

Control of Spermatogenesis in Mice by the Cyclin D-Dependent Kinase Inhibitors p18^{Ink4c} and p19^{Ink4d}

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Male mice lacking both the *Ink4c* and *Ink4d* genes, which encode two inhibitors of D-type cyclin-dependent kinases (Cdks), are infertile, whereas female fecundity is unaffected. Both p18^{Ink4c} and p19^{Ink4d} are expressed in the seminiferous tubules of postnatal wild-type mice, being largely confined to postmitotic spermatocytes undergoing meiosis. Their combined loss is associated with the delayed exit of spermatogonia from the mitotic cell cycle, leading to the retarded appearance of meiotic cells that do not properly differentiate and instead undergo apoptosis at an increased frequency. As a result, mice lacking both *Ink4c* and *Ink4d* produce few mature sperm, and the residual spermatozoa have reduced motility and decreased viability. Whether or not *Ink4d* is present, animals lacking *Ink4c* develop hyperplasia of interstitial testicular Leydig cells, which produce reduced levels of testosterone. The anterior pituitary of fertile mice lacking *Ink4c* or infertile mice doubly deficient for *Ink4c* and *Ink4d* produces normal levels of luteinizing hormone (LH). Therefore, the failure of Leydig cells to produce testosterone is not secondary to defects in LH production, and reduced testosterone levels do not account for infertility in the doubly deficient strain. By contrast, *Ink4d*-null or double-null mice produce elevated levels of follicle-stimulating hormone (FSH). Because *Ink4d*-null mice are fertile, increased FSH production by the anterior pituitary is also unlikely to contribute to the sterility observed in *Ink4c/Ink4d* double-null males. Our data indicate that p18^{Ink4c} and p19^{Ink4d} are essential for male fertility. These two Cdk inhibitors collaborate in regulating spermatogenesis, helping to ensure mitotic exit and the normal meiotic maturation of spermatocytes.

Spermatogenesis in mammals is characterized by a well-defined sequence of mitotic and meiotic divisions that lead to the production of mature spermatozoa (27). In newborn mice, male germ cell precursors undergo self-renewal in the testis between days 1 and 7 postpartum (pp) (Fig. 1). From day 7 pp onward, inception of spermatogenesis begins synchronously in a cohort of precursors, starting with at least two mitotic divisions followed by one round of meiosis. The early cell divisions lead to the development of type A and type B spermatogonia, the latter of which undergo premeiotic replication and enter meiosis as primary spermatocytes. Meiosis I is characterized by a prolonged prophase that allows chromatid exchange through crossing over. Segregation of homologous chromosomes occurs at the end of meiosis I, and resulting secondary spermatocytes then proceed through a second meiotic division in which haploid germ cells are generated. These differentiate to form round spermatids and, eventually, mature elongated spermatozoa (spermogenesis). The first round of spermatogenesis is followed by additional waves, enabling continuous sperm production throughout the life of the animals.

Spermatogenesis is regulated hormonally through the pituitary-gonadal axis. The anterior lobe of the pituitary gland produces the gonadotropins follicle-stimulating hormone (FSH) and luteinizing hormone (LH). In males, FSH stimulates Sertoli cells, whose number determines the thickness of the seminiferous epithelium and, in turn, the size of the testis (36). LH induces interstitial Leydig cells to produce testosterone, a gonadal steroid necessary for spermatogenesis (19).

Cyclin-dependent kinases (Cdks) likely govern both the mitotic and meiotic divisions that characterize spermatogenesis, but it remains unclear which classes of enzymes are required for the various processes. Using immunohistochemical methods, cyclins D2 and D3 and their catalytic partner Cdk4 were seen to be expressed at the periphery of the seminiferous tubules between days 1 and 13 pp in spermatogonia undergoing mitosis (7, 22, 28, 33, 34, 46). By contrast, little cyclin D2 and Cdk4 expression was observed later in differentiated spermatocytes and spermatids (7, 28), although cyclin D3 expression was maintained (33, 46). *cyclin D2*-null male mice are fertile but have reduced testicular size and low sperm counts (39), whereas *Cdk4*-null mice are for the most part sterile at birth or become so at an early age (32, 42). Atrophic seminiferous tubules observed within the testes of *Cdk4*-null mice showed reduced numbers of spermatogonia and spermatocytes, suggesting that their proliferative capacity was reduced. Cdk6, the other known catalytic partner of D-type cyclins, was

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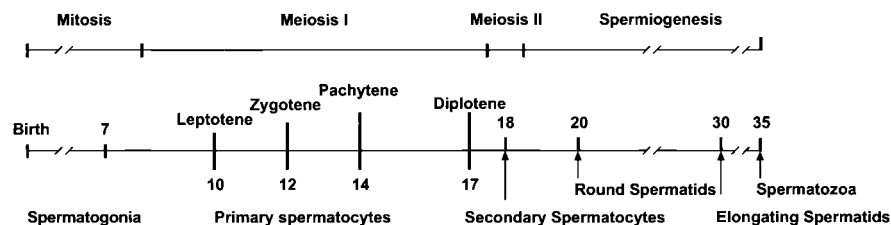


FIG. 1. Idealized timing of the first wave of spermatogenesis. The time line from birth onward indicates the temporal sequence of events in the first 35 days pp (27). Intervals in which mitotic cell division, meiosis I, meiosis II, and spermiogenesis occur are indicated above the time line, noting different stages during prophase of meiosis I. Spermatogonia populate the seminiferous tubules after birth, giving rise to spermatocytes, spermatids, and spermatozoa, as indicated below the time line.

expressed at lower levels in testes of *Cdk4*-null animals (42). These findings together suggest that cyclins D2 and D3 in combination with Cdk4, and possibly Cdk6, may regulate G_1 progression in spermatogonia in the postnatal testis and that cyclin D3 may play an additional role later in germ cell development.

The cyclin D-dependent kinases are negatively regulated by small polypeptide Cdk inhibitors encoded by four distinct *Ink4* genes (38). Two of the *Ink4* gene products, p16^{Ink4a} and p15^{Ink4b}, are not detectably expressed during mouse fetal development and are first observed in tissues of young adult animals (48). Disruption of either *Ink4a* or *Ink4b* leads to no developmental defects, and the young animals are healthy and fertile (25, 35). By contrast, the other *Ink4* family members, p18^{Ink4c} and p19^{Ink4d}, are expressed during mouse embryogenesis and into adult life, particularly in the central nervous system and testis (48–50). *Ink4c*-null male mice are fertile and manifest no apparent testicular abnormalities, but they exhibit organomegaly and eventually develop midlobe pituitary tumors and hyperplasia of Leydig cells (16, 17, 25). Mice expressing a Cdk4 knock-in allele encoding a mutant form of the kinase (R24C) that cannot be inhibited by *Ink4* proteins are fertile and also manifest Leydig cell hyperplasia (32). *Ink4d*-null mice are fertile and enjoy a normal life span, but they do not manifest Leydig cell abnormalities and instead display marked testicular atrophy associated with increased apoptosis in seminiferous tubules and reduced sperm counts (50). These findings suggest that expression of unrestrained cyclin D-dependent kinases can not only increase Leydig cell proliferation but can also have additional effects on male germ cell development, whether their expression is cell autonomous or not.

To address possible redundant roles for p18^{Ink4c} and p19^{Ink4d} in testicular development, we bred mice lacking *Ink4c* (25) with *Ink4d*-null mice (50) to derive animals lacking both genes. We show that deletion of both *Ink4c* and *Ink4d* (referred to here as *Ink4cd* double-null mice) results in complete infertility in males but has no effect on female reproductive function. Our data suggest that inappropriate regulation of cyclin D-dependent kinases in male germ cell progenitor cells inhibits them from undergoing meiosis.

MATERIALS AND METHODS

Generation of mouse strains and mouse embryo fibroblasts. Mouse strains (C57BL/6 \times 129Svj) deficient in *Ink4c* (25) or *Ink4d* (50) were intercrossed. *Ink4d*-deficient females were bred with *Ink4c*-null males to yield compound heterozygotes. Interbreeding generated wild-type and single- and double-null

animals at the expected Mendelian frequencies. Genotyping was performed by Southern blotting with mouse tail DNA from weanling pups, as previously described (25, 50). Primary mouse embryonic fibroblasts (MEFs) were isolated from 13.5- to 14.5-day-old embryos as described previously and were propagated on a 3T9 schedule (48).

Sperm counts. Two-cauda epididymides from 10- to 14-week-old male mice were harvested at the same time of day (between 12:00 and 2:00 p.m.). The sperm-containing fluid was squeezed out of the cauda, which was later cut into pieces. The sperm fluid and the pieces of cauda were suspended in 1 ml of Dulbecco's modified Eagle medium containing 25 mM HEPES buffer (pH 7.5) and 4 mg of bovine serum albumin per ml and were incubated at 37°C for 20 min. Suspensions of spermatozoa (20 μ l) were fixed in 480 μ l of 10% formalin. We used a hemacytometer to determine the number of spermatozoa.

Histologic and immunohistochemical methods. Testes were placed in Bouin's fixative (Electron Microscopy Sciences, Fort Washington, Pa.) for a period between 6 h and overnight, depending on the size of the testis. After fixation, the testes were stored in 70% ethanol at 4°C until processed. Tissues were dehydrated in increasing concentrations of ethanol, embedded in paraffin wax, and sectioned at a thickness of 5 μ m. Sections were stained with hematoxylin and eosin or were processed for immunohistochemical analysis with antibodies to either germ cell nuclear antigen 1 (GCNA1; a kind gift of George Enders, University of Kansas Medical Center) or the steroidogenic enzyme P450 side chain cleavage (P450scc; Chemicon International Inc., Temecula, Calif.). The anti-P450scc antibody was used at a dilution of 1:100. Detection of the proliferating-cell nuclear antigen (PCNA) was performed with 12- μ m frozen sections of testis fixed in 4% paraformaldehyde in 0.1 M sodium phosphate buffer, pH 7.6. PCNA was detected as described previously (21).

In situ hybridization and apoptosis assay. Male mice were anesthetized with intraperitoneal injections of ketamine and xylazine (ratio of 0.4 to 0.6 [vol/vol]) and perfused intracardially with 4% paraformaldehyde in 0.1 M sodium phosphate buffer (pH 7.6) for 20 min. Isolated testes were postfixed for 4 h at 4°C, transferred to 25% sucrose solution in 0.1 M sodium phosphate buffer (pH 7.6), and incubated at 4°C for an additional 24 h. A microtome cryostat (Microm, Walldorf, Germany) was used to section tissue into 12- μ m cryosections, which were mounted on Fischerbrand Superfrost Plus slides (Fischer Scientific, Pittsburg, Pa.) and stored at -20°C. In situ hybridization was performed as previously described (49). Briefly, sections were hybridized with radiolabeled antisense and sense riboprobes generated from mouse *Ink4c* and *Ink4d* cDNAs (49), from a *PstI* fragment containing the 3' end of the *Cdk4* gene (nucleotides 588 to 867), and from a *KpnI-SacI* fragment containing the 3' end of the *Cdk6* gene (nucleotides 549 to 1000). Apoptotic cells were visualized by the terminal deoxynucleotidyltransferase-mediated dUTP-biotin nick 3'-end labeling (TUNEL) assay (TdT FragEL DNA fragmentation detection kit; Calbiochem, La Jolla, Calif.).

Protein analysis. Sequential immunoprecipitation and immunoblotting were performed as described previously (49). Analyses of mouse p18^{Ink4c} and p19^{Ink4d} were performed with commercially available polyclonal antisera: M-167 for immunoprecipitation and immunoblotting of p19^{Ink4d}, M-20 for immunoprecipitation of p18^{Ink4c}, and M-168 for immunoblotting of p18^{Ink4c} (Santa Cruz Biotechnologies, Inc., Santa Cruz, Calif.). Mouse Cdk4 was precipitated from cell lysates with antiserum (Rz) to the C-terminal peptide of mouse Cdk4 (26) and was immunoblotted with antibody C-22 (Santa Cruz Biotechnologies, Inc.). Mouse Cdk6 was detected with antiserum to a C-terminal peptide (49).

Hormone measurements. Blood samples were collected from 10- to 14-week-old male mice (between 12:00 and 2:00 p.m.) and were kept on ice. Serum was separated and stored at -80°C. The amounts of serum testosterone were measured by a standard coated-tube radioimmunoassay kit (Diagnostic Systems

TABLE 1. Phenotypes of *Ink4c*-null, *Ink4d*-null, and *Ink4cd* double-null male mice

Strain genotype	Fertility	Testis wt (mg) ^a	Sperm count ^b	Seminiferous tubule histochemistry	Testosterone (ng/ml) ^b	Leydig cells	FSH (ng/ml) ^b
Wild type	Fertile	124 ± 10.3	3.0 × 10 ⁷ ± 0.8 × 10 ⁷	Normal	13.5 ± 2.8	Normal	18.4 ± 3.6
<i>Ink4c</i> ^{-/-}	Fertile	149 ± 12.7	4.5 × 10 ⁷ ± 1.4 × 10 ⁷	Normal	3.52 ± 1.3	Hyperplastic ^c	19.1 ± 5.1
<i>Ink4d</i> ^{-/-}	Fertile	101 ± 18.4	2.0 × 10 ⁷ ± 0.7 × 10 ⁷	Meiotic delay, increased apoptosis	9.89 ± 2.8	Normal	35.8 ± 7.5
<i>Ink4cd</i> ^{-/-}	Infertile	84.2 ± 9.9	0.44 × 10 ⁷ ± 0.3 × 10 ⁷	Pronounced meiotic delay, greatly increased apoptosis	2.84 ± 0.56	Hyperplastic ^c	45.0 ± 11

^a Testes from 6-month-old mice (wild type, *n* = 10; *Ink4c* null, *n* = 18; *Ink4d* null, *n* = 14; *Ink4cd* double null, *n* = 15). Data are averages ± standard deviations.

^b Total number recovered per mouse using 10- to 14-week-old males. Numbers represent averages ± standard deviations from five or more animals of each genotype.

^c Hyperplastic Leydig cells did not appear to differentiate based on the relatively low levels of histochemically detectable P450scc, the rate-limiting enzyme for steroid biogenesis.

Laboratories, Webster, Tex.). Interassay and intra-assay controls were included. Serum FSH measurements were carried out by radioimmunoassays (8) performed with antibodies and standards generously provided by the National Hormone and Pituitary Program of the National Institute of Diabetes and Digestive and Kidney Diseases (Washington, D.C.).

RESULTS

Phenotypic features of mice lacking *Ink4c* and *Ink4d*. Mice containing disrupted *Ink4c* alleles were bred with *Ink4d*-null animals, and intercrosses between heterozygotes yielded mice lacking both genes at the expected Mendelian frequency. *Ink4c*-deficient mice exhibit organomegaly (16, 25), whereas *Ink4d*-null animals do not differ from their wild-type littermates in size (50). Female mice doubly deficient for both *Ink4c* and *Ink4d* exhibited a mean increase in body weight relative to wild-type and *Ink4d*-null female littermates but still remained 8% smaller than *Ink4c*-null animals (data not shown). In contrast, *Ink4cd* double-null males were similar in size to *Ink4c*-deficient males. Despite expression of high levels of *Ink4c* and *Ink4d* in the developing mouse brain (49), the doubly deficient mice displayed no overt neurological defects. However, we recently recognized that *Ink4d*-null mice are deaf (B. Chen, F. Zindy, M. F. Roussel, and N. Segil, unpublished observations). Other Cdk inhibitors may well compensate for the lack of these Ink4 proteins in the brain, as animals lacking p19^{Ink4d} and p27^{Kip1}, but neither alone, exhibit profound neurologic defects leading to the death of the animals in the first few weeks of life (47).

Ink4c-null animals develop progressively expanding adenomas of the intermediate lobe of the pituitary gland with nearly complete penetrance by 8 to 10 months of age, and by 14 months at least half of the animals die as a consequence (16, 25). *Ink4c*-null mice also develop pheochromocytomas at a low but significant frequency (ca. 10%) (17, 25). At least one independently derived strain exhibits hyperplasia of pancreatic β-islet cells (25), mirroring observations with mice expressing an Ink4-resistant mutant Cdk4 allele in place of the wild-type gene (32). *Ink4cd* double-null mice exhibited each of these features of the *Ink4c*-null strain with similar types, frequencies, and rates of tumor development. The fact that *Ink4d* loss did not synergize with *Ink4c* deficiency in this respect is consistent

with the previous finding that p19^{Ink4d} is not a tumor suppressor in mice (50).

To determine whether the combined loss of *Ink4c* and *Ink4d* affects cell proliferation, we explanted MEFs into culture from 13.5-day-old single- and double-null embryos that normally express both proteins (48). Primary MEFs from the various knockout strains could not be distinguished from those derived from wild-type littermates with respect to their proliferative rates, their ability to arrest in medium containing limiting concentrations of serum, or their rate of entry into S phase following serum withdrawal and restimulation. The cells underwent replicative senescence after 20 to 30 population doublings, associated with the induction during progressive cell culture of various stress response proteins, including p16^{Ink4a}, p15^{Ink4b}, p21^{Cip1}, and p19^{Arf} (reviewed in reference 37). Like nonimmortalized MEF strains, *Ink4cd* double-null cells were resistant to transformation by oncogenic Ras. In short, the combined loss of *Ink4c* and *Ink4d* had no overt effects on the fibroblast cell cycle or replicative senescence.

Sterility in *Ink4cd* double-null males. *Ink4cd* double-null females exhibited no obvious fertility defects, but all doubly deficient males were sterile, and test matings with wild-type females failed to induce pregnancy. Atrophy of the testes in 6-month-old *Ink4cd* double-null males was more pronounced than that in males lacking *Ink4d* alone (*P* < 0.005) (Table 1). In contrast, the testes of organomegalic *Ink4c*-null mice were larger than those of their wild-type littermates (*P* < 0.0001). The number of spermatozoa in fluid extruded from the epididymides of *Ink4d*-null males was significantly lower than that of both wild-type (*P* = 0.03) and *Ink4c*-null (*P* < 0.001) males (Table 1). More strikingly, the sperm counts of double-null males were 85% lower than those of wild-type littermates (*P* < 0.001) (Table 1). In addition, the morphology of residual epididymal spermatozoa in the double-null mice was abnormal and more than 60% of spermatozoa were nonviable (data not shown).

Histologic comparison of transverse and sagittal sections of testes from 3-month-old and 6-month-old *Ink4cd* double-null and wild-type mice revealed that steroidogenic Leydig cells were abnormal in doubly deficient males. Hyperplasia of the Leydig cells was already apparent at 3 months of age (Fig. 2B

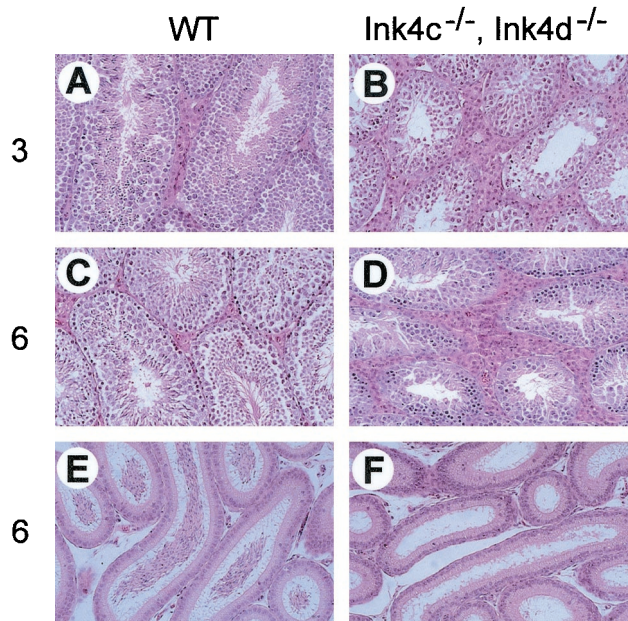


FIG. 2. Leydig cell hyperplasia and reduced numbers of sperm in *Ink4cd* doubly deficient mice. Hematoxylin and eosin staining of sections of testes from 3-month-old (A) and 6-month-old (C) wild-type (WT) mice show normal seminiferous tubule architecture and interstitial Leydig cells. In the testes of 3-month-old (B) and 6-month-old (D) *Ink4cd* double-null males, Leydig cell hyperplasia is apparent. Although the 6-month-old wild-type epididymis (E) was replete with mature spermatozoa, the epididymis of the 6-month-old *Ink4cd* double-null mouse (F) had few mature spermatozoa. Magnification, $\times 200$.

versus A) and was even more obvious at 6 months of age (Fig. 2D versus C). The excessive proliferation of Leydig cells in *Ink4cd* double-null males seemed somewhat more severe than that seen in *Ink4c*-null males (data not shown). Consistent with the sterility of *Ink4cd* double-null males, the number of mature sperm observed in the epididymal lumina of 6-month-old an-

imals was much less than that seen in sections of wild-type testes (Fig. 2F versus E).

Apoptosis normally plays a role in regulating production of mature functional sperm. However, the decreased number of sperm in *Ink4d*-null mice is associated with an increased number of apoptotic germ cells; this is reflected by increases in the number of multinucleated giant cells, of cells with condensed nuclei, and of DNA free ends detected by TUNEL (50). In testes from wild-type mice, as the first cohort of germ cells reaches the pachytene stage at day 14 or 15 pp (Fig. 1), apoptosis of tubular cells increases (Fig. 3A) but then declines by day 19 pp as some cells complete meiosis I (Fig. 3E). Apoptosis was also observed at day 15 pp in testes from *Ink4c*-null mice (Fig. 3B), but unlike that of testes from wild-type mice, was even more evident at day 19 pp (Fig. 3F). The tubules in testes of *Ink4d*-null (Fig. 3C) and *Ink4cd* double-null mice (Fig. 3D) also revealed elevated apoptosis through day 19 pp (Fig. 3G and H) and even into adulthood (data not shown). Analysis of multiple sections from at least five animals of each genotype suggested that the apoptotic indices in testes from mice at day 19 pp lacking *Ink4d* were higher than those of animals lacking *Ink4c* (Fig. 3G and H versus F), although we did not quantify these effects by grain counts. Therefore, despite Leydig cell hyperplasia, the *Ink4cd* doubly deficient males had small testes with tubular atrophy, germ cell apoptosis, reduced sperm production, and loss of viability of the remaining spermatozoa, leading to male infertility.

Altered hormone levels in males lacking *Ink4c*, *Ink4d*, or both. A complex network of hormonal signals orchestrates spermatogenesis. Gonadotrophs in the anterior lobe of the pituitary gland secrete LH and FSH, which stimulate Leydig cells to produce testosterone. To investigate the possible contribution of the pituitary-gonadal axis to male sterility, we measured the levels of testosterone, FSH, and LH in 10- to 14-week-old males of all genotypes (Table 1).

The onset of mature Leydig cell function at puberty is characterized by their ability to terminally differentiate and pro-

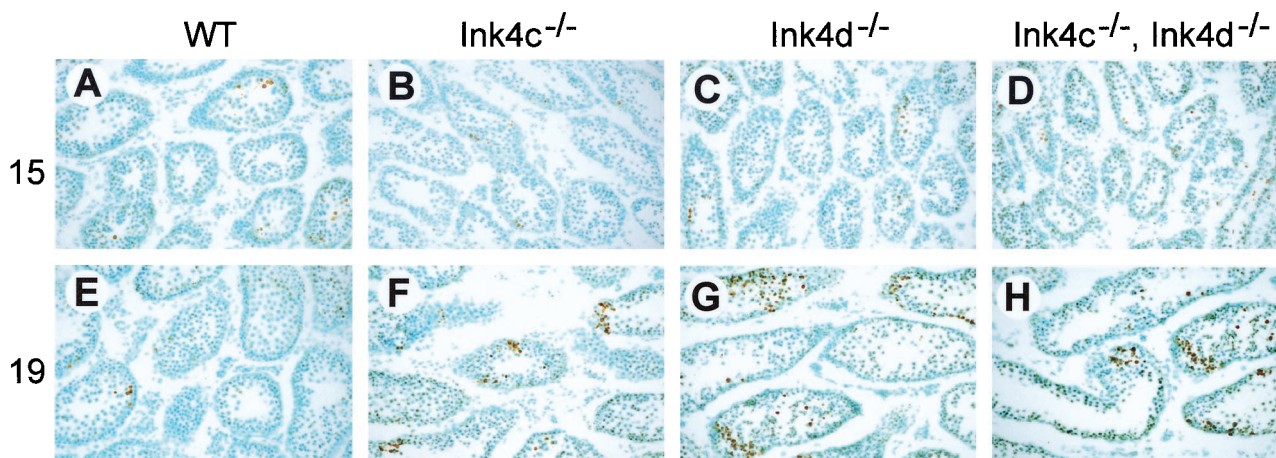


FIG. 3. Increased apoptotic index in germ cells from *Ink4cd* doubly deficient mice. Sections of testes of 15-day-old (A through D) and 19-day-old (E through H) mice of indicated genotypes were subjected to TUNEL assay to score apoptotic cells. As visualized by the brown precipitate, testes from *Ink4d*-null (G) and *Ink4cd* double-null (H) mice at day 19 pp showed increased numbers of apoptotic cells within the seminiferous tubules. WT, wild type.

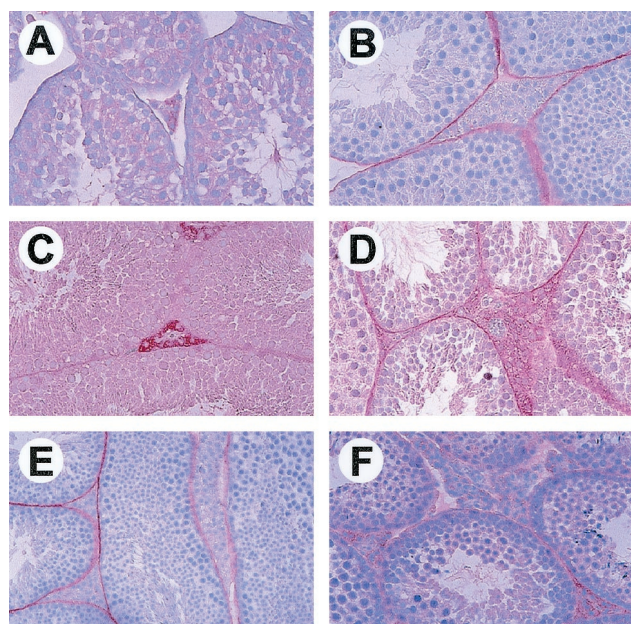


FIG. 4. Impaired differentiation of Leydig cells from *Ink4c*-null and *Ink4cd* double-null mice. P450scc staining of Leydig cells is indicated by red precipitate. (A) Immature Leydig cells from 1-month-old wild-type mice lack P450scc staining. (B) Absence of staining of 3-month-old wild-type testis with preimmune serum used as negative control. P450scc staining in Leydig cells from 3-month-old wild-type (C) and *Ink4d*-null (D) mice. Testes from 3-month-old *Ink4c*-null (E) and *Ink4cd* doubly deficient (F) mice failed to reveal significant P450scc staining.

duce testosterone and other androgens at levels seen in adult males. As noted above, however, spermatogenesis begins earlier in life, arguing that adequate levels of testosterone to support this process are available soon after birth. One marker of Leydig cell differentiation is P450scc, the steroidogenic enzyme responsible for conversion of cholesterol to pregnenolone, the first and rate-limiting step in the steroidogenic pathway (29). As expected, in testis sections from 1-month-old wild-type males, Leydig cells failed to show P450scc immunoreactivity (Fig. 4A) above background levels (Fig. 4B). By 3 months of age, however, Leydig cells of wild-type males showed strong P450scc immunoreactivity, as demonstrated by the red precipitate within the interstitial space of the testis (Fig. 4C). Similarly, Leydig cells of testes from adult *Ink4d*-null males also showed up-regulated levels of P450scc (Fig. 4D) correlating with nearly normal levels of testosterone in their serum (Table 1). However, the levels of P450scc immunoreactivity was severely reduced in Leydig cells of *Ink4c*-null (Fig. 4E) and *Ink4cd* double-null testes (Fig. 4F), suggesting a failure of Leydig cell differentiation. In turn, levels of serum testosterone were reduced by approximately 75% in *Ink4c*-null and *Ink4cd* doubly deficient males (Table 1). The levels of LH, released by the anterior lobe of the pituitary, were equivalent in males of all genotypes (data not shown), suggesting that the apparent failure of Leydig cell differentiation in mice lacking *Ink4c* was not secondary to pituitary dysfunction. We presume that the hyperplasia of Leydig cells observed somewhat later in life in the two mouse strains lacking *Ink4c* reflects a consequence of the same pathologic process; namely, their failure to

differentiate and produce testosterone. However, despite the decline in serum testosterone levels, *Ink4c*-null mice remain fertile.

Normal levels of FSH were present in the serum of wild-type and *Ink4c*-null males, whereas FSH levels were increased in both *Ink4d*-null and *Ink4cd* double-null males (Table 1). The physiologic basis or consequence, if any, of the increased FSH levels remains unclear but occurs independently of production of the other gonadotroph product, LH. Normal levels of other pituitary hormones secreted by the anterior lobe, including corticotropin and thyroid-stimulating hormone, were produced in males of all genotypes (data not shown).

Expression of *Ink4* proteins, *Cdk4*, and *Cdk6* in testes. To understand the nature of sterility in *Ink4cd* double-null males, we first studied the expression of *Ink4* proteins and cyclin D-dependent kinases during testicular development in wild-type mice. We isolated testes from animals between 7 days and 1 month pp, the period that marks the first wave of spermatogenesis (Fig. 1), and also from mice at 5 months of age. We then used sequential immunoprecipitation and immunoblotting to score for the presence of the various proteins in testicular tissue and to look for complexes between p18^{Ink4c} or p19^{Ink4d} with *Cdk4* and *Cdk6* (Fig. 5). As controls for antibody specificity, we used testes from the *Ink4cd* doubly deficient animals (Fig. 5, lanes C). Testes from at least five mice were pooled to make lysates, and analysis of the same lysates was repeated twice with similar results.

As previously described (34, 46), *Cdk4* protein was most abundant in the immature testis (days 7 to 11 pp), after which its levels declined (Fig. 5A). This may reflect the fact that *Cdk4* expression in Sertoli cells and spermatogonia is diluted by an increasing proportion of postmeiotic germ cells as the organ develops postnatally. *Cdk6* expression was also maximal in the immature testis but, in contrast to that of *Cdk4*, was undetectable after day 17 pp (Fig. 5A). Since our antibodies detect recombinant *Cdk4* and *Cdk6* with nearly equal efficiency, we can conclude that the levels of *Cdk4* eclipse those of *Cdk6* at all times during maturation of the organ. The pattern of expression of p18^{Ink4c} was biphasic (Fig. 5A): the first burst of expression occurred between days 7 and 11 pp, which marks the mitotic phase of spermatogenesis, and the second phase occurred between days 15 and 25 pp, peaking at about day 17 pp, which roughly corresponds to the end of meiosis I (Fig. 1). p18^{Ink4c} was detected in complexes with both *Cdk4* and *Cdk6* at days 7 to 11 pp but was more difficult to visualize in complexes with *Cdk4* at later times (Fig. 5B). In contrast, the level of p19^{Ink4d} progressively increased from day 7 to day 17 pp and then slowly declined (Fig. 5A). p19^{Ink4d} failed to bind to *Cdk6* but formed complexes with *Cdk4* from day 7 pp through adulthood (Fig. 5B).

Because our antibodies proved inadequate for examining protein expression by immunohistochemistry in fixed tissue sections, we performed *in situ* hybridization using antisense probes (50) with sections of testes from wild-type mice in an attempt to localize cells expressing *Ink4* and *Cdk4* transcripts during the first wave of spermatogenesis (Fig. 6). At each stage, we never obtained specific hybridization with sense probes used as controls (Fig. 6, Ctl panels). At day 9 pp, *Ink4c* RNA detected with the antisense probe was expressed in a diffuse pattern throughout the testis (Fig. 6B), although for

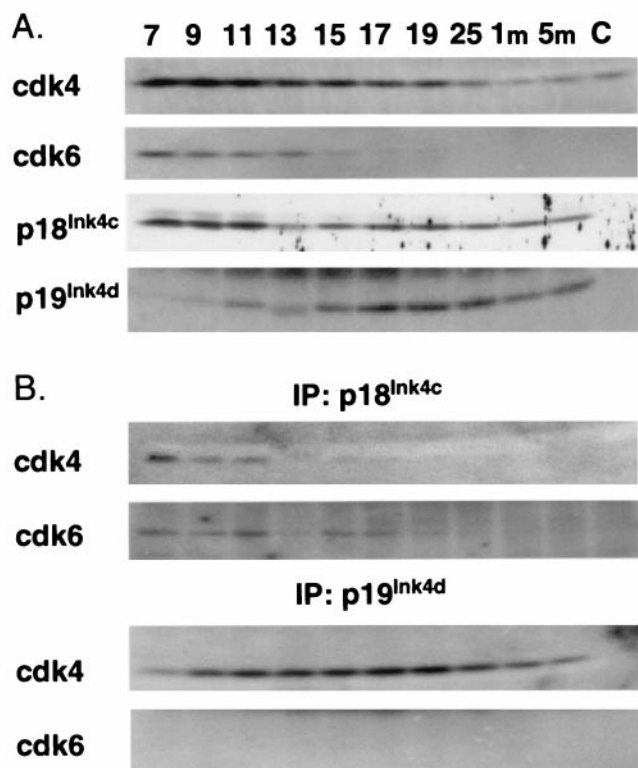


FIG. 5. Expression of Ink4 proteins and Cdk4 during testis development in wild-type mice. (A) Lysates from whole testes isolated from wild-type males at 7, 9, 11, 13, 15, 17, 19, and 25 days pp and at 1 and 5 months (m) of age were normalized for protein content and precipitated with antibodies to Ink4 proteins or Cdk4. Denatured precipitates were electrophoretically separated on denaturing gels and immunoblotted with antisera directed to the same proteins, as indicated at the left of the panels. (B) Lysates were precipitated with antibodies directed to either p18^{Ink4c} or p19^{Ink4d}, as indicated at the top of the panels. Immunoprecipitates (IP) were denatured, separated electrophoretically, and immunoblotted with antisera to the Cdk4 noted at the left of the panels. Controls performed with cells of *Ink4cd* double-null mice are in lanes C.

unknown reasons the relative intensity of RNA staining at this time was significantly weaker than what might have been predicted based on parallel protein expression studies (Fig. 5A). This underscores a potential shortcoming of this approach since disparities between mRNA expression and protein turnover could well reflect translational or posttranslational controls. Immunohistochemical staining of sections of human testes performed with a monoclonal antibody to p18^{Ink4c} detected this protein in interstitial Leydig cells and tubular Sertoli cells as well as in germ cells (4).

Only during later periods of development was *Ink4c* RNA clearly localized to the mouse seminiferous tubules, appearing most evident by day 17 pp (Fig. 6B). Higher power resolution of grains scored at day 17 pp (Fig. 6A and B, bright field, bottom) demonstrated selective staining of intratubular cells corresponding to spermatocytes while sparing the more immature cells at the periphery. *Ink4d* RNA expression increased with the age of the testis (Fig. 6D), consistent with the graded expression of the protein observed by immunoblotting (Fig. 5A). *Ink4d* transcripts were detected in tubules at day 13 pp

before the appearance of *Ink4c* RNA. By day 17 pp, intratubular spermatocytes were highly labeled (Fig. 6C and D, bright field, bottom). The temporal and physical localization of *Ink4c* and *Ink4d* RNAs within the tubules indicates that both Cdk inhibitors are primarily expressed in postmitotic spermatocytes, in agreement with immunohistochemical studies performed on human testes (4, 5). *Cdk4* staining was detected in seminiferous tubules during the first meiotic wave (Fig. 6F, bright field, bottom), but the hybridization signal decreased in intensity as spermatogenesis progressed, in agreement with results of protein immunoblotting (Fig. 5A). These data suggest that, unlike the *Ink4* proteins, *Cdk4* is primarily expressed during earlier stages of spermatogenesis at the time when spermatogonia undergoing mitosis predominate in the tubules.

Impaired meiotic progression in *Ink4cd* doubly deficient mice. Because p18^{Ink4c} and p19^{Ink4d}, when expressed at high levels, were expected to inhibit *Cdk4* activity and to block entry of cells into S phase, we anticipated that the loss of both inhibitors might increase premeiotic spermatogonial proliferation. To detect mitotically active spermatogonia, sections of testes from males of all genotypes were stained with an antibody to PCNA, a protein expressed in S phase (6). Up to day 15 pp, PCNA-positive cells were detected in testicular tubular cells of all genotypes (Fig. 7A through D). By day 19 pp, the number of PCNA-positive cells in tubules from wild-type and *Ink4c*-null mice was diminished, suggesting that many cells had exited the mitotic cycle (Fig. 7E and F versus A and B, respectively). In contrast, more PCNA-positive cells remained in some but not all tubules of *Ink4d*-deficient mice and were most evident in those of the doubly null strain (Fig. 7G and H). By adulthood, however, germ cell proliferation was restricted to a monolayer of cells along the basement membrane of the seminiferous tubules in males of all genotypes (Fig. 7I through L). The relative persistence of mitotic activity of the spermatogonial population in 19-day-old *Ink4d*-null and *Ink4cd* double-null males might therefore reflect a delay in the onset of the first wave of meiosis I.

To determine if and when meiotic progression was impaired in *Ink4d* and *Ink4cd* double-null males, we compared the progress of germ cell differentiation through meiosis I using testis sections made from male mice of various genotypes sacrificed between days 9 and 25 pp. Sections were stained with an antibody against GCNA1, a germ cell nuclear antigen that is specifically expressed in premeiotic cells up to the pachytene stage of meiosis I (43). At early stages, densely stained premeiotic cells are localized close to the basement membranes of seminiferous tubules (Fig. 8A). Passage through meiosis I is marked by a progressive loss of intensity of GCNA1-positive red precipitate during early prophase I, resulting in a weaker punctate staining pattern in pachytene cells (Fig. 8E, I, and M).

At day 9 pp, the patterns of GCNA1 localization were similar in all genotypes (Fig. 8A through D), with at least one layer of stained spermatogonia observed on the tubular basement membrane, and by day 13 pp cells showed signs of entry into meiosis I (Fig. 8E through H). Whereas pachytene cells were detected at day 17 pp in *Ink4c*-null (Fig. 8J) and *Ink4d*-null testes (Fig. 8K), cells in many tubules of *Ink4cd*-deficient mice continued to stain brightly with GCNA1 (Fig. 8L), suggesting retarded progression through meiosis concomitant with increased apoptosis (see Fig. 3H above). By day 25 pp, many

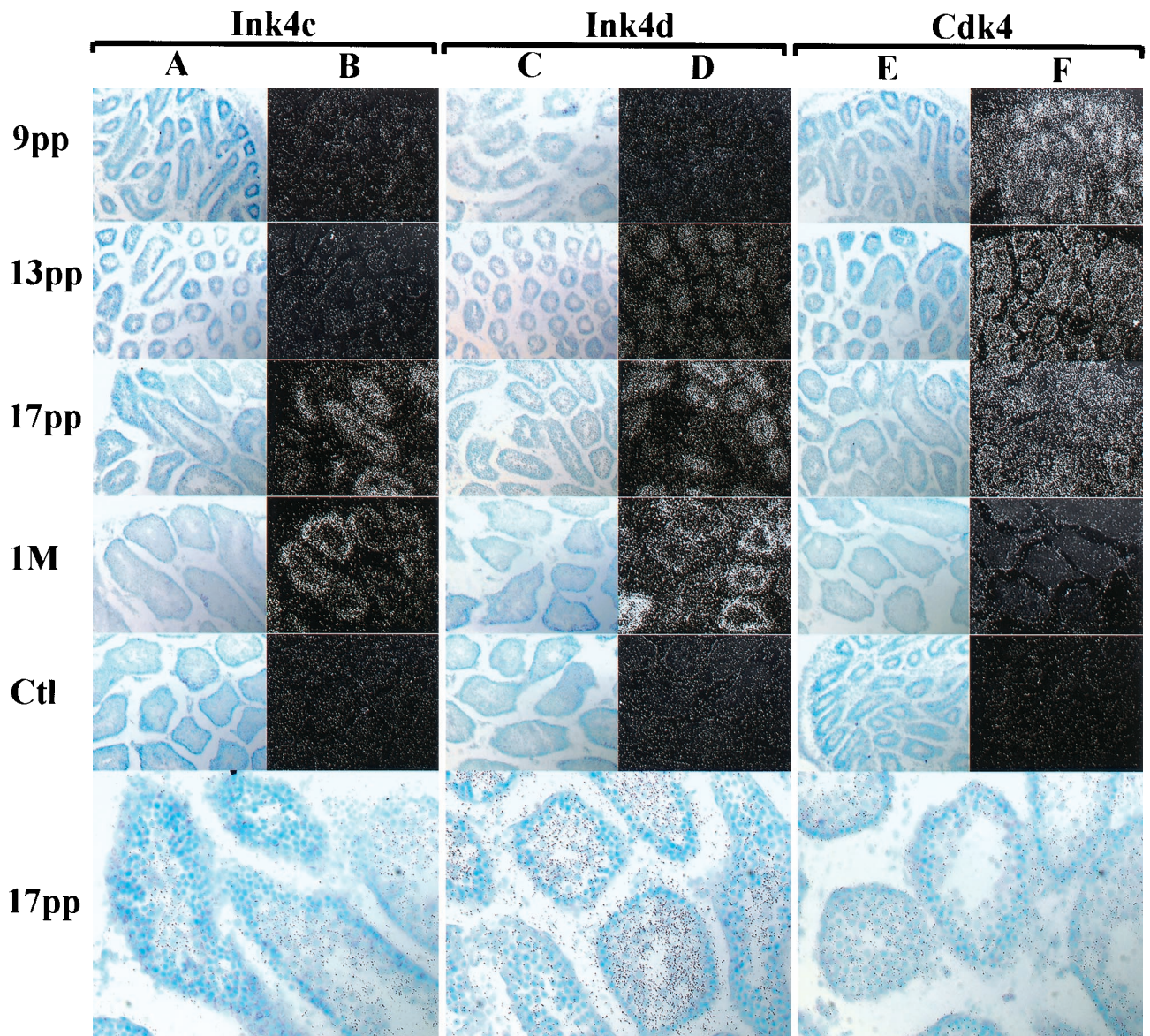


FIG. 6. *Ink4c*, *Ink4d*, and *Cdk4* mRNAs expressed during normal testis development. In situ hybridizations were performed on sections of testes taken from wild-type animals at 9, 13, and 17 days and 1 month (M) pp as indicated at the left of the panels. Sections were hybridized with the antisense probes indicated at the top. Morphology of the tubules is revealed in bright-field images (A, C, and E), and the hybridization intensity is revealed by dark-field images (B, D, and F). Control hybridizations (Ctl) performed with sense strand probes at 1 month pp (*Ink4c* and *Ink4d*) and at day 9 pp (*Cdk4*) are shown. All magnifications are $\times 200$. The three bottom bright-field panels indicate the localization of grains within tubules, as photographed at higher power (magnification, $\times 400$).

apoptotic cells were present within the tubular lumina of *Ink4cd*-deficient mice (Fig. 8P; confirmed by hematoxylin and eosin staining and TUNEL assays). In summary, these results show that $p18^{\text{Ink4c}}$ and $p19^{\text{Ink4d}}$ are required during the first wave of spermatogenesis to regulate the differentiation of mature viable sperm.

DISCUSSION

Impaired spermatogenesis in *Ink4cd* double null males. Females lacking both *Ink4c* and *Ink4d* exhibit no reproductive abnormalities, whereas males lacking both Cdk inhibitors are infertile. *Ink4cd* double-null males produce low numbers of

mature sperm, the majority of which exhibited abnormal morphology and reduced motility. Male sterility correlated with an ineffective exit of spermatogonia from the mitotic cycle and with a failure of residual postmitotic spermatocytes to correctly undergo the meiotic divisions required to generate haploid gametes. The temporally delayed entry into meiosis of *Ink4cd*-null germ cells was also associated with increased apoptosis of spermatocytes, leading ultimately to the production of mostly nonviable sperm (genotypes are summarized in Table 1).

By day 19 pp during the first wave of spermatogenesis, mitotic cell divisions in the seminiferous tubules of wild-type mice, marked by PCNA-stained S-phase cells, become largely confined to spermatogonial cells that reside close to the base-

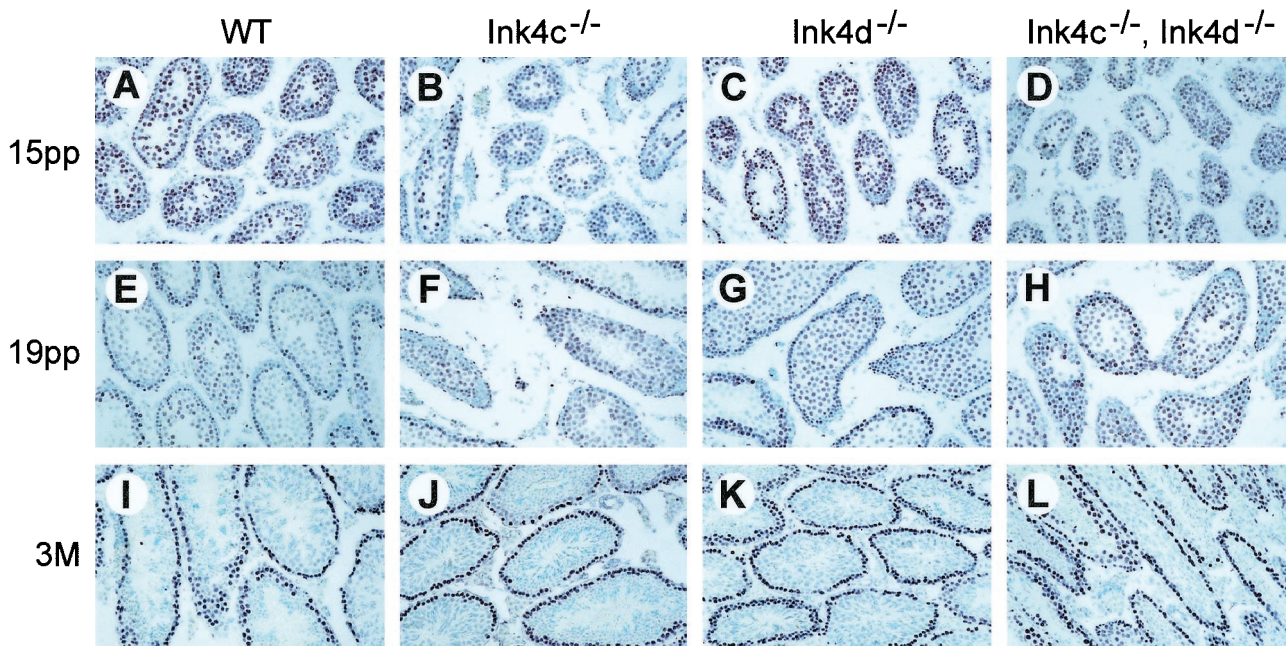


FIG. 7. PCNA-positive cells in seminiferous tubules of mice of different genotypes. Sections of testes from mice of the indicated genotypes (top) taken at day 15, day 19, and 3 months pp (indicated at the left) were immunostained with an antibody against PCNA. A deep purple precipitate indicates cellular proliferation that, in adult males, is eventually restricted to a single or double layer of premeiotic germ cells on the periphery of the seminiferous tubules (I through L). At day 19 pp, more widespread proliferation is observed in the tubules of single *Ink4*-null (F and G) and *Ink4cd* double-null males (H) as compared to the patterns in wild-type (WT) mice (E).

ment membrane. In contrast, intraluminal spermatocytes at a further distance from the basement membrane have exited the cell cycle to undergo meiotic division and subsequent differentiation. Upon completion of meiosis, the latter cells differentiate to spermatids and mature spermatozoa that can be most easily visualized histologically in the epididymis. GCNA staining of cells in seminiferous tubules can distinguish the densely stained premeiotic cells at the tubular periphery from the more punctately stained cells in the pachytene stage of meiosis I within the lumen. These patterns of staining were distorted in the tubules of *Ink4cd* double-null mice (and, to a lesser extent, in *Ink4d*-null males), in which an expanded intraluminal population of mitotic S-phase cells was detected and fewer pachytene cells appeared, the latter only after a significant delay. Abnormalities in stages of meiotic progression were associated with an increased apoptotic index, and in turn, spermatids and spermatozoa were not readily seen in the seminiferous tubules of the *Ink4cd* doubly deficient mice.

Both p18^{Ink4c} and p19^{Ink4d} are expressed at relatively high levels in testes compared to those in other organs of neonatal mice (48). *Ink4d* RNA transcripts were first readily detected by in situ hybridization in the seminiferous tubules of wild-type mice by day 13 pp, consistent with the increase in overall testicular p19^{Ink4d} protein expression observed by immunoblotting. This corresponds to the time of entry into meiosis by the first wave of developing germ cells. At later stages, *Ink4d* RNA expression remains restricted to tubular cells, while the overall p19^{Ink4d} protein level in the testis is maintained as additional waves of spermatogenesis ensue. In agreement with these findings, immunohistochemical studies of human p19^{INK4d} in testes revealed maximal levels of protein expres-

sion in primary and secondary spermatocytes (5). This argues that *Ink4d* is normally expressed in postmitotic spermatocytes, where it presumably functions to extinguish the activity of cyclin D-dependent kinases, thereby helping to prevent further mitotic divisions.

The overall pattern of p18^{Ink4c} protein expression was more complex, with an earlier wave of protein synthesis (days 7 to 11 pp) followed by a nadir at day 13 pp and a return of expression from day 15 pp onward. By use of in situ hybridization, we were unable to clearly localize the earliest round of *Ink4c* expression to tubular germ cells and instead saw a more diffuse pattern of staining throughout the testis. However, *Ink4c* expression eventually became restricted to seminiferous tubular cells by day 17 pp onward, implying that the protein is expressed at high levels in spermatocytes. Similar results were obtained by others who applied immunohistochemical techniques to study p18^{INK4c} in human testes (4). Loss of *Ink4d* leads to modest but still clearly defined inhibition of mature sperm cell production, but so far *Ink4c* has not been implicated in this process. Our data now suggest that p18^{Ink4c} collaborates with p19^{Ink4d} in helping to ensure the transition from mitotic to meiotic cell cycles during male germ cell maturation.

Evidence is accumulating that progression of meiosis I is monitored to eliminate germ cells that have sustained unrepaired DNA damage. For example, defects in genes, such as *Atm* (2, 12, 20, 44), *ligase-4* (3, 15), and *MSH 4* and *MSH 5* (11, 23), that monitor or effect DNA repair routinely lead to male sterility. One possibility, then, is that *Ink4* genes have effector functions in cell cycle checkpoint control, arresting cells with broken chromatids until DNA damage is repaired. TUNEL assays indicated that loss of *Ink4d*, or more dramatically, loss

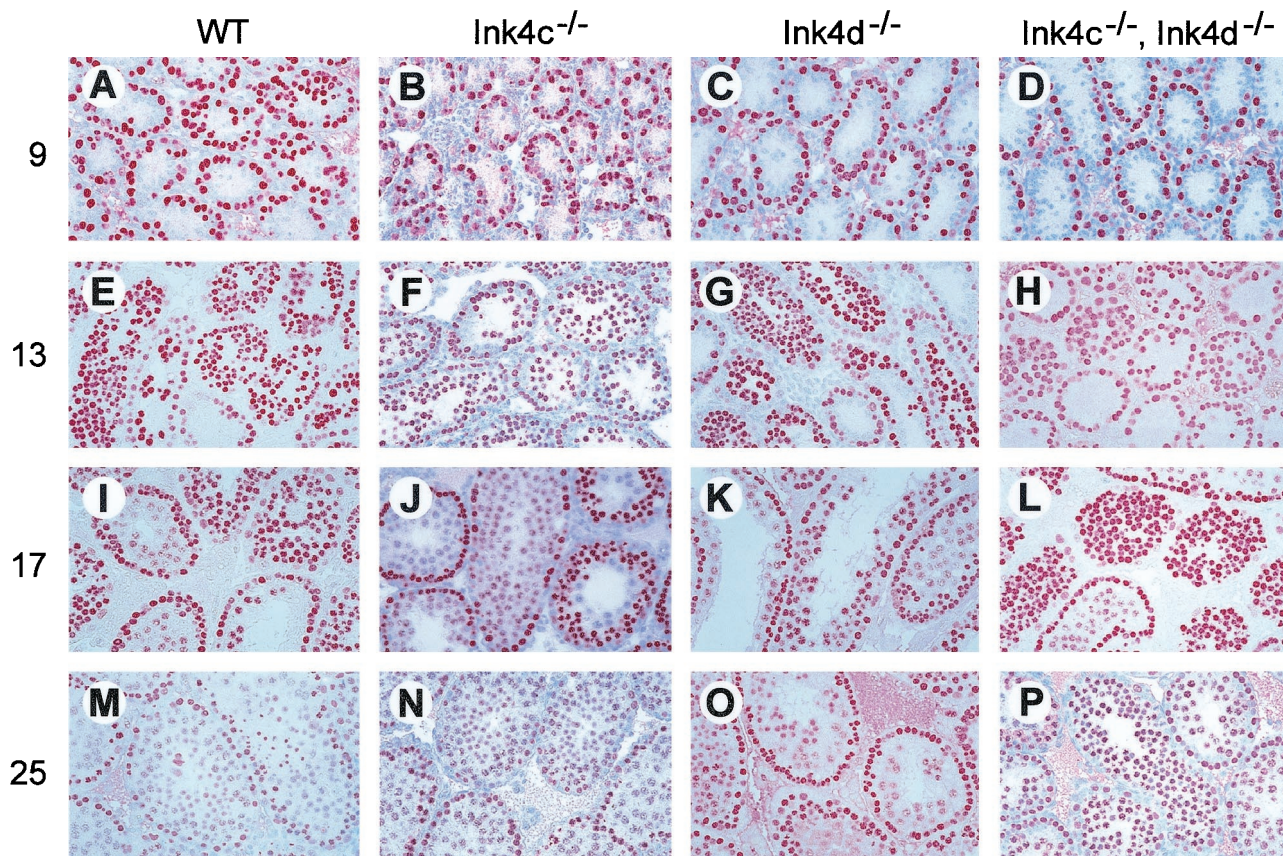


FIG. 8. Meiosis I is delayed in germ cells from *Ink4cd* doubly deficient mice. Testis sections from mice of the indicated genotypes (top) were immunostained using an antibody against GCNA1 (red precipitate). Testes were removed at the times (days pp) indicated at the left. GCNA1-positive germ cells are apparent within the testes as early as day 9 pp in all genotypes and are located at the periphery of the tubules (A through D). Their numbers increase through day 13 pp (E through H), although the intensity of staining of some cells starts to decrease. By day 17 pp, many cells within the lumina of the seminiferous tubules of wild-type (WT) (I) and *Ink4c*-null (J) males have lost their bright GCNA1 signal, indicative of progression of cells toward meiosis II. The proportion of GCNA1-positive cells continues to drop in these mice through day 25 pp (M and N). In *Ink4d*-null (K and O) and *Ink4cd* double-null (L and P) males, however, the number of GCNA1-positive cells remains higher from day 17 to day 25 pp, indicative of reduced progression through meiosis I.

of *Ink4d* in collaboration with disabled *Ink4c*, led to increased apoptosis once spermatocytes entered prophase I. The increased number of apoptotic cells coincided with the progression of spermatocytes up to and through pachynema (31). In contrast, the germ cells of *Ink4c*-null mice, like those of wild-type littermates, did not undergo significantly increased apoptosis. Hence, lack of *Ink4d* had a greater impact than loss of *Ink4c*, although both genes can contribute.

Unlike the two Ink4 proteins, Cdk4 is expressed at maximal levels at earlier stages of spermatogenesis, where spermatogonia undergoing mitotic cell cycles predominate. Although its level declined as the animals aged, Cdk4 was still readily detected at reduced levels thereafter. Lower levels of Cdk6 relative to those of Cdk4 were also observed from day 7 to day 15 pp during the first wave of spermatogenesis but fell below the limit of detectability of our assays at later times. This indicates that Cdk4 is the major cyclin D-dependent catalytic subunit in the mouse testis but does not formally preclude the participation of Cdk6 in spermatogenesis in adult mice. Our results demonstrate that the effects of codeletion of *Ink4c* and *Ink4d* on spermatogenesis are not equivalent to those observed in the

Ink4-resistant *Cdk4 R24C* knock-in strain, which is fertile (32). The mechanistic basis for these differences remains unclear. Under normal circumstances, p19^{Ink4d} interacts preferentially with Cdk4 (data shown here), whereas p18^{Ink4d} binds more selectively to Cdk6 (18). Possibly, the presence of the Cdk4 R24C mutant in place of wild-type Cdk4 might enhance the association of p19^{Ink4d} with Cdk6, leading to a compensatory decrease in overall cyclin D-dependent kinase activity. A detailed analysis of spermatogenesis in the *Cdk4 R24C* knock-in strain has not been undertaken, and it is conceivable that the mice manifest subtle defects similar to those seen in males lacking *Ink4d* alone.

Disruption of the *cyclin D2* gene, which is normally expressed in spermatogonia (7, 22, 28, 33, 34, 46), leads to testicular atrophy and reduced sperm counts (39), while loss of *Cdk4* ultimately results in decreased numbers of spermatogonia and male infertility (32, 42). Together, these results indicate that cyclin D-dependent kinases are required for progression through the early mitotic divisions that characterize postnatal male germ cell development. However, their unopposed expression at later stages due to *Ink4cd* loss also pre-

vents proper germ cell maturation, underscoring a requirement for precise temporal regulation of these enzymes.

In humans, testicular germ cell tumors are derived from primordial germ cells. Both seminomas and other germ cell tumor types appear to arise from cytogenetically identical carcinomas in situ that progress to invasive lesions (7). p19^{INK4d} is undetectable in fetal germ cells that are thought to be the precursors of testicular tumors, and as might be expected, deletion of *INK4d* has not been seen in such tumors (5). Unlike p19^{INK4d}, however, p18^{INK4c} expression is abundant in carcinomas in situ, and loss of *INK4c* is frequently associated with progression from carcinomas in situ to invasive germ cell tumors in human testes (4). These findings point to the possibility for additional roles for p18^{INK4c} in fetal germ cell development.

Effects of *Ink4c* and *Ink4d* loss on the pituitary-gonadal axis.

Although the noted defects in spermatogenesis in *Ink4cd* double-null mice are likely to be germ cell autonomous, the loss of *Ink4c* and *Ink4d* also affects the production of hormones that regulate germ cell development. The pituitary tumors observed in *Ink4c*-null or *Ink4cd* double-null mice arise gradually throughout life but with complete penetrance and primarily involve the intermediate lobe of the gland. Pituitary tumorigenesis is unlikely to be responsible for the observed early postnatal effects on germ cell production in *Ink4cd* double-null animals, because mice lacking *Ink4c* alone are fertile and produce normal numbers of sperm and their pituitary tumors arise much later in life and occur in a region of the gland that does not produce LH or FSH.

Whether *Ink4d* is present or not, *Ink4c*-null animals, like mice expressing the Ink4-resistant mutant form of Cdk4 (32), exhibit Leydig cell hyperplasia. However, these cells produced only low levels of P450scc, the rate-limiting enzyme for steroid biogenesis (29). In turn, serum testosterone levels in *Ink4c*-null and *Ink4cd* double-null mice at 10 to 14 weeks of age were equally low. Because *Ink4c*-null males had normal sperm counts and were fertile, decreased testosterone production in *Ink4cd* double-null males was not strictly associated with infertility. In fact, previous studies have demonstrated that the amount of testosterone synthesized by the rodent testis far exceeds that required for normal spermatogenesis (1, 51). Moreover, the initial failure of male gametogenesis in *Ink4cd* double-null mice was manifested before puberty during the very first wave of spermatogenesis, so the effects of *Ink4c* loss on Leydig cell function are unlikely to contribute to the reduced meiotic progression. In agreement, we were unable to rescue defective spermatogenesis in *Ink4cd* double-null males by enforced administration of testosterone (data not shown). Because *Ink4c*-null and *Ink4cd* double-null animals synthesized normal levels of LH, the failure to produce normal levels of testosterone does not reflect pituitary LH insufficiency. Instead, *Ink4c* expression appears to control the ability of pubertal Leydig cells to properly exit the cell cycle, up-regulate steroidogenic enzyme expression, and increase testosterone production. The fact that *Ink4d*-null mice produced near normal levels of testosterone connotes a distinct role for p18^{INK4c}, but not p19^{INK4d}, within Leydig cells.

Regardless of whether or not *Ink4c* was present, mice lacking *Ink4d* produced elevated levels of FSH. It has been suggested that FSH is required not for the initiation of spermatogenesis

but rather for the maintenance of sperm viability and motility (24). Male mice lacking the FSH receptor (10) or lacking the FSH- β subunit (24) and men with a mutation in the FSH receptor (40) are fertile despite decreased testicular mass and reduced sperm motility and numbers. However, FSH treatment of oligospermic men results in increased spermatogenesis rather than improved spermiogenesis, arguing that FSH might increase the entry of spermatogonia into meiosis (14). Despite high levels of FSH, however, mice lacking *Ink4d*, or both *Ink4d* and *Ink4c*, have small testes and a reduction in sperm counts and viability, suggesting that the testis is unable to up-regulate spermatogenesis in the absence of the Ink4 proteins. We saw good correlation between testis size and sperm counts, a phenomenon also found in mice lacking other genes involved in testicular development, including *cyclin D2* (39), *Cdk4* (32, 42), *E2F-1* (13, 45), and *Egr4* (41).

In males, FSH enhances functions of Sertoli cells that, in turn, provide support for spermatogonial maturation (10, 36). In females, FSH plays a similar role in sustaining granulosa cell development, and disruption of *cyclin D2*, an FSH-responsive gene in these cells, leads to faulty granulosa cell function and female infertility (39). Disruption of *Cdk4* also results in female infertility, but there is no apparent defect in granulosa cell development. Instead, female infertility appears to be due to defects in the hypothalamic-pituitary axis, where Cdk4 loss leads to a significant reduction in FSH production (32). Notably, this contrasts with the elevated FSH levels observed in both *Ink4d*-null and *Ink4cd* double-null animals. Although we have no clear explanation for the increased FSH production by the pituitary gland in these mice, this could well reflect a role of Cdk4 in regulating FSH levels in males. In this regard, *Ink4d* appears to play a more prominent role than *Ink4c*.

The rate of meiotic progression in the testis is governed by an inhibin-mediated feedback loop that is initiated in the Sertoli cells. Under normal conditions, the accumulation of pre- and perimeiotic germ cells stimulates Sertoli cells to produce inhibin, which selectively suppresses FSH production by gonadotrophs in the anterior pituitary (9, 10, 30). In light of the reduced rate of meiotic entry and progression, we might expect that inhibin levels would be reduced in *Ink4cd* double-null mice. This feedback mechanism would explain the selective increase in FSH seen in *Ink4cd*-null mice, with serum LH concentrations remaining unaffected. We attempted to analyze this possibility by staining for inhibin in the testes of these mice, but available commercial antibodies did not yield specific immunohistochemical signals in Sertoli cells. This formally leaves open the possibility that a surfeit of FSH produced by the anterior pituitary gland of *Ink4cd*-null mice acts through Sertoli cells to drive premeiotic hyperproliferation of *Ink4cd*-null spermatogonia. However, given that FSH levels in fertile *Ink4d*-null mice and infertile *Ink4cd* double-null mice are 2.0- and 2.5-fold greater, respectively, than those in wild-type animals, and that FSH is used to improve spermatogenesis in men with low sperm counts (14, 36), we believe that elevated FSH levels are unlikely to explain the abnormalities in meiotic progression observed in the *Ink4cd* double-null strain.

In summary, our results demonstrate that p18^{INK4c} and p19^{INK4d} are crucial for the progression of germ cells past the pachytene stage of spermatogenesis and are most consistent with the idea that this defect in progression is intrinsic rather

than hormonally driven. Human *INK4d* and *INK4c* may be candidate markers for some forms of male-only infertility, including those caused by autosomal recessive disorders in which early meiotic arrest is a hallmark.

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F.Z. and W.D.B. contributed equally to this work.

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