, which have been studied more extensively than most other species, 5α -androstan- 3α , 17β -diol (3α -DIOL) and androsterone (AO) are abundant in the circulation between days 20-40 postpartum when T is still low (1). Testicular tissue of rats aged 15-40 days metabolizes radiolabeled progesterone and T to 3α -DIOL and, to a lesser extent, AO (2–5). Testicular enzyme activities of 3β -hydroxysteroid dehydrogenase $(3\beta$ HSD), 17 α -hydroxylase/C17–20 lyase (P450_{c17}), and 17 β hydroxysteroid dehydrogenase (17BHSD) increase gradually between the ages of 20 and 60 days and plateau thereafter (6, 7). The amount of testicular 5α -reductase (5α R) activity, on the other hand, sharply increases between days 20 and 40,

Address all correspondence and requests for reprints to: Dr. Matthew P. Hardy, The Population Council, 1230 York Avenue, New York, New York 10021. E-mail: m-hardy@popcbr.rockefeller.edu. $(\text{mean} \pm \text{se})$ vs. 330.13 \pm 44.19 in immature Leydig cells and 523.23 \pm 67.29 in adult Leydig cells. The predominant androgen produced by progenitor, immature, and adult Leydig cells differed, with AO being released by progenitor cells (72.08 \pm 9.02% of total androgens), 3α -DIOL by immature Leydig cells (73.33 \pm 14.52%), and T by adult Leydig cells (74.38 \pm 14.73%). Further examination indicated that changes in the predominant androgen resulted from differential gene expression of T biosynthetic and metabolizing enzymes. Low levels of type III 17\beta-hydroxysteroid dehydrogenase (17\betaHSD) mRNA and enzyme activity were present in progenitor cells compared with immature and adult Leydig cells. In contrast, levels of type I 5 α -reductase $(5\alpha R)$ and 3α -hydroxysteroid dehydrogenase $(3\alpha HSD)$ mRNA and enzyme activities were dramatically lower in adult Leydig cells compared with those in progenitor and immature Leydig cells. Several T biosynthetic enzymes attained equivalent levels in immature and adult Leydig cells, but T was rapidly metabolized in the former to 3α -DIOL by high 5α R and 3α HSD activities, which were greatly reduced in the latter. Therefore, declines in $5\alpha R$ and $3\alpha HSD$ activities are hypothesized to be a major cause of the ascendancy of T as the predominant androgen end product produced by adult Leydig cells. These results indicate that steroidogenic enzyme gene expression is not induced simultaneously, but through sequential changes in T biosynthetic and metabolizing enzyme activities, resulting in different androgen end products being secreted by Leydig cells during pubertal development. (Endocrinology 139: 3787-3795, 1998)

and then falls between days 40 and 60 (7). Further analysis demonstrated that type I 5 α R and 3 α -hydroxysteroid dehydrogenase (3 α HSD) messenger RNA (mRNA) and protein are abundantly present in progenitor and immature Leydig cells during days 15–35 (8, 9). These findings suggest that the Leydig cell itself is a metabolizing site for androgens during puberty, at least via the 5 α -reduction pathway. However, technical barriers prevented the testing of this hypothesis, because steroidogenic enzymes such as 17 β HSD (10–12), 5 α R (13, 14), and 3 α HSD (14, 15) are present in other cell types in the testis. Therefore, purified Leydig cells were examined in the present study to delineate T biosynthesis and metabolism occurring in this cell type during pubertal differentiation.

Androstenedione (DIONE; a precursor of T), T, and the 5α -reduced metabolites, AO and 3α -DIOL, were measured in purified progenitor Leydig cells on day 21, in immature Leydig cells on day 35, and in adult Leydig cells on day 90. The predominant androgen end product varied depending on the stage of differentiation; it was AO in progenitor Leydig cells, 3α -DIOL in immature Leydig cells, and T in adult Leydig cells. These results led to a further examination of T biosynthetic [cholesterol side-chain cleavage enzyme

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(P450_{scc}), 3β HSD, P450_{c17}, and 17β HSD] and metabolizing (5α R and 3α HSD) enzyme activities in Leydig cells. The results showed that activities of T biosynthetic and metabolizing enzymes in developing Leydig cells determine not only their capacity for T production but also the predominant androgen end product that is secreted.

Materials and Methods

Chemicals

25-[26,27-³H]Hydroxycholesterol, [7-N-³H]pregnenolone, [1,2-N-³H]17α-hydroxyprogesterone, [1 β ,2 β -N-³H]androst-4-ene-3,17-dione, [1,2,6,7-N-³H]T, [1,2-N-³H]dihydrotestosterone, 5α-[9,11-N-³H]androstane-3α,17 β -diol, and [9,11-N-³H]AO were purchased from DuPont-New England Nuclear (Boston, MA). [1,2,6,7-N-³H]Progesterone was purchased from Amersham International (Aylesbury, UK). Nonradio-active steroids were purchased from Sigma Chemical Co. (St. Louis, MO) or Steraloids (Wilton, NH). 4-Methyl-aza-3-oxo-5α-pregnan-20(5)-carboxylate, an inhibitor of 5αR, was provided by Merck (Rahway, NJ). The antibodies for 3α-DIOL and AO in RIA were provided by Dr. D. T. Armstrong (Department of Obstetrics and Gynecology, University of Western Ontario, London, Ontario, Canada).

Animals

Sprague-Dawley rats (dams with litters of male pups, immature males, and adult males) were purchased from Charles River Laboratories (Wilmington, MA). The males rats were 21, 35, and 90 days of age on the day of Leydig cell isolation. The animals were killed by asphyxiation with CO_2 . The animal protocol was approved by the institutional animal care and use committee of the Rockefeller University (Protocol 91200). A complete description of the procedure that was used to isolate each of the three stages of Leydig cell differentiation has been published (16, 17). The purity of cell fractions was evaluated by histochemical staining for 3 β HSD activity with 0.4 mM etiocholanolone as the steroid substrate (18). Enrichment of the three fractions was typically more than 95%. The absence of androgen-binding protein mRNA in the cell fractions demonstrated that there was no appreciable contamination by Sertoli cells.

Androgen production

Isolated progenitor Leydig cells, immature Leydig cells, and adult Leydig cells were incubated at a concentration of $0.1-0.25 \times 10^6$ cells/ml in Leydig cell culture medium consisting of DMEM and Ham's F-12 medium (D2906, Sigma Chemical Co.) buffered with 15 mM HEPES and $14\,mM\,NaHCO_3$ and containing 1% BSA for 3 h at 34 C in a shaking water bath. Incubations of triplicate samples were conducted in medium alone (basal) or in medium plus a maximally stimulating dose of ovine LH (100 ng/ml). At the end of 3 h, the samples were centrifuged at $500 \times g$. Supernatants were extracted with 2 ml ethyl acetate twice, and the organic layer was dried under nitrogen gas. Steroids in the samples were fractionated using Sephadex LH-20 (Pharmacia Biotech, Uppsala, Sweden) column chromatography as previously described (19). The elution system was chloroform-butane-ethanol (50:50:1, vol/vol/vol) saturated with distilled water. Clear separation of DIONE, AO, T, and 3α -DIOL in this system was confirmed using radiolabeled steroids (19). The recovery rates following extraction and column separation (DIONE, 90.7 \pm 5.8%; AO, 76.2 \pm 2.0%; T, 73.9 \pm 3.7%; 3 α -DIOL, 86.0 \pm 1.2%) were used to correct the final concentration measured by RIA. RIAs of DIONE, AO, T, and 3α -DIOL were performed as previously described (20–22). The results of three separate experiments were averaged for statistical analysis.

Enzyme assay

With the exception of P450_{scc}, steroidogenic enzyme activities were measured by incubation of purified Leydig cells with radiolabeled substrates and separation of products by TLC as previously described (23–25). The substrate concentrations used for each enzyme were maximal to ensure that the concentration of substrate was not rate limiting.

Control samples of culture medium alone were run in parallel with each enzyme assay. Briefly, reaction mixture (0.5 ml) was prepared in Leydig cell medium that contained 1 μ M substrate (1 μ Ci) in medium. As T undergoes 5α-reduction in immature Leydig cells, 4-methyl-aza-3-oxo- 5α -pregnan-20(S)-carboxylate (2 μ M) was used to inhibit 5α R when P450_{c17} or 17β HSD was measured. The reaction mixture was maintained at pH 7.2. Reactions were initiated by adding to the reaction medium an aliquot of $0.1-0.2 \times 10^6$ Leydig cells. The reaction mixtures, conducted in triplicate, were maintained at 34 C in a shaking water bath for 10 min. Reactions were terminated by adding ice-cold ethyl acetate, and steroids were rapidly extracted. The organic layer was dried under nitrogen. The radioactivity was measured using a radiometric scanner (System 200/ AC3000, Bioscan, Washington DC). The activity of 3βHSD was determined by measuring conversion of pregnenolone to progesterone. P450_{c17} catalyzes two mixed function oxidase reactions: 17α -hydroxylation and C17–20 cleavage. The activity of 17α -hydroxylation was determined by measuring conversion of progesterone to 17α-hydroxyprogesterone, DIONE, and T. The activity of C17-20 cleavage was determined by measuring conversion of 17α -hydroxyprogesterone to DIONE and T. The activity of 17β HSD was determined by measuring the conversion of DIONE to T. The activity of $5\alpha R$ was determined by measuring the conversion of T to dihydrotestosterone, 3α -DIOL, and 5α -androstane- 3β ,17 β -diol. The activity of 3α HSD was determined by measuring the conversion of dihydrotestosterone to 3α -DIOL. The steroids were separated on TLC plates in chloroform-methanol (97:3) for 3β HSD, 17β HSD, and 5α R assays; chloroform-ether (7:1, vol/vol) for P450_{c17} 17α-hydroxylation and C17–20 cleavage assays; and diethyl ether-acetone (98:2) for the 3α HSD assay.

Activity of P450_{scc} was determined by measuring the conversion of side-chain labeled 25-[26,27-³H]hydroxycholesterol to radioactive 4-hydroxyl-4-methyl-pentanoic acid as previously described (26). Leydig cells were incubated in a total volume of 0.5 ml medium containing 1 μ Ci 25-[26,27-³H]hydroxycholesterol (1 μ M 25-hydroxycholesterol). Incubations were performed for 30 min at 34 C, and at the end of the incubation 0.5 ml NaOH (0.5 m) was added. The mixture was extracted twice with 2 ml chloroform and mixed with neutral alumna to remove nonmetabolized substrate (26), and an aliquot was removed for measurement by liquid scintillation counting.

RT-PCR

Within the cytochrome P450 enzyme superfamily, P450_{scc} is encoded by a single P450_{scc} gene (designated CYP11A1), and P450_{c17} is encoded by a single P450_{c17} gene (designated CYP17). However, the existence of multiple isoforms of hydroxysteroid dehydrogenases and 5 α R have been demonstrated. In the rat, four distinct 3 β HSD complementary DNAs (cDNAs) have been identified (27). These four isoforms share 76–94% identity in their amino acid sequences (28, 29). Types I and II are the only isoforms that are expressed in rat testis (28).

Four distinct 17 β HSD cDNAs have been cloned in several species (reviewed in Ref. 30). Although three isoforms have been identified in the rat (30), it is believed that the type III 17 β HSD isoform is the major protein in the testis responsible for the conversion of DIONE to T (31). In the rat, however, type III 17 β HSD cDNA has not been cloned. Therefore, we designed a pair of primers to amplify the common sequence of human and mouse type III 17 β HSD (31, 32). The primers amplified a 360-bp product from Leydig cell cDNA. The PCR product was partially sequenced and analyzed with respect to the known isoforms (rat types I, II, and IV 17 β HSD, type III human and mouse 17 β HSD) (31–35) using the LFASTA program (36).

At least three isoforms of 3α HSD have been cloned in several species (37), and the type I isoform has been cloned from rat liver cDNA (38–40). This isoform has been demonstrated to be present in Leydig cells (7). The two isoforms of 5α R that have been cloned in the rat are present in rat testis (41, 42).

Therefore, using the published sequences of the rat P450_{scc} (43), types I and II 3 β HSD (28), P450_{c17} (44), types I and II 5 α R (41, 42), type I 3 α HSD (39), and ribosomal protein S16 (RPS16) (45), primers were selected using PRIMER software (Whitehead Institute of Biomedical Research, Cambridge, MA) and synthesized on an oligonucleotide synthesizer (Gene Assembler Special, LKB, Rockville, MD). Table 1 summarizes the properties of the primers.

Total RNA was extracted from isolated Leydig cells by a single step

Target mRNA (Ref.)	Localization and sequence of pair	GenBank or EMBL	Size of PCR product (bp)	Identification
P450scc (43) Forward Reverse	5'-aggtgtagctcaggacttca-3' 5'-aggaggctataaaggacacc-3'	gb (J05156) ^a	399	RE $(Fok \ I)^b$
3β-HSD (I, II) (28) Forward Reverse	5'-TCACATGTCCTACCCAGG-3' 5'-ATTTTTCAGGATGCTCCC-3'	gb (M38178) gb (M38179)	264	RE $(Taq I)^b$
P450c17 (44) Forward Reverse	5'-tcatcaagaagggaaaagaa-3' 5'-tgaagcagatagcacagatg-3'	gb (M31681)	294	RE $(Fok \ I)^b$
17β-HSD (III) (32) Forward Reverse	5'-tcaatgggacaatgggcagt-3' 5'-gcygygycaycyygacyacg-3'	gb (U66827) gb (U05659)	363	Sequencing
5α-R (I) (41) Forward Reverse	5'-CAATCCTGCAAGATTCCACC-3' 5'-ATTGGTCCTTGGGTGCATTC-3'	gb (J05035)	380	RE $(EcoR I)^b$
5α-R (II) (42) Forward Reverse	5'-accaccacacaggttctacctt-3' 5'-ctgagtcaggattaggttgc-3'	gb (M95058)	519	RE $(EcoR I)^b$
3α-HSD (39) Forward Reverse	5'-CCCATCGATGCGTAACAAG-3' 5'-GCCGTGCTTCAATCTTGCTT-3'	gb (M61937)	237	$\operatorname{RE} (Fok \ \mathbf{I})^b$
RPS16 (45) Forward Reverse	5'-aagtcttcggacgcaagaaa-3' 5'-gacaagacgaagacccgtt-3'	emb (X17665)	148	RE $(Pst \ I)^b$

TABLE 1. Parameters of oligonucleotide primer pairs for seven target genes

^{*a*} The sequence accession number from GenBank (gb) or EMBL (emb).

^b The restriction endonuclease (RE) was listed.

method, using phenol and guanidinium thiocyanate (Trireagent, Molecular Research Center, Cincinnati, OH) according to the manufacturer's instructions. Leydig cell total RNA (400 ng) was reverse transcribed with avian myeloblastosis virus reverse transcriptase according to a previously described procedure for RT-PCR (46). Briefly, total RNA samples were reverse transcribed using random hexamers plus deoxy-NTPs at 42 C for 75 min, and the reaction was terminated by heating at 95 C for 5 min. Target cDNA was coamplified with RPS16 cDNA, an internal standard. PCR was initiated by addition of Taq DNA polymerase in the presence of $[\alpha^{-32}P]$ deoxy-CTP, and proceeded for 30 cycles at an annealing temperature of 52 C. Radiolabeled PCR bands were visualized by exposure to autoradiographic film. Quantitative analysis of mRNA levels was performed by scanning the films in a laser densitometer (Ultroscan, LKB). The signal intensities for steroidogenic enzyme mRNAs were normalized to RPS16. Preliminary studies showed that the targeted cDNAs were amplified linearly between 15-35 cycles of PCR.

Statistics

All measurements were repeated at least three times. The data were analyzed by the Kruskal-Wallis ANOVA, followed by multiple comparisons testing to identify significant differences between groups (47).

Results

Androgens produced by Leydig cells

Stage-dependent differences were observed in Leydig cell capacity for T production. The basal T production was, respectively, 4 and 30 times higher in immature and adult Leydig cells compared with that in progenitor Leydig cells. When the total androgen profile was measured, however, the stage-dependent differences were less pronounced (Table 2). The three Leydig cell preparations showed, respectively, 2-, 6-, and 6-fold increases in total androgen production in response to LH stimulation compared with basal control values (Table 2). These results indicated that progenitor Leydig cells were less responsive to LH stimulation than immature and adult Leydig cells.

Further analysis of the total androgen profile showed that the primary androgen end products produced by progenitor, immature, and adult Leydig cells were different. AO was the primary androgen secreted by progenitor Leydig cells because this steroid constituted 72.08 \pm 9.02% of the total androgens released under LH-stimulated conditions. Parallel measurements showed that 3 α -DIOL was the primary androgen for immature Leydig cells (73.33 \pm 14.52% of total androgens), and T was primary androgen for adult Leydig cells (74.38 \pm 14.73%; Fig. 1).

Synthesis of T and 3α -DIOL requires 17β HSD activity, and the low release rates for these steroids from progenitor cells suggest that the amount of 17β HSD is low in these cells. AO and 3α -DIOL require the activities of 5α R and 3α HSD, and the high release rates of these steroids by progenitors and immature Leydig cells indicate that androgen-metabolizing enzymes are most highly expressed in Leydig cells before the completion of puberty and then decline in adulthood.

Steroidogenic enzyme activities

The underlying basis for the different profiles of androgen release from three distinct stages of Leydig cell differentiation was examined further by measuring T biosynthetic and metabolizing enzyme activities in purified cells. Steroidogenic enzyme activities were measured in intact Leydig cells, because homogenization can change the relative rates of oxidative and reductive activities in most hydroxysteroid dehydrogenases. The first step of T biosynthesis is the con-

Conditions	Age group	Total androgen $(ng/10^6 \text{ cells} \cdot 3 \text{ h})$	Ratio of 17-hydroxyl to 17-keto androgen	Ratio of 5α -reduced to 5α -oxidized androgen
Basal	Progenitor Leydig cells Immature Leydig cells Adult Leydig cells	$egin{array}{rl} 36.77 \pm 5.32^a\ 55.10 \pm 10.41^a\ 89.33 \pm 4.13^b \end{array}$	$egin{array}{l} 0.17 \pm 0.05^a \ 4.97 \pm 0.76^b \ 5.71 \pm 0.87^b \end{array}$	$egin{array}{r} 8.45 \pm 1.49^a \ 2.69 \pm 0.57^a \ 0.12 \pm 0.02^b \end{array}$
+LH	Progenitor Leydig cells Immature Leydig cells Adult Leydig cells	$\begin{array}{c} 84.33\pm8.74^b\\ 330.13\pm44.19^c\\ 523.23\pm69.29^c\end{array}$	$egin{array}{l} 0.20 \ \pm \ 0.03^a \ 4.88 \ \pm \ 1.33^b \ 4.55 \ \pm \ 1.33^b \end{array}$	$4.92 \pm 0.08^a \ 5.41 \pm 0.94^a \ 0.09 \pm 0.02^b$

TABLE 2. Leydig cell capacity for androgen production during postnatal development

The incubations were performed in DMEM/F-12 alone (basal) or in medium plus ovine LH (+LH).

All data were expressed as the mean \pm SE from four separate experiments.

 $^{a-c}$ Groups in the same column sharing the same letter were not significantly different at P < 0.05.

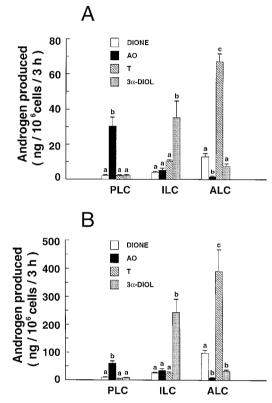


FIG. 1. Androgens produced by Leydig cells during pubertal development. Progenitor Leydig cells (PLC), immature Leydig cells (ILC), and adult Leydig cells (ALC) were isolated from 21-, 35-, and 90-dayold rats, respectively. The incubations were performed in DMEM-Ham's F-12 medium alone (basal; A) or in medium plus 100 ng/ml ovine LH (+ LH; B) with $0.1-0.25 \times 10^6$ cells. Measurement of DI-ONE, AO, T, and DIOL was performed as described in *Materials and Methods*. The levels of different androgens were expressed as the mean \pm SE in duplicate samples from four sets of separate experiments. Groups sharing the same alphabet letter were not significantly different at P < 0.05.

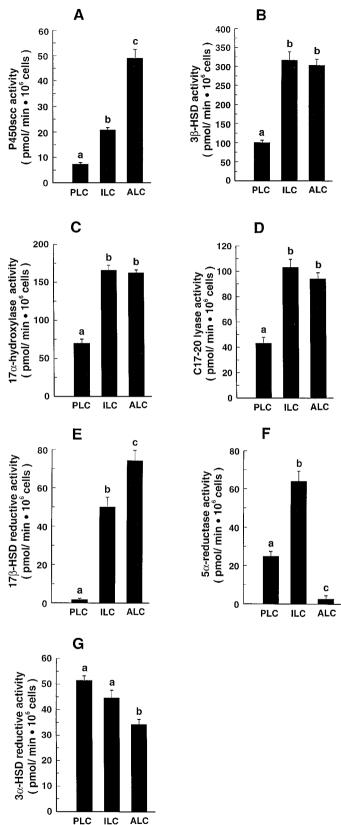
version of cholesterol to pregnenolone, which is catalyzed by P450_{scc}. The conversion of Δ^5 -3 β -hydroxysteroids to Δ^4 -3-ketosteroids is catalyzed by 3 β HSD. P450_{c17} catalyzes both the 17 α -hydroxylase and C17–20 lyase reactions to produce DIONE (48). The final step of T biosynthesis is catalyzed by 17 β HSD reductive activity. DIONE and T can be metabolized by 5 α R and 3 α HSD into AO and 3 α -DIOL, respectively. As shown in Fig. 2, the level of P450_{scc} was low in progenitor Leydig cells and increased during Leydig cell differentiation. 3 β HSD and P450_{c17} (17 α -hydroxylation and C17–20 cleav-

age) attained half of their mature activity levels in progenitor cells, and levels in immature and adult Leydig cells were statistically equivalent. Progenitor cells had negligible 17β HSD reductive activity, which greatly increased in immature and adult Leydig cells.

There was a significant increase in 5α R activity during the transition from progenitor to immature Leydig cell, followed by a sharp decline to almost undetectable levels in adult Leydig cells. The levels of 3α HSD activity were highest in progenitor Leydig cells and lowest in adult Leydig cells (Fig. 2). These results indicate that changes in the activities of both T biosynthetic and metabolizing enzymes catalyze the formation of distinct androgen end products during Leydig cell development.

Steady state mRNA levels of steroidogenic enzymes

Differential changes in T biosynthetic and metabolizing enzyme activities could result from variation in the steady state levels of mRNAs that encode these enzymes. Therefore, the steady state mRNA levels of Leydig cell steroidogenic enzymes were evaluated. With the exception of P450_{scc} and P450_{c17}, multiple isoforms of hydroxysteroid dehydrogenases and $5\alpha R$ have been identified. Four distinct $3\beta HSD$ cDNAs have been found in the rat (27), and rat Leydig cells are known to express 3β HSD isoforms I and II (28). Five distinct isoforms of 17βHSD have been identified in several species. Type III 17β HSD is thought to be responsible for catalyzing the conversion of DIONE to T in Leydig cells (31). Although it has not been cloned in the rat, type III 17β HSD is the predicted isoform for rat Leydig cells. A pair of primers was designed to identify sequences common to human and mouse type III 17βHSD cDNAs (31, 32). A 0.36-kb PCR product was identified in immature Leydig cells and adult Leydig cells (Fig. 3). Partial sequence analysis of the first 200 bp of the 0.36-kb PCR product showed that it had a 75% sequence similarity with mouse type III 17βHSD compared with only 35%, 40%, and 35% similarity with rat type I, II, and IV isoforms. Accordingly, this cDNA fragment was identified as type III. Two distinct isoforms of $5\alpha R$ have been identified in the rat testis (9, 41). In the testis, type I is present exclusively in Leydig cells (9). Type I $5\alpha R$ mRNA was abundant in progenitor and immature Leydig cells and almost undetectable in adult Leydig cells. No type II $5\alpha R$ mRNA was detected in any stage of Leydig cell despite a clear signal in the positive control, the prostate (data not shown). Of the three isoforms of 3α HSD that have been identified, type I is known to be expressed in the testis (8). As shown in Fig. 3, when the



steady state mRNA levels of these genes were evaluated, the trends for mRNAs corresponded closely with the trends defined for enzyme activities, suggesting that these genes transcriptionally regulate the levels of steroidogenic enzyme activity.

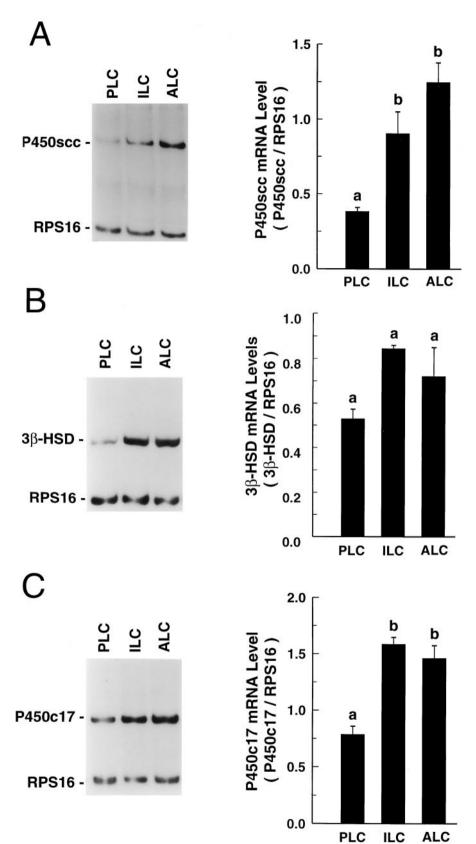
Discussion

From the data reported herein, it is apparent that development of the capacity for T production is achieved not only by increases in T biosynthetic enzymes but also by concurrent declines in T-metabolizing enzymes. Studies of serum and testicular and rogen levels have shown that 5α -reduced androgens are high in immature males (1-5). In the rat, 3α -DIOL is the predominant and rogen in circulation during days 20-40 (1). A distinction must therefore be drawn between steroidogenic ability and the capacity for T production. Steroidogenic ability is defined as the ability of cells to produce steroids, whereas the capacity for T production reflects the balance of both T biosynthetic and metabolizing enzyme activities. In the present study, using purified intact Leydig cells, we measured a 46-fold increase in basal T production during the transition from 21-day-old progenitor to 90-day-old adult Levdig cell under LH-stimulated conditions. In contrast, when the release rate was totaled for the four androgens, DIONE, T, AO, and 3α -DIOL, it only increased by 5-fold during the transition between progenitor and adult Leydig cell. This indicated that steroidogenic ability and capacity for T production are distinct measures of Leydig cell function. The steroidogenic ability of the progenitor Leydig cells, although lower than that of the more mature Leydig cell stages, is notable because these cells are devoid of smooth endoplasmic reticulum membranes that are the sites of steroidogenic enzyme localization. It is possible, therefore, that the steroidogenic ability of progenitor Leydig cells results from cytosolic, as opposed to membranebound, isoforms of steroidogenic enzymes. This is supported by the presence of type I 3α HSD, a cytosolic isoform, which attained its highest levels in progenitor Leydig cells. Another possible explanation is that mitochondrial isoforms of steroidogenic enzymes may be present in progenitor Leydig cells before the development of membrane-bound isoforms in the smooth endoplasmic reticulum. In this regard, a mitochondrial form of 3βHSD has been found to exist in bovine adrenal cortical cells (49). The primary and rogen end product was different for each of the three stages of Leydig cell differentiation, with AO being the primary androgen released by progenitor Leydig cells, and 3a-DIOL and T, respectively, the primary androgens for immature and adult Levdig cells.

Synthesis of T and 3α -DIOL require 17β HSD reductive activity, and their low release rates from progenitor cells

tively. Determination of enzyme activity was performed as described in *Materials and Methods*. The enzyme activities of P450_{scc} (A), 3 β HSD (B), 17 α -hydroxylase (C), C17–20 lyase (D), 17 β HSD (E), 5 α -reductase (F), and 3 α HSD (G) were evaluated in PLC, ILC, and ALC. Values represent the mean \pm sE from 9–15 different samples in 3–5 independent experiments. Shared alphabet letters denote groups that were not significantly different at P < 0.05.

FIG. 3. Quantification of steady state mRNA levels for T biosynthetic and metabolizing enzymes in Leydig cells during pubertal development. Progenitor Leydig cells (PLC), immature Leydig cells (ILC), and adult Leydig cells (ALC) were isolated from 21-, 35-, and 90-dayold rats, respectively. Steady state mRNA levels of enzymes were measured by RT-PCR and normalized to RPS 16, an internal control. The steady state mRNA levels for $P450_{scc}$ (A), $_{3\beta}$ HSD (B), P450_{c17} (C), type III 17 β HSD (D), type I 5 α -reductase (E), and type I 3α HSD (F) were evaluated in PLC, ILC, and ALC. Values represent the mean \pm se from three different assays in three independent experiments. Shared alphabet letters denote groups were not significantly different at P <0.05.



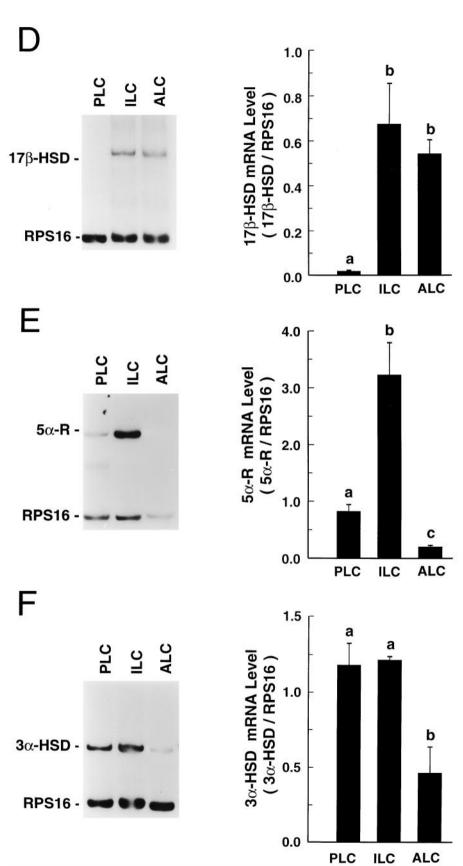


FIG. 3. Continued

suggest that 17β HSD activity is low in these cells. AO and 3α -DIOL require the activities of 5α R and 3α HSD, and the high release rates for these steroids by progenitors and immature Leydig cells indicate that androgen-metabolizing enzymes are abundantly expressed in Leydig cells before the completion of puberty, with declining levels in adulthood. Direct measurements of enzyme activities in purified cells explained the changes in end products released during Leydig cell differentiation. In progenitor cells, several androgen biosynthetic enzyme activities attained half of the more mature values, with the exception of 17β HSD, which was low, resulting in negligible conversion of DIONE to T. However, progenitor Leydig cells readily converted DIONE to 5α and rost ane dione by $5\alpha R$ and ultimately to the and rogen end product, AO, through 3α HSD, because the activities of both $5\alpha R$ and $3\alpha HSD$ were high. Although T biosynthetic enzymes nearly all attained adult values in immature Leydig cells, T formed in these cells was rapidly metabolized to another and rogen end product, 3α -DIOL. Adult Leydig cells contain significant amounts of 3α HSD activity. However, the absence of 5α -reductase in adult Leydig cells eliminates the catabolism of T, making this steroid the primary androgen end product, as T is not a substrate for 3α HSD. The 3α HSD activity present in adult Leydig cells may act to metabolize dihydrotestosterone, which is still produced at trace levels in the mature testis.

Steroidogenic enzyme activities have been measured in the developing testis (5–7). However, these studies report that 3α -DIOL is the primary androgen secreted from the immature testis on day 21 postpartum, when progenitor Leydig cells are present (2–5). Given the negligible 3α -DIOL production that was observed in progenitor Leydig cells, the predominance of this steroid in immature rat testis indicates that high levels of 17 β HSD are expressed by other cell types in the testis. The seminiferous tubules and Sertoli cells in particular contain 17 β HSD activity (10, 11). Indeed, the testis remains capable of 17 β HSD activity when Leydig cells are completely destroyed by the Leydig cell toxicant ethylene-1,2-dimethanesulfonate, with only a 50% decline in conversion of DIONE to T (12).

The present data indicate that the development of steroidogenic enzyme activities results from sequential, rather than simultaneous, induction of steroidogenic enzyme gene transcription. The steady state mRNA levels of steroidogenic enzymes were measured in Leydig cells to evaluate the relationship between mRNA levels and enzyme activity. We demonstrated that mRNA levels for T biosynthetic enzymes were significantly lower in progenitor Leydig cells than in immature and adult Leydig cells. Of the T biosynthetic enzymes, type III 17βHSD was lowest in progenitor cells, consistent with the enzyme activity data. The presence of type III 17βHSD mRNA in rat Leydig cells indicated that this isoform encodes Leydig cell-specific 17βHSD, which is also expressed in human and mouse testes (31, 32). Levels of type I $5\alpha R$ were highest in immature Leydig cells, followed by progenitor and adult Leydig cells, confirming the transitory expression of this enzyme activity during pubertal development. Although type II $5\alpha R$ is reported to be in rat testis (8, 42), this isoform was undetectable in Leydig cells.

In summary, these data demonstrate that biosynthetic and

androgen-metabolizing capacities are separately modulated in Leydig cells during pubertal differentiation and that this has significant consequences for the overall rate of T production. Progenitor Leydig cells primarily produce AO, with 3α -DIOL predominant in immature Leydig cells and T predominant in adult Leydig cells. During the transition from progenitor to immature Leydig cells, increased type III 17β HSD mRNA and enzyme activities confer adult levels of T biosynthetic capacity. During the later transition from immature to adult Leydig cell, loss of type I 5α R and decline of type I 3α HSD lower androgen metabolism, making T the primary androgen end product.

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