

Tumorigenesis and Neoplastic Progression

# Inactivation of $p21^{WAF1/cip1}$ Enhances Intestinal Tumor Formation in $Muc2^{-/-}$ Mice

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**In the  $Apc1638^{+/-}$  mouse model of intestinal tumorigenesis, targeted inactivation of the cyclin-dependent kinase inhibitor  $p21^{WAF1/cip1}$  is highly effective in enhancing Apc-initiated tumor formation in the intestine. Because  $p21^{WAF1/cip1}$  plays a critical role in regulating intestinal cell proliferation, maturation, and tumorigenesis, we examined whether its inactivation would enhance tumor formation in a different mouse model of colon cancer. Therefore, we mated  $p21^{-/-}$  mice with mice carrying a genetic deficiency of the  $Muc2$  gene, which encodes the major gastrointestinal mucin.  $Muc2^{-/-}$  mice develop tumors in the small and large intestine and the rectum, but in contrast to tumors in  $Apc1638^{+/-}$  mice, this does not involve increased expression or nuclear localization of  $\beta$ -catenin. We found that inactivation of  $p21^{WAF1/cip1}$  significantly increased the frequency and size of intestinal tumors in  $Muc2$  knockout mice and also led to development of more invasive adenocarcinomas. This enhanced tumorigenesis significantly decreased mouse life span. Further, inactivation of  $p21^{WAF1/cip1}$  increased cell proliferation, decreased apoptosis, and decreased intestinal trefoil factor expression in the mucosa of both the small and large intestine. Surprisingly, reduced expression of  $p27^{kip1}$  was also observed in the  $Muc2^{-/-}$ ,  $p21^{+/-}$ , and  $p21^{-/-}$  mice. In contrast, the expression of  $c-myc$  was significantly elevated. Thus,  $p21$  modulates the formation of tumors whose initiation does (Apc) or does not ( $Muc2$ ) involve altered  $\beta$ -catenin-Tcf4 signaling, but which may converge on common elements downstream of this signaling pathway. (*Am J Pathol* 2005, 166:1239–1246)**

$p21^{WAF1/cip1}$  is an inhibitor of cyclin-dependent kinase (cdk) activity and is therefore an important regulator of cell cycle progression, and overall cell maturation, including

differentiation and apoptosis.<sup>1–4</sup> Despite this apparently central role of p21, mice with a targeted, homozygous inactivation of the p21 gene are essentially normal, although embryonic fibroblasts from such mice are defective in cell-cycle arrest at the G<sub>1</sub> checkpoint.<sup>5</sup> Thus, either p21 is not critical in cell maturation or its loss can be compensated for by other alterations that preserve pathways necessary for normal development and tissue homeostasis.

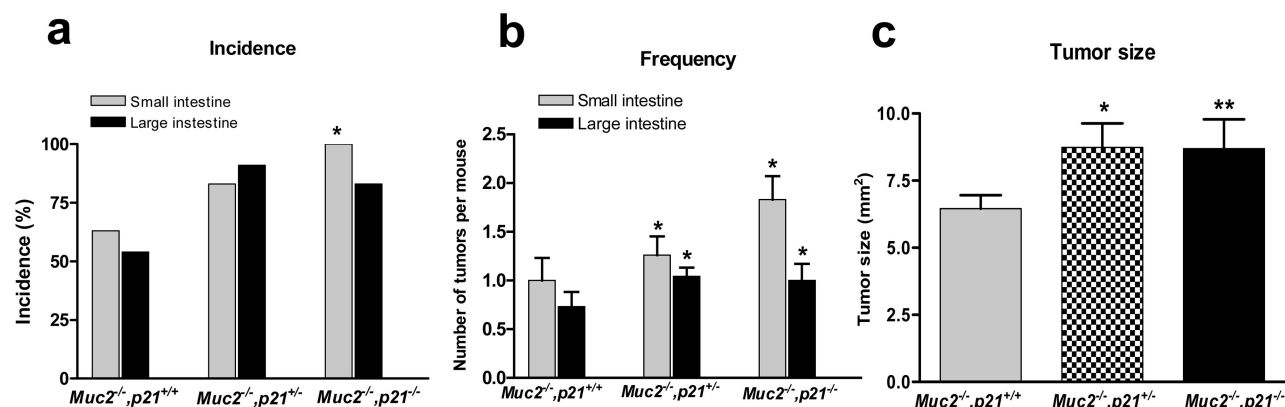
In the intestinal tract, p21 appears to play a key role in the maturation of cells as they migrate from the crypt toward the lumen.<sup>6</sup> p21 is expressed as cells exit the proliferative compartment. This pattern of expression, as well as the known functions of p21, suggested to us that p21 levels would modulate tumor formation in the intestine, and indeed, the introduction of a targeted inactivation of p21 into the  $Apc1638^{+/-}$  mouse, in which intestinal tumors form when the wild-type Apc gene is lost or inactivated, substantially increased intestinal tumor formation.<sup>3</sup> More recently, the importance of p21 in homeostasis of the intestinal mucosa has been emphasized and clarified by the finding that expression of p21 is repressed by c-MYC protein,<sup>7–9</sup> a target of APC- $\beta$ -catenin-Tcf signaling.<sup>8,10–12</sup> c-MYC sequesters the transcription factor MIZ in a MIZ/MYC complex,<sup>7,9,13,14</sup> and data have been presented that the down-regulation of  $c-myc$  gene expression, and consequent activation of p21 expression by MIZ, is a critical event in triggering intestinal cell differentiation.<sup>8</sup>

Interestingly, analysis of gene expression profiles in colonic tumor cells in which  $\beta$ -catenin-Tcf signaling is abrogated by expression of a dominant-negative Tcf-4 demonstrated up-regulation of gene markers that are characteristic of either the mucosecretory or the absorptive cell lineages in the intestinal mucosa,<sup>15–18</sup> consistent with our report that down-regulation of  $\beta$ -catenin-Tcf signaling accompanied, and was mechanistically linked to, colonic cell differentiation in tissue culture.<sup>12</sup> This therefore suggests that loss of normal cell differentiation patterns in the intestinal mucosa, possibly attributed to the

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**Figure 1.** The incidence (a), frequency (b), and size (c) of intestinal tumors in the *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>*, *Muc2<sup>-/-</sup>, p21<sup>+/-</sup>*, or *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice. **a:** \**P* = 0.025, in comparison to *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* mice by Fisher's exact test. **b:** \**P* = 0.02. **c:** \**P* = 0.03 and \*\**P* = 0.02 in comparison to *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* mice by Mann-Whitney test.

regulation of p21 expression through effects of  $\beta$ -catenin-Tcf signaling and *c-myc* expression, is a key event in tumor formation, and indeed, introduction of the targeted inactivation of p21 in the *Apc1638<sup>+/-</sup>* mouse, which increased tumor formation, decreased the number of mucin-expressing goblet cells in the intestinal mucosa.<sup>3</sup>

We also showed that genetic inactivation of the *Muc2* gene, which encodes the principal colonic mucin,<sup>19–23</sup> was sufficient to cause tumor formation.<sup>24</sup> However, unlike the loss of *Apc* function, this loss of *Muc2* expression was not accompanied by elevated  $\beta$ -catenin expression, or the relocalization of  $\beta$ -catenin to the nucleus.<sup>24</sup> Thus, there was no evidence that  $\beta$ -catenin-Tcf signaling was aberrant in this model, although *c-myc* expression was elevated in the tumors.<sup>24</sup> Therefore, although the mechanism of tumorigenesis in the *Muc2* and the *Apc* models may converge, the initial events seem to be distinct, and differ in their overt effects on  $\beta$ -catenin-Tcf signaling. The question therefore arises whether the targeted inactivation of p21 would also be effective in augmenting tumor formation in the *Muc2<sup>-/-</sup>* mouse model. Such data are essential for interpretation of the role that loss of p21 plays in the central pathways involved in intestinal tumorigenesis. We here report that introduction of a targeted inactivation of *p21* increased tumor formation in the small intestine, colon, and rectum, of mice with a targeted inactivation of *Muc2*, accompanied by increased cell proliferation, decreased apoptosis, and decreased differentiation in the intestinal mucosa, which was associated with down-regulation of p27kip and up-regulation of *c-myc*.

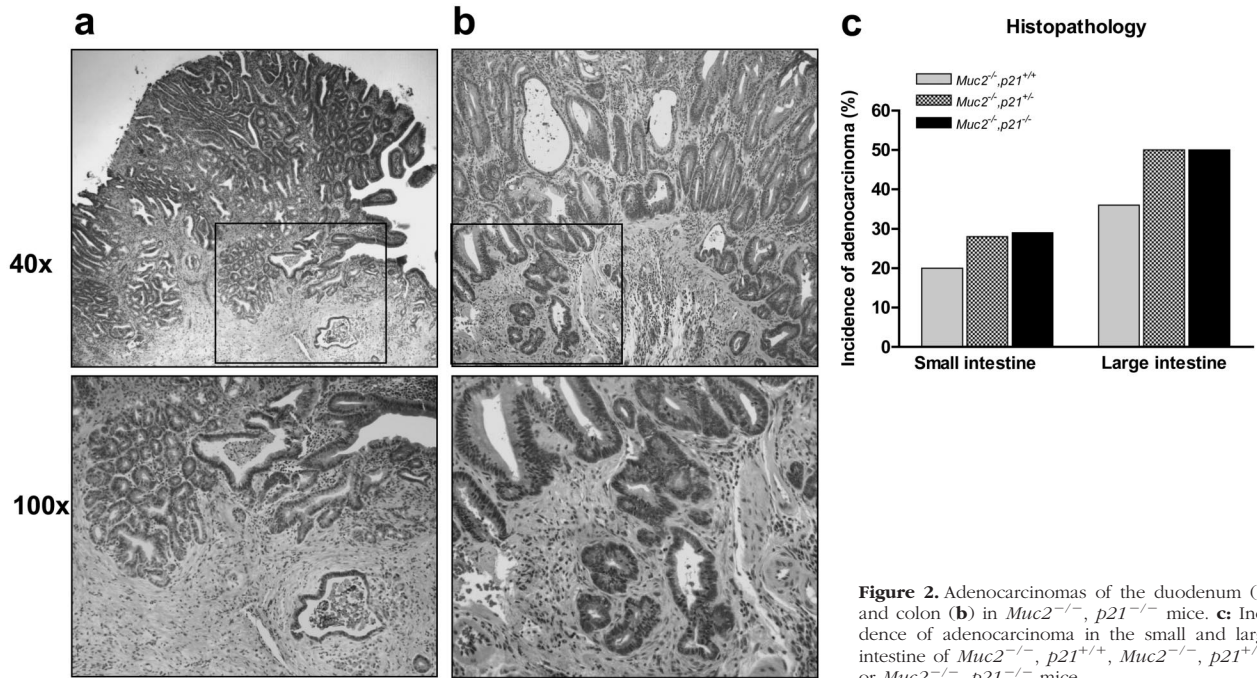
### Materials and Methods

The *p21* and *Muc2* mouse models, and methods for genotyping, have been reported.<sup>3,24</sup> *Muc2<sup>-/-</sup>* mice (mixed C57BL/6J and 129/SvOla background) were mated with *p21<sup>-/-</sup>* mice (mixed 129S6/SvEvTac and NIH Black Swiss background) to generate *Muc2<sup>+/-</sup>, p21<sup>+/-</sup>* offspring (F<sub>1</sub>). F<sub>1</sub> mice were mated to produce desired genotypes: *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>*, *Muc2<sup>-/-</sup>, p21<sup>+/-</sup>*, or *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>*. At weaning (~3 to 4 weeks), littermates were fed AIN-76A diet (Teklad, Madison, WI), *ad libitum*.

Mice were weighed weekly and maintained on diet for 36 weeks, or until they exhibited significant weight loss or other signs of extensive tumor formation. Mice were killed by CO<sub>2</sub> overdose and cervical dislocation, and then rapidly dissected for evaluation of tumors and fixation of tissues, as described previously.<sup>3,4,24,25</sup> Proliferation and apoptosis were evaluated by staining for proliferating cell nuclear antigen (Zymed, South San Francisco, CA) or terminal dUTP nick-end labeling (TUNEL) assay (Trevigen, Gaithersburg, MD), as described.<sup>3,26</sup>

Total RNA and protein were isolated from the frozen tissues using TRIzol reagent (Invitrogen Life Technology, Carlsbad, CA), as described.<sup>25</sup> The quantity of RNA and protein were measured spectrophotometrically. As we described,<sup>25</sup> cDNA was synthesized from DNase-treated total RNA using TaqMan Multiscribe reverse transcriptase (Applied Biosystems, Inc., Foster City, CA). Quantitative polymerase chain reaction analysis was done using the ABI Prism 7900-HT sequence detection system (96-well, Applied Biosystems, Inc., Foster City, CA). The primers for *p21*, *p27*, *c-myc*, and  $\beta$ -actin, the amplification conditions for the quantitative real-time polymerase chain reaction, and data analysis, were reported previously.<sup>25</sup>

Western blot analyses of steady-state levels of specific proteins were done by standard methods, as described,<sup>25</sup> using the following primary antibodies for detection: anti-p21, anti-p27, anti-*c-myc* (Santa Cruz Biotechnology, Santa Cruz, CA); and anti- $\beta$ -actin (Sigma, St. Louis, MO). Signal was detected by the enhanced chemiluminescence technique (Amersham Life Science, Piscataway, NJ). Immunohistochemical staining for intestinal trefoil factor (ITF) was previously reported in detail.<sup>24</sup> Briefly, 4- $\mu$ m formalin-fixed and paraffin-embedded sections were deparaffinized and rehydrated, quenched with 1.5% H<sub>2</sub>O<sub>2</sub>, blocked with 10% normal goat serum, and probed with rabbit anti-ITF polyclonal antibody (kindly provided by Catherine Tomasetto, Strasbourg, France). Detection was with biotinylated anti-rabbit IgG (Santa Cruz Biotechnology), followed by incubation with avidin-biotin complex (Vector Labs, Burlingame, CA) and the substrate 3',5'-diaminobenzidine, combined with hematoxylin counterstaining.



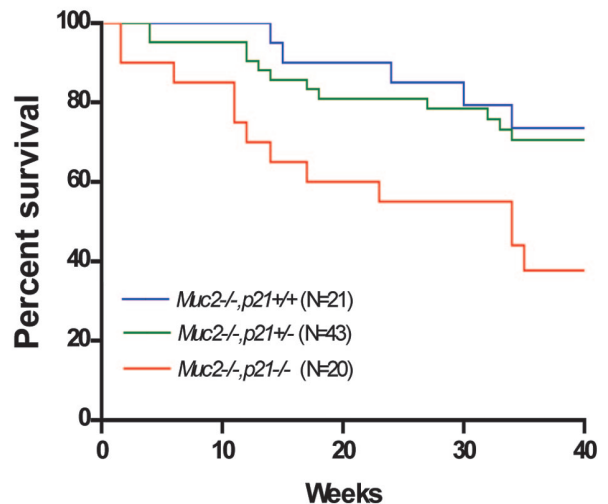
**Figure 2.** Adenocarcinomas of the duodenum (a) and colon (b) in *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice. c: Incidence of adenocarcinoma in the small and large intestine of *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup>, *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup>, or *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice.

## Results

Tumors developed throughout the intestinal tract in the *Muc2*<sup>-/-</sup>, *p21* wild-type mice. Small intestinal tumors developed in ~60% of the *Muc2*<sup>-/-</sup>, *p21* wild-type mice at a frequency of 1.0 tumor per mouse at an age of 36 weeks (Figure 1, a and b). This is similar to the incidence and frequency of tumors we previously reported for the *Muc2*<sup>-/-</sup> mice.<sup>24</sup> However, littermates that were *Muc2*<sup>-/-</sup> and either *p21*<sup>+/-</sup> or *p21*<sup>-/-</sup> had a significantly higher tumor incidence of 83% and 100% in small intestinal tumors, respectively ( $P = 0.025$ ) (Figure 1a). In addition, the small intestinal tumor frequency per mouse was increased by 30% in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> mice and by >80% in the *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice (Figure 1b) ( $P = 0.029$ ). The effect on tumor size was also striking: small intestinal tumors in *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> or *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> mice were 35% larger than the tumors in *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice ( $P = 0.02$  and  $P = 0.03$ , respectively, compared to the *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice) (Figure 1c). Most important, 28% of the small intestinal tumors in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> and *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice (8 of 29 and 8 of 28, respectively) were invasive adenocarcinomas (Figure 2a), more than the 20% in the *Muc2*<sup>-/-</sup>, *p21* wild-type mice (3 of 15) (Figure 2, a and c). Because *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice died earlier (Figure 3), the *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice that were sacrificed at 36 weeks, which provided the histopathological data, may overrepresent those with less aggressive phenotype, accounting for the similarity in the incidence, frequency, size, and pathology of these mice to the *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> mice.

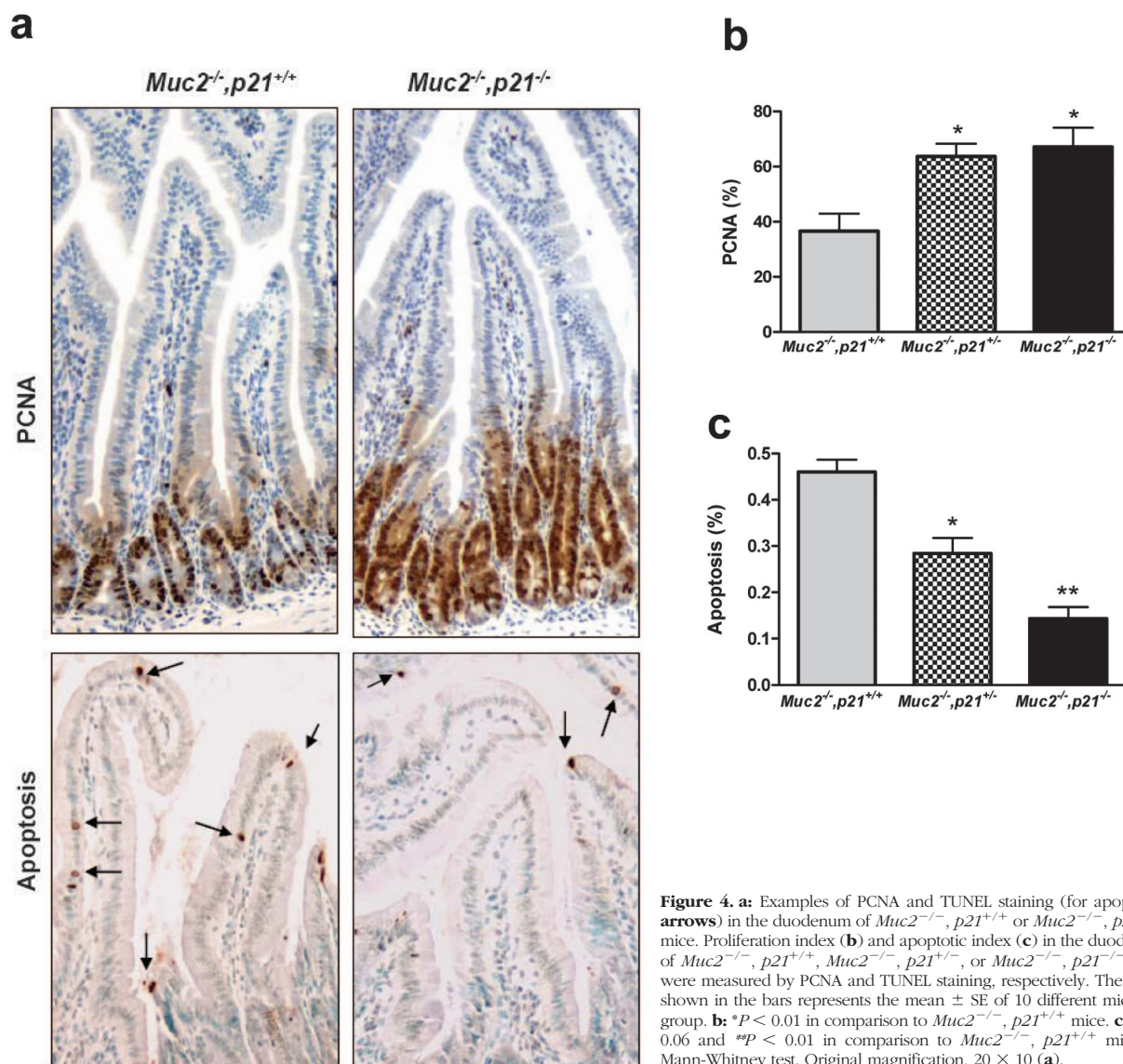
The inactivation of p21 also increased tumor formation in the colon and rectum of the mice. Although 53% of *Muc2*<sup>-/-</sup>, *p21* wild-type mice developed large intestinal tumors, in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> or *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice, this increased to 91% and 83%, respectively (Figure 1a),

and again the frequency was also significantly increased from 0.63 large intestinal tumors per mouse in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice to 1.04 and 1.00 in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> or *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice (Figure 1b). Analysis of histopathology revealed that 50% of the large intestinal tumors were early or advanced invasive adenocarcinomas (Figure 2b) in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> or *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice (13 of 26 and 8 of 16, respectively) (Figure 2c), whereas 36% (4 of 11) of large intestinal tumors in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice were invasive adenocarcinoma. More interesting, 33% (5/15) of *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice developed rectal tumors. The incidence of this pathology was somewhat higher in the p21 heterozygous mice (43%, 10 of 23), and increased further, to 60% (9 of 15), in the *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice. All of the rectal tumors were either adenomas



**Figure 3.** Survival of *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup>, *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup>, or *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice.





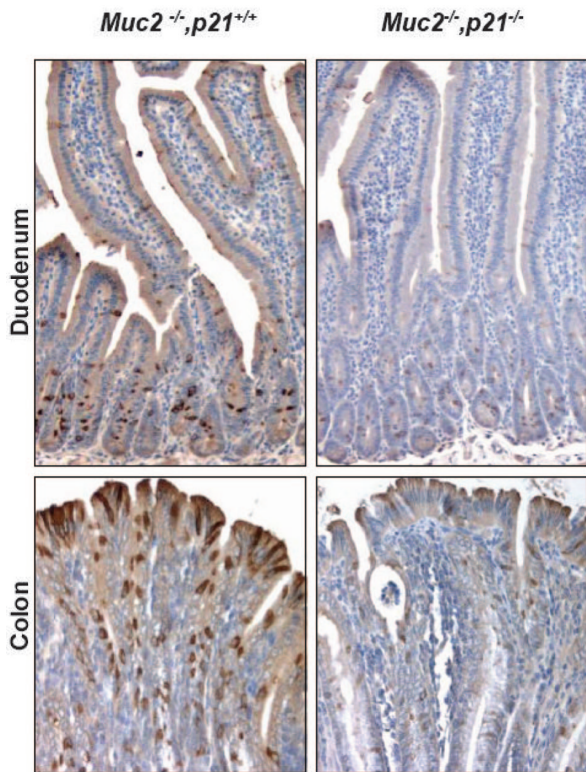
**Figure 4. a:** Examples of PCNA and TUNEL staining (for apoptosis, **arrows**) in the duodenum of *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> or *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice. Proliferation index (**b**) and apoptotic index (**c**) in the duodenum of *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup>, *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup>, or *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice were measured by PCNA and TUNEL staining, respectively. The value shown in the bars represents the mean  $\pm$  SE of 10 different mice per group. **b:** \**P* < 0.01 in comparison to *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice. **c:** \**P* = 0.06 and \*\**P* < 0.01 in comparison to *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice by Mann-Whitney test. Original magnification, 20  $\times$  10 (**a**).

or invasive adenocarcinomas. As for the small intestinal tumors, mice with less aggressive disease may be under-represented in the data because of early death of the *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice (Figure 3).

In the *Apc* mouse model of tumorigenesis, the increased tumor formation caused by the introduction of the targeted inactivation of p21 was accompanied by a disruption in homeostasis of the intestinal mucosa, demonstrated by increased proliferation, decreased apoptosis, and a decrease in recognizable goblet cells in the intestinal mucosa.<sup>3</sup> All of these changes were already present in the *Muc2*<sup>-/-</sup> model that was wild-type for p21.<sup>24</sup> However, because inactivation of p21 was effective in increasing tumor formation in the *Muc2*<sup>-/-</sup> mice, we investigated whether there were further perturbations in these three aspects of cell maturation.

Examples of the assays for proliferation [proliferating cell nuclear antigen (PCNA) staining] and apoptosis (TUNEL staining) are shown in Figure 4a. In the duodenum, the proliferation index (percent PCNA-positive cells)

of *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice was 38% (Figure 4b). Using the same assay, with mice wild-type at both loci and of similar age maintained on the same diet, we reported that the proliferation index was 19%.<sup>24</sup> The twofold increase is consistent with the increase we have reported for the *Muc2*<sup>-/-</sup> mice.<sup>24</sup> However, the introduction of one inactivated p21 allele increased the proliferation index even further to ~60% (Figure 4b; *P* < 0.01), and inactivation of the second p21 allele resulted in an increased proliferation index of ~70%. Similarly, the *Muc2*<sup>-/-</sup>, *p21*<sup>+/+</sup> mice had an apoptotic index in the duodenum of 0.45 (Figure 4c), lower than the 0.90 level that we reported for mice wild-type at both loci, and consistent with the reduced apoptosis we reported in the *Muc2*<sup>-/-</sup> mice.<sup>24</sup> Again, however, introduction of the targeted inactivation of p21 further reduced the apoptotic index, in this case more clearly in a p21 gene dose-dependent manner: by 38% in the *Muc2*<sup>-/-</sup>, *p21*<sup>+/-</sup> mice and by 60% in the *Muc2*<sup>-/-</sup>, *p21*<sup>-/-</sup> mice (Figure 4c, *P* = 0.06 and <0.01, respectively). The increased proliferation and decreased apo-



**Figure 5.** ITF expression in the duodenum and colon of *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* or *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice. Original magnification, 20 × 10.

ptosis were also seen in the flat mucosa of the colons from the *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* or *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice compared with *Muc2<sup>-/-</sup>* littermates wild-type for p21 (data not shown). Therefore, changes in proliferation and apoptosis were consistently associated with the increased tumor formation because of p21 inactivation.

The most interesting finding regarded differentiation of the goblet cell lineage. The intestinal mucosa of the *Muc2<sup>-/-</sup>* mouse is characterized by the absence of recognizable goblet cells.<sup>24</sup> However, this cell lineage is not completely ablated, because cells in the mucosa still stain immunohistochemically for ITF, another principal secreted product of goblet cells and component of intestinal mucus.<sup>27,28</sup> This can be clearly seen in the duodenum and colon of the *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* mice (Figure 5). However, on homozygous inactivation of the p21 gene, the goblet cell lineage is further perturbed as indicated by the significant reduction in ITF staining (Figure 5). Thus, the loss of p21 expression in the intestinal mucosa of the *Muc2<sup>-/-</sup>* mice, which accelerates and enhances tumor formation, alters cell maturation—increasing cell proliferation, decreasing cell differentiation and apoptosis—in the intestinal mucosa.

Gene expression was investigated in the mucosa of these mice. As expected, the levels of p21 mRNA decreased in a gene dosage-dependent manner in the *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* and *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice compared to the *Muc2<sup>-/-</sup>, p21* wild-type mice (Figure 6a), and this was also reflected in the protein levels of p21 (Figure 6d). We have previously shown that the inactivation of *p27<sup>kip1</sup>* in mice maintained on the AIN-76A diet was sufficient to

cause tumor formation (W. Yang, et al, manuscript in preparation).<sup>25</sup> It was therefore of interest that the inactivation of p21 in the *Muc2<sup>-/-</sup>* mice led to a concurrent reduction in p27 mRNA by 30% (Figure 6b) and protein expression by 50% (Figure 6, d and e).

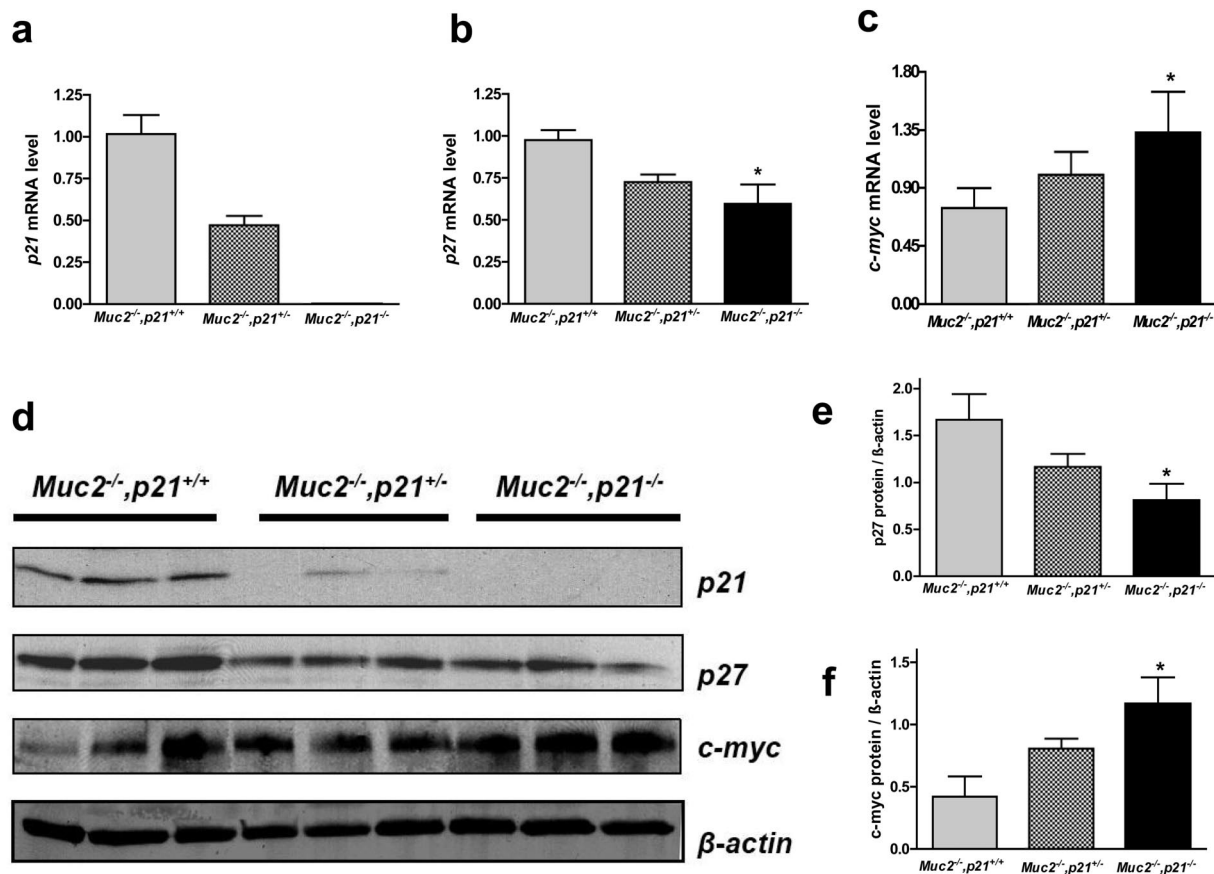
As regards the reduction in p27 expression, it may be specific for the compound knockout (ie, *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>*) mice, because preliminary data indicated that there is no reduction in p27 expression in the *Muc2<sup>-/-</sup>* mice compared to *Muc2<sup>+/+</sup>* mice when both are wild-type for p21 (N. Popova and A. Velcich, personal communication), and it has been reported that p27 expression is not reduced in *p21<sup>-/-</sup>* mice compared to wild-type mice.<sup>29–31</sup>

Finally, as outlined in the Introduction, p21 is under negative control of myc expression in the intestinal mucosa. We therefore investigated whether there is a feedback, and thus whether the decreased p21 expression alters *c-myc* expression. Figure 6c illustrates that *c-myc* mRNA levels indeed rose in the intestinal mucosa of the *Muc2<sup>-/-</sup>* mice in conjunction with the targeted inactivation of p21. This increase was ~60% in the *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice compared to the *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* mice ( $P < 0.05$ ) (Figure 6c). Further quantifying Western blot signals demonstrated that *c-myc* protein level in *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice was elevated by 1.78-fold greater than the *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* mice ( $P < 0.05$ ) (Figure 6, d and f).

## Discussion

There is considerable evidence that *p21<sup>WAF1/cip1</sup>* plays a fundamental role in pathways that regulate intestinal cell maturation and homeostasis of the intestinal mucosa.<sup>1,3,4,6,32</sup> In the intestinal tract, *p21* is expressed as cells exit the proliferative compartment, and loss of both expression and topological regulation is detected early in colon tumor formation.<sup>1,6</sup> Absence of *p21* is linked to inability of colon tumor cells to arrest in the G<sub>1</sub> phase of the cell cycle,<sup>13,33,34</sup> when stimulated by the nonsteroidal anti-inflammatory drug sulindac,<sup>4,34,35</sup> or by radiation.<sup>36</sup> However, despite this evidence for a key role of p21, its targeted inactivation does not grossly perturb the mucosa and does not cause tumor formation.<sup>5</sup>

Loss of p21 in mice that have an initiating mutation in *Apc* does cause marked enhancement of tumor formation.<sup>3</sup> This report extends this observation to another initiator of intestinal tumorigenesis: the inactivation of *Muc2*, the gene that encodes the major intestinal mucin. Inactivation of *Muc2* initiates tumor formation by a pathway that is distinct from that of loss of APC function, in that loss of *Muc2* expression does not involve elevation of  $\beta$ -catenin expression or accumulation in the nucleus, both of which are characteristic of APC-initiated tumors.<sup>24</sup> The difference in mechanism is confirmed by more recent evidence that combining the *Apc* and *Muc2* mutations in the mouse is synergistic in terms of tumor formation initiated by either mutation alone (A. Velcich, manuscript in preparation). Thus, the fact that inactivation of p21 enhances tumor formation in these two different



**Figure 6.** p21, p27 and *c-myc* expression in the colon of *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>*, *Muc2<sup>-/-</sup>, p21<sup>+/-</sup>*, or *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice. **a–c:** Relative level of mRNA of *p21* (**a**), *p27* (**b**), and *c-myc* (**c**) in the colon, assayed by quantitative real-time reverse transcriptase-polymerase chain reaction. The value shown in the bars represents the mean ± SE of five different mice per group. **d:** Protein expression of p21, p27, and *c-myc*, assayed by Western blot. The quantification of p27 (**e**) and *c-myc* (**f**) protein were normalized to β-actin. \**P* < 0.05 in comparison to *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* mice by Mann-Whitney test.

mouse models underlines the importance of p21 in intestinal homeostasis and tumorigenesis.

As in the *Apc1638<sup>+/-</sup>* model, the targeted inactivation of p21 causes a disruption in cell maturation pathways that includes further elevation in proliferation, depressed apoptosis, and perturbation of differentiation lineages. In the *Apc1638<sup>+/-</sup>* model, the perturbation of differentiation by inactivation of p21 was manifest as a decrease in the number of mucin-containing goblet cells.<sup>3</sup> However, in the *Muc2<sup>-/-</sup>* mouse, this cell phenotype is already not detectable in the intestinal mucosa, but the loss of p21 appears to further perturb development of the lineage because it results in decreased expression of ITF, another marker of this cell type, which instead persists in the *Muc2<sup>-/-</sup>, p21<sup>+/+</sup>* mice (Figure 5).<sup>24</sup> In this regard, it is of interest that in the *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice, there is down-regulation of another cdk inhibitor, *p27<sup>kip1</sup>*. Although the inactivation of p27 is able to initiate tumor formation in mice maintained on the AIN-76A diet,<sup>25</sup> like *Muc2<sup>-/-</sup>* mice, the *p27<sup>-/-</sup>* mice still express ITF, albeit assayed at the mRNA level, while the mucin-expressing goblet cells are reduced.<sup>25</sup> Therefore, it may be that in the *Muc2<sup>-/-</sup>, p21<sup>-/-</sup>* mice, the combined inactivation of p21, and down-regulation of p27, has a more extensive affect on the development of the goblet cell lineage than does

inactivation and/or down-regulation of only one of these cdk inhibitors.

In the intestinal mucosa, tumor initiation by loss of function of APC, either through mutation, deletion, or epigenetic events, is most often because of the effects of APC on β-catenin/Tcf signaling, and indeed, evidence that altered β-catenin-Tcf signaling is sufficient to initiate tumor formation is very strong.<sup>15,18,37–39</sup> One of the direct, key targets of β-catenin-Tcf signaling is *c-myc*. Brief inactivation of *c-myc* is sufficient to induce a sustained loss of the transformed phenotype.<sup>40</sup> The importance of *c-myc* is also supported by the fact that *Muc2<sup>-/-</sup>* initiated tumor formation, although not targeting β-catenin-Tcf signaling, still caused elevation of *c-myc* expression,<sup>24</sup> and that loss of p21, which increases tumor initiation and progression, further elevates *c-myc* expression (Figure 6). Thus, these data reinforce a key role for *c-myc* in intestinal tumor formation. This is consistent with our recent report that *c-myc* likely plays a key role in regulating the maturation pathway as cells migrate from the proliferative compartment toward the lumen in the intestinal mucosa.<sup>41</sup>

In summary, our data demonstrated that the loss of p21 in the *Muc2<sup>-/-</sup>* mouse model of intestinal tumor formation is linked to further perturbation in cellular mech-



anisms of intestinal homeostasis. This is coincident with alterations of expression of at least two molecules critical in the maturation of intestinal epithelial cells: down-regulation of p27 and up-regulation of *c-myc*. Each has multiple targets and pathways that it can modulate. The alterations of these targets and pathways by gene and proteomic profiling of cells as they migrate along the crypt-villus axis of the intestinal tract in mouse genetic models, with and without modulation of tumorigenesis by other loci and environmental factors, will be fundamental in defining the key events in establishing probability of tumor formation and progression.

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