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Deletion of p37^{Ing1} in Mice Reveals a p53-Independent Role for Ing1 in the Suppression of Cell Proliferation, Apoptosis, and

Tumorigenesis

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Abstract

ING proteins have been proposed to alter chromatin structure and gene transcription to regulate numerous aspects of cell physiology, including cell growth, senescence, stress response, apoptosis, and transformation. *ING1*, the founding member of the *in*hibitor of growth family, encodes p37^{Ing1}, a plant homeodomain (PHD) protein that interacts with the p53 tumor suppressor protein and seems to be a critical cofactor in p53-mediated regulation of cell growth and apoptosis. In this study, we have generated and analyzed p37^{Ing1}-deficient mice and primary cells to further explore the role of Ing1 in the regulation of cell growth and p53 activity. The results show that endogenous levels of p37^{Ing1} inhibit the proliferation of p53-wild-type and p53-deficient fibroblasts, and that p53 functions are unperturbed in p37^{Ing1}-deficient cells. In addition, loss of p37^{Ing1} induces *Bax* expression and increases DNA damage–induced apoptosis in primary cells and mice irrespective of p53 status. Finally, p37^{Ing1} suppresses the formation of spontaneous follicular B-cell lymphomas in mice. These results indicate that p53 does not require p37^{Ing1} to negatively regulate cell growth and offers genetic proof that Ing1 suppresses cell growth and apoptosis in a p53-independent manner.

Introduction

ING1 was initially isolated in a screen to identify genes that were underexpressed in transformed versus normal human epithelium (1). Subsequently, ING1 was identified as a member of a five-gene ING family that encode plant homeodomain (PHD)-finger domain proteins conserved in yeast and vertebrates (2). Forced overexpression of ING1 in normal human diploid fibroblast cells was found to inhibit cell growth by inducing a G_1 -phase cell cycle arrest. Conversely, down-regulation of ING1 in cell lines by antisense RNA increased cell proliferation and cell transformation, as assayed by colony formation in soft agar (1,3), and inhibited Myc-induced apoptosis (4). The ability of ING1 to inhibit cell growth and

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promote apoptosis in these various assays indicated that ING1 might function as a tumor suppressor.

The murine *Ing1* gene produces several spliced isoforms of Ing1 mRNA that generate two distinct Ing1 proteins (5). The larger 37-kDa protein ($p37^{Ing1}$) contains all of the residues present in the smaller 31-kDa ($p31^{Ing1}$) protein, as well as additional sequences at the NH₂ terminus that interact with the p53 tumor suppressor protein (5). Both mouse $p37^{Ing1}$ and the human orthologue p33 ^{ING1} have been shown to coimmunoprecipitate with p53 in transfected cells (5,6). Furthermore, p33 ^{ING1} has been reported to stabilize p53 levels in transfected cells by binding with p53 and blocking murine double minute-2 (Mdm2)-p53 interactions and by inhibiting hSir2 deacetylation of p53 (6–8). Functional data linking Ing1 with regulation of p53 activity were provided by transfection studies in which forced overexpression of ING1 in cell lines was found to both increase cell sensitivity to DNA double-strand breaks in a p53-dependent manner (9) and induce the expression of certain p53 target genes such as *p21* following DNA damage (7). In addition, the ability of exogenous p53 to inhibit cell growth in BALB/c 3T3 cells seems to be compromised in cells partially depleted for *ING1* by antisense RNA (6).

These cell-based studies suggest that the negative effects of Ing1 on cell growth are mediated through p53. Furthermore, mutation of ING1 has also been detected in human primary tumors and in tumor-derived cell lines, further underscoring the possibility that ING1 functions as a tumor suppressor (2,10). In support of this model, mice deleted for Ing1 via gene targeting experiments in embryonic stem cells were recently found to be smaller in size, have reduced viability following whole body irradiation, and display a slight increase in the rate of spontaneous tumor formation relative to wild-type (Wt) mice, particularly in lymphomagenesis (11). However, these Ing1-deficient mouse primary fibroblasts displayed little or no differences in replicative life span or in sensitivity to various forms of genotoxic stress. Thus, the link between Ing1, p53 activity, and tumor suppression remains unclear.

To determine if Ing1 regulates p53 functions in cell growth and tumorigenesis, we generated p37^{Ing1}-deficient mice by using mouse embryonic stem cells bearing a gene trap mutation that specifically ablates p37^{Ing1}. Analysis of p37^{Ing1}-deficient mice and mouse embryonic fibroblasts (MEF) revealed that loss of p37^{Ing1} increased the growth rate of MEFs, providing direct genetic evidence in primary cells that the endogenous level of Ing1 negatively regulates cell proliferation. However, deletion of p37^{Ing1} also increased the proliferation of p53-deficient MEFs, showing a p53-independent role for p37^{Ing1} in the control of cell growth. Furthermore, loss of p37^{Ing1} failed to alter numerous other cell characteristics normally governed by p53, including the rate of primary cell immortalization, oncogene-induced cell senescence, or cell growth arrest following DNA damage by known inducers of p53 activity. In addition, p37^{Ing1} deletion did not affect the p53-induced embryonic lethality of Mdm2-null mice. These data indicate that p53 remains fully functional in cells lacking p37^{Ing1}, indicating that p37^{Ing1} down-regulates cell growth in a p53-independent manner.

Although loss of p37^{Ing1} did not alter the levels of the proapoptotic p53 response gene *Puma*, p37^{Ing1} deletion was found to significantly increase *Bax* gene expression and promote apoptosis following DNA damage in thymocytes irrespective of p53 status, showing that p37^{Ing1} also has a p53-independent, prosurvival role in apoptosis. Whereas DNA damage– induced apoptosis was up-regulated in p37^{Ing1}-deficient primary cells, mice lacking p37^{Ing1} developed spontaneous follicular B-cell lymphomas, indicating that p37^{Ing1} functions as a tumor suppressor *in vivo*.

Materials and Methods

Generation of p37^{Ing1b}-deficient mice

Mouse (129 strain) embryonic stem cells (Omnibank no. OST206270, Lexicon Genetics, The Woodlands, TX) containing a retroviral promoter gene trap inserted into one allele of *Ing1* were used to generate chimeric mice, which were bred to generate p37^{Ing1}-heterozygous mice and subsequent p37^{Ing1}-homozygous mice. Mice were genotyped by PCR analysis of genomic DNA using the following primer sequences: Wt allele, ING1g6673SN: 5'-

TCCCCCGTGGAATGTCCTTATC-3', ING1g7259ASN: 5'-

CTGCCAACAGTAGTTCTGAGCAAG-3'; targeted allele, ING1gLTR: 5'-

AAATGGCGTTACTTAAGCTAGCTTGC-3', INGg1874ASN: 5'-

CTTGTGGAGAAAAATGCC-3'. A Southern blot strategy using a *BgI*II digest of genomic DNA and a probe to the $p31/p37^{Ing1}$ common exon was used to confirm the PCR genotyping results. Ing1b-null mice were crossed to Mdm2^{+/-} or p53-null mice and genotyped as previously described (12,13). All mice were maintained and used in accordance with both federal guidelines and the University of Massachusetts Animal Care and Use Committee.

Cell culture and proliferation assay

MEFs were generated from E13.5 day embryos as previously described (14). To determine the rates of cell proliferation, multiple lines of Wt, p53-null, p37^{Ing1}-heterozygous, p37^{Ing1}-null, or p37^{Ing1}/p53-double null MEFs were seeded at 2×10^5 per 60-mm plate or 1×10^5 per well of a six-well plate. Triplicate plates of each line were harvested and counted every 24 h using a Z1 Coulter Particle Counter (Beckman Coulter, Miami, FL). Duplicate gelatinized 10-cm plates with 10,000 cells per plate were seeded in MEF medium to assay for cell survival and growth at low plating density. At 8 to 12 days postplating, cells were fixed with methanol and stained with 0.1% crystal violet to visualize colony formation.

Cell immortalization assay

A 3T9 assay was done as described (15) to examine the rate of spontaneous immortalization of Wt, p37^{Ing1}-null, or p53-null MEFs. Briefly, 3×10^6 cells were plated into MEF medium on 10-cm plates 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 before counting and replating at a density of 3×10^6 per 10-cm plate every 3 days.

Cell senescence assay

MEFs were transduced with a recombinant *Ras* retrovirus as previously described (16) and plated at 2×10^5 per 60-mm plate. Triplicate wells of each of two lines of cells per genotype were harvested and counted every 48 h following initial plating using a Z1 Coulter Particle Counter (Beckman Coulter). Data were expressed as a ratio of the cell number after 6 days in culture versus the cell number at initial plating.

Cell growth arrest studies

MEFs from three independent lines of each genotype were seeded onto 10-cm plates at a density of 8×10^5 per plate in 0.1% fetal bovine serum for 3 days. Cells were then fed with medium supplemented with 10% serum for 4 h, and the cultures left untreated or exposed to 8 Gy of γ -radiation in a cesium irradiator, 300 ng/mL Adriamycin (Sigma-Aldrich, St. Louis, MO), or 60 J/cm² UVC in a UV Crosslinker (Stratagene, Cedar Creek, TX). At 15 h posttreatment, MEFs were pulse labeled with 60 µmol/L bromodeoxyuridine (BrdUrd) for 3 h, harvested by trypsinization into PBS, and fixed in 70% ethanol overnight at 4°C. Flow cytometric analysis of DNA synthesis and total DNA content was done using an anti-BrdUrd antibody, propidium iodide staining, and Flowjo software. Data are presented as a ratio of percentage of cells in S phase for treated versus untreated cells.

Apoptosis studies

MEFs transduced with a recombinant retrovirus encoding the adenovirus EIA gene (17) were plated at 10⁶ per 10-cm plate and treated with 0.3 µg/mL of Adriamycin for 24 h. Adherent and nonadherent cells were collected and assayed either for trypan blue exclusion or for propidium iodide staining by fluorescence-activated cell sorting (FACS) analysis. Wt, p37^{Ing1}-heterozygous, p37^{Ing1}-null, or p53-null mice that were 4 to 6 weeks old were wholebody irradiated with 10 Gy of ionizing radiation, and thymi harvested at 12 h post ionizing radiation. Thymi were ground between frosted glass slides and cells were assayed for viability by trypan blue exclusion. Approximately 10⁶ thymocytes were stained with CD4-phycocyanin (BD PharMingen, San Jose, CA; diluted 1:250) and CD8-allophycocyanin (BD PharMingen, diluted 1:100) for 30 min before FACS analysis. For spontaneous apoptosis experiments, single-cell suspensions of thymocytes were plated in RPMI medium with 5% FCS at 10^6 per 60-mm plate and incubated at 37°C, 5% CO₂ for 96 h. Plates were harvested, fixed in 70% ethanol, propidium iodide stained, and analyzed by FACS. Ex vivo thymocyte apoptosis experiments were done by plating 10⁶ thymocytes onto 60-mm plates in RPMI medium with 5% FCS and either mock treating or irradiating the plates with 2.5-Gy ionizing radiation. After 4 h, cells were harvested, stained with propidium iodide, and analyzed by FACS.

Quantitative real-time reverse transcription-PCR

Relative levels of mRNA expression were analyzed by quantitative reverse transcription-PCR (RT-PCR) as previously described (18). cDNA was generated using Invitrogen SuperScript first-strand synthesis system for RT-PCR. The following primer sequences (shown 5'–3') were used in the PCR reactions: *Bcl-2*, CTCGTCGCTACCGTCGTGACTTCG and GTGGCCCAGGTATGCACCCAG; *Puma*, CCTGGAGGGTCATGTACAATCT and TGCTACATGGTGCAGAAAAAGT; and *Bax*, CTGAGCTGACCTTGGAGC and GACTCCAGCCACAAAGATG. Samples were normalized to EF1α levels present in each tissue as previously described (18).

Tumor assays

A tumor cohort was established for p37^{Ing1}-heterozygous and p37^{Ing1}-homozygous null mice. Moribund mice or those that reached 22 months of age were sacrificed for necropsy, and select tissues harvested for DNA, RNA, and protein. A portion of each tissue was fixed in 10% phosphate-buffered formalin, paraffin embedded, and stained with H&E. Tumors were classified by morphology and by immunostaining with antibodies against surface antigen B220 (BD PharMingen, diluted 1:50) or CD-3 (DAKO, Denmark; diluted 1:400).

Results

Generation of p37^{Ing1}-deficient mice

A mouse embryonic stem cell line bearing a retroviral promoter trap inserted into one allele of the *Ing1* gene was used in standard blastocyst injection experiments to generate chimeric mice (19). DNA sequence analysis of the gene-trapped locus revealed that the retrovirus inserted into exon C, the third exon of Ing1 (5), interrupting the coding of the longer Ing1b transcript that encodes p37^{Ing1} (Fig. 1*A*). Several high-degree chimeric mice were bred with C57Bl/6 mice to generate agouti offspring, and genomic DNA was harvested from tail biopsies and analyzed by Southern blot hybridization (Fig. 1*B*) and PCR (Fig. 1*C*) to identify mice that inherited the *Ing1*-targeted allele. Heterozygous F₁ generation mice were intercrossed to obtain mice homozygous for the mutant *Ing1* allele, which were recovered with normal Mendelian

frequency. Similar to a recent report of mice mutated at the *Ing1* locus (11), homozygous mutant mice weighed ~15% less at weaning but were otherwise indistinguishable from the Ing1-heterozygous and Wt littermates. Western blot analysis of Ing1 proteins harvested from the spleens and thymi of Wt and homozygous mutant mice revealed that the homozygous mutant mice retained expression of the p31^{Ing1} protein in the presence (4 Gy whole body) or absence of DNA damage, but displayed loss of p37^{Ing1} protein (Fig. 1*D*). Thus, retroviral insertion into the Ing1 locus in the embryonic stem cells resulted in a p37^{Ing1}-specific knockout allele.

Negative regulation of cell growth by p37^{Ing1}

We generated primary fibroblasts from Wt, heterozygous $p37^{Ing1}$ -deficient, and homozygous $p37^{Ing1}$ -deficient embryos to characterize and compare the growth rates of these cells with p53-null MEFs (Fig. 2*A*). The results indicate that the proliferation rate of $p37^{Ing1}$ -heterozygous MEFs is similar to Wt MEFs. However, $p37^{Ing1}$ -null MEFs proliferate significantly faster than Wt MEFs, although not as fast as MEFs lacking p53. Although $p37^{Ing1}$ -null MEFs displayed Wt plating efficiencies under normal conditions, these cells showed increased survival and growth relative to Wt MEFs when plated at low density (Fig. 2*B*), similar to results obtained with p53-null MEFs (15). To determine if the down-regulation of cell proliferation by $p37^{Ing1}$ is mediated through p53, we bred the $p37^{Ing1}$ -deficient mice with p53-null mice (12) and generated $p37^{Ing1}/p53$ -double null MEFs. Analysis of the growth rate of these-double null cells revealed that deletion of $p37^{Ing1}$ enhanced the growth of p53-deficient primary fibroblasts (Fig. 2*C*), indicating that the increased proliferation rate of $p37^{Ing1}$ -null MEFs was not due to alteration of p53 activity in these cells.

Immortalization of p37^{Ing1} null MEFs

Spontaneous immortalization of MEFs is dependent, in part, on p53 functions. To analyze further the role of $p37^{Ing1}$ in cell growth, we did a modified 3T3 (3T9) cell immortalization assay. MEFs were generated from Wt, $p37^{Ing1}$ -null, or p53-null embryos and passed continuously in culture following a standard protocol (15). In agreement with our previous reports (15,20), p53-null MEFs displayed a rapid rate of cell growth throughout the course of the assay relative to Wt MEFs (Fig. 2*D*). In contrast, $p37^{Ing1}$ -deficient MEFs continued to divide more slowly than p53-null MEFs and never exhibited robust induction of cell growth. Thus, unlike deletion of p53, loss of $p37^{Ing1}$ does not alter the immortalization rate of primary cultured cells.

Ras-induced cell senescence in p37^{Ing1}-deficient MEFs

Oncogene-induced stress effects a p53-dependent, replicative senescent phenotype in MEFs (16). To determine if forced expression of an activated *ras* oncogene induces senescence in p37^{Ing1}-deficient fibroblasts, MEFs that were either Wt, p37^{Ing1} null, or p53 null were infected with pBABE retrovirus (vector) or with pBABE bearing an activated H-*ras* gene (vector + ras). After puromycin selection for 72 h, the transduced cells were plated at equal densities and scored for cell growth at day 6 postinfection. Vector alone (no *ras*) had little effect on the growth of MEFs regardless of genotype (Fig. 3A). As expected, exogenous *ras* suppressed the growth of Wt MEFs whereas the growth of p53-null MEFs was not affected by activated *Ras* expression. Similar to Wt MEFs, the growth of p37^{Ing1}-null MEFs was suppressed following oncogene transduction. These data indicate that p53-mediated cell senescence induced by inappropriate oncogene expression in MEFs is unaffected by the presence or absence of p37^{Ing1}.

Response of p37^{Ing1}-deficient MEFs to DNA damage

Induction of DNA double-strand breaks in MEFs by treatment with ionizing radiation or Adriamycin induces stabilization of p53 and subsequent inhibition of cell growth. To determine

if loss of p37^{Ing1} alters the ability of p53 to arrest cells following DNA damage, Wt, p37^{Ing1}null, or p53-null MEFs were synchronized in their cycling and mock treated or treated with either 8-Gy ionizing radiation or 0.3 µg/mL Adriamycin. The results of this experiment (Fig. *3B*) are expressed as the ratio of treated cells in S phase to untreated cells in S phase. As expected, Wt MEFs treated with either ionizing radiation or Adriamycin had a large reduction in the numbers of cells present in the S phase of the cell cycle, whereas p53-null MEFs displayed far less cell growth inhibition following treatment with ionizing radiation or Adriamycin. However, MEFs lacking p37^{Ing1} were indistinguishable from Wt MEFs in their response to these DNA-damaging agents, indicating that the ability of p53 to induce cell growth arrest in response to DNA double-strand breaks is unaltered in p37^{Ing1}-null MEFs. In addition, Wt, p37^{Ing1}-null, or p53-null MEFs were treated with 60 J/cm² UVC and assayed for a damageinduced reduction in S phase. Although UVC treatment strongly inhibits cell growth in Wt MEFs, this effect requires functional c-jun NH₂-terminal kinase signaling and is not p53 dependent in MEFs (21,22). Similar to Wt and p53-deficient MEFs, cells lacking p37^{Ing1} were also growth inhibited by UVC treatment.

Deletion of p37^{Ing1} does not alter p53-induced lethality in Mdm2-null mice

Mdm2 is a well-established regulator of p53 activity and p53 stability in cells (23,24), and mice deleted for Mdm2 display an early embryonic lethal phenotype that is rescued by deletion of p53 (13). Recently, the NH₂ terminus of p33^{ING1}, the human orthologue of p37^{Ing1}, has been found to complex with the p19-ADP ribosylation factor (ARF) tumor suppressor (25). In cells, p19-ARF regulates p53 activity by binding with Mdm2 and inhibiting Mdm2 ubiquitination of p53 (26). Interaction of p19-ARF and p33^{ING1} suggests that Ing1 may play a role in p19-Mdm2-p53 signaling, and Ing1 has previously been proposed to alter p53 stability, possibly by interfering with Mdm2-p53 interactions (25).

To explore whether deletion of $p37^{Ing1}$ alters p53 activity and Mdm2-p53 signaling during development, we bred $p37^{Ing1}$ -null mice with mice bearing a mutated Mdm2 allele (13). Intercrosses of the resulting Mdm2^{+/-}, $p37^{Ing1}$ -null mice were done, and DNA isolated from the offspring (n = 40) of these matings was genotyped to determine Mdm2 status. This experiment failed to generate any Mdm2-null mice (expected n = 10), indicating that the embryonic lethality induced by p53 in the absence of Mdm2 was not dependent on $p37^{Ing1}$ function. This result is in keeping with our finding that $p37^{Ing1}$ -null MEFs are not compromised in p53-mediated cell growth arrest following DNA damage.

Antiapoptotic role of p37^{Ing1} in DNA damage response

MEFs transduced with adenovirus EIA undergo p53-induced apoptosis following treatment with Adriamycin (24). To determine if p53 apoptosis was compromised in p37^{Ing1}-null cells, MEFs that were Wt, p53 null, or p37^{Ing1} null were infected with a recombinant retrovirus encoding both the adenovirus E1A gene and a puromycin resistance gene. Following a 2-day selection in puromycin, E1A-transduced MEFs were recovered, plated in triplicate at equal cell density, and mock treated or treated with 0.3 µg/mL Adriamycin for 24 h. Cell viability was scored by trypan blue exclusion. As expected, Wt MEFs displayed reduced cell viability on treatment with Adriamycin ($35.3 \pm 4\%$ viable cells), whereas p53-null MEFs were much less susceptible to treatment with this DNA-damaging agent ($64.7 \pm 4\%$ cell viability). Surprisingly, p37^{Ing1}-null MEFs showed far greater sensitivity to Adriamycin-induced death $(9.7 \pm 1\%)$ viable cells) than Wt cells. To confirm that cell death was due to apoptosis, the experiment was repeated and the percentage of cells undergoing apoptosis was determined by FACS analysis of the sub- G_1 DNA content for each genotype (Fig. 3C). As before, p53-null MEFs were compromised in their ability to undergo apoptosis relative to Wt MEFs, whereas p37^{Ing1}-null MEFs were highly sensitive to the effects of Adriamycin with most of the p37^{Ing1}-null MEFs undergoing apoptosis.

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To further explore aberrant apoptosis in p37^{Ing1}-null mice, we isolated thymocytes from 6-week-old Wt, p53-null, or p37^{Ing1}-null mice. The total number of thymocytes recovered from the thymi of p37^{Ing1}-null mice was reduced 2-fold relative to the cell numbers recovered from Wt mice, whereas no differences were observed in the cellularity of the spleens or peripheral lymph nodes in these mice (data not shown). However, Wt, p53-null, and p37^{Ing1}-null thymocytes displayed similar kinetics of cell death when plated in culture (Fig. 4A), suggesting that the spontaneous, non-DNA damage–induced thymocyte death that occurs under these culture conditions is independent of either p53 or p37^{Ing1} status.

Mouse immature thymocytes (CD4/CD8 double-positive T cells) will undergo p53-dependent apoptosis in response to ionizing radiation (27,28). To determine if DNA damage-induced cell death is up-regulated in thymocytes lacking p37^{Ing1}, three 4- to 6-week-old mice that were Wt, p53 null, p37^{Ing1} null, or p37^{Ing1}/p53 double null were whole-body treated with 10-Gy ionizing radiation and thymocytes were recovered at 12 h posttreatment from these mice and from agematched, nontreated mice. FACS analysis of thymocyte populations stained with CD4 and CD8 antibodies revealed that $45.5 \pm 3.1\%$ of the Wt CD4/CD8 double-positive cells survived ionizing radiation treatment, whereas only $31.4 \pm 2.1\%$ of the nondamaged levels of p37^{lng1}null double-positive T cells were recovered after ionizing radiation treatment. As expected, far more p53-null double-positive T cells survived ionizing radiation damage ($78.5 \pm 1.4\%$). However, deletion of p37^{Ing1} also significantly reduced the survival of p53-null doublepositive T cells (56.3 \pm 0.3%). These data suggest that p37^{Ing1} plays a prosurvival role in thymocytes following DNA damage irrespective of p53 status. To confirm that p37^{Ing1} deletion alters thymocyte apoptosis, thymocytes were harvested from Wt, p53-null, p37^{Ing1}-null, or p53/p37^{Ing1}-double null mice and either untreated or treated with 2.5 Gy of ionizing radiation. Cells were collected 4 h after treatment or mock treatment and analyzed by propidium iodide staining and FACS to measure sub-G₁ DNA content (Fig. 4B). In agreement with the findings of the previous E1A-MEF experiments, deletion of p37^{Ing1} led to increased DNA damageinduced apoptosis in Wt thymocytes. Furthermore, although deletion of p53 correlated with reduced thymocyte apoptosis following ionizing radiation damage, loss of p37^{Ing1} also resulted in increased apoptosis following DNA damage in p53-null cells, indicating that p37^{Ing1} has a p53-independent, antiapoptotic role in thymocytes.

To determine if increased apoptosis in p37^{Ing1}-null thymocytes reflected up-regulated expression of the proapoptotic p53 target gene *Puma*, an important regulator of DNA damage–induced apoptosis in lymphocytes and MEFs (29), the levels of *Puma* RNA transcripts were measured by quantitative PCR before and after DNA damage. Total RNA was harvested at 4 h posttreatment from the thymi, spleens, livers, and brains of mock-treated or ionizing radiation–treated (10 Gy) mice that were Wt, p53 null, p37^{Ing1} null, or p53/p37^{Ing1} double null, and real-time PCR analysis was done to determine the expression levels. As expected, *Puma* expression was increased in the thymi of Wt mice in response to DNA damage, but not in mice lacking functional p53 (Fig. 4*C*). However, there was no significant difference in the levels of *Puma* induction in thymi of Wt mice and p37^{Ing1}-null mice, or in p53-null mice in the presence or absence of p37^{Ing1}.

Bax is also a critical inducer of apoptosis in T cells and in other cell types, but it is not thought to be an important transcriptional target of p53 (30,31). In keeping with this model, Bax RNA levels were only slightly up-regulated in Wt thymus, spleen, and liver following DNA damage and were unchanged or slightly decreased following ionizing radiation treatment in p53-null samples. In contrast, *Bax* mRNA levels were induced ~8- to 10-fold after DNA damage in both $p37^{Ing1}$ -null and $p37^{Ing1}/p53$ -double null thymi (Fig. 4*D*). Induction of *Bax* expression following DNA damage was also elevated 4- to 6-fold in the spleens, livers, and brains of $p37^{Ing1}$ -null mice and ~2-fold in these tissues in $p37^{Ing1}$ -null mice lacking p53. In addition,

levels of *Bcl-2* RNA, an antiapoptotic gene, were not significantly altered in these tissues by the presence or absence of either $p37^{Ing1}$ or p53 (data not shown).

These data suggest that loss of $p37^{Ing1}$ does not promote thymocyte apoptosis by altering p53 transactivation of *Puma* following DNA damage. Instead, loss of $p37^{Ing1}$ leads to up-regulated *Bax* gene expression following DNA damage of thymocytes irrespective of p53 status.

Tumorigenesis in p37^{Ing1}-deficient mice

To determine if mice specifically deleted for $p37^{Ing1}$ are tumor prone, we generated cohorts of Wt, $p37^{Ing1}$ -heterozygous ($p37^{Ing1+/-}$), and $p37^{Ing1}$ -null ($p37^{Ing1-/-}$) mice and monitored these mice for the development of spontaneous tumors. Approximately 40% of the $p37^{Ing1+/-}$ mice and half of the $p37^{Ing1-/-}$ mice formed tumors by 22 months of age, whereas spontaneous tumorigenesis was not observed in the Wt (C57Bl/ 6×129 -SV hybrid) mice during this time (Fig. 5*A*). All of the tumors arose in the spleen or peripheral lymph nodes in these mice and were classified as follicular B-cell lymphomas. Staining of tumor tissues with antibody against B220 or CD3 confirmed that these tumors originated in the B-cell compartment (Fig. 5*B*) and not in T cells. Thus, although deletion of $p37^{Ing1}$ does not alter p53 growth control and greatly up-regulates apoptosis in response to DNA damage, $p37^{Ing1}$ clearly functions as a tumor suppressor in B cells in mice.

Discussion

ING family members are highly conserved, PHD-containing proteins that are linked to chromatin remodeling and transcriptional regulation through their association with histone acetyl transferase and histone deacetylase complexes (2,32–34). Previous studies have implicated Ing1 (or human ING1) as an important regulator of numerous aspects of mammalian cell physiology, including cell growth, apoptosis, senescence, and tumorigenesis (2). The *Ing1* gene encodes for three alternatively spliced isoforms of Ing1 message: Ing1a and Ing1c encode a 31-kDa protein and Ing1b encodes for a 37-kDa protein in mice (p33 kDa in human). Forced Ing1b overexpression or antisense RNA knockdown experiments in cultured cells have indicated that p37^{Ing1} (or p33^{ING}) and p53 coimmunoprecipitate and can function in a synergistic manner to inhibit cell growth or promote apoptosis following DNA damage (5–7). Furthermore, some ING proteins, including p33^{ING}, have been proposed to regulate cellular p53 levels by modifying p53 interactions with components of the p19-Mdm2-p53 signaling pathway (2,8). In addition to altering p53 activity, transfection experiments have also suggested possible p53-independent roles for ING1b in the regulation of cell proliferation and apoptosis (4,35).

In this study, we generated p37^{Ing1}-deficient mice and primary cells to examine the role of this p53-binding protein in the regulation of cell growth, cell death, and tumorigenesis. Specific deletion of p37^{Ing1} in MEFs increases the rate of cell proliferation and growth at low plating densities, confirming that physiologic levels of an Ing protein can function to inhibit cell growth in primary cells. However, this regulation does not seem to involve p53 because deletion of p37^{Ing1} also increases the proliferation of MEFs lacking p53. Furthermore, p37^{Ing1}-deficient MEFs fail to exhibit perturbation of other cell growth parameters governed by p53, including spontaneous cell immortalization, oncogene-induced senescence, or cell cycle arrest following DNA damage. In addition, deletion of p37^{Ing1} failed to rescue the p53-dependent embryonic lethality seen in Mdm2-null mice, offering further support that loss of p37^{Ing1} does not seem to compromise p53 activities in cells or in mice.

Human ING1b (p33^{ING1}) has previously been proposed to play a proapoptotic role in fibroblasts (4,36). In contrast, analysis of apoptosis in p37^{Ing1}-deficient MEFs indicates that physiologic levels of p37^{Ing1} strongly inhibit apoptosis in these cells following DNA damage.

To confirm a prosurvival role for p37^{Ing1} following DNA damage, we analyzed ionizing radiation–induced apoptosis in primary thymocytes and in thymus tissue *in vivo*. As expected, apoptosis was decreased in cells lacking p53 following DNA damage. However, apoptosis was increased in thymocytes both *in vitro* and *in vivo* in p37^{Ing1}-deficient cells, in keeping with our previous results in primary fibroblasts. Notably, apoptosis was dramatically increased in thymocytes isolated from both p37^{Ing1}-deficient mice and p37^{Ing1}/p53-double null mice following DNA damage. Thus, in contrast to the proapoptotic role of p53 in these cells, p37^{Ing1} plays an antiapoptotic role in thymocytes following DNA damage that is clearly p53 independent.

Puma is a p53 transcriptional target validated *in vivo* as important regulator of apoptosis following DNA damage (29). In thymocytes, p53-mediated apoptosis is governed primarily through Puma activation because mice deleted for either Puma or p53 are equally deficient in ionizing radiation–induced thymocyte apoptosis. However, Puma expression was equally upregulated in Wt and p37^{Ing1}-null thymocytes following DNA damage, further supporting a p53-independent role for p37^{Ing1} in the suppression of apoptosis. Bax, another critical regulator of mitochondrial-associated apoptosis, functions by altering permeabilization of the mitochondrial outer membrane. Deletion of p37^{Ing1} dramatically increased up-regulation of Bax expression *in vivo* in both Wt and p53-null thymocytes following DNA damage. Loss of p37^{Ing1} also up-regulated Bax levels to a lesser extent in spleen, liver, and brain tissues. Increased levels of Bax induction in p37^{Ing1}-deficient mice following DNA damage occurred independent of p53 status in all tissues, confirming a p53-independent role for p37^{Ing1} in suppressing Bax expression following DNA damage.

The results of our study indicate that $p37^{Ing1}$ suppresses cell proliferation in the presence or absence of p53 and does not alter p53-mediated cell growth arrest following DNA damage. In addition, $p37^{Ing1}$ negatively regulates Bax levels *in vivo* and functions as a prosurvival molecule following DNA damage in thymocytes regardless of p53 status. These data suggest that $p37^{Ing1}$ might also suppress tumor formation in a p53-independent manner. Interestingly, all of the tumors that arose in the $p37^{Ing1}$ -deficient colonies were classified as follicular B-cell lymphoma. This tumor type is rarely observed in p53-null mice (12), which present mainly with thymic lymphomas and osteosarcomas, offering further support that p53 and p37^{Ing1} may suppress tumorigenesis through different mechanistic pathways. Furthermore, whereas there is a significant acceleration of tumorigenesis in mice haploinsufficient or deleted for $p37^{Ing1}$, the onset of tumors in heterozygous or homozygous $p37^{Ing1}$ -null mice is similar, suggesting that any reduction in $p37^{Ing1}$ levels might promote spontaneous tumor formation.

Recently, mice lacking both $p37^{Ing1}$ and $p31^{Ing1}$ have been analyzed (11). The $p37^{Ing1}/p31^{Ing1}$ -deficient mice undergo normal development but have reduced body size and present with spontaneous tumors later in life. In addition, MEFs derived from $p37^{Ing1}/p31^{Ing1}$ -deficient mice display little or no changes in cell cycling after treatment with Taxol or other DNA-damaging agents. As these characteristics are also observed in $p37^{Ing1}$ -deficient mice, it is possible that loss of $p37^{Ing1}$ is the underlying cause of the defects observed in the $p37^{Ing1}/p31^{Ing1}$ -deficient mice and cells. However, in addition to B-cell lymphomas, histiocytic sarcomas and myeloid leukemias were also observed in $p37^{Ing1}/p31^{Ing1}$ -deficient mice, suggesting that the $p31^{Ing1}$ protein may also have unique tumor-suppressing properties. Interestingly, expression of $p31^{Ing1}$ seems to be slightly elevated in mice lacking the $p37^{Ing1}$ -ablated cells and mice. However, any contribution of increased $p31^{Ing1}$ levels to the phenotype of $p37^{Ing1}$ -deficient mice and cells must also be p53 independent.

The results of this study show a p53-independent role for $p37^{Ing1}$ in the negative regulation of cell proliferation and apoptosis and highlight a role for $p37^{Ing1}$ in the suppression of B-cell lymphomagenesis. Furthermore, the data indicate that loss of $p37^{Ing1}$ correlates with increased Bax expression in thymocytes following DNA damage. Although up-regulation of Bax has been found to also increase cell proliferation in some settings (37,38), it is possible that increased apoptosis in $p37^{Ing1}$ null thymocytes might prevent tumor formation in this tissue. Further work will be needed to better characterize the role of Bax in the $p37^{Ing1}$ regulation of cell proliferation, apoptosis, and B-cell lymphomagenesis.

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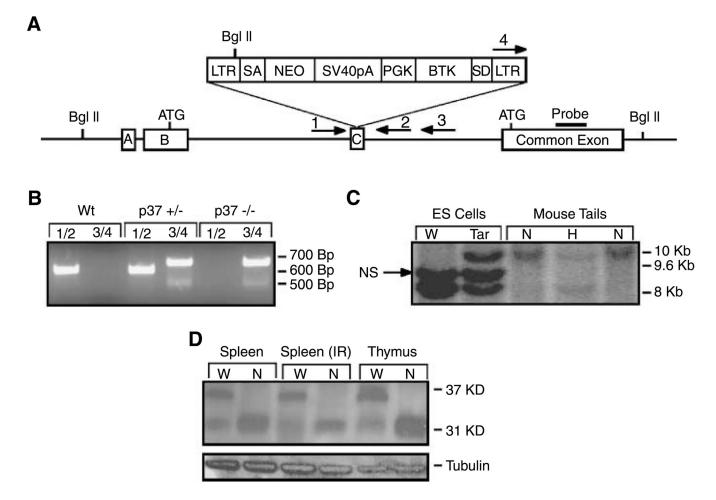


Figure 1.

Generation of $p37^{Ing1}$ -deficient mice. *A*, schematic of the gene-trapped locus showing the gene trap inserted into exon C. The trap interrupts the Ing1b message isoform and corresponding p37 protein but encodes the shorter Ing1c isoform and p31 protein. *B*, PCR genotyping of the embryonic stem cells and mouse tail biopsies showing the presence of the gene trap. A 650-bp fragment is generated from Wt Ing1 using primers 1 and 2 and a 700-bp fragment is generated Ing1 allele using primers 3 and 4. *C*, a Southern blot strategy using a *Bgl*II digest and a probe to the common exon was used to confirm germ-line transmission of the gene trap. The Wt fragment is ~8 kb and the mutant fragment is ~10 kb in length. A nonspecific (*NS*) band corresponding to an Ing1 pseudogene present in 129 strain mouse DNA is also observed. *D*, to confirm that the longer p37^{Ing1} form was specifically deleted, a Western blot was done with an Ing1 antibody that recognizes sequences encoded in the common exon. The presence of the shorter form (p31) and the absence of the longer form (p37) were observed. Tubulin was used as a loading control.

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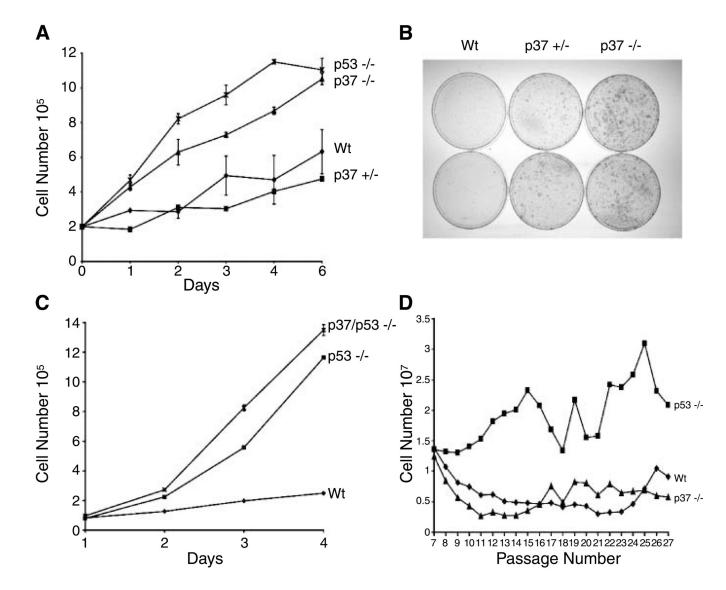


Figure 2.

Growth regulation byp37^{Ing1}. *A*, proliferation of p37^{Ing1}-deficient cells. Two independent MEF cell lines of p37^{Ing1}-Wt, p37^{Ing1}-heterozygous, p37^{Ing1}-null, or p53-null genotypes were plated in triplicate in 60-mm plates and growth was analyzed over the course of 7 d. Statistical analysis between Wt and p37^{Ing1}-heterozygous curves shows no significant difference (P < 1.00), but there is statistical difference between Wt and p37^{Ing1}-null curves (P < 0.005). *B*, growth of p37^{Ing1}-deficient cells at low density. Two lines of Wt, p37^{Ing1}-heterozygous, or p37^{Ing1}-heterozygous genotype were seeded at 10⁴ cells per 10-cm plate and incubated for 8 to 12 d before staining with crystal violet. *C*, growth inhibition by p37^{Ing1} is independent of p53 status. Three MEF cell lines deficient for p53 or for p53 and p37^{Ing1} were plated in triplicate in six-well plates and counted over a 7-d period. Wild-type MEFs were also similarly plated as a control. A statistical difference exists between the proliferation rates of p53-null and p37^{Ing1}/p53-double null MEFs (P < 0.0005). *D*, immortalization of p37^{Ing1}-deficient cells. A 3T9 assay was done using two independent MEF cell lines of p37^{Ing1}-Wt, p37^{Ing1}-null, or p53-null MEFs.

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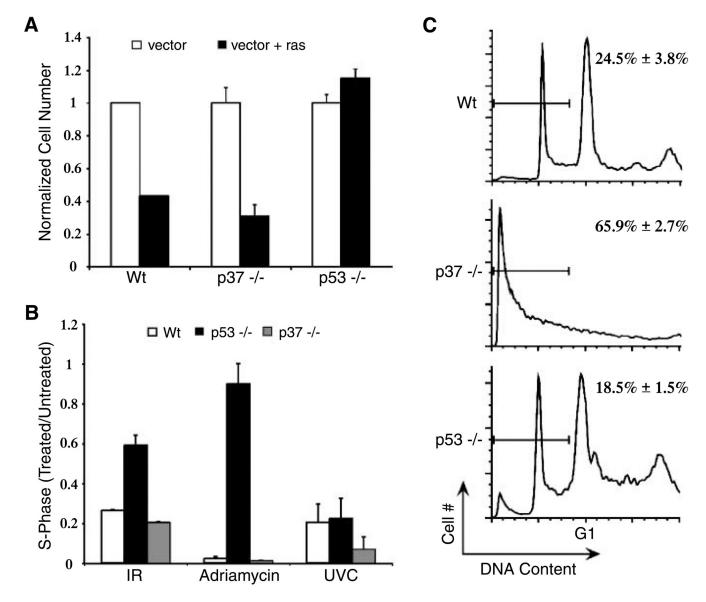


Figure 3.

Cell cycle arrest and *ras*-induced senescence are normal in $p37^{Ing1}$ -deficient MEFs, but apoptosis is elevated. *A*, oncogene-induced senescence in $p37^{Ing1}$ -deficient cells. Triplicate experiments were done wherein multiple lines each of Wt, $p37^{Ing1}$ -null, or p53-null MEFs were transduced with recombinant virus with or without H-*ras*, and cells were plated at equal densities and counted. *Columns*, ratio of cell number after 6 d in culture versus the cell number at initial plating. *B*, absence of $p37^{Ing1}$ does not alter cell cycle arrest due to DNA-damaging agents. Three lines each of wt, $p37^{Ing1}$ -null, or p53-null MEFs were plated in duplicate and synchronized in their growth before mock treatment or treatment with Adriamycin, ionizing radiation, or UVC. Cells were stained with BrdUrd and propidium iodide and analyzed by FACS. *Columns*, ratio of treated cells in S phase versus untreated cells in S phase. *C*, triplicate experiments of multiple lines of E1A-transduced MEFs were mock treated or treated with Adriamycin for 24 h. Cell viability was determined by staining with propidium iodide before FACS analysis. Representative result and sub-G₁ content derived from three separate experiments. *Columns*, mean; *bars*, SD.

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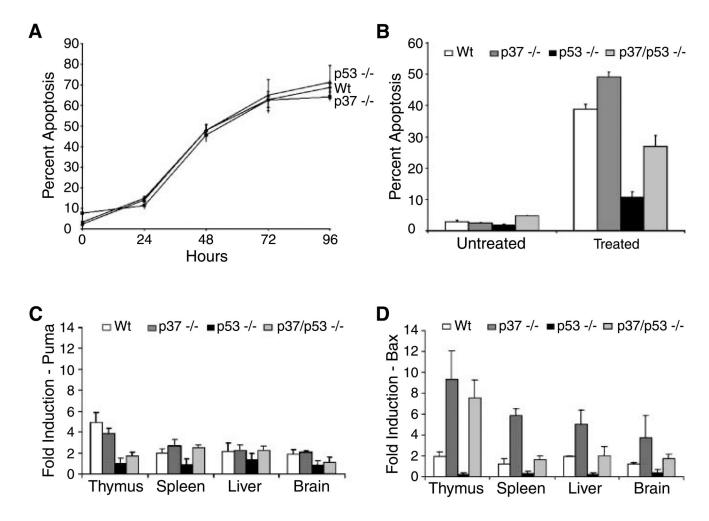


Figure 4.

Apoptosis is altered in p37^{Ing1}-deficient thymocytes. A, spontaneous apoptosis is unchanged in p37^{Ing1}-deficient thymocytes. Thymocytes were isolated from two Wt, p37^{Ing1}-null, or p53null mice, plated in triplicate, and cultured for 4 d ex vivo. Cells were harvested at 24-h intervals, fixed in 70% ethanol, and stained with propidium iodide and analyzed by FACS. B, thymocytes were isolated from Wt, p37^{Ing1}-null, p53-null, or p37^{Ing1}/p53-double null mice and either mock treated or irradiated with 2.5 Gy of ionizing radiation, fixed in 70% ethanol after 4 h, and stained with propidium iodide for FACS. Columns, average of three separate experiments. The difference in apoptosis between Wt and p37^{Ing1}-null or between p53-null and p37^{Ing1}/p53double null thymocytes is statistically significant (P < 0.005). C and D, real-time PCR was done with primers to Puma or Bax on tissues harvested from mice that were either mock treated or whole-body treated with 10 Gy of ionizing radiation and harvested 4 h later. Columns, average values (relative fold induction) of three mice for each genotype. The difference in *Bax* expression is statistically significant between Wt and p37^{Ing1}-null tissues for thymus and spleen (P < 0.05), but not for brain and liver (P < 0.07). The difference in Bax expression between p53-null and p37^{Ing1}/p53-double null tissues is statistically significant for thymus, spleen, and brain (P < 0.05), but not for liver (P < 0.06).

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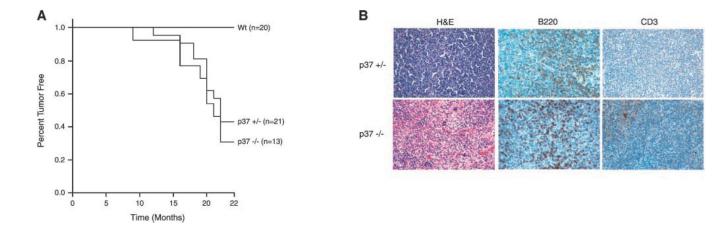


Figure 5.

Tumorigenesis in mice deficient for p37^{Ing1}. *A*, Kaplan-Meier survival curves of cohorts of Wt, p37^{Ing1}-heterozygous, or p37^{Ing1}-null mice. Mice that were either moribund or reached 22 mo of age were sacrificed for necropsy and fixed in 10% phosphate-buffered formalin. The rate of tumor incidence between Wt and either p37^{Ing1}-heterozygous or p37^{Ing1}-null mice is statistically significant (P < 0.005 or P < 0.05, respectively). *B*, tumors arising in mice were paraffin embedded and H&E stained for pathologic analysis. Tumor sections were also stained with either B220 or CD3 antibodies to determine tissue of tumor origin.