

# Tumorigenesis in the multiple intestinal neoplasia mouse: Redundancy of negative regulators and specificity of modifiers

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**The interaction between mutations in the tumor-suppressor genes *Apc* and *p53* was studied in congenic mouse strains to minimize the influence of polymorphic modifiers. The multiplicity and invasiveness of intestinal adenomas of *Apc*<sup>Min/+</sup> (Min) mice was enhanced by deficiency for *p53*. In addition, the occurrence of desmoid fibromas was strongly enhanced by *p53* deficiency. The genetic modifier *Mom1* and the pharmacological agents piroxicam and difluoromethylornithine each reduced intestinal adenoma multiplicity in the absence of *p53* function. *Mom1* showed no influence on the development of desmoid fibromas, whereas the combination of piroxicam and difluoromethylornithine exerted a moderate effect. The ensemble of tumor suppressors and modifiers of a neoplastic process can be usefully analyzed in respect to tissue specificity and synergy.**

Genes of several kinds regulate normal and neoplastic growth in the mammal. Positive and negative regulators can act either cell autonomously or nonautonomously to alter the proliferative potential of the stem cell that establishes the normal or neoplastic lineage. Individual genes in the ensemble can affect the transition between normal and neoplastic growth and/or the maintenance of one of the growth states. One important class of genes has been identified by loss-of-function alleles transmitted through the germline in heterozygous form, each predisposing a carrier to a particular spectrum of neoplasms. In these tumors, it is commonly found that the remaining wild-type allele has been lost or inactivated. The gene in question, commonly called a “tumor suppressor,” is formally a cell-autonomous negative regulator of the neoplastic state.

Humans and mice heterozygous for germline mutations that inactivate the *Adenomatous polyposis coli* (*APC/Apc*) gene develop only a limited range of neoplasms, including intestinal adenomas and desmoid fibromas (1). Because this gene is widely expressed (2), this restriction in neoplastic histotype is surprising. Does the limited range of neoplasia reflect cooperation between *APC/Apc* and other negative regulators?

The *p53* gene is a negative regulator that is mutated in a broad range of human neoplasms, but its gene product is constitutively active in only a few cell types (3, 4). For example, the loss of *p53* adenocarcinomas in the human intestine, but the protein becomes detectable in this tissue after stress such as ionizing radiation (5). Does an interaction between *Apc* and *p53* affect intestinal neoplasia in experimental models of cancer (6)? Preliminary studies tested whether a lack of *p53* activity affected neoplasia in the intestine of Min (multiple intestinal neoplasia) mice heterozygous for the *Min* nonsense allele of *Apc*, but no significant effect was observed (7–10). Those studies utilized populations of mice with heterogeneous genetic backgrounds. In this report, we reinvestigate whether a tissue-specific or stage-specific interaction between *Apc* and *p53* affects the spectrum of neoplasms in mice, by using a homogeneous genetic background.

A central issue addressed in these experiments is whether the major negative regulators each act singly in controlling neoplasia of a particular histotype, as implied by the “gatekeeper hypothesis” (11).

Beyond these major negative regulators, a number of other factors have been found to influence intestinal neoplasia in humans and mice. Such modifiers can act to either promote or retard tumor growth, thereby affecting tumor multiplicity in experimental models. The genetic modifier *Mom1* encodes a secretory phospholipase, *Pla2g2a*, expressed throughout the intestinal tract. The active allele of *Pla2g2a* leads to a reduction in the growth rate and multiplicity of intestinal adenomas in the Min mouse. Similarly, pharmacological agents can affect tumorigenesis. Piroxicam, a nonsteroidal antiinflammatory agent that inhibits cyclooxygenase-1 and -2, and difluoromethylornithine (DFMO), a suicide substrate of ornithine decarboxylase, each reduce the multiplicity of intestinal adenomas. The strongest inhibition is seen with a combination of piroxicam and DFMO (12). In the present study, we have asked whether the action of either *Mom1* or the piroxicam/DFMO combination is tissue specific. Further, we have asked whether either of these growth-inhibitory actions depends on *p53* activity.

## Materials and Methods

**Mice.** The C57BL/6-*p53* congenic mouse strain (B6-*p53*) was founded by a (129/Sv × B6) F2 female that carried a targeted disruption of the *p53* gene (13). Designating the founder female as N1, an expanded N10 intercross population was produced by mating *Apc*<sup>+/+</sup> *Mom1*<sup>R/S</sup> *p53*<sup>+/-</sup> females to *Apc*<sup>Min/+</sup> *Mom1*<sup>R/S</sup> *p53*<sup>+/-</sup>, *Apc*<sup>Min/+</sup> *Mom1*<sup>R/S</sup> *p53*<sup>-/-</sup>, and *Apc*<sup>Min/+</sup> *Mom1*<sup>R/S</sup> *p53*<sup>-/-</sup> males. Mice were housed as described previously (14).

**Drug Treatment.** After weaning at approximately 30 days of age, animals were housed in groups of one to five in microisolator cages under fluorescent lighting on a 12-hr cycle and weighed once per week. Tap water was available *ad libitum* for the duration of the experiment and was replaced weekly. The mice were treated with the appropriate drug or control vehicle mixed in the defined synthetic AIN-93G diet (Dyets, Bethlehem, PA), then killed after the specified duration of treatment.

The chemoprotective agents were stable for at least 7 days in a standard rodent diet at the concentrations used in these

Abbreviation: DFMO, difluoromethylornithine.

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**Table 1. Effect of *p53* on intestinal tumor multiplicity in *Min* mice**

	<i>Mom1<sup>S/S</sup></i>			<i>Mom1<sup>R/S</sup></i>			<i>Mom1<sup>R/R</sup></i>			Treated*		
	N	Tumor count	P-value <sup>†</sup>	N	Tumor count	P-value <sup>†</sup>	N	Tumor count	P-value <sup>†</sup>	N	Tumor count	P-value <sup>‡</sup>
<i>p53<sup>+/+</sup></i>	20	32 ± 11	NA	20	13 ± 5	NA	19	5 ± 4	NA	5	5 ± 2	0.0007
<i>p53<sup>+/-</sup></i>	19	37 ± 9	0.072	21	15 ± 7	0.403	20	8 ± 4	0.053	10	6 ± 4	0.00001
<i>p53<sup>-/-</sup></i>	21	45 ± 13	0.0014	14	26 ± 8	0.00002	16	13 ± 6	0.00008	4	12 ± 8	0.003

Mice were killed at 90 days of age. Tumor counts are shown as means ± SD.

\**Mom1<sup>S/S</sup>* mice were treated with piroxicam and DFMO (see *Materials and Methods*).

<sup>†</sup>P-values were calculated by using two-sided Wilcoxon rank sum tests compared with *p53<sup>+/+</sup>* mice of the same *Mom1* genotype.

<sup>‡</sup>P-values were calculated by using two-sided Wilcoxon rank sum tests compared with untreated mice of the same *p53* genotype.

studies. New batches were prepared weekly by thorough mixing of the diet with the indicated doses of drug and were stored until use in sealed containers at 4°C. Fresh diet was added to protected feeders three times weekly and was completely changed after the feeders were emptied once weekly.

Piroxicam (CAS no. 36322–90-4) was purchased from Sigma, and DFMO (CAS no. 70052–12-9) was a gift from Ajit Verma (University of Wisconsin–Madison). Piroxicam at the intended concentration was mixed in the diet beginning at approximately 30 days of age; the mice were killed at age 90 days after 2 mo of treatment. Mice treated with DFMO were given water mixed with the intended concentration of that drug from age 30 to 90 days. When administered to mice after weaning, these cyclooxygenase and decarboxylase inhibitors were not overtly toxic at the doses used.

**DNA Preparation.** Mice were anesthetized with ether, and 250 μl of blood was collected from the retroorbital sinus. DNA was isolated from blood as described previously (14).

**Genotyping.** Mice were genotyped to identify carriers of the *Min* allele of *Apc*, the resistance allele of *Mom1*, and the knockout allele of *p53* with PCR assays as described (14–16). In some cases, tail DNA was prepared from neonates to verify independently the *p53* genotype by Southern blot analysis, as described previously (13).

**Tumor Counts.** All mice were killed at 90 days by CO<sub>2</sub> asphyxiation. The intestinal tract was removed, washed with PBS, opened longitudinally, and laid out as described previously (6). In this method, 4-cm sections from the proximal, medial, and distal regions of the small intestine and the entire colon were examined. The number of tumors was scored with a dissecting microscope by a single observer blind to the genotype of the mice or their treatment group. The samples were then fixed in 10% buffered formalin, washed in 70% ethanol, and stored in this solution. Tumor counts were verified by a second observer in a subset of the postfixed samples.

At sacrifice, the number of desmoid fibromas was scored by a single observer blind to the genotype of the mice or their treatment group. The abdominal body wall was then removed, stretched, fixed overnight in 10% formalin, washed in 70% ethanol, and stored in this solution. Tumor counts were verified by a second observer for a subset of the postfixed samples. To facilitate scoring, the tissue was stained with Fast green, a dye that detects collagen. Samples were incubated for 30 min each in the following series: distilled water; acetate buffer (1 M acetic acid/7.2 mM sodium acetate/0.25% formalin); 5% (wt/vol) phosphotungstic acid in distilled water; Fast green [1 M acetic acid/110 mM sodium acetate/0.05% (wt/vol) Fast green]; acetate buffer; 10 mM sulfuric acid; 50% ethanol; and 70% ethanol.

**Histological Analysis of Intestinal Tumors.** The largest tumors were isolated from sections of the small intestine and the entire colon. After fixation in 10% buffered formalin, the tumors were embedded in paraffin, sectioned, and stained with hematoxylin and eosin. Sections were analyzed by light microscopy for signs of progression by H. C. Pitot (McArdle Lab), also blind to the genotype of the samples.

**Histological Analysis of Desmoid Fibromas.** Desmoid fibromas were isolated from postfixed abdominal body walls. The tumors were embedded in paraffin, sectioned, and stained with either hematoxylin and eosin or Masson's trichrome.

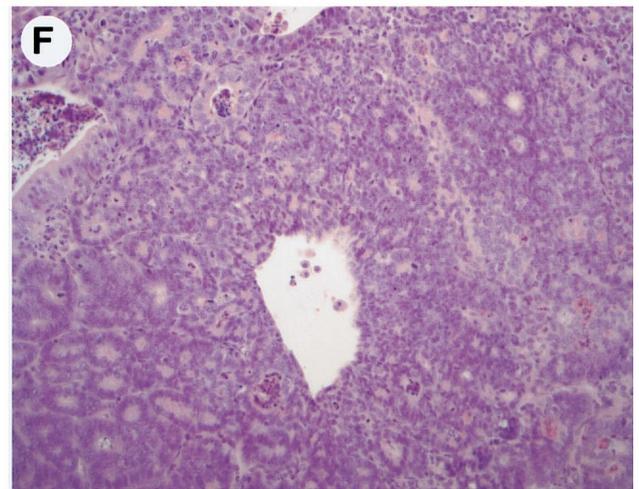
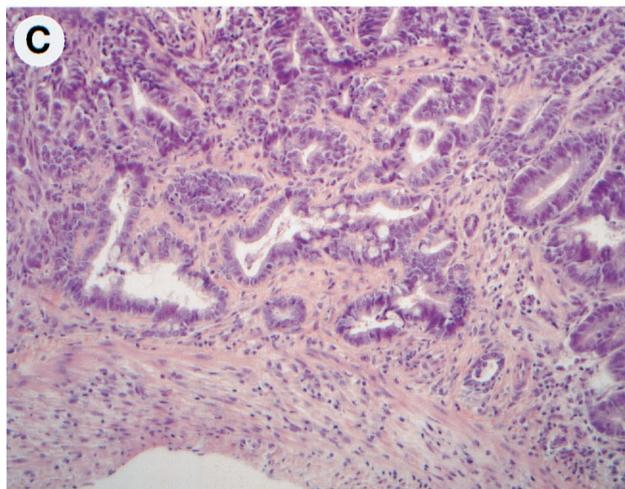
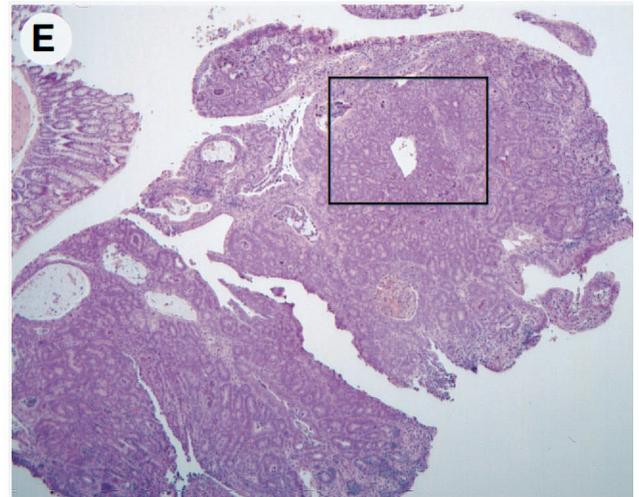
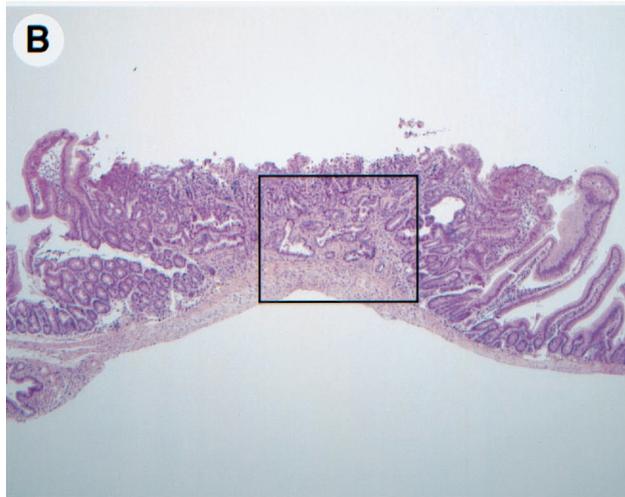
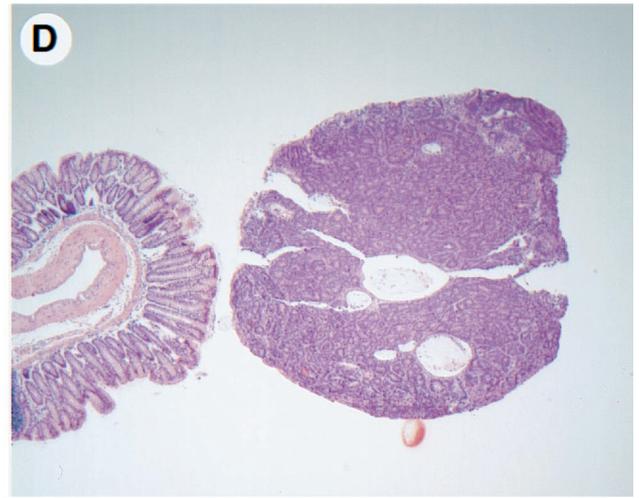
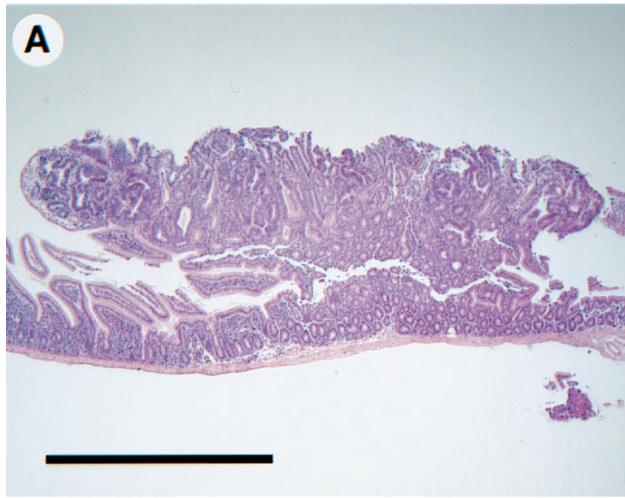
## Results

***p53* Effect on Multiplicity of Intestinal Adenomas.** Does a lack of *p53* activity affect the multiplicity of adenomas in the intestine of the *Min* mouse? *Min* mice homozygous for a null allele of *p53* developed significantly more intestinal adenomas than those homozygous for the wild-type allele of *p53*, regardless of the *Mom1* genotype (Table 1). Thus, *p53* negatively regulates the development of adenomas in the intestine of the *Min* mouse.

*Min* mice heterozygous for a null allele of *p53* (*Apc<sup>Min/+</sup> p53<sup>+/-</sup>*) also develop more tumors than those homozygous for the wild-type allele of *p53* (Table 1). The difference is at the borderline of statistical significance for two of the *Mom1* classes.

One interpretation for the effect of the *p53* genotype is that a polymorphic modifier, linked to the *p53* locus, affects the development of intestinal adenomas in the *Min* mouse (14). In deriving the B6-*p53* congenic animals, a region from chromosome 11 of the 129 genome was moved onto the B6 genetic background. We have tested this hypothesis and found no evidence for such a modifier. The extent of the 129/SvEv genome carried by the *p53* mutant line was assessed by genotyping *Min* mice heterozygous for a null allele of *p53* with markers distributed along chromosome 11. This analysis indicated that the introgressed region of the 129/SvEv genome in the B6-*p53* congenic strain extends at least from 19.7 cM to the *p53* locus at 39 cM on chromosome 11. To test whether a polymorphic modifier was present in this region of the 129 genome, B6 females were mated to (129/SvPas × B6)F1 *Min* males. At least 20 resulting progeny were scored for the number of adenomas and genotyped with markers from chromosome 11. Tumor multiplicity was comparable between *Min* mice that were heterozygous for the 129/SvPas allele and those that are homozygous for the B6 allele at each locus (data not shown). Thus, the heterozygous effect of the *p53* genotype cannot be explained by a dominant polymorphic modifier of intestinal tumorigenesis that is linked to the *p53* locus.

***p53* Effect on the Progression of Intestinal Adenomas.** Mutation of *p53* has been correlated with the progression of tumors in the human colon (17). To investigate whether a lack of *p53* function enhances tumor progression in *Min* mice, the largest tumors



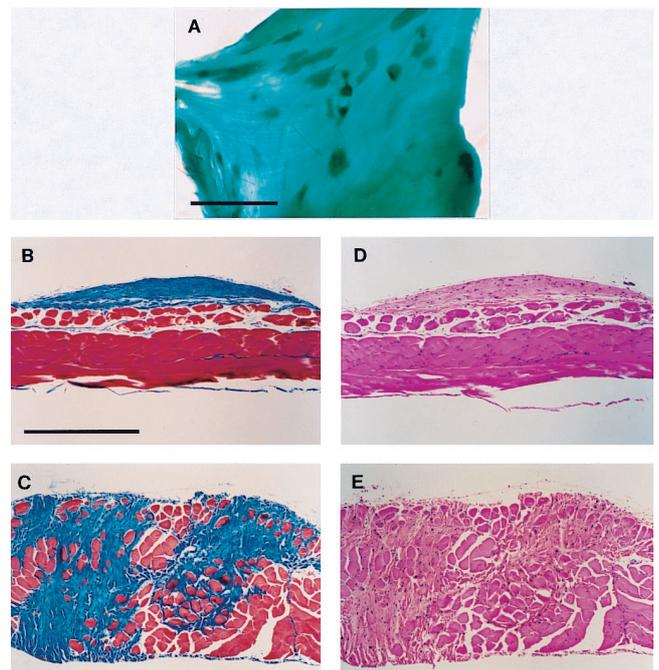
**Fig. 1.** Intestinal tumors exhibit signs of progression in *Apc<sup>Min/+</sup> p53<sup>-/-</sup>* mice. Tumors were isolated from *Apc<sup>Min/+</sup> Mom1<sup>S/S</sup>* mice carrying zero, one, or two copies of the *p53* knockout allele and stained with hematoxylin and eosin. Most tumors from the small intestine and colon of *p53*-deficient mice were typical adenomas (A and D, respectively). However, signs of progression were exhibited by some of the tumors isolated from the proximal region of the small intestine and the colon (B and E, respectively). The regions that exhibited signs of progression are boxed in B and E and are shown at higher magnification in C and F. The magnification was the same for A, B, D, and E, with the scale bar in A representing 1 mm. C and F are  $\times 4$  of B and E, respectively.

from three regions of the small intestine and the entire colon were analyzed for signs of morphological progression in Min mice of each *p53* genotype. Two of the 42 tumors that were isolated from the small intestines of *Apc*<sup>Min/+</sup> *p53*<sup>-/-</sup> mice were locally invasive with tumors invading the underlying *muscularis mucosae* (Fig. 1 *B* and *C*), whereas two of the nine tumors isolated from the colons of these mice exhibit features characteristic of *in situ* carcinomas (Fig. 1 *E* and *F*). By contrast, all 78 tumors isolated from *Apc*<sup>Min/+</sup> *p53*<sup>+/+</sup> and *Apc*<sup>Min/+</sup> *p53*<sup>+/-</sup> mice were benign. This difference in the incidence of invasive tumors lies at the borderline of statistical significance [ $P = 0.05$ , Fisher's exact test, one sided]. Thus, the lack of *p53* activity may affect the Min phenotype by permitting the progression of adenomas to adenocarcinomas. Alternatively, *p53* deficiency may permit invasive tumors to arise from a stem cell population distinct from that giving rise to adenomas. Would a stronger effect be observed if the tumors could be analyzed in older mice? Note that mice were killed at 90 days of age in this study. Unfortunately, the average lifespan of *Apc*<sup>Min/+</sup> *p53*<sup>-/-</sup> mice is 122 days, regardless of the *Mom1* genotype. Furthermore, the loss-of-function *p53* allele used in these studies may not fully simulate any gain-of-function *p53* allele found in human tumors (18). Other experimental designs are necessary to explore more fully the extent to which alterations in *p53* can lead to more advanced tumors in the Min mouse.

**Function of *p53* on the Multiplicity of Desmoid Fibromas.** Min mice and humans carrying germline defects in *APC* develop desmoid fibromas. This type of tumor is characterized by nodules or penetrating masses of unusually dense connective tissue that result from the active proliferation of fibroblasts. Some of these features are evident in sections of tumors stained with dyes that detect collagen (Fig. 2).

The multiplicity of these lesions in Min mice is dramatically affected by the lack of *p53* activity (Table 2). These lesions formed on the musculature of the entire body but were scored only in the abdominal region. *Apc*<sup>Min/+</sup> *p53*<sup>-/-</sup> mice develop on average 52 desmoid fibromas, whereas *Apc*<sup>+/+</sup> *p53*<sup>+/-</sup>, *Apc*<sup>Min/+</sup> *p53*<sup>+/+</sup>, and *Apc*<sup>Min/+</sup> *p53*<sup>+/-</sup> mice develop on average less than a single tumor per animal. This difference indicates that both *Apc* and *p53* must be altered for maximal transformation of normal fibroblasts into a neoplastic counterpart. A germline mutation in one of these genes is almost completely without effect in the presence of a wild-type form of the other. These results contrast with the observation by Smits *et al.* (19) of desmoid tumors in an *Apc* knockout strain wild type for *p53*. It is possible that the neomycin-resistance insertion in the knockout allele of *Apc* exerted a position effect on a gene adjacent to *Apc* (20).

**Effects of *Mom1* and Pharmacological Agents on the Multiplicity of Intestinal Adenomas and Desmoid Fibromas.** The resistance allele of *Mom1* (*Mom1*<sup>R</sup>) and the pharmacological agents piroxicam and DFMO slow the net growth rate of intestinal adenomas in the Min mouse and reduce multiplicity. Do these factors act by stimulating a *p53*-dependent activity, such as cell cycle arrest or apoptosis? The number of adenomas was scored in *Apc*<sup>Min/+</sup> *p53*<sup>-/-</sup> mice carrying zero, one, or two copies of *Mom1*<sup>R</sup>. A single copy of this allele reduced the number of tumors by a factor of 1.7, two copies reduced the number by a factor of 3.5 (Table 1). The semidominant effect of *Mom1* in *p53*-deficient animals was indistinguishable from its effect in mice heterozygous or wild type for *p53*. Similarly, the effect on tumor multiplicity of the piroxicam/DFMO combination was unaffected by the lack of *p53* activity (Table 2). Thus, *Mom1* and these pharmacological agents act independently of any *p53*-dependent processes. Furthermore, the growth inhibition exhibited by the piroxicam/



**Fig. 2.** Desmoid fibromas develop on the abdominal body wall of *Apc*<sup>Min/+</sup> *p53*<sup>-/-</sup> mice. The abdominal body wall was isolated and fixed in 10% formalin. The postfixated tissue was either stained with Fast green (*A*) or further dissected to isolate individual tumors. Isolated tumors were sectioned and stained either with Masson's Trichrome (*B* and *C*), in which muscle stains red and collagen blue, or with hematoxylin and eosin (*D* and *E*). These lesions can form on the surface of the abdominal wall muscle (*B* and *D*) or infiltrate and disrupt the organization of this tissue (*C* and *E*). The scale bar in *A* represents 5 mm, whereas that in *B* represents 0.5 mm. The magnification in *B*–*E* was the same.

DFMO combination does not depend on the active *Mom1* resistance allele encoding the Pla2 g2a secretory phospholipase.

Do the resistance allele of *Mom1* and the piroxicam/DFMO combination affect the development of desmoid fibromas? Tumors were scored on the abdominal body wall of *Apc*<sup>Min/+</sup> *p53*<sup>-/-</sup> mice either carrying *Mom1*<sup>R</sup> or treated with both piroxicam and DFMO (Table 2). All *Mom1* classes of mice were indistinguishable in desmoid multiplicity, indicating that the resistance allele of *Mom1* does not suppress the development of desmoid fibromas. By contrast, mice treated with the piroxicam/DFMO combination showed a significant reduction in desmoid multiplicity. Additional mice need to be studied to confirm this effect. Thus, *Mom1* action is tissue specific, affecting tumorigenesis only in the intestine of the *Apc*<sup>Min/+</sup> mouse, whereas the action of the piroxicam/DFMO combination appears not to be restricted to the intestine.

## Discussion

We are analyzing the genetic network controlling intestinal neoplasia in the laboratory mouse to further a fundamental understanding of human colon cancer. One of the mysteries in the development of human colon cancer is its nonlinear age dependence, fitted to a six-hit process (21). One strategy to identify the elements contributing to such a multihit process is to generate mice carrying germline mutations of candidate tumor-suppressor genes and to observe strong enhancement of neoplasia. The candidates currently available for such a test are the genes of the mouse homologous to those mutated in familial or sporadic colon cancer in the human. An impediment to a rigorous test of cooperation between negative regulators is the creation of heterogeneous genetic backgrounds when the distinct mutations are carried on different inbred backgrounds. This

**Table 2. Tests for effects of *p53* and *Mom1* on the development of desmoid fibromas in Min mice**

	<i>Mom1<sup>S/S</sup></i>			<i>Mom1<sup>R/S</sup></i>			<i>Mom1<sup>R/R</sup></i>			Treated*		
	N	Tumor count	P-value†	N	Tumor count	P-value†	N	Tumor count	P-value†	N	Tumor count	P-value†
<i>p53<sup>+/+</sup></i>	27	0.5 ± 0.7	0.69	19	0.7 ± 1.0	0.69	19	0.6 ± 1.0	0.89	2	0.5 ± 0.7	0.84
<i>p53<sup>+/-</sup></i>	40	0.8 ± 1.3	0.11	28	1.6 ± 2.2	0.11	26	1.1 ± 1.7	0.16	3	0 ± 0	0.25
<i>p53<sup>-/-</sup></i>	23	53 ± 11	0.12	17	58 ± 10	0.12	14	51 ± 12	0.66	2	34 ± 9	0.04

Tumor counts are shown as means ± SD. Mice were killed at 90 days of age, and the number of desmoid fibromas in the abdominal region were scored. Using two-sided Wilcoxon rank sum tests, the results for *Mom1<sup>S/S</sup>* mice were compared with those for *Mom1<sup>R/S</sup>* ( $P = 0.122$ ) and *Mom1<sup>R/R</sup>* mice ( $P = 0.661$ ). Desmoid fibromas were not observed in *Apc<sup>+/+</sup> p53<sup>+/+</sup>* mice, regardless of *Mom1* type.

\**Mom1<sup>S/S</sup>* mice were treated with piroxicam and DFMO (see *Materials and Methods*).

† $P$ -values were calculated by using two-sided Wilcoxon rank sum tests compared with *Mom1<sup>S/S</sup>* mice of the same *p53* genotype.

‡ $P$ -values were calculated by using two-sided Wilcoxon rank sum tests compared with untreated *Mom1<sup>S/S</sup>* mice of the same *p53* genotype.

heterogeneity creates genetic variation for polymorphic modifiers of the phenotype (7, 15, 22) that can obscure any cooperation. This study has utilized congenic derivatives of a single inbred strain, B6, to detect cooperation between *p53* and *Apc* in suppressing intestinal adenoma formation (Table 1) and progression (Fig. 1). More impressively, strong cooperation has been detected between these negative regulators in suppressing desmoid fibromas of the skeletal muscle wall (Fig. 2; Table 2).

Strong cooperation between distinct negative regulators of neoplasia (“tumor suppressors”) can provide a basis for multihit cancer development with a kinetic order higher than the two-hit process hypothesized by Knudson (23). Such a higher-order process may contain within it the canonical two-hit/one-locus process developed for retinoblastoma and the retinoblastoma gene in the human (24) and intestinal adenoma formation in the mouse (25). These simpler cases may fit the “Gatekeeper Hypothesis,” in which a single negative regulator controls the neoplastic potential of a particular cell type (11). However, a more general understanding of neoplasia in mammals must recognize instances of redundant regulation, such as that between *p53* and *Apc* in controlling desmoid fibromas (Table 2) and that between *p53* and *Rb* in controlling endocrine tumors (26).

The *APC/Apc* gene is expressed very broadly throughout mammals (27). The restricted spectrum of neoplasms found in humans and mice carrying germline mutations in *APC/Apc* could be explained by cooperation with other negative regulators. However, the *p53* gene is not an obvious candidate for fulfilling such a role, because it appears to be constitutively expressed in only a few cell types (3, 4), becoming more broadly active under several conditions of stress (28, 29). Several possibilities can be considered for the involvement of *p53* function in the development of invasive intestinal tumors and desmoid fibromas. The *p53* gene may be constitutively active in the precursors of these neoplasms. Note that Li-Fraumeni patients carrying germline mutations in the *p53* gene usually develop soft tissue sarcomas of mesenchymal origin (30). Alternatively, the *p53* gene may be induced as the neoplasm is initiated. The loss of *APC/Apc* activity in the tumor might lead to the activation of the *c-Myc* protooncogene (31), which in turn might activate *p53* via sequential steps of negative regulation from *p19ARF* to *Mdm2* to *p53* (32, 33). If this process activates *p53*-dependent apoptosis, then a deficiency of *p53* would enhance tumor formation.

Other testable mechanisms for interaction between *Apc* and *p53* can be proposed. The *p53* transcription factor is required for *Apc* expression *in vitro* (34), so a deficiency of *p53* may lead to silencing of *Apc* expression (35) and enhanced adenoma formation. A molecular analysis of the status of the wild-type *Apc* allele in the tumors that arise in Min mice lacking *p53* may be informative. Indirect modes of interaction are also conceivable and testable, involving any of the polypeptides known

to interact with *APC/Apc* (36) or any of the genes known to be regulated by *p53* activity or polypeptides known to interact with *p53* (28, 29, 37).

Beyond the negative regulatory genes of neoplasia, *APC/Apc* and *p53*, the modifier gene *Mom1* encoding the secretory phospholipase *Pla2g2a* and the pharmacological agents piroxicam and DFMO must be placed in the network of interactions controlling neoplasia. The activity of each of these factors leads to a reduction in net growth rate of the adenomas of Min mice (12, 38). The modifying effect of *Mom1* is tissue specific, affecting only the intestinal phenotype of the Min mouse. Early insight into the selective action of modifiers in pleiotropic cancer syndromes has been reported for familial neurofibromatosis type I (39).

Two questions are important in assessing these modifying agents for chemoprevention or therapy of tumors. Is the effect on growth rate lost in *p53*-deficient tumors? Clearly both *Mom1* and the piroxicam/DFMO combination are active against intestinal adenomas under *p53*-deficient conditions (Tables 1 and 2). Is there synergistic action between modifiers? It has been found previously that the treatment of Min mice with the combination of piroxicam plus DFMO is much more effective than either agent alone, resulting in a significant number of tumor-free mice (12). These findings of successful combination chemoprevention pertain to Min mice on the B6 genetic background, deficient in the secretory phospholipase *Pla2g2a*. Finding that *Mom1* resistance is also independent of *p53* function points to the importance of investigating whether the secretory phospholipase or one of the products of its action will synergize with piroxicam/DFMO to give complete chemoprevention and/or therapy for adenomatous polyps. A parallel study has found *p53* independence and strong synergy between the modifiers of intestinal tumors, *Mom1* and *Dnmt* (R. T. Cormier and W.F.D., unpublished work). Single factors, each with a small effect on a disease process, can be combined usefully.

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