

# Androgen Receptor Roles in Spermatogenesis and Fertility: Lessons from Testicular Cell-Specific Androgen Receptor Knockout Mice

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Androgens are critical steroid hormones that determine the expression of the male phenotype, including the outward development of secondary sex characteristics as well as the initiation and maintenance of spermatogenesis. Their actions are mediated by the androgen receptor (AR), a member of the nuclear receptor superfamily. AR functions as a ligand-dependent transcription factor, regulating expression of an array of androgen-responsive genes. Androgen and the AR play important roles in male spermatogenesis and fertility. The recent generation and characterization of male total and conditional AR knockout mice from different laboratories demonstrated the necessity of AR signaling for both external and internal male phenotype development. As expected, the male total AR knockout mice exhibited female-typical external appearance (including a vagina with a blind end and a clitoris-like phallus), the testis was located abdominally, and germ cell development was severely disrupted, which was similar to a human complete androgen insensitivity syndrome or testicular feminization mouse. However, the process of spermatogenesis is highly dependent on autocrine and paracrine communication among testicular cell types, and the disruption of AR throughout an experimental animal cannot answer the question about how AR in each type of testicular cell can play roles in the process of spermatogenesis. In this review, we provide new insights by comparing the results of cell-specific

AR knockout in germ cells, peritubular myoid cells, Leydig cells, and Sertoli cells mouse models that were generated by different laboratories to see the consequent defects in spermatogenesis due to AR loss in different testicular cell types in spermatogenesis. Briefly, this review summarizes these results as follows: 1) the impact of lacking AR in Sertoli cells mainly affects Sertoli cell functions to support and nurture germ cells, leading to spermatogenesis arrest at the diplotene primary spermatocyte stage prior to the accomplishment of first meiotic division; 2) the impact of lacking AR in Leydig cells mainly affects steroidogenic functions leading to arrest of spermatogenesis at the round spermatid stage; 3) the impact of lacking AR in the smooth muscle cells and peritubular myoid cells in mice results in similar fertility despite decreased sperm output as compared to wild-type controls; and 4) the deletion of AR gene in mouse germ cells does not affect spermatogenesis and male fertility. This review tries to clarify the useful information regarding how androgen/AR functions in individual cells of the testis. The future studies of detailed molecular mechanisms in these *in vivo* animals with cell-specific AR knockout could possibly lead to useful insights for improvements in the treatment of male infertility, hypogonadism, and testicular dysgenesis syndrome, and in attempts to create safe as well as effective male contraceptive methods. (*Endocrine Reviews* 30: 119–132, 2009)

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Abbreviations: ACTB,  $\beta$ -Actin; AIS, androgen insensitivity syndrome; AMH, anti-Müllerian hormone; AR, androgen receptor; AR<sup>-/-</sup>, AR knockout male; AR<sup>+/+</sup>, wild-type AR male; AR<sup>+/-</sup>, wild-type AR female; CAIS, complete androgen insensitivity syndrome; Cre, cre recombinase; DHT, 5 $\alpha$ -dihydroxytestosterone; floxed, flanked by lox site; G-AR<sup>-/-</sup>, germ cell-specific AR knockout; 3 $\beta$ -HSD, 3 $\beta$ -hydroxysteroid dehydrogenase; 17 $\beta$ -HSD, 17 $\beta$ -hydroxysteroid dehydrogenase; L-AR<sup>-/-</sup>, Leydig cell-specific AR knockout; LHR, LH receptor; PM, peritubular myoid; PM-AR<sup>-/-</sup>, PM cell-specific AR knockout; P450c17, 17 $\alpha$ -hydroxylase or 17,20 lyase; P450scc, P450 side-chain cleavage; RLF, relaxin-like factor; S-AR<sup>-/-</sup>, Sertoli cell-specific AR knockout; StAR, steroidogenic acute regulatory protein; T-AR<sup>-/-</sup>, total AR knockout; *Tfm*, testicular feminization; TSP-2, thrombospondin type 2.

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## I. Introduction

**S**PERMATOGENESIS (EXOCRINE) and androgen biosynthesis (endocrine) are the major functions of mammalian testis. Both functions are complicated and highly regulated. Spermatogenesis is a process of generating mature sperm with half the number of chromosomes (haploid) produced from germ cell precursors (diploid). Androgens, by signaling through the androgen receptor (AR), mediate a wide range of physiological responses and developmental processes, involving both reproductive and nonreproductive systems in the male (1–3). The appropriate regulation of androgen activity via the hypothalamic-pituitary-testis axis is necessary for development of the male phenotype, as well as for initiation and maintenance of spermatogenesis (2, 4).

AR, which has been localized to the long arm of the X chromosome (at Xq11-12), is a member of the nuclear receptor superfamily and acts as a ligand-inducible transcription factor to modulate expression of target genes (5–7). The binding of testosterone or its metabolite 5 $\alpha$ -dihydroxytestosterone (DHT) to AR induces receptor dimerization, facilitating the ability of AR to bind to its cognate response elements and recruit coregulators to promote the expression of target genes (4, 8). Failure of the mutated receptor to activate its target genes causes a spectrum of hereditary disorders of androgen insensitivity syndrome (AIS) or testicular feminization (*Tfm*) mutation (9–12). The humans with the most severe type of AIS, presumably complete AIS (CAIS), display feminized external genitalia and intraabdominal testis and do not undergo development of secondary sex characteristics at puberty. In less severe cases, where some level of androgen responsiveness is maintained, there are a wide range of phenotypes. In addition to men, the inherited syndrome of *Tfm* has been described in several species, including the dog (13), the rat (14), the mouse (15, 16), and the cat (17). In those species, the affected male pseudohermaphrodites manifest several common features, such as feminized external genitalia, an absence of internal genitalia other than testis (13–16), and marked resistance to endogenous and exogenous androgens (14, 16).

The mammalian testis has two distinct functional compartments known as the seminiferous tubules and the interstitium, with spermatogenesis arising in the seminiferous tubules and androgen biosynthesis in the interstitial Leydig cells. Both testicular compartments contain a variety of different cell types, and the spermatogenesis is highly dependent on autocrine and paracrine communication among all cell types (18). The interstitial space, or space between the seminiferous tubules, consists of the testosterone-producing Leydig cells (19), macrophages, perivascular smooth muscle cells, and vascular endothelial cells. The Sertoli cells comprise the main structural component of the seminiferous tubules. They are responsible for the structural support for germ cell development (20, 21), facilitating germ cell movement and mature germ cell release (18, 22), and secretion of diverse functional glycoproteins and peptides to nourish

germ cells (23, 24), as well as maintenance of the blood-testis barrier, and secretion of seminiferous tubular fluid (25).

The blood-testis barrier is located in the basal third of the seminiferous tubules and segregates the seminiferous tubules into the basal compartment (containing spermatogonia, preleptotene, and leptotene spermatocytes) and the adluminal compartment (containing different stages of meiotic spermatocytes, round spermatids, elongated spermatids, and spermatozoa). The blood-testis barrier is known to function as a natural barrier to regulate the passage of various molecules into and out of the adluminal compartment of seminiferous tubules, and an immunological barrier to create a specialized environment for the differentiation and movement of developing germ cells (22).

Peritubular myoid (PM) cells are mesenchymal cells that form the outer border of the seminiferous tubules and, in conjunction with Sertoli cells, produce the basement membrane required to maintain normal tubule morphology. Several important functions of PM cells have been proposed: 1) they can synthesize several secretory products, such as P-Mod-S (peritubular factors that modulate Sertoli cell function), to modulate a number of Sertoli cell functions (26, 27); and 2) they have the ability to contract, thus inducing peristalsis-like waves and impulses in the seminiferous tubule, which will help the transport of spermatozoa through the tubular lumen and into the epididymis to control testicular output of both fluid and sperm (28). PM cells have also been indicated as one of the principal target cells in testis that respond to androgens and participate in the androgenic control of spermatogenesis (29).

There is a general agreement that AR can be detected in Sertoli cells, PM cells, and cells in the interstitial spaces including Leydig cells and perivascular smooth muscle cells in testis (30–35). However, using antibody staining, the localization of the AR in male germ cells remains controversial. Several studies indicated that AR is present in germ cells in different species (32, 34–41), but other reports show that there is no AR staining in the germ cells (30, 33, 42–46). Interestingly, some previous studies have revealed that the AR expression in male germ cells is stage specific, only expressing in elongated spermatid at spermatogenic stage XI in rat testis (34). Furthermore, the presence of testosterone binding sites has been detected on monkey sperm and correlated with motility (41). The presence of the AR in human sperm has been demonstrated by Western blot and by immunofluorescence assay (38). These studies suggest that AR might play a direct role in germ cells.

The recent generation and characterization of male total AR knockout (T-AR<sup>-/y</sup>) mice, which confirmed the similar phenotype to a human with CAIS or *Tfm* mouse, exhibited female-typical external appearance (including a vagina with a blind end and a clitoris-like phallus). Testes were located abdominally, and germ cell development was severely disrupted. These observations further emphasized the necessity of AR signaling for both external and internal male phenotype development (47, 48). However, a gene alteration in the whole body might cause a complex phenotype in which it is hard to differentiate direct effects in a particular tissue or particular cell type from those secondary effects arising from a gene change in other cell types. In view of the fact that the

AR has developmental roles in establishing the male phenotype and the process of spermatogenesis is highly dependent on autocrine and paracrine communication among testicular cell types, the disruption of AR throughout an experimental animal cannot answer the question about how AR in each cell type within the testis can play roles in the spermatogenesis. Therefore, it is necessary to use a cre-lox strategy in transgenic mice with cell-specific expression of the cre recombinase (Cre) to generate a mouse model in which disruption of AR function is exclusively in a particular cell type in the testis. Here, we summarize the results of cell-specific AR knockout ( $AR^{-/y}$ ) in germ cells (G- $AR^{-/y}$ ), peritubular myoid cells (PM- $AR^{-/y}$ ), Leydig cells (L- $AR^{-/y}$ ), and Sertoli cells (S- $AR^{-/y}$ ) mouse models to understand the spermatogenic consequences of AR loss in different testicular cell types.

## II. Generation of Various Testicular Cell-Specific Androgen Receptor (AR) Knockout Mice

To generate tissue-specific  $AR^{-/y}$  mice, a Cre-loxP strategy for conditional knockout is necessary. The Cre-loxP system utilizes the expression of P1 phage Cre to catalyze the excision of DNA located between flanking loxP sites (49). This strategy differs from the standard targeted disruption procedure in that embryonic stem cells are generated in which the target segment is not disrupted but is flanked by loxP sites (floxed). The target gene thus functions normally, and mice can be bred to homozygosity for the targeted locus. The C57-B6/129/SvEv loxP-floxed AR mice were first generated (47, 48) and then mated with different testicular cell-specific Cre mice for the study of total or cell-specific  $AR^{-/y}$  mice (Fig. 1). T- $AR^{-/y}$  mice were generated by mating the floxed AR mice with  $\beta$ -actin (*ACTB*)-Cre (*ACTB*-Cre is under the control of the  $\beta$ -actin promoter) transgenic mice (47); S- $AR^{-/y}$  mice were generated by mating the floxed AR mice with anti-Müllerian hormone (*AMH*)-Cre (*AMH*-Cre is under the control of the anti-Müllerian hormone promoter) transgenic mice (50–53); L- $AR^{-/y}$  mice were generated by

mating the floxed AR mice with *AMHRII*-Cre (*AMHRII*-Cre is under the control of the anti-Müllerian hormone receptor II promoter) transgenic mice (54); PM- $AR^{-/y}$  mice were generated by mating the floxed AR mice with *Transgelin*-Cre (*Transgelin*-Cre is under the control of the transgelin promoter) transgenic mice (55); and G- $AR^{-/y}$  mice were generated by mating the floxed AR mice with *Sycp1*-Cre [*Sycp1*-Cre is under the control of the synaptonemal complex protein 1 promoter, expressed at an early stage of the male meiosis (leptotene to zygotene)] transgenic mice (56). Male wild-type AR ( $AR^{+/y}$ ) or floxed AR littermate mice without Cre transgene were used as control  $AR^{+/y}$  to study the AR roles in individual cells within the testis.

## III. Serum Testosterone Levels in Various Testicular Cell-Specific AR Knockout Mice

Serum testosterone levels are influenced by two key factors: testosterone biosynthesis rate in the Leydig cells, and number of functional Leydig cells available in the testis.

### A. Testosterone biosynthesis in the Leydig cells

The conversion of cholesterol to testosterone involves a number of steps that are catalyzed by enzymes, including hydroxylase [17 $\alpha$ -hydroxylase (P450c17)], two cleavage enzymes [P450 side-chain cleavage (P450scc) and 17,20 lyase (P450c17)], and one isomerization  $\Delta^5$ - $\Delta^4$  enzyme [3 $\beta$ -hydroxysteroid dehydrogenase (3 $\beta$ -HSD), conversion of  $\Delta^5$ -3 $\beta$ -hydroxysteroids to  $\Delta^4$ -3-ketosteroids], predominantly belonging to the cytochrome P450 family. LH, through the LH receptor (LHR) on the surface of Leydig cells, directly stimulates the synthesis of a steroidogenic acute regulatory (StAR) protein and the outer mitochondrial membrane translocator protein. StAR and outer mitochondrial membrane translocator protein are key regulators of cholesterol transport from the outer to the inner mitochondrial membrane (57, 58). Once delivered to the inner mitochondrial membrane, cholesterol is enzymatically converted to pregnenolone by the enzyme cytochrome P450scc (59). Pregnenolone diffuses across the mitochondrial membranes and might progress to testosterone production through either the  $\Delta^4$  or  $\Delta^5$  pathway (60) by enzymes associated with the smooth endoplasmic reticulum. Pregnenolone can be converted to progesterone through the enzyme 3 $\beta$ -HSD (the  $\Delta^4$  pathway) or can be hydroxylated at the 17 $\alpha$  position by the enzyme 17 $\alpha$ -hydroxylase to form 17 $\alpha$  hydroxypregnenolone (the  $\Delta^5$  pathway). The  $\Delta^5$  pathway predominates in human Leydig cells (61), whereas in the rodent, both pathways are catalyzed, although the  $\Delta^4$  pathway is preferred (62). The further conversion of 17 $\alpha$  hydroxypregnenolone (the  $\Delta^5$  pathway) involves the formation of the 19C steroid dehydroepiandrosterone catalyzed by the enzyme 17,20 lyase. Both 17 $\alpha$ -hydroxylation and cleavage of the 17,20 carbon-carbon bond are catalyzed by a single microsomal enzyme cytochrome P450c17, which is encoded by a single gene on chromosome 10 (63, 64). The final step in the production of testosterone in the testis is via conversion of androstenedione (a weak androgen) to testosterone in the  $\Delta^4$  pathway or dehydroepiandrosterone to androstenediol in the  $\Delta^5$  pathway by the

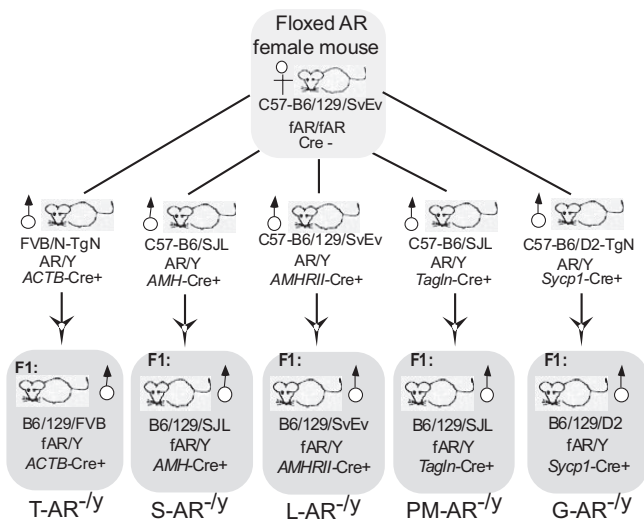


FIG. 1. The mating strategy to generate T- $AR^{-/y}$ , S- $AR^{-/y}$ , L- $AR^{-/y}$ , PM- $AR^{-/y}$ , and G- $AR^{-/y}$  mice.

enzyme 17 $\beta$ -hydroxysteroid dehydrogenase (17 $\beta$ -HSD)/17-ketosteroid reductase (65). Biosynthesis of all biologically active steroid hormones requires conversion of  $\Delta^5$ -3 $\beta$ -hydroxysteroids to  $\Delta^4$ -3-ketosteroids, and this reaction is catalyzed by the enzyme 3 $\beta$ -HSD (66). Notably, the 3 $\beta$ -HSD type I isoform is expressed at significant levels in fetal-type Leydig cells, whereas the expression of 3 $\beta$ -HSD type VI isoform is specific to the adult population of Leydig cells in the mouse testis (67). Testosterone is the principal steroid produced and secreted by the testis, although numerous other steroids (C18, C19, and C21) are also produced (68, 69).

### B. Leydig cell development and maturation

In mammals, there are two distinct populations of Leydig cells arising in a sequential manner during normal testicular development (70, 71). In the mouse testis, the first population, fetal Leydig cells, develops at around 12.5 d after coitum after normal testicular differentiation (71, 72) and starts to produce testosterone at around 13 d after coitum for masculinization of the male urogenital system (71, 73). The fetal Leydig cell population persists after birth but becomes ancillary to the newly developed secondary population, adult Leydig cells, which originate from undifferentiated mesenchymal spindle-shaped precursors in the testicular interstitium (71, 74). The intermediates of adult Leydig cells first become apparent by around postnatal d 7–11 (67, 75, 76). Subsequently, the total number of Leydig cell precursors increases rapidly in a LH-dependent manner between postnatal d 10 and d 20, finally reaching a maximal number at about postnatal d 35 (77–79). Earlier studies reviewing the morphological change of adult Leydig cell population showed that the mesenchymal spindle-shaped precursor cells are mainly found in the peritubular region in the interstitium and are transformed to the progenitor Leydig cells in the same region. Subsequently, the progenitor Leydig cells are transformed from spindle-shaped to round immature adult Leydig cells, most commonly seen in the testis during postnatal d 28 to d 56 (76). The immature adult Leydig cells contain numerous lipid droplets and abundant smooth endoplasmic reticulum and are moved to the central interstitial region. An increase in cell size, the volume of smooth endoplasmic reticulum, and the decline in cytoplasmic lipid droplets characterize the transition between immature and mature adult Leydig cells by postnatal d 56 (76, 79). The factors regulating the proliferation and differentiation of the two populations of Leydig cells are still largely unknown, but gonadotropins such as LH (80, 81) and FSH (82, 83), thyroid hormone (84, 85), adrenocorticotrophic hormone (81), estrogens (86), and androgens (87) are suggested to play an important role. In addition, a body of evidence either from *in vitro* studies using testicular-derived mesenchymal cells in culture or from *in vivo* studies using the *Tfm* mice have demonstrated that the development of adult Leydig cell function definitely requires the action of androgens (88, 89).

### C. *Tfm* mice and humans with AIS

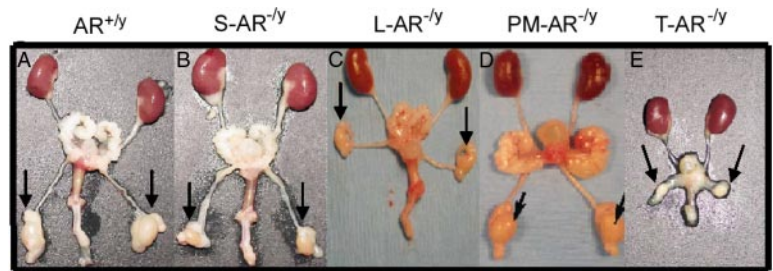
The serum testosterone levels in *Tfm* mice are reduced ( $1.8 \pm 0.3$  vs.  $9.3 \pm 2.0$  nmol/liter;  $P < 0.05$ ) (90), and serum

LH levels are increased as compared with AR<sup>+/y</sup> controls (89, 91). In addition, examination of Leydig cell-specific steroidogenic enzyme and functional gene expressions revealed that in adult *Tfm* mice, the 17 $\beta$ -HSD III, 3 $\beta$ -HSD VI, and prostaglandin synthetase gene expressions are barely detectable, and the P450c17 and relaxin-like factor (RLF) gene expressions are largely reduced compared with AR<sup>+/y</sup> controls. However, the gene expression levels of P450scc, StAR, LHR, 3 $\beta$ -HSD I, and thrombospondin type 2 (TSP-2) are normal or increased in adult *Tfm* mice compared with AR<sup>+/y</sup> controls. The highly abnormal patterns of Leydig cell-specific gene expression in *Tfm* mice shows that the initial differentiation of adult Leydig cells occurs, but there is a developmental failure of adult Leydig cell maturation, which leads to the adult Leydig cells in *Tfm* mice only acquiring partial maturational characteristics of the adult Leydig cell (92). These findings are consistent with the earlier studies in which the *Tfm* mice showed elevated plasma LH levels, increased Leydig cell oxidative enzyme activities indicating that Leydig cells were hyperstimulated, and reduced testosterone production and secretion (93). Earlier studies showed that the Leydig cell number increased 26-fold from postnatal d 12 to d 140, wherein most increases in Leydig cell number occurred during the period between postnatal d 12 and d 50 (94). The evidence from *Tfm* mice showed that the Leydig cell number and function are normal up to postnatal d 5, indicating that fetal Leydig cell proliferation, development, and function are not dependent on androgens before postnatal d 5 (92). The normal prepubertal rise in Leydig cell number is significantly reduced and only reaches 30% of AR<sup>+/y</sup> control values on postnatal d 20. Between postnatal d 20 and adulthood, Leydig cell number shows a similar trend of increase in both AR<sup>+/y</sup> control and *Tfm* mice, indicating that only the early part of the developmental phase is associated with AR function (92). In contrast to rodents, wherein the AR might play a role in the differentiation of Leydig cell precursors into mature adult Leydig cells, the humans with CAIS have relatively normal or slightly increased levels of plasma testosterone, elevated LH levels, and normal FSH levels (12, 95, 96), and Leydig cells are shown to have hyperplastic changes (97). In consideration of the wide variation of clinical characteristics in AIS and the wide variation of serum testosterone levels between individuals, there are still debates existing in regard to this issue in human patients.

### D. T-AR<sup>-/y</sup> mice

The serum testosterone levels in T-AR<sup>-/y</sup> mice are decreased ( $7.9 \pm 0.8$  vs.  $192 \pm 13.1$  ng/dl;  $P < 0.05$ ), and serum LH levels are increased compared with AR<sup>+/y</sup> controls in our study (Fig. 2F) (50, 56). These results are similar to *Tfm* mouse (90). Another study also showed the similar trend of serum testosterone level changes in T-AR<sup>-/y</sup> mice compared with AR<sup>+/y</sup> controls, although this was not statistically significant due to a wide variation of serum testosterone levels between animals within the same genotype groups (94). The results of quantitative RT-PCR for Leydig cell-specific steroidogenic enzymes and functional gene expressions revealed that adult T-AR<sup>-/y</sup> mice have reduced P450c17 gene expression and increased 3 $\beta$ -HSD I and P450scc gene expressions compared

FIG. 2. The internal genitalia of 14-wk-old male  $AR^{+/y}$  (A),  $S-AR^{-/y}$  (B),  $L-AR^{-/y}$  (C),  $PM-AR^{-/y}$  (D), and  $T-AR^{-/y}$  (E). Arrows indicate the testis. [Sections of figures were reproduced from Refs. 50 and 55. Copyright 2004 and 2006, respectively, Proceedings of the National Academy of Sciences.] F, Body weight, testis weight, and serum testosterone levels (mean  $\pm$  SEM) of 14-wk-old male  $AR^{+/y}$ ,  $G-AR^{-/y}$ ,  $PM-AR^{-/y}$ ,  $S-AR^{-/y}$ ,  $L-AR^{-/y}$ , and  $T-AR^{-/y}$  mice. <sup>a</sup> Significant difference ( $P < 0.05$ ; *t* test) compared with  $AR^{+/y}$ . All of these results originated from our previous publications. The other studies also showed the similar trend of serum testosterone level changes in  $T-AR^{-/y}$  mice and  $S-AR^{-/y}$  mice compared with  $AR^{+/y}$  controls, although this was not statistically significant due to a wide variation of serum testosterone levels between animals within the same genotype groups.



F Body weight, testis weight and serum testosterone level (Mean $\pm$ SEM) at 14 weeks old mice.

Group	N	BW (gm)	Testis Weight (mg)	Serum testosterone level (ng/dl)
$AR^{+/y}$ (B6)	10	29.91 $\pm$ 0.85	110.6 $\pm$ 5.04	192.03 $\pm$ 13.1
$G-AR^{-/y}$	10	27.94 $\pm$ 0.78	105.6 $\pm$ 3.48	186.10 $\pm$ 15.3
$PM-AR^{-/y}$	10	28.43 $\pm$ 0.38	87.7 $\pm$ 1.47 <sup>a</sup>	159.99 $\pm$ 17.9
$S-AR^{-/y}$	10	26.84 $\pm$ 0.78	23.8 $\pm$ 0.64 <sup>a</sup>	48.48 $\pm$ 13.8 <sup>a</sup>
$L-AR^{-/y}$	10	28.52 $\pm$ 0.49	35.3 $\pm$ 1.71 <sup>a</sup>	20.14 $\pm$ 1.5 <sup>a</sup>
$T-AR^{-/y}$	10	27.76 $\pm$ 0.50	7.7 $\pm$ 0.13 <sup>a</sup>	7.90 $\pm$ 0.8 <sup>a</sup>

with  $AR^{+/y}$  controls (94). Moreover, the result from  $T-AR^{-/y}$  mice showed that the Leydig cell number reached 37% of  $AR^{+/y}$  values on postnatal d 12 and reached only 22% of  $AR^{+/y}$  values on postnatal d 140. Similarly, the Leydig cell cytoplasmic volume in  $T-AR^{-/y}$  mice achieved only 60% of  $AR^{+/y}$  values on postnatal d 140 (94). Overall, the result from the above studies clearly demonstrated that loss of AR in whole testis, including Leydig cells, PM cells, and Sertoli cells, will cause a profound functional defect in Leydig cells. There are reduced Leydig cell numbers, size, and steroidogenic enzyme gene expressions in  $T-AR^{-/y}$  mice compared with  $AR^{+/y}$  controls, but these findings in  $T-AR^{-/y}$  mice are not explained by the coincidental cryptorchidism (92).

#### E. $S-AR^{-/y}$ mice

The serum testosterone levels in  $S-AR^{-/y}$  mice are decreased by 74.7% of  $AR^{+/y}$  values (48.48  $\pm$  13.8 vs. 192  $\pm$  13.1 ng/dl;  $P < 0.05$ ) (Fig. 2F), and the serum LH levels are increased compared with  $AR^{+/y}$  controls in our study (50, 53). The other studies also showed the similar trend of serum testosterone level changes in  $S-AR^{-/y}$  mice compared with  $AR^{+/y}$  controls, although this was not statistically significant (51, 94). But one independent study has shown almost a 40-fold increase in serum testosterone levels in both *loxP*-floxed AR mice and  $S-AR^{-/y}$  mice compared with  $AR^{+/y}$  controls (52). In view of the wide variation of serum testosterone levels between animals within genotype groups, this issue is still debatable. The results of quantitative RT-PCR for Leydig cell-specific steroidogenic enzyme and functional gene expressions also revealed that adult  $S-AR^{-/y}$  mice have significant increases of P450<sub>scc</sub>, P450<sub>c17</sub>, and 3 $\beta$ -HSD I gene expressions compared with  $AR^{+/y}$  controls (94). Our studies also found that adult  $S-AR^{-/y}$  mice have significant increases of P450<sub>c17</sub>, 3 $\beta$ -HSD I, 3 $\beta$ -HSD VI, 17 $\beta$ -HSD III, and LHR compared with  $AR^{+/y}$  controls (data not shown). The evidence from  $S-AR^{-/y}$  mice showed that the Leydig cell numbers are comparable with  $AR^{+/y}$  controls up to postnatal d 12, and progressive decreases in Leydig cell numbers are evident at all later ages; for example, the Leydig cell number is reduced by 42% of  $AR^{+/y}$  values on postnatal d 20 and by 54% of  $AR^{+/y}$  values on postnatal d 140. In contrast, the

Leydig cell cytoplasmic volume in  $S-AR^{-/y}$  mice is 23% greater than  $AR^{+/y}$  controls on postnatal d 140 (94). Based on above  $T-AR^{-/y}$  and  $S-AR^{-/y}$  mice data, it is suggested that the androgen/AR signals in Sertoli cells contribute to the establishment of approximately 50% of adult Leydig cell number in testes, whereas the establishment of normal Leydig cell size is not dependent on such action. In addition, the evidence indicated that androgen/AR signals on testicular cell types other than Sertoli cells are essential for the establishment of approximately 25% of the final adult Leydig cell number and the development of normal adult Leydig cell size (94).

Despite increased Leydig cell cytoplasmic volume and increased Leydig cell steroidogenic activity in the testis of  $S-AR^{-/y}$  mice, our study showed that the serum testosterone level is lower than  $AR^{+/y}$  control (Fig. 2F). The possible explanation for these discrepancies could be that, although the Leydig cells might have gain-of-function in the environment of low serum testosterone levels in  $S-AR^{-/y}$  mice, they cannot restore the normal serum testosterone values due to the 78.5% decline in testicular size (Fig. 2F and Table 1) and the 50% decrease in Leydig cell numbers.

#### F. $L-AR^{-/y}$ mice

The serum testosterone levels in  $L-AR^{-/y}$  mice are decreased (20.14  $\pm$  1.5 vs. 192  $\pm$  13.1 ng/dl;  $P < 0.05$ ) (only reached 10.5% of  $AR^{+/y}$  values) (Fig. 2F), and the serum LH levels are increased compared with  $AR^{+/y}$  controls (54, 56). The results of quantitative RT-PCR for Leydig cell-specific steroidogenic enzyme and functional gene expressions revealed that adult  $L-AR^{-/y}$  mice have significant decreases in P450<sub>c17</sub>, 3 $\beta$ -HSD VI, 17 $\beta$ -HSD III, and RLF, as well as significant increases in LHR and TSP-2 gene expressions compared with  $AR^{+/y}$  controls (54). Earlier reports have shown that RLF, 3 $\beta$ -HSD VI, and 17 $\beta$ -HSD III are mainly expressed in the adult Leydig cells and barely detectable in the fetal Leydig cells; therefore they act as markers for adult Leydig cell differentiation. In contrast, the expression pattern of TSP-2 is predominantly in the fetal Leydig cell population, and the expression levels are gradually decreased after puberty due to the rapid increase of adult Leydig cell popula-

TABLE 1. Comparison of reproductive phenotypes in T-AR<sup>-/-</sup>, G-AR<sup>-/-</sup>, S-AR<sup>-/-</sup>, L-AR<sup>-/-</sup>, and PM-AR<sup>-/-</sup> mice

	T-AR <sup>-/-</sup>	G-AR <sup>-/-</sup>	S-AR <sup>-/-</sup>	L-AR <sup>-/-</sup>	PM-AR <sup>-/-</sup>
H-P-T axis					
Serum FSH	Elevated	Normal	Normal	Elevated	Normal
Serum LH	Elevated	Normal	Elevated	Elevated	Normal
Serum testosterone	Decreased	Normal	Decreased	Decreased	Normal
Testicular function					
Testis size	Decreased	Normal	Decreased	Decreased	Decreased
Testis size/WT testis size (%)	7	Normal	23.40	31.10	79.00
Epididymal sperm count	No epididymis	Within normal range	No sperm	No sperm	Decreased by ~57% of WT sperm count
General histology observation	No lumen formation; germ cell hypoplasia and development to pachytene spermatocyte	Normal full range of spermatogenesis	Decrease of lumen formation; germ cell development to diplotene spermatocyte	Decrease of lumen formation; germ cell development to round spermatid	Normal full range of spermatogenesis
Overall fertility	Infertile	Normal fertility	Infertile	Infertile	Normal fertility

H-P-T, Hypothalamic-pituitary-testis; WT, wild-type. All of these results originated from our previous publications. The other studies also showed the similar trend of serum testosterone level changes in T-AR<sup>-/-</sup> mice and S-AR<sup>-/-</sup> mice as compared to AR<sup>+/+</sup> controls, although this was not statistically significant due to a wide variation of serum testosterone levels between animals within the same genotype groups.

tion (78). Looking at the results of L-AR<sup>-/-</sup> mice, the markers of adult Leydig cell differentiation are all significantly decreased, but they are still detectable, and the TSP-2 gene expression levels are significantly increased compared with AR<sup>+/+</sup> controls (54). In contrast, we found that the Leydig cell number and size in L-AR<sup>-/-</sup> mice are comparable with AR<sup>+/+</sup> controls. These findings fit the hypothesis that the early initiation of adult Leydig cell differentiation and prepubertal rise in adult Leydig cell numbers are normal, but the later differentiation of adult Leydig cells to establish the full capacity of steroidogenic function is affected by loss of functional AR in the Leydig cells.

#### G. PM-AR<sup>-/-</sup> mice and G-AR<sup>-/-</sup> mice

The serum testosterone levels in PM-AR<sup>-/-</sup> and G-AR<sup>-/-</sup> mice are comparable to AR<sup>+/+</sup> controls (Fig. 2F), which implies that the functional AR in PM cells and germ cells are not essential for Leydig cell development, differentiation, and steroidogenesis.

#### IV. Phenotypes of External Genitalia and Internal Male Accessory Genital Organ Size in Various Testicular Cell-Specific AR Knockout Mice

It is generally agreed that the testicular differentiation from the bipotential gonad in early embryonic development is not an androgen/AR-dependent process. However, the early differentiation of male accessory sex organs during embryonic development is dependent on testosterone when the enzymes necessary to convert testosterone to DHT do not yet exist. The major effect of testosterone at early embryonic stages involves directing the differentiation of male-specific internal genital structures derived from the Wolffian ducts, including the epididymis, vas deferens, and seminal vesicles

(98–100). During later embryonic developmental stages, when the enzymes responsible for mediating testosterone to DHT conversion became available, DHT-dependent development of the prostate and prostatic urethra occurs internally, and the differentiation of genital structures externalizes (4). Failure of the AR to activate its target genes in the presence of androgen during these critical stages will result in severe defects in induction of male sex differentiation and development of the male phenotype.

The results from recent publications showed that T-AR<sup>-/-</sup> male mice, with AR knockout in all cell types, exhibited female-like external genitalia, such as shorter genito-anal distance (the distance from the sex papilla to the anus; 0.59 cm in T-AR<sup>-/-</sup> compared with 1.12 cm in AR<sup>+/+</sup> 8-wk-old mice), a vagina with a blind end, and a microphallus. The male internal reproductive organs originating from the Wolffian ducts (such as epididymis, vas deferens, and seminal vesicles) or urogenital sinus (prostate) were absent in T-AR<sup>-/-</sup> male mice (Fig. 2E) (47, 48, 51), which was similar to *Tfm* male mice and humans with CAIS. In contrast, results from various testicular cell-specific AR knockout mice showed that the external genitalia and internal reproductive organs are normally developed (Fig. 2, B–D). However, the epididymis size in S-AR<sup>-/-</sup> mice is decreased by 63% of AR<sup>+/+</sup> size (53), and in L-AR<sup>-/-</sup> mice is decreased by 44% of AR<sup>+/+</sup> size (54). This decrease in epididymis size in S-AR<sup>-/-</sup> and L-AR<sup>-/-</sup> mice could be due to the lower serum testosterone levels and no mature sperm production in these mice (details will be discussed in Section VII). These *in vivo* results display the clear evidence that androgen/AR signals in Wolffian ducts, genital tubercle, and urogenital sinus during embryonic developmental stage play a critical role in the differentiation and development of internal and external male genitalia.

## V. Testis Position in Various Testicular Cell-Specific AR Knockout Mice

The testis position of T-AR<sup>-/-</sup> mice is similar to *Tfm* male mice and men with CAIS and is located anywhere in the abdomen along the pathway from the normal position of the ovaries in female mice to the inguinal region (47, 48). The developing gonads are initially attached to the posterior abdominal wall around the pararenal position by two ligamentous structures derived from the genital mesenteries. One is the cranial suspensory ligament, a ligamentous structure connecting the upper tip of the testis to the posterior abdominal wall. The other is the gubernaculum, a ligamentous structure of condensed mesenchymal cells connecting the caudal pole of the testis to the precursor structure of the scrotum. Regulation of transabdominal descent centers on the control of gubernacular enlargement and regression of the cranial suspensory ligament (98, 101, 102). Testicular descent is suggested to be mediated by the gubernaculum in that androgen/AR activity promotes outgrowth of the gubernaculum to allow caudal testis migration and is believed to be required for the inguinoscrotal phase of testicular descent (103). Androgen/AR action also drives regression of the cranial suspensory ligament, another process necessary for testis migration. Notably, both T-AR<sup>-/-</sup> male mice and *Tfm* male mice, as well as the humans with CAIS, present an intraabdominal migration of testis but a failure of inguinoscrotal migration of testis (104). The lack of an androgenic effect might affect the normal involution of the cranial suspensory ligament and enlargement of the gubernaculum, causing the testis to remain in the peritoneal cavity (101, 102). In contrast, the testis of S-AR<sup>-/-</sup>, L-AR<sup>-/-</sup>, PM-AR<sup>-/-</sup>, and G-AR<sup>-/-</sup> mice were normally descended compared with AR<sup>+/+</sup> control mice. This *in vivo* evidence further emphasizes that the extratesticular mesenchymal (gubernaculum and cranial suspensory ligament) androgen/AR signals are important in normal testis descent, especially in the phase of inguinoscrotal migration.

## VI. Testis Size in Various Testicular Cell-Specific AR Knockout Mice

### A. S-AR<sup>-/-</sup> mice

Recent studies point out that the Sertoli cell number is the main contributory factor for the capacity of sperm production per testis and the final testis size in adulthood (105–107). The exact regulatory mechanism of Sertoli cell proliferation is not clear yet, but several hormones are involved in these processes. These include FSH, which stimulates Sertoli cell proliferation (108, 109), and thyroid hormones, which regulate the cessation of Sertoli cell proliferation (105, 110). Several earlier studies showed that androgens/AR play an important role in the Sertoli cell maturation process (105, 111), rather than Sertoli cell proliferation, because the Sertoli cells do not express AR during fetal and early neonatal time periods of Sertoli cell proliferation (33, 35, 112). The expression of AR is first detectable in fetal rat testis at 14–15 d after coitus (113, 114), coincident with the time that the fetal Leydig cells begin to produce testosterone (115, 116). The

positive AR immunostaining in the fetal testis is mostly localized in the interstitial cells and PM cells. In contrast, the Sertoli cells do not obviously express AR until postnatal d 5 (114). However, recent studies using the transgenic *Tfm* mouse model (109), gene expression profiling of androgen-regulated transcripts in neonatal mice testis (117), and combinations of hormonal manipulations in neonatal rat (118) suggested that androgens/AR might play a role in Sertoli cell proliferation, especially during early postnatal testis development. Another study using gene expression profiling of AR antagonists in the fetal rat testis suggested that the fetal testis is not a major target for AR activity during this stage of testis development, and results indicated that there are no overt histology changes and no common set of gene targets among various treated groups and the control group (119).

The size of testis in adult S-AR<sup>-/-</sup> mice is decreased by 21.5% of AR<sup>+/+</sup> testis size (Fig. 2F and Table 1) (50–52, 56); however, there are studies (51, 120) to show that adult S-AR<sup>-/-</sup> mice develop nearly normal numbers of Sertoli cells as compared with AR<sup>+/+</sup> control. In agreement with the earlier studies in which androgens/AR play an important role in the Sertoli cell maturation rather than Sertoli cell proliferation, other studies also suggest that the decrease in testis size in S-AR<sup>-/-</sup> mice can be mainly due to the functional defect of Sertoli cells to support late meiotic and post-meiotic germ cells (50, 53).

### B. *Tfm* mice and T-AR<sup>-/-</sup> mice

Earlier studies indicated that *Tfm* mice have significant decreases in testis size as early as postnatal d 5 compared with AR<sup>+/+</sup> control (108, 109), and the testis of adult T-AR<sup>-/-</sup> mice were markedly reduced in size (only reached 7% of AR<sup>+/+</sup> testis size at 14 wk old; Fig. 2F and Table 1) and cryptorchid (47, 56). The testis sizes of AR<sup>+/+</sup> and T-AR<sup>-/-</sup> are comparable up to postnatal d 2 despite significant decreases in Sertoli cell numbers in T-AR<sup>-/-</sup> testis (120), but by postnatal d 5, the testis size of *Tfm* mice was significantly smaller as compared with AR<sup>+/+</sup> testis (109). The reasons for testis hypoplasia in these mice could be the result of multiple factor effects, including loss of AR function in whole testicular cell types and the effect of elevated testicular temperature due to the intraabdominal position of testis.

### C. L-AR<sup>-/-</sup> mice, PM-AR<sup>-/-</sup> mice, and G-AR<sup>-/-</sup> mice

The size of testis in adult L-AR<sup>-/-</sup> mice is decreased to 32.0% of AR<sup>+/+</sup> testis size (Fig. 2F and Table 1). The result showed that the decrease in testis size in L-AR<sup>-/-</sup> mice is mainly due to the defect of Leydig cell steroidogenic function subsequently affecting Sertoli cell functions to support post-meiotic germ cells (54, 56). The size of testis in adult PM-AR<sup>-/-</sup> mice is decreased to 79.3% of AR<sup>+/+</sup> testis size (Fig. 2F and Table 1). The result suggested that loss of functional AR in PM cells might impair the paracrine effect from PM cells to Sertoli cells, resulting in the influence of normal Sertoli cell nourishing function and reduction in testis germ cell number (55). Interestingly, the testis of G-AR<sup>-/-</sup> mice showed similar testis size compared with AR<sup>+/+</sup> testis (Fig. 2F and Table 1) (56). This result clearly demonstrated that

androgen/AR signals in germ cells do not have significant effects on sperm maturation and testis development.

### VII. Testis Morphology, Epididymal Sperm Count, and Fertility Test in Various Testicular Cell-Specific AR Knockout Mice

Qualitatively complete spermatogenesis is defined as the presence of all germ cell types, but the cell number might be subnormal, whereas quantitatively complete spermatogenesis is the presence of a full complement of germ cells both in type and number (121). Androgen replacement alone has been shown to initiate qualitatively complete spermatogenesis in the gonadotropin-deficient mice (122), but FSH replacement alone fails to rescue spermatogenesis beyond the meiotic stages in these mice (123, 124). Moreover, earlier studies using hypophysectomy to remove androgens from adult rats demonstrated an initial display of loss of midstage round spermatids and elongated spermatids (125). After long-term hypophysectomy and elimination of residual testosterone activity by flutamide or ethane dimethanesulphonate treatment, spermatogenesis rarely proceeds beyond meiosis and only a few round spermatids can be observed as well as the absence of elongated spermatids (126, 127). These results provide additional supportive evidence that androgens/AR are necessary for the completion of meiosis and the differentiation of round spermatids into the spermatozoa.

#### A. Humans with AIS, *Tfm* mice, and *T-AR*<sup>-/-</sup> mice

The histology of testis in humans with CAIS showed that a majority of the seminiferous tubules were dysgenic. Some

seminiferous tubules contained only spermatogonia and occasional meiotic spermatocytes. There were no mature germ cells in any seminiferous tubules (128).

The testicular histology of 14-wk-old *T-AR*<sup>-/-</sup> mice showed that a majority of seminiferous tubules lack germ cells, whereas others contained few germ cells, had no lumen formation in seminiferous tubules as well as decreases in diameter, and germ cell development stopped at the pachytene primary spermatocyte stage of the first meiosis division (47, 56), which was similar to *Tfm* mice (15, 16). The epididymis in *T-AR*<sup>-/-</sup> and *Tfm* mice were absent. As mentioned previously, there are reduced Leydig cell numbers, size, and steroidogenic enzyme gene expressions in *T-AR*<sup>-/-</sup> mice compared with *AR*<sup>+/-</sup> controls. But these findings in *T-AR*<sup>-/-</sup> mice are not explained by the coincidental cryptorchidism (92).

#### B. *S-AR*<sup>-/-</sup> mice

The histology of *S-AR*<sup>-/-</sup> testis showed a decrease of lumen formation as well as a decrease in diameter of seminiferous tubules and reduced germ cell complement as compared with *AR*<sup>+/-</sup> littermates. The seminiferous tubules of *S-AR*<sup>-/-</sup> mice displayed poor germ cell differentiation, with the majority of germ cell maturation ceasing at the diplotene primary spermatocyte stage (Fig. 3G) (50). Occasionally, secondary spermatocytes occur, but in greatly reduced numbers, and very few round spermatids exist (51, 52). Consequently, the results from epididymal sperm count indicated that there was no sperm existing in the epididymis of *S-AR*<sup>-/-</sup> mice, and the mating tests showed that *S-AR*<sup>-/-</sup> mice failed to impregnate the wild-

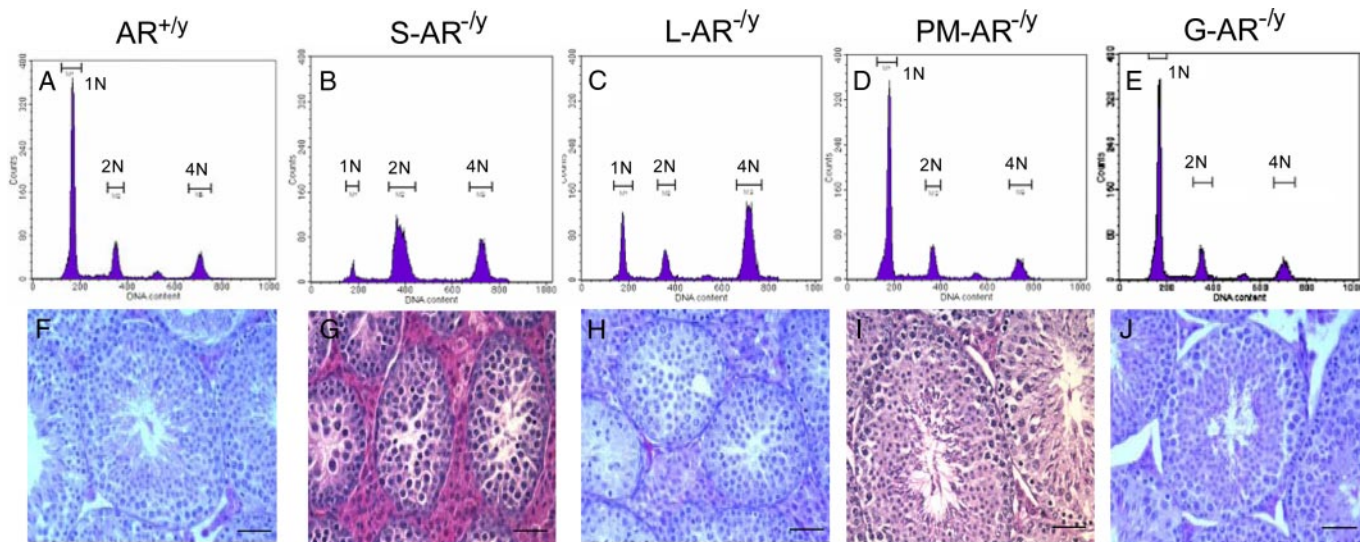


FIG. 3. Analysis of germ cell DNA content of 14-wk-old male *AR*<sup>+/-</sup> (A), *S-AR*<sup>-/-</sup> (B), *L-AR*<sup>-/-</sup> (C), *PM-AR*<sup>-/-</sup> (D), and *G-AR*<sup>-/-</sup> (E) mice by using flow cytometry. 1N represents haploid cells, 2N represents diploid cells, and 4N represents tetraploid cells. Compared with *AR*<sup>+/-</sup> testis (A), *S-AR*<sup>-/-</sup> testis showed 3-fold increase in diploid cells, 2-fold increase in tetraploid cells, and 11-fold reduced haploid cells (B); *L-AR*<sup>-/-</sup> testis showed 4-fold increase in tetraploid cells and 2.8-fold reduced haploid cells (C). There were similar distributions of DNA content histogram picks between *PM-AR*<sup>-/-</sup> (D), *G-AR*<sup>-/-</sup> (E), and *AR*<sup>+/-</sup> testis. Histology of testis by hematoxylin and eosin staining in testicular sections from 14-wk-old *AR*<sup>+/-</sup> (F), *S-AR*<sup>-/-</sup> (G), *L-AR*<sup>-/-</sup> (H), *PM-AR*<sup>-/-</sup> (I), and *G-AR*<sup>-/-</sup> (J) mice. Four to six 14-wk-old mice from individual groups were killed, and testes were excised for histology section. Compared with *AR*<sup>+/-</sup> testis (F), *S-AR*<sup>-/-</sup> testis showed that decrease of lumen formation in seminiferous tubules as well as germ cell development stopped at diplotene primary spermatocyte (G); *L-AR*<sup>-/-</sup> testis showed that decrease of lumen formation in seminiferous tubules as well as germ cell development stopped at round spermatid and no further differentiated elongated spermatid or released spermatozoa can be found (H); *PM-AR*<sup>-/-</sup> (I) and *G-AR*<sup>-/-</sup> (J) testis showed relatively comparable seminiferous tubule diameters and full range of germ cell development. [Sections of figures were reproduced from Refs. 55 and 56. Copyright 2006, Proceedings of the National Academy of Sciences.] All of these results originated from our previous publications.

TABLE 2. Fertility test (pup number per litter; mean  $\pm$  SEM) and epididymal sperm content analysis in AR<sup>+/y</sup>, G-AR<sup>-/y</sup>, PM-AR<sup>-/y</sup>, L-AR<sup>-/y</sup>, S-AR<sup>-/y</sup>, and T-AR<sup>-/y</sup> mice

Genotype	Mate no.			Vaginal plug	Sperm count/epididymis (n = 5)	Motility
	1	2	3			
AR <sup>+/y</sup>	8.5 $\pm$ 0.5	7.5 $\pm$ 0.5	7.2 $\pm$ 1.0	+	26 $\times$ 10 <sup>6</sup> /ml	Normal
G-AR <sup>-/y</sup>	7.5 $\pm$ 0.5	7.8 $\pm$ 0.5	8.2 $\pm$ 0.5	+	25 $\times$ 10 <sup>6</sup> /ml	Normal
PM-AR <sup>-/y</sup>	9.0 $\pm$ 2.6	8.6 $\pm$ 1.5	8.7 $\pm$ 2.1	+	12 $\times$ 10 <sup>6</sup> /ml <sup>a</sup>	Normal
L-AR <sup>-/y</sup>	0	0	0	+	0	0
S-AR <sup>-/y</sup>	0	0	0	+	0	0
T-AR <sup>-/y</sup>	0	0	0	+	No epididymis	

All of these results originated from our previous publications.

<sup>a</sup> Significant difference ( $P < 0.05$ ;  $t$  test) as compared to AR<sup>+/y</sup>.

type (AR<sup>+/+</sup>) C57BL/6 female mice (Table 2) (50–52). Using flow cytometric scanning of propidium iodide-labeled cells, S-AR<sup>-/y</sup> testes showed a 3-fold increase in diploid cells, 2-fold increase in tetraploid cells, and 11-fold reduction in haploid cells compared with AR<sup>+/y</sup> testes (Fig. 3B *vs.* Fig. 3A) (50). Moreover, recent studies suggest that androgen, acting through Sertoli cells AR, regulates the microenvironment of seminiferous epithelium by influencing a broad spectrum of gene changes in Sertoli cells (53, 129, and 130). The results showed that loss of AR specifically in Sertoli cells could affect the following: 1) structural support elements of Sertoli cells leading to impaired normal supportive function for movement of developing germ cells; 2) junction complex formation and basement membrane development of Sertoli cells leading to impaired functional integrity of the blood-testis barrier; and 3) Sertoli cell-specific proteases, transport proteins, and paracrine factor production and/or secretion, leading to impaired Sertoli cell nursery functions for developing germ cells (53, 129). The results of S-AR<sup>-/y</sup> mice studies (50–53, 120, 129, 130) clearly demonstrated that AR function in Sertoli cells is essential for the maintenance of fully competent Sertoli cell functions as well as the appropriate hormone levels that support the completion of meiosis I during spermatogenesis.

### C. L-AR<sup>-/y</sup> mice

Testicular testosterone is produced by the Leydig cells and is essential for qualitatively and quantitatively complete spermatogenesis and development of the male phenotype. Earlier studies have demonstrated that long-term intratesticular testosterone withdrawal causes the failure of progression of round spermatid to elongated spermatids, particularly during stages VII and VIII of the spermatogenic cycle (131, 132). The histology of L-AR<sup>-/y</sup> testis revealed a decrease of lumen formation as well as a decrease in diameter of seminiferous tubules, and germ cell development stopped at the round spermatid stage as compared with AR<sup>+/y</sup> littermates. There are no further differentiated mature elongated spermatids or spermatozoa throughout the L-AR<sup>-/y</sup> testis (Fig. 3H). Consequently, the results from epididymal sperm count indicated that no sperm exists in the epididymis of L-AR<sup>-/y</sup> mice, and the mating tests showed that L-AR<sup>-/y</sup> mice failed to impregnate the AR<sup>+/+</sup> C57BL/6 female mice (Table 2). The flow cytometric scanning showed that L-AR<sup>-/y</sup> testes have 4-fold increased tetraploid cells and

2.8-fold reduced haploid cells compared with AR<sup>+/y</sup> testes (Fig. 3C *vs.* Fig. 3A) (54). The results from L-AR<sup>-/y</sup> mice studies illustrated that AR function in Leydig cells is essential for the maintenance of appropriate Leydig cell steroidogenic function and subsequent affect on Sertoli cell functions to support the final differentiation of round spermatids to mature elongated spermatids, the so-called process of spermiogenesis, in testis (54). The results from L-AR<sup>-/y</sup> mice also confirmed previous *in vitro* findings (133) demonstrating that androgen/AR signaling in Leydig cells displays in an autocrine regulation manner.

### D. PM-AR<sup>-/y</sup> mice

The testicular histology of 14-wk-old PM-AR<sup>-/y</sup> mice showed relatively comparable seminiferous tubule diameters and the full range of germ cell development (Fig. 3I), but the epididymal sperm count in PM-AR<sup>-/y</sup> mice revealed 57% decreases compared with AR<sup>+/y</sup> mice (Table 2). Despite decreased epididymal sperm count in PM-AR<sup>-/y</sup> mice, the mating test showed that there were no significant differences in the litter number and the pup number per litter compared with AR<sup>+/y</sup> male mice (Table 2). The flow cytometric scanning showed that there were similar distributions of DNA content histogram peaks between PM-AR<sup>-/y</sup> and AR<sup>+/y</sup> controls (Fig. 3D *vs.* Fig. 3A) (55). In addition, the result of selective gene expression changes of PM-AR<sup>-/y</sup> testis, such as the genes related to smooth muscle contractions (endothelin, endothelin receptor A and B, adrenomedullin, adrenomedullin receptor, and vasopressin receptor 1a) or genes related to Sertoli cells' functional genes and cell junction genes (transferrin, epidermal fatty acid binding protein, tissue-type plasminogen activator, urokinase-type plasminogen activator, androgen-binding protein, occludin, testin, nectin, zyxin, vinculin, laminin  $\gamma$ 3, gelsolin, connexin43, and N-cadherin), indicated that there might be impaired smooth muscle contractility, Sertoli cell nourishing function, and the integrity of Sertoli cell junctions. All of these consequently affect testis sperm production and slow down the germ cell movement (55). The results in PM-AR<sup>-/y</sup> mice clearly showed that lack of functional AR in PM cells might directly cause functional impairment of PM cell contractility and indirectly cause some functional defects of Sertoli cells to support germ cell development and spermatozoa output, which subsequently lead to decreased epididymal sperm count.

### E. *G-AR*<sup>-/-y</sup> mice

In comparison with AR<sup>+/-y</sup> mice, testes from G-AR<sup>-/-y</sup> mice have normal spermatogenesis at every spermatogenic stage (Fig. 3J). There are no significant differences in the litter number and the pup number per litter between G-AR<sup>-/-y</sup> and AR<sup>+/-y</sup> controls (Table 2). The flow cytometric scanning also showed that there were similar distributions of DNA content histogram peaks between G-AR<sup>-/-y</sup> and AR<sup>+/-y</sup> controls (Fig. 3E *vs.* Fig. 3A) (56). Although it was suggested that the presence of functional AR in germ cells was not essential for development into sperm in a chimera study (134) and in a spermatogonial stem cell transplantation study (135), studies from G-AR<sup>-/-y</sup> mice clearly demonstrated that the quality and the quantity of spermatogenesis, as well as fertility, were almost normal if AR gene is deficient in nearly all of germ cells beyond the pachytene stage (56). The major concern about the chimera study (134) is that many germ cells from the pachytene stage to spermatids (the obvious androgen-dependent period during spermatogenesis) still contain normal AR that might support spermatogenesis through autocrine or paracrine effects. Another potential argument is that the floxed AR allele might not be deleted in spermatogonia and early spermatocytes in *Sycp1-Cre* (+) G-AR<sup>-/-y</sup> mice. However, little AR expression was detected in spermatogonia and spermatocytes in any of the previous studies (34). The testicular morphology from T-AR<sup>-/-y</sup> and *Tfm* mice also suggests that AR has little effect on this stage of spermatogenesis because their testis contained spermatogonia and the spermatocytes that reached the pachytene stage (47, 91). However, the animal model might not be exactly the same as the human condition. The testicular morphology of humans with CAIS showed only Sertoli cells and very scant spermatogonia in seminiferous tubules (136, 137). In the mouse counterpart, the testicular histology of T-AR<sup>-/-y</sup> revealed maturation arrest, with the development of germ cells stopping at the pachytene stage of spermatocytes (47). Thus, the role of AR in the human spermatogenesis might be more critical than in mice. Earlier studies using immunofluorescence labeling and confocal microscopy localized AR in the midpiece and mitochondria of human sperm (38) and demonstrated that AR in human sperm have the ability to modulate the phosphatidylinositol-3-kinase/AKT pathway (40). Therefore, we cannot completely rule out the possibility of the presence of the AR nongenomic signaling within the germ cells. Manipulating the AR in the laboratory in developed human embryonic gonadal cells (138) or cultured human spermatogonia might help us to further clarify the AR role in human germ cells during spermatogenesis.

### VIII. Concluding Remarks and Future Directions

The powerful new techniques of genetic manipulations have provided us new animal models with precisely defined gene ablations in specific cell types in the body. These animal models along with emerging phenotypes have led to the accumulation of a great deal of useful information on less well-known mechanisms involved in pathological effects of androgen/AR function in male reproductive health. This review clearly indicates that AR has differential roles in the

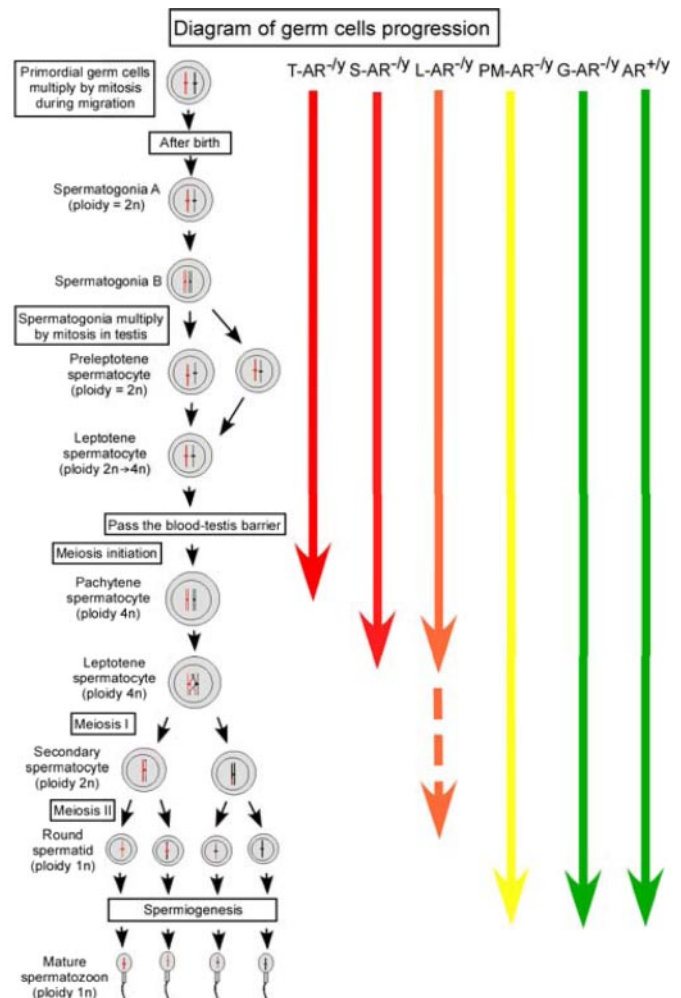


FIG. 4. Diagram of germ cell progression in T-AR<sup>-/-y</sup>, S-AR<sup>-/-y</sup>, L-AR<sup>-/-y</sup>, PM-AR<sup>-/-y</sup>, G-AR<sup>-/-y</sup>, and AR<sup>+/-y</sup> testis. AR<sup>+/-y</sup>, G-AR<sup>-/-y</sup>, and PM-AR<sup>-/-y</sup> testis can achieve full germ cell progression. However, spermatogenesis in the T-AR<sup>-/-y</sup>, S-AR<sup>-/-y</sup>, and L-AR<sup>-/-y</sup> testis ceases predominately at the pachytene, diplotene, and round spermatid stages, respectively.

different testicular cells responsible for spermatogenesis and male fertility (Fig. 4). The androgen/AR signaling in Sertoli cells plays the most important role in the process of meiosis I during spermatogenesis, and the testis from S-AR<sup>-/-y</sup> mice shows the most detrimental phenotype in that the spermatogenesis arrests predominantly at the diplotene primary spermatocyte stage before the accomplishment of first meiotic division (50, 51, 53). The lack of functional AR in Leydig cells has a major influence on Leydig cells steroidogenic functions leading to spermatogenesis arrest predominately at the round spermatid stage (54). Male mice lacking functional AR only in the smooth muscle cells and testicular PM cells (PM-AR<sup>-/-y</sup>) have similar fertility despite decreased sperm output compared with AR<sup>+/-y</sup> controls (55). The functional AR in germ cells is not essential for spermatogenesis and male fertility in mice (56). However, there are still many questions that need to be answered regarding the molecular mechanisms for androgen/AR regulation of Sertoli cell, Leydig cell, and PM cell proliferation and/or differentiation. The

exact mechanism(s) of stage-specific regulation of Sertoli cells via androgen/AR signals in testis are still largely unknown. It would also be of interest to determine the role of AR in the modulation of the complex paracrine signaling between different testicular cell-cell communications during testis development and spermatogenesis, or whether androgens/AR can regulate some aspects of spermatogonia stem cell proliferation or differentiation through the PM cell AR. The future studies towards understanding detailed molecular mechanisms in these *in vivo* animals with cell-specific AR knockout could possibly lead to useful insights for improvements in the treatment of male infertility, hypogonadism, testicular dysgenesis syndrome, and in the attempts to create a safe as well as effective male contraceptive method.

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