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Functional Interaction of the Retinoblastoma and Ini1/Snf5 Tumor Suppressors in Cell Growth and Pituitary Tumorigenesis

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Abstract

The Ini1 subunit of the SWI/SNF chromatin remodeling complex suppresses formation of malignant rhabdoid tumors in humans and mice. Transduction of Ini1 into Ini1-deficient tumor-derived cell lines has indicated that Ini1 arrests cell growth, controls chromosomal ploidy, and suppresses tumorigenesis by regulating components of the retinoblastoma (Rb) signaling pathway. Furthermore, conditional inactivation of Ini1 in mouse fibroblasts alters the expression of various Rb-E2F-regulated genes, indicating that endogenous Ini1 levels may control Rb signaling in cells. We have reported previously that loss of one allele of Ini1 in mice fibroblasts results only in a 15% to 20% reduction in total Ini1 mRNA levels due to transcriptional compensation by the remaining Ini1 allele. Here, we examine the effects of Ini1 haploinsufficiency on cell growth and immortalization in mouse embryonic fibroblasts. In addition, we examine pituitary tumorigenesis in Rb-Ini1 compound heterozygous mice. Our results reveal that heterozygosity for Ini1 up-regulates cell growth and immortalization and that exogenous Ini1 down-regulates the growth of primary cells in a Rb-dependent manner. Furthermore, loss of Ini1 is redundant with loss of Rb function in the formation of pituitary tumors in Rb heterozygous mice and leads to the formation of large, atypical Rb⁻/⁻ tumor cells lacking adrenocorticotropic hormone expression. These results confirm in vivo the relationship between Rb and Ini1 in tumor suppression and indicate that Ini1 plays a role in maintaining the morphologic and functional differentiation of corticotrophic cells. (Cancer Res 2006; 66(16): 8076-82)

Introduction

SWI/SNF chromatin remodeling complexes hydrolyze ATP to alter nucleosome-DNA contacts to activate or repress eukaryotic gene expression (reviewed in refs. 1, 2). The subunit composition of SWI/SNF complexes varies among cell types; however, all human SWI/SNF complexes include either the catalytic ATPase subunit BRM (Brahma) or BRG1 (Brahma-related gene 1), as well as Ini1 (3, 4). The Ini1 subunit (also known as Snf5) was originally isolated as a cellular host protein that binds specifically to HIV-1 integrase and as a human homologue to the yeast Snf5 component of yeast SWI/SNF complex (5, 6).

Chromosomal abnormalities at the Ini1 locus have linked Ini1 mutation with early childhood cancers. Known broadly as rhabdoid predisposition syndrome, these cancers include atypical teratoid and rhabdoid tumors of the central nervous system and malignant rhabdoid tumors (MRT). Germ-line mutations in Ini1 have been detected in MRT of the kidney and brain, with subsequent deletion or mutation of the wild-type (WT) Ini1 allele in these tumors (7). Biallelic Ini1 mutations have also been detected in sporadic renal rhabdoid tumors, choroid plexus carcinomas, medulloblastomas, and central primitive neuroectodermal tumors (8–10). The link between Ini1 mutation and cancer and the loss of heterozygosity (LOH) for Ini1 observed in human familial MRT strongly suggests that Ini1 functions to suppress tumorigenesis.

The tumor-suppressing properties of Ini1 were confirmed by studies of mice bearing mutations in Ini1. Embryos homozygous for disruption of Ini1 (Ini1⁻/⁻) fail to develop beyond the peri-implantation stage of gestation (11, 12). Mice haploinsufficient for Ini1 (Ini1⁻/+⁻) develop normally, but approximately a quarter of these mice will develop rhabdoid tumors and undifferentiated sarcomas primarily on the head and neck, many of which are highly aggressive and metastatic (11–13). Further analysis indicated that tumor tissues showed loss of the remaining Ini1 locus or were deficient for Ini1 protein, confirming that the tumors underwent LOH for Ini1 (11, 12). More recently, transient inactivation of Ini1 using a reversible conditional Ini1 allele in mice led to rapid bone marrow failure and death of the mice, whereas sporadic inactivation of Ini1 in hematopoietic tissues and in other organs resulted in an extremely rapid onset of CD8⁺ lymphomas and rhabdoid tumors in all mice (14).

Reintroduction of Ini1 into Ini1-deficient human tumor cell lines induces a G1 arrest in these cells, a characteristic flat cell formation, and, in some cases, cell senescence or apoptosis (15–19). Ini1-deficient tumor cell lines also exhibit chromosomal instability and polyploidy, both of which can be restored to more normal conditions by transduction of Ini1 (20). Recent work has begun to investigate the underlying mechanisms of Ini1-mediated effects on cell growth and tumor suppression. The SWI/SNF ATPase subunits BRG1 and BRM are implicated in control of cell growth and proliferation through their direct interaction with the retinoblastoma (Rb) tumor suppressor protein (21–24). Rb functions as a key regulator of the G1-S transition during cell cycling by binding and inhibiting the activity of E2F transcription factors. Phosphorylation of Rb by cyclin-dependent kinases (CDK) interrupts Rb-E2F interactions, leading to E2F-induced expression of genes involved in DNA replication and progression of the cell cycle into S phase. The link between SWI/SNF ATPases and Rb has promoted investigations into the effects of Ini1 loss on the Rb signaling pathway. Induction of Rb signaling was found to restore the G1 arrest in Ini1-deficient cells, suggesting that Ini1 acts upstream of Rb in regulating the G1 checkpoint (16, 18). In support of this hypothesis, reintroduction of Ini1 into MRT cells was reported to activate the CDK inhibitor p16INK4a (25) and to induce senescence in cultured
rhabdoid cells by binding directly to the p16INK4a and p21WAF/CIP promoters and up-regulating the expression of these CDK inhibitors (26). In addition, exogenous Ini1 induces HDAC-dependent repression of the cyclin D1 promoter (17), and RNA interference inhibition of cyclin D1 activity was sufficient to restore a G1 arrest and apoptosis in Ini1-deficient MRT cells (26). Finally, gene expression profiling indicates that Ini1 activates a mitotic checkpoint and ploidy control through regulation of E2F target genes (20).

These tumor cell–based studies suggest that Ini1 functions by altering CDK phosphorylation of the Rb tumor suppressor. Studies in Drosophila also implicate the Ini1 homologue Snr1 in controlling cell growth via the Rb-E2F pathway (27, 28). However, less evidence is available about the effects of Ini1 loss in normal mammalian cells. Very recently, Ini1 was conditionally deleted in mouse embryonic fibroblasts (MEF) by use of the Cre-lox system. In one report, loss of Ini1 led to aberrant regulation of certain E2F target genes whose expression levels are also altered in MRT cells. In addition, inactivation of Ini1 induced a G2 cell cycle arrest, polyploidy, and apoptosis, with concomitant increases in levels of p53 and p21 (29). However, in another report, inactivation of Ini1 impaired cell proliferation and survival and induced p53 levels without affecting E2F-responsive genes (30). These data suggest that Ini1 loss may perturb both Rb and p53 tumor suppression pathways. Conditional inactivation of either Rb or p16INK4a did not alter the rate of tumorigenesis in mice, in which Ini1 was also conditionally inactivated by induction of Cre expression from an MX1-Cre transgene (29), whereas coinactivation of both Ini1 and p53 led to a dramatic increase in the rate of tumor onset (29, 30). Therefore, apparently, mutation of Rb, but not p53, might be redundant with Ini1 loss of function in vivo.

We have generated previously Ini1-heterozygous mice and MEFs and observed that the Ini1 levels in these Ini1-haploinsufficient cells are similar to WT Ini1 levels due to an increase in the activity of the remaining Ini1 promoter (12, 31). However, this transcriptional compensation for loss of one allele of Ini1 is not complete and results in a 15% to 20% reduction in total Ini1 levels in the Ini1-heterozygous cells. Thus, individuals heterozygous for Ini1 may be susceptible to the consequences of reduced Ini1 levels. To better understand the functional relationship between Ini1 and the Rb tumor suppressor and to determine if haploinsufficiency of Ini1 is sufficient to alter cell growth, we examined the growth characteristics of Ini1-heterozygous MEFs. Our results indicate that haploinsufficiency of Ini1 increases the rate of cell proliferation and cell immortalization and that Ini1 requires functional Rb to induce a senescent-like phenotype in primary cells. Moreover, mice heterozygous for Rb or for both Rb and Ini1 form highly penetrant pituitary tumors at similar rates. Although adrenocorticotropic hormone (ACTH)–positive tumor cells displaying LOH for the remaining WT Rb allele were present in tumors from both cohorts of mice, large atypical tumor cells lacking ACTH expression were also readily detected solely in pituitary tumors in the Rb+/−, Ini1+/− mice. Laser capture microdissection (LCM) revealed that the small, ACTH-positive cells underwent LOH for the WT Rb allele while retaining heterozygosity for Ini1, whereas the large atypical tumor cells underwent LOH for Ini1 but not for Rb. These findings indicate that Ini1 and Rb have redundant functions in suppressing pituitary tumorigenesis in Rb-heterozygous mice and that Ini1 is important for maintaining the morphologic and functional differentiation of normal corticotropic cells as well as those undergoing neoplastic transformation.

Materials and Methods

Cells and mice. Ini1-heterozygous mice and Rb-heterozygous mice have been described previously (12, 32). MEFs were generated from E12 to E14 embryos as described (33). Genomic DNA was prepared from the hind limb of each embryo for PCR analysis to identify the genotype of the resulting fibroblast line. All studies were initiated using low-passage embryonic fibroblasts (passages 2-4). Ini1+/−, Rb+/−, and Ini1−/−, Rb−/− mice were maintained on a C57Bl/6×129 genetic background. All mice were maintained and used in accordance with the University of Massachusetts Animal Care and Use Committee.

Cell studies. To determine the rates of cell proliferation, three different lines of WT MEF or Ini1-heterozygous MEFs were plated at 5 × 10^4 cells per 10-cm dish in MEM media (DMEM, with 10% fetal bovine serum, 0.37% sodium bicarbonate, penicillin, and streptomycin). Triplicate plates of cells for each line were harvested and counted at various times following initial plating using a Z1 Coulter Particle Counter (Beckman Coulter, Miami, FL.). A 3T9 assay was done as described (34) to examine the rate of spontaneous immortalization of WT MEF and Ini1-heterozygous MEFs. Briefly, 3 × 10^6 cells were plated into MEM media in 10-cm dish every 3 days. A total of three plates (9 × 10^6 cells) were maintained for two separate lines of fibroblasts for each genotype. Triplicate plates for each line were trypsinized and mixed before counting and replating. Recombinant Ini1 retrovirus was generated as described previously (31) and used to transduce Ini1 into WT and Rb-null MEFs. Cells were plated onto 10-cm dishes in MEM media at 5 × 10^5 per plate and infected with either an Ini1-flag recombinant retrovirus or an empty vector recombinant virus for 24 hours in the presence of polybrene. Infected cells were transiently selected in blastcicidin (10 μg/mL) for 60 hours to identify cells transduced by the recombinant Ini1 virus, harvested by trypsinization, and counted. Triplicate platings of transduced cells for each genotype were replated at a density of 4 × 10^6 cells per 10-cm dish, and plates were harvested at various time points following initial plating and counted using a Z1 Coulter Particle Counter.

Reverse transcription-PCR to detect Ini1 transcripts. Reverse transcription-PCR (RT-PCR) was done on reverse-transcribed total RNA isolated from MEFs infected with either pBABE-flag-Ini1 or pBABE-no insert (31). The primers for the flag-Ini1 cDNA were 5’-GTCACCACCATCGCATACAG-3’ (F922) and 5’-TCATATTGTGTCTCTGCTTTTC-3’ (R1188). The primers for the 36B4 control cDNA were 5’-CAGGCCCCTTGACCTC-TGGTCTGTTG-3’ (F699) and 5’-TTGTGGTGTGGTGTTGCCGGGATTAGTCT-3’ (R1023).

Western protein. Protein was extracted from 3T9 cells at various passages and quantified by Bradford analysis, and 50 μg of each sample were used for Western analysis with an Ini1 antibody as described (12).

Tumor assays. Ini1-heterozygous mice were crossed with Rb heterozygous mice to establish a cohort of Rb-Ini1 compound heterogeneous mice. Tumor assays were done using these mice and Rb heterozygous mice. Mice were inspected every other day for morbidity or tumor formation. Necropsies were done and tumors were harvested and fixed in 10% buffered formalin phosphate and then processed for paraffin embedding and sectioning using standard methods. Sections were mounted on a glass slide and stained with H&E. Sections of adenomas and normal pituitaries from WT mice were also immunostained for ACTH. Before LCM, coverslipped, H&E-stained, and ACTH-immunostained sections were studied microscopically. Areas of interest were then identified in replicate deparaffinized, noncovered sections, and LCM was done using an Arcturus (Mountain View, CA) Pixcell 2 instrument.

Genotyping of tumor samples. Following microdissection of select areas of tumor slides, the samples were placed into a small Eppendorf tube and genotyped by PCR for Ini1 and Rb status as described previously (12, 32).

Results

Increased proliferation of Ini1-heterozygous fibroblasts. Multiple lines of Ini1+/− MEFs or WT MEFs were generated by standard methods from E13.5 embryos isolated from pregnant...
Ini1<sup>+/−</sup> female mice intercrossed with WT male mice. Three separate lines of Ini1<sup>+/−</sup> or WT MEFs were harvested, plated, and counted at various times after initial plating to characterize the growth rate of these early-passage fibroblasts. The results indicate that Ini1<sup>+/−</sup> primary fibroblasts show an increase in their rate of proliferation relative to WT fibroblasts (Fig. 1A). This assay was repeated twice with similar results.

Cell immortalization studies. To explore the effects of Ini1 haploinsufficiency on cell immortalization, we did a 3T9 assay on two lines each of MEFs that were either WT or heterozygous for Ini1 (Fig. 1B). WT MEFs failed to immortalize during the course of this 3T9 assay, whereas MEFs that were Ini1<sup>+/−</sup> underwent rapid spontaneous immortalization by the fifth passage. Western analysis of Ini1<sup>+/−</sup> MEFs at various passages during the 3T9 assay revealed that these cells did not require Ini1 LOH to immortalize, as Ini1 expression was detected throughout various time points in Ini1<sup>+/−</sup> cells during the course of the 3T9 assay, including as late as passage 30 (Fig. 1C).

Cell transduction studies. Forced overexpression of Ini1 in MRT cells induces a flattened cell morphology and reduced cell growth, and several reports have implicated involvement of the Rb signaling pathway in Ini1-induced senescence (19, 26). To document the role of Rb in primary cells in Ini1-mediated cell senescence, we infected multiple lines of WT MEFs or MEFs derived from Rb-null embryos with a recombinant retrovirus that encodes a flag-tagged Ini1 and a blasticidin drug selection marker. Following infection and transient selection in blasticidin, the cells were harvested and plated in 10-cm dishes, and a proliferation assay was done using the transduced cells (35). WT MEFs infected with an empty vector virus continued to proliferate slowly throughout the assay (Fig. 2A). In contrast, WT MEFs transduced with virus encoding Ini1 assumed a flattened cell appearance and divided only minimally postinfection. As expected, Rb-null MEFs infected with an empty vector proliferated far faster than WT MEFs infected with empty vector virus. Interestingly, transduction of Ini1 failed to alter the morphology or growth characteristics of Rb-null MEFs, indicating that the replicative senescence-like effects of Ini1 expression are Rb-dependent. RT-PCR analysis of total RNA harvested from mock or Ini1-tranduced cells confirmed expression of exogenous, flag-tagged Ini1 in Rb and WT MEFs infected with the recombinant Ini1 virus (Fig. 2B).

Tumorigenesis in Ini1-Rb compound heterozygous mice. We have described previously Ini1 as a tumor suppressor in mice (12). To explore further genetic interactions between the Ini1 and Rb tumor suppressors in vivo, we generated Rb-Ini1 compound heterozygous mice by breeding Ini1<sup>+/−</sup> mice with Rb<sup>+/−</sup> mice. Rb-null mice develop normally until E14 but fail to develop thereafter due to massive cell death in the nervous system and defects in erythropoesis (32, 36, 37). Rb<sup>+/−</sup> mice are viable and appear normal but form foci of atypical melanotroph precursor cells in the intermediate lobe of the pituitary as early as 30 days after birth (38). Eventually, adenocarcinomas develop in the anterior pituitary lobe as do medullary thyroid adenomas and neuroendocrine tumors (38–41). Ultimately, 100% of Rb<sup>+/−</sup> mice succumb to pituitary tumors, and LOH for the WT Rb allele is observed in all late-stage tumors and in some of the microdissected atypical cells of early proliferates (38, 40).

To determine whether haploinsufficiency for Ini1 alters tumorigenesis in Rb heterozygous mice, a cohort of Rb<sup>+/−</sup>, Ini1<sup>+/−</sup> mice was generated and analyzed for the formation of spontaneous tumors. The rate and tissue spectrum of tumor formation in these compound heterozygous mice was compared with tumorigenesis in mice heterozygous for either Rb or Ini1. Mice heterozygous for

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**Figure 1.** Haploinsufficiency for Ini1 alters the proliferation and spontaneous immortalization of MEFs. A, analysis of the growth rate of (WT) MEFs and Ini1-heterozygous MEFs reveals that Ini1 haploinsufficiency induces a faster rate of proliferation in cells. B, immortalization of WT MEFs and Ini1-heterozygous MEFs. WT MEFs failed to immortalize following a 3T9 protocol to assess spontaneous immortalization, whereas cells haploinsufficient for Ini1 readily immortalized. C, Western analysis of Ini1 protein in the immortalized Ini1<sup>+/−</sup> cells at various passage numbers reveals that the Ini1-heterozygous cells retain Ini1 expression. The Ini1 antibody detects two Ini1 proteins generated through differential splicing of the Ini1 transcript (50).
both Rb and Ini1 developed pituitary tumors at a rate and penetrance indistinguishable from tumor formation in Rb\(^{-/-}\) mice (Fig. 3). The spectrum of tumors observed in Rb\(^{-/-}\) mice WT or heterozygous for Ini1 was also quite similar, except for one compound heterozygous mouse that developed thymic lymphoma, only pituitary tumorigenesis was detected Rb\(^{-/-}\) mice and in Rb\(^{-/-}\), Ini1\(^{+/-}\) mice. Interestingly, no rhabdoid-like tumors or poorly differentiated sarcomas were observed on the head and neck of Ini1\(^{-/-}\) Rb\(^{-/-}\) mice during this assay, although \(\sim 20\%\) of Ini1\(^{-/-}\) mice present with these cancers in the same time frame. Thus, haploinsufficiency for Ini1 does not alter the onset, penetrance, or spectrum of tumor formation in Rb\(^{-/-}\) mice. Furthermore, deletion of WT Rb or WT Ini1 did not rescue the embryonic lethality of either Ini1-null mice or Rb-null mice, respectively, because intercrossing of the compound heterozygous mice did not cause Ini1-null or Rb-null mice.

Histologic features and genotype of tumors arising in Rb\(^{-/-}\), Ini1\(^{+/-}\) mice. The size of pituitary tumors in the Rb-Ini1 compound heterozygous mice ranged from 4 to 7 mm in diameter, similar to what was observed in pituitary tumors arising in Rb\(^{-/-}\) mice in this and in other published studies (39, 42).

Pituitary tumors in Rb\(^{-/-}\) mice contained a uniform population of cells that morphologically resemble corticotrophic cells normally present in the intermediate and anterior pituitary lobes. Analysis of tumor sections revealed that the tumor cells in the pituitary of Rb heterozygous mice consistently immunostained strongly for ACTH (Fig. 4A). ACTH is a marker for anterior lobe corticotrope pituitary cells (Fig. 4B) and is expressed in both benign and malignant pituitary tumors that arise in Rb\(^{-/-}\) mice (43).

Tumors from Rb\(^{-/-}\), Ini1\(^{+/-}\) compound mice also exhibited a similar morphology and robust ACTH staining pattern. However, large ACTH-negative cells could also be found in more than half of the Rb\(^{-/-}\), Ini1\(^{+/-}\) tumors. These atypical cells were usually polygonal, displayed large round vesicular nuclei, and abundant amphophilic cytoplasm and present in aggregates throughout the tumor tissue (Fig. 4C-D). Notably, these large ACTH-negative cells were observed only in pituitary tumors that arose in the Rb-Ini1 compound heterozygous mice.

Figure 2. Overexpression of Ini1 in MEFs induces cell growth arrest.
A, retroviral-mediated transduction of Ini1 into WT MEFs retards the growth of these cells and induces a senescence-like state. However, transduction of Ini1 into Rb-null MEFs does not alter the growth of these cells, showing that effects of Ini1 overexpression on cell growth are Rb-dependent. B, transduced Ini1 (flag-tag) expression is detected by RT-PCR using primers to Ini1 and the flag sequences in WT and Rb\(^{-/-}\) MEFs infected with Ini1 recombinant retrovirus. RT-PCR against 36B4 cDNA was done as a control for RNA quality.

Discussion
Evidence in the literature indicates that the Ini1 tumor suppressor plays a role in regulating cell growth and chromosomal
...by regulating components of the Rb signaling pathway, including CDK inhibitors and/or cyclin D1 (reviewed in ref. 44). However, most of these studies were done using forced overexpression of Ini1 into Ini1-deficient malignant rhabdoid tumor-derived cell lines. More recently, analysis of tumorigenesis in mice revealed tumorigenesis induced by deletion of Ini1 was not altered on codeletion of either Rb or p16ink4, whereas loss of cyclin D1 activity inhibited tumorigenesis in Ini1-null mice (29, 45). These findings suggest that the Rb and Ini1 may possess redundant tumor-suppressing capabilities. Less data are available about the role of Ini1 in regulating p53 functions in cells or mice. Conditional deletion of Ini1 was reported to increase p53 activity in MEFs, suggesting that Ini1 negatively regulates p53 functions (29, 30). However, deletion of p53 also dramatically increased the onset of tumor formation following conditional deletion of Ini1 in mice in these studies, suggesting that Ini1 and p53 may also cooperate in preventing oncogenesis.

In this study, we have examined the growth characteristics of Ini1-haploinsufficient MEFs. The results reveal that loss of one allele of Ini1 increases the rate of cell proliferation, in keeping with the findings of earlier reports, indicating that Ini1 overexpression negatively regulates cell growth (15–19, 31). These data also indicate that haploinsufficiency of Ini1 can have functional ramifications on cell proliferation despite the compensatory effects of increased Ini1 expression from the remaining WT Ini1 allele (31). Spontaneous immortalization of Ini1+/− MEFs was also readily achieved in the 3T3 assay. Interestingly, cells haploinsufficient for Ini1 spontaneously immortalized without undergoing loss for Ini1, whereas WT cells failed to immortalize in this assay, again suggesting that primary cells haploinsufficient for Ini1 have altered growth characteristics.

To explore the Rb-mediated effects of Ini1 in primary cells, we transduced Ini1 expression into WT and Rb-deficient MEFs. Expression of exogenous Ini1 in MEFs induced a cell growth arrest and flat cell morphology, similar to previous findings of the effect of forced Ini1 expression in MRT cells (15–19). However, Ini1 transduction had no effect on MEFs lacking functional Rb, indicating that the senescence-inducing properties of Ini1 are Rb-dependent in these cells.

The development of pituitary adenomas in Rb+/− mice was accelerated on crossing the Rb-mutant mice with mice deficient for p27, p53, or p19ARF tumor suppressors (46–48). However, LOH of the WT Rb allele was observed in all tumors arising in these studies. To explore further the relationship between Rb and the Ini1 tumor suppressor in vivo, we generated Ini1-Rb compound heterozygous mice and compared tumor formation in these mice with tumorigenesis in Ini1+/− or Rb+/− mice. The rate and spectrum of tumor formation in Rb-heterozygous mice was unaltered by the presence of one or two copies of functional Ini1. Analysis of the genotypes of tumor cells determined that most tumor cells displayed LOH for the WT Rb allele as expected. However, not all pituitary adenoma cells lost functional Rb; the large atypical cells that were ACTH negative retained the WT Rb allele and displayed LOH for the functional Ini1 allele. These results indicate that loss of Rb or Ini1 is redundant in...
some pituitary tumors in mice, suggesting that these two tumor suppressors function similarly in preventing this type of cancer. However, loss of Ini1 in these tumors also seemed to alter the morphology of the tumor cell, with Ini1-null cells appearing larger and lacking ACTH expression. Based on these observations, we hypothesize that loss of Rb or Ini1 occurs early in cells of the intermediate and anterior pituitary lobes in the Ini1+/C0, Rb+/C0 mice and that complete loss of one tumor suppressor eliminates the need for LOH of the other suppressor gene in tumor formation. Surprisingly, mice haploinsufficient for both Rb and Ini1 failed to form any sarcomas or rhabdoid-like tumors of the head and neck in contrast to those observed in Ini1+/C0 mice. Therefore, although Rb and Ini1 appear redundant in suppressing pituitary tumorigenesis in these mice, the lack of MRT tumors in the Rb+/C0, Ini1+/C0 mice suggests that Rb is epistatic to Ini1 in tumor suppression.

The appearance of Ini1-deficient pituitary tumor cells that were enlarged relative to the Ini1 heterozygous, Rb-deficient tumor cells is reminiscent of the behavior of proliferating immortalized cells that express a dominant-negative version of the Brg1 ATPase. These cells were larger than counterpart cells not expressing the mutant Brg1, showing increased area of surface attachment, increased cell volume, and increased levels of proteins associated with cell adhesion (49). In both studies, the results suggest that interference with SWI/SNF enzyme function affects pathways controlling cell size and shape without blocking cell proliferation. Additional analyses will be needed to identify the specific genes involved and to address mechanisms by which chromatin remodeling enzymes affect the physical variables controlling cell size and shape. We note that these observations differ from previous documentation of a "flat cell" phenotype when Ini1 or BRG1 are ectopically expressed in Ini1- or BRG1-deficient tumor cells (15–19, 21, 22), as these cells are inevitably growth arrested.

The results of our study indicate that Ini1 regulates Rb functions in murine primary cells in vitro and in vivo and that Ini1 haploinsufficiency has functional consequences on cell growth. These results suggest that there might be clinical ramifications to Ini1 haploinsufficiency in members of MRT families. In addition, Ini1 seems to have morphologic consequences in pituitary cells and to regulate ACTH expression. Further analysis of Rb functions in Ini1+/C0 and Ini1-null cells and in mice should help elucidate the precise interplay of these signaling pathways in regulating cell proliferation and in tumor suppression.

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