Activation of retinoblastoma protein in mammary gland leads to ductal growth suppression, precocious differentiation, and adenocarcinoma

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The retinoblastoma (Rb) tumor suppressor controls cellular proliferation, survival, and differentiation and is functionally inactivated by mutations or hyperphosphorylation in most human cancers. Although activation of endogenous Rb is thought to provide an effective approach to suppress cell proliferation, long-term inhibition of apoptosis by active Rb may have detrimental consequences in vivo. To directly test these paradigms, we targeted phosphorylation-resistant constitutively active Rb alleles, Rb\DeltaKs, to the mouse mammary gland. Pubescent transgenic females displayed reduced ductal elongation and cell proliferation at the endbuds. Postpuberty transgenic mice exhibited precocious cellular differentiation and

β-casein expression and extended survival of the mammary epithelium with a moderate but specific effect on the expression of E2F1, IGF1Rα, and phospho–protein kinase B/AKT. Remarkably, ~30% RbΔK transgenic females developed focal hyperplastic nodules, and ~7% exhibited fullblown mammary adenocarcinomas within 15 mo. Expression of the RbΔK transgene in these mammary tumors was reduced greatly. Our results suggest that transient activation of Rb induces cancer by extending cell survival and that the dual effects of Rb on cell proliferation and apoptosis impose an inherent caveat to the use of the Rb pathway for long-term cancer therapy.

Introduction

The retinoblastoma (Rb)* tumor suppressor exerts diverse effects on cell growth and differentiation by modulating the activity of transcription factors such as members of the E2F protein family (Dyson, 1998). Rb itself is regulated at the phosphorylation level by G1 cyclins, their associated cyclin-dependent kinases (Cdks), and specific inhibitors (e.g., p16^{lnk4a}) (Sherr, 2000). During most of the G1 phase of the cell cycle, hypophosphorylated Rb interacts with E2F on cognate binding sites and actively represses transcription of

Key words: Rb; breast cancer; mammary gland; apoptosis; transgenic mice

© The Rockefeller University Press, 0021-9525/2002/1/185/14 \$5.00 The Journal of Cell Biology, Volume 156, Number 1, January 7, 2002 185–198 http://www.jcb.org/cgi/doi/10.1083/jcb.200106084

genes required for cell cycle progression and DNA replication. Active repression involves, at least in part, the recruitment by Rb of chromatin-modifying enzymes such as the histone deacetylase HDAC1 (Lai et al., 1999; Harbour and Dean, 2000a). Mitotic signals propagated through G1 cyclins induce sequential phosphorylation and inactivation of Rb (Lundberg and Weinberg, 1998). Early in G1, Cdk4/6-cyclin D1 phosphorylates Rb at phosphoacceptor sites just downstream of the Rb pocket domain, which is involved in protein-protein interaction (Harbour et al., 1999). Phosphorylation of these sites induces intramolecular interaction between the negatively charged phosphate groups and basic residues in the pocket. This intramolecular interaction induces a conformation change that expels HDAC1 from Rb, thereby relieving active transcriptional repression. Later in G1, Cdk2–cyclin E phosphorylates Rb at Serine-567, resulting in dissociation of Rb from E2F, transcriptional depression of target genes, and cell cycle progression.

In addition to keeping the cell cycle in check, Rb also controls cell differentiation and survival. There is evidence that Rb can bind a plethora of differentiation factors and cooper-

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^{*}Abbreviations used in this paper: Cdk, cyclin-dependent kinase; H-E, hematoxylin and eosin; IHC, immunohistochemistry; MMTV, mouse mammary tumor virus; PCNA, proliferation cell nuclear antigen; PKB, protein kinase B; Rb, retinoblastoma; RT, reverse transcribed; TEB, terminal endbud; TUNEL, TdT-mediated dUtp-biotin nick end labeling; WAP, whey acidic protein.

Figure 1. Targeted expression of constitutively active Rb to the mammary gland. (A) Schematic structure of Rb showing the pocket domain (A and B), relative location of 16 Cdk phosphoacceptor sites, and mutated residues in RbAp34 and $Rb\Delta K11$. (B) Schematic presentation of the MMTV-Rb Δ p34 and MMTV-Rb Δ K11 transgenes. (C) Detection of Rb Δ K transgene by Northern blot analysis. The transgene has a shorter 3' UTR and migrates faster. (D and F) Western blots of protein lysates (50 μ g) from mammary glands of (D) 13-wk-old or (F) 14-mo-old transgenic and wildtype females were developed with anti-HA monoclonal antibody. (E) Protein lysates (1.2 mg) from 14-mo-old transgenic and control mammary glands were immunoprecipitated with anti-HA antibodies and Western blotted with anti-Rb antibody. (G and H) RT-PCR analysis of the MMTV-RbΔp34#24 and MMTV-RbΔK11#29 transgenes relative to keratin 18 in virgin (V), pregnant (P), lactating (Lac), and involuting (Inv) glands. w, weeks. + and - signs indicate the presence or absence of reverse transcriptase.



atively induce terminal differentiation and transcriptional activation of differentiation genes in vitro (Macleod, 1999; DiCiommo et al., 2000). In addition, Rb mutant embryos exhibit incomplete differentiation in certain tissues and aberrant expression of specific markers, for example, NGF receptors during neurogenesis (Lee et al., 1994), muscle creatine kinase during myogenesis (Zacksenhaus et al., 1996b), and filensin/crystallin γ B during lens development (Liu and Zacksenhaus, 2000).

A role for Rb as a survival factor is evident from the massive cell death observed in Rb-deficient mice in tissues where Rb is normally highly expressed (Lee et al., 1994; Zacksenhaus et al., 1996b; Jiang et al., 1997; Macleod, 1999). Furthermore, in vitro, loss of Rb increases cell susceptibility to cytotoxic drugs, whereas ectopic expression of Rb inhibits cell death (Almasan et al., 1995; Haas-Kogan et al., 1995; Kranenburg et al., 1996). The dual effects of Rb on cell proliferation and apoptosis are mediated in some cell types by E2F1, which controls the expression of genes required for cell cycle progression and cell death. For example, deregulated E2F1 can induce the transcription of apoptosis protease-activating factor (Apaf)-1 (Moroni et al., 2001) and p14/19ARF, which binds and inactivates MDM2, a p53 inhibitor, thereby linking loss of Rb to p53-induced apoptosis (Sherr, 2000). Consistent with these in vitro studies, apoptosis in Rb-deficient mice is mediated in certain tissues by an E2F1–p53-Apaf-1–dependent pathway (Morgenbesser et al., 1994; Macleod et al., 1996; Tsai et al., 1998; Guo et al., 2001).

Although germ-line mutations in Rb predispose individuals to retinoblastoma, somatic mutations in Rb are often associated with cancer progression (DiCiommo et al., 2000). However, in many types of cancer, such as breast cancer (Buckley et al., 1993), other components of the Rb pathway (i.e., Cdk4, cyclin D1, or p16^{lnk4a}) are mutated preferentially or deregulated (Weinberg, 1995). In such tumors, Rb becomes hyperphosphorylated and inactive, but the protein is intact and amenable to therapeutical activation. Reactivation of endogenous Rb in such tumors (e.g., by Cdk inhibitors) may provide an effective approach to suppress cell proliferation. In support of this notion, introduction of phosphorylation-resistant constitutively active Rb alleles, Rb Δ Ks, into many cell types in vitro (Mittnacht, 1998) and in vivo (Chang et al., 1995) efficiently suppresses cell proliferation. However, the inherent ability of Rb to suppress both cell division and apoptosis points to a potential caveat. Activation of Rb may inadvertently protect cells, which have acquired oncogenic alterations or are programmed to die in response to normal developmental signals, from apoptosis and lead to detrimental consequences in vivo. Of relevance here is the observation that both overexpression and inactivation of E2F1 induce cancer in mouse models presumably by disrupting different aspects (proliferation versus apoptosis, respectively) of cell physiology (Field et al., 1996; Yamasaki et al., 1996; Pan et al., 1998; Pierce et al., 1998, 1999).

To directly address the consequences of activating Rb in vivo, we have targeted Rb Δ K alleles to the mammary gland of transgenic mice. We show that transgenic females initially display ductal growth suppression but later exhibit precocious differentiation, extended survival of the mammary epithelium, and ultimately develop hyperplastic lesions and mammary adenocarcinomas. The implications of these results to cancer therapy are discussed.

Results

Targeted expression of phosphorylation-resistant Rb alleles in the mammary gland

The interaction of Rb with transcription factors is dictated by cyclin D– and cyclin E–associated kinases that phosphorylate Rb in a cell cycle–dependent manner (Lundberg and Weinberg, 1998; Mittnacht, 1998). Rb contains 16 potential phosphoacceptor sites that are distributed along the protein (Lundberg and Weinberg, 1998; Mittnacht, 1998) (Fig. 1 A). Recent evidence indicates that a cluster of serine and threonine in exon 23 of Rb just downstream of the pocket plays a major role in regulating Rb activity (Harbour et al., 1999). In our study, we used two Rb Δ K alleles: Rb Δ p34 contains mutations in eight Cdk sites, four of which are within exon 23 (Hamel et al., 1992) (Fig. 1 A). Using RbAp34 as a backbone, we generated three additional serine/threonine to alanine substitutions in exon 23, yielding Rb Δ K11, in which all phosphoacceptor sites in exon 23 are protected completely from Cdk-mediated phosphorylation (see Materials and methods) (Fig. 1 A). Due to their resistance to phosphorylation by Cdks, the RbAKs outperform wild-type Rb in multiple assays and are viewed as constitutively active, underphosphorylated, native Rb (Hamel et al., 1992; Geng et al., 1996; Knudsen et al., 1998; Sellers et al., 1998; Brown et al., 1999; Harbour et al., 1999). To determine the effects of constitutive activation of Rb on mammary gland development and cancer progression, the two $Rb\Delta K$ alleles were targeted to the mammary gland under control of the mouse mammary tumor virus (MMTV) long terminal repeat (Matsui et al., 1990; Fig. 1 B). Four independent MMTV-Rb Δ p34 transgenic lines in mixed C57BL/6×SJL background and three MMTV- $Rb\Delta K11$ lines in pure FVB background were established (Fig. 1 and Table I).

Transgenic transcripts were found in the mammary gland and the salivary gland and spleen but not in the brain, kidney, and liver (Fig. 1, C, G, and H; data not shown). The HA-tagged Rb Δ K transgenic protein was readily detected by immunoprecipitations and Western blots with antibodies specific to the HA epitope (Fig. 1, D–F). However, we have been unable so far to detect the transgenes by immunohistochemistry (IHC) with our anti-HA antibodies. Reverse transcribed (RT)-PCR analysis showed that the MMTV-Rb Δ p34 and MMTV-Rb Δ K11 transgenes (collectively re-

Table I. Incidence of focal hyperplastic nodules and adenocarcinomas in transgenic mice expressing constitutively active Rb in the mammary gland

Transgenic line	Number of mammary glands analyzed ^b	Number of mammary glands with tumors ^c	Incidence per line	Incidence combined
			%	
MMTV-Rb∆p34(K8)				
#24	38	8^{d}	21	
#16	7	4	57	
#8	16	1	6.25	
#27	2	1 ^d	50	
Total	63	14		22
Control littermates ^a	50	0	0	0
MMTV-Rb∆K11				
#29	18	4 ^{3d}	22	22
Control littermates ^a	7	0	0	0
WAP-Rb∆p34(K8)				
# 9	7	2	28.6	
#44	5	4^{d}	80	
#61	9	2 ^d	22.2	
Total	21	8		38
Control littermates ^a	21	0	0	0

^aControl littermates were caged together with transgenic females and analyzed in parallel.

^bFocal hyperplastic nodules were detected by whole-mount staining of the right inguinal mammary glands of 10–15-mo-old females. Palpable adenocarcinomas were confirmed by histopathology.

Numbers include both hyperplastic nodules and adenocarcinomas.

^dIndicates one full-blown mammary adenocarcinoma in this group.



Figure 2. Temporal expression of endogenous Rb during mammogenesis in wild-type females. (A–F) RNA In situ hybridization analysis of endogenous Rb. Shown are bright field (A, C, and E) and dark field (B, D, and F) micrographs of representative sections. (G–I) IHC of endogenous Rb in wild-type mammary glands. Arrows indicate Rb-positive nuclei.

ferred to as MMTV-Rb Δ K) were expressed throughout mammary gland development, pregnancy, lactation, and involution (Fig. 1, G and H).

Regulated expression of endogenous Rb during mammogenesis in wild-type mice

To relate the expression of the Rb Δ K transgenes to endogenous Rb, we performed in situ hybridization and IHC analyses on wild-type mammary glands (Fig. 2). Both methods of detection revealed that endogenous Rb expression was below detection level or sporadic in nulliparous females, induced at mid-pregnancy, and peaked at lactation and involution (Fig. 2). Thus, compared with endogenous Rb the Rb Δ K transgenes are expressed at a relatively high level in nulliparous females (Fig. 1, G and H). Accordingly, we observed the strongest phenotypes in nulliparous MMTV-Rb Δ K transgenic mice as described below.

MMTV-Rb∆K transgenes suppress ductal growth in pubescent females

During the ductal stage, the mammary epithelium infiltrates the fat pad by proliferation at the terminal endbuds (TEBs) and dichotomous side branching (Medina, 1996; Hennighausen and Robinson, 1998; Cardiff et al., 2000; Silberstein, 2001). Whole-mount analysis of pubescent MMTV-Rb Δ K transgenic females revealed a moderate delay in TEB progression (Fig. 3, A–D). The suppression of ductal growth coincided with reduced expression of proliferation cell nuclear antigen (PCNA), a marker for cell proliferation (Fig. 3, F–I). On average, MMTV-Rb Δ K TEBs exhibited three- to fourfold less PCNA-positive nuclei compared with control wild-type littermates (Fig. 3 J). The suppression of ductal growth was most evident at 5–7 wk of age and reduced thereafter (Fig. 3 E). Accordingly, there were no obvious histological differences between the mammary glands of transgenic and wild-type littermates during pregnancy and lactation, and transgenic females were able to nurse their young and raise normal size litters (unpublished data).

MMTV-Rb∆K transgenes induce alveolar outgrowth and precocious differentiation in nulliparous females

After puberty, the mammary epithelium undergoes cyclic proliferation, limited differentiation, and cell death during the estrous cycle (Robinson et al., 1995, 2000). Interestingly, we found that fully developed nulliparous MMTV-RbAK transgenic females exhibited consistent alveolar outgrowth relative to control littermates (Fig. 4, A and B). PCNA-staining revealed very few cycling cells in both wildtype and transgenic mice (Fig. 4, C and D), indicating that enlargement of the alveolar compartment was not due to overt cellular proliferation. To test whether the alveolar outgrowth was indicative of precocious differentiation of the mammary epithelium, we determined the expression of β -casein, a milk protein. Remarkably, β -casein transcripts were detected by RT-PCR in \sim 28% (10 of 35) of 9–20-wkold nulliparous transgenic females but only rarely and at lower levels in wild-type littermates (Fig. 4 Q). Furthermore, IHC and Western blot analysis revealed that this milk protein was expressed in some nulliparous transgenic but in none of the wild-type females (Fig. 4, E, F, and R).

The alveolar outgrowth and precocious expression of β -casein became significantly more pronounced in 12–14mo-old nulliparous MMTV-Rb Δ K mice. Some transgenic females exhibited lactating-like glands that appeared distended with milk (unpublished data). Western blot analysis revealed a high level of β -casein in these glands (Fig. 4 R). Upon histological examination, some Rb Δ K transgenic females exhibited hyperplastic alveoli, dilated ducts, and enlarged lobules that resembled secretory lobuloalveoli in pregnant females with characteristic vacuoles and acinar morphology (Fig. 4, G–M).



Figure 3. **Suppression of ductal growth in pubescent MMTV-RbΔK transgenic mice.** Whole-mount staining of 5- and 7-wk-old pubescent MMTV-RbΔp34#27 (A and B) and MMTV-RbΔK11#29 (C and D) transgenic and control females. LN, lymph node. (E) The distance from the endbuds to the lymph nodes in 7-wk-old wild-type females was set at 100%. The average growth rates of endbuds in three transgenic females and three wild-type littermates were plotted against this value for each time point. PCNA staining of sections through endbuds of 6-wk-old MMTV-RbΔp34#24 (F and G) and MMTV-RbΔK11#29 (H and I) transgenic females and control

IHC analysis demonstrated high levels of β -casein expression in the mammary gland of four out of five transgenic females but not in control littermates (Fig. 4, N–P). RT-PCR analysis of another group of three 18-mo-old nulliparous transgenic and three control females revealed β -casein expression exclusively in the transgenic females; whey acidic protein (WAP), a later differentiation marker, was detected in two of the three transgenic mice (unpublished data). Thus, persistent expression of the Rb Δ K transgenes in nulliparous transgenic females induces precocious differentiation of the mammary epithelium both at the molecular and cellular levels.

MMTV-Rb∆K transgenic females develop hyperplastic nodules

Intriguingly, whole-mount analysis of 10–15-mo-old MMTV-Rb Δ K transgenic females revealed the presence of focal hyperplastic nodules (Fig. 5). These nodules appeared in 22% of the transgenic females (18 out of 81) from independent MMTV-Rb Δ p34 and MMTV-Rb Δ K11 lines (Table I). Both nulliparous and multiparous transgenic females developed these lesions, indicating that pregnancy was not required for neoplastic transformation. In many cases there were multifocal lesions per gland (Fig. 5, B, E, and F), suggesting independent stochastic transformation of the mammary epithelium. The incidence of hyperplastic nodules may be underestimated because only one mammary gland per animal (of the ten glands in a female mouse) was subject to whole-mount analysis. No such lesions were observed in over 57 wild-type littermate females (Fig. 5, A and C; Table I).

WAP-Rb∆p34 transgenic females also develop hyperplastic nodules

To further corroborate these observations, we targeted $Rb\Delta p34$ to the mammary gland under control of the WAP promoter/3' UTR (Fig. 6 A). The WAP regulatory unit directs linked transgenes exclusively to differentiated mammary epithelial cells during the estrous cycle and pregnancy, and WAP transgenes are expressed and exert phenotypic changes in both nulliparous and multiparous females (Jhappan et al., 1993; Sympson et al., 1994; Robinson et al., 1995). Three WAP- $Rb\Delta p34$ transgenic lines were established in C57BL/ $6 \times$ SJL background. Expression of the WAP-Rb Δ p34 transgenes was low in all three lines, and transcripts could only be detected during pregnancy (Fig. 6 B). Further analysis revealed normal ductal elongation, pregnancy, and lactation in WAP-Rb Δ p34 transgenic females (unpublished data). However, remarkably 38% (8 of 21) of the transgenic females from the three independent WAP-Rb Δ p34 lines developed focal hyperplastic nodules within 10-15 mo (Fig. 6, D and E; Table I). As with the MMTV-Rb Δ K lines, both nulliparous and parous females developed these micro lesions. No similar lesions were found in >21 wild-type littermate fe-

wild-type littermates. Arrows indicate PCNA-positive nuclei. (J) Quantitative analysis of the relative proliferation (PCNA-positive nuclei) in TEBs at the indicated stages. For each time point, two sections each from three transgenic and three wild-type mice were analyzed. The average number of PCNA-positive nuclei in the control wild-type females for MMTV-Rb Δ K11 was set at 100%. Bars in E and J indicate standard deviation.

Figure 4. Precocious differentiation of the mammary epithelium in nulliparous MMTV-RbAK transgenic females. Whole-mount analyses of 13-wk-old nulliparous transgenic and control females (A and B); PCNA staining (C and D); IHC for β-casein (E and F). Whole-mount hematoxylin staining of 14-mo-old control (G) and two nulliparous transgenic (H and I) females $(40 \times)$. Note the dilated ducts (arrow). (J) Higher magnification $(70\times)$ of the mammary gland in I, showing a lobuloalveolar-like structure (arrow). (K and M) H-E staining of mammary glands of 14-mo-old transgenic (M) and wild-type (K) littermate female. (L) H-E staining of a mammary gland from a 17.5-d-old pregnant wild-type female (P17.5). Note the similarity between nulliparous transgenic (M) and pregnant wild-type glands (L). (N-P) IHC of β-casein in mammary glands of 14-mo-old nulliparous wild-type (N) and two transgenic (O and P) females. (Q) RT-PCR analysis of β-casein expression in 9–20-wk-old nulliparous transgenic and control females. (R) Western blot analysis of β-casein expression in transgenic (TG) and wildtype (WT) females.



males (Fig. 6 C; Table I). Thus, targeted activation of Rb by two independent promoters, MMTV and WAP, induces hyperplastic lesions in the mammary epithelium.

MMTV-Rb Δ K and WAP-Rb Δ p34 transgenic females develop full-blown mammary adenocarcinomas

A fraction (7 in 102 is \sim 7%) of the MMTV-Rb Δ p34, MMTV-Rb Δ K11, and WAP-Rb Δ p34 transgenic females developed visible masses of 1–2 cm in diameter in their mammary glands after 10–15 mo (Table I). Pathology analysis revealed that these masses represented full-blown mammary adenocarcinomas (Fig. 7; see legend for pathology). None of

>100 wild-type littermate females in our colony developed mammary tumors (Table I). The incidence of tumors was highest in the MMTV-Rb Δ K11#29 line (Table I and unpublished data). We have analyzed recently two retired MMTV-Rb Δ K11#29 transgenic and two wild-type breeders at 20 mo of age (not included in Table I). Both transgenic females but none of the control mice developed palpable tumors in several mammary glands. Histological analysis revealed that each of these mammary glands contained one to four tumors. Fig. 7 G shows one such gland with a small tumor (Fig. 7 G, letter H) and several additional lesions (Fig. 7 G, letter I–K). Taken together, the results indicate that some



Figure 5. Hyperplastic nodules in the mammary gland of MMTV-Rb Δ K transgenic mice. Low power views (8×) of wholemount staining of #4 right inguinal mammary glands from 13.5-mo-old nulliparous wild-type female (A) and MMTV-Rb Δ p34#16 transgenic littermate (B). Arrows point to hyperplastic nodules. (C and D) 13.5-mo-old multiparous wild-type and MMTV-Rb Δ K11#29 transgenic littermate gland with a hyperplastic nodule. (E and F) High power views of representative hyperplastic nodules (arrows) in independent MMTV-Rb Δ p34 transgenic lines.

hyperplastic nodules in $Rb\Delta K$ transgenic mice progress into full-fledged mammary adenocarcinomas.

Transgene expression is lost in mammary adenocarcinomas from MMTV-Rb∆K transgenic mice

IHC analysis for PCNA revealed a relatively high proliferation index in the mammary adenocarcinomas derived from MMTV-Rb Δ K and WAP-Rb Δ p34 mice (Fig. 7, L–M). This observation has prompted us to determine whether expression of Rb Δ K was lost during the progression of these tumors. To this end, we analyzed the expression of the Rb Δ K transgene in two tumor biopsies and tumor cells derived from the mammary adenocarcinoma shown in Fig. 7 F and successfully propagated in vitro. Remarkably, Rb Δ K transcripts were virtually lost in both the tumor biopsies and the cultured tumor cells (Fig. 7, O and P). These results suggest that downregulation of the transgene with its inhibitory effect on cell proliferation may precede tumor progression in MMTV-Rb Δ K transgenic mice.

MMTV-Rb∆K transgenes extend the survival of differentiated mammary epithelial cells after single and multiple pregnancies

To directly test the effect of $Rb\Delta K$ on cell survival, we examined the first postlactational involution in MMTV- $Rb\Delta K$ transgenic mice. During this process, the extracellular matrix



Figure 6. **WAP-Rb** Δ **p34 transgenic mice also develop hyperplastic nodules.** (A) Schematic structure of the WAP-Rb Δ p34 transgene. (B) RT-PCR analysis of WAP-Rb Δ p34 expression in an 18.5-d-old pregnant transgenic female. (C and D) Whole-mount analysis of 12-mo-old nulliparous wild-type (C) and WAP-Rb Δ p34#44 transgenic (D) glands (8×). (E) Representative high power view of hyperplastic nodules in a WAP-Rb Δ p34#61 transgenic female (40×). (F) H-E staining of cross-section through a hyperplastic nodule. Arrows in D and E point to hyperplastic nodules.

is degraded and the mammary epithelium undergoes massive apoptosis and remodeling back to a virgin-like state. After 6 d, most of the differentiated epithelial cells are eliminated, and the expression of milk genes is greatly reduced (Strange et al., 1992; Lund et al., 1996). At this stage, there were remarkably more β -casein– expressing lobuloalveoli in MMTV-Rb Δ K11 females compared with wild-type littermates (Fig. 8 D; unpublished data). RT-PCR and Western blot analyses demonstrated a modest yet consistent elevation in the expression of WAP and β-casein in the transgenic MMTV-RbΔK11 females relative to control littermates (Fig. 8, F and G). In situ apoptosis by the TdT-mediated dUtp-biotin nick end labeling (TUNEL) assay revealed that collapsed lobules that did not express β -casein exhibited some cell death (Fig. 8, A and B), whereas lobules that maintained β-casein expression contained very few apoptotic nuclei (Fig. 8, C and D). Overall, there was a 44% reduction in apoptosis in the involuting glands of transgenic females compared with wild-type littermates (Fig. 8 E). Combined, the results suggest that the persistent expression of milk genes at day 6 of involution was due to inhibition of lobuloalveolar cell death by activated Rb.

Although Rb Δ K exerted only a moderate suppression of apoptosis during the first involution, we asked whether it might have a compounded effect after multiple pregnancies. To address this possibility, we performed serial cross-section analysis on multiparous, retired, transgenic MMTV-Rb Δ K breeders. Interestingly, nonregressing areas with enlarged and aberrant lobuloalveolar-like structures were detected in each of five transgenic females (Fig. 8, I and K) but not in five wild-type control



Figure 7. **Mammary adenocarcinomas in MMTV-Rb** Δ K and WAP-Rb Δ p34 transgenic mice and loss of transgene expression. Histology of mammary adenocarcinomas. (A) Whole-mount staining of a palpable mammary adenocarcinoma in a 13.5-mo-old nulliparous MMTV-Rb Δ p34#27 transgenic female, showing the periphery of a tumor with invasion into the ducts. (B) Mammary adenocarcinoma with multiple cystic areas in a 12-mo-old multiparous MMTV-Rb Δ p34#24 transgenic female. (C) Well-differentiated tubulo-acinar mammary adenocarcinoma in a 12-mo-old nulliparous WAP-Rb Δ p34#44 transgenic female. (D) Well-differentiated papillary mammary adenocarcinoma with peripheral growth of basaloid cells and radiating keratinization in a 14-mo-old multiparous WAP-Rb Δ p34#61 transgenic female. (E and F) Papillary mammary adenocarcinoma and mammary adenocarcinoma in the first and third glands, respectively, from a multiparous 18-mo-old MMTV-Rb Δ K11#29 transgenic female. (G) Multiple lesions in an MMTV-Rb Δ K11#29 reired breeder. (H and K) High power magnifications of the lesions shown in (G). H–J are mammary adenocarcinomas; K is an enlarged lobuloalveolar-like structure. Tumors in A–F were ~1–2 cm in length and significantly larger than H. Analysis of mammary adenocarcinomas. (L–M) PCNA staining demonstrating high mitotic index (M, arrows) in a tumor relative to a tumor-free gland in the same transgenic female (L). (N) Positive staining of mammary adenocarcinomas with antikeratin antibody, an epithelial marker. (O and P) RT-PCR analysis of transgene (O) and keratin 18 (P) expression in tumor biopsies and a derived cell line. Adenocarcinoma biopsies from the tumors shown in E and F (1 and 2); mammary gland from a MMTV-Rb Δ K11 transgenic female (3); cell line derived from the mammary adenocarcinoma in F (4); mammary gland from a wild-type female (5). The + and – signs indicate the presence or absence of reverse transcriptase.

breeders (Fig. 8, H and J). These lobuloalveolar-like structures expressed extracellular matrix as revealed by Gomori-trichrome, a specific stain for collagen (Fig. 8, L and M), and were often observed in multiparous transgenic females alongside hyperplastic nodules and mammary adenocarcinomas (Fig. 7 K).

Aberrant expression of survival factors in multiparous MMTV-Rb∆K transgenic breeders

We next determined whether the expression pattern of known survival factors was altered in multiparous MMTV-

Rb Δ K transgenic mice. Western blot analyses of whole mammary glands revealed that E2F1 was reduced moderately; IGFR-1R α and phospho–protein kinase B (PKB)/ AKT were elevated slightly; p21^{waf1/cip1} and BCl-XI exhibited additional faster migrating forms of unknown functions (Fig. 8, N and P). Expression of cyclin D1 and BCl2 was not obviously affected (Fig. 8 N; unpublished data). The aberrant expression of E2F1, IGFR-1R α , phospho-PKB/AKT, p21^{waf1/cip1}, and BCl-XL was specific to the multiparous transgenic females and was not observed in 4-mo-old nullip-



Figure 8. **Extended survival of the mammary epithelium in MMTV-Rb** Δ K transgenic mice. (A–G) Effects of activated Rb on day 6 of the first postlactational involution. TUNEL analysis (A and C) and β -casein IHC (B and D) were performed on adjacent sections (A and B and C and D) from MMTV-Rb Δ K11#29 transgenic females at day 6 of first involution. Note the inverse correlation between the presence of apoptotic nuclei (A, black arrows) and the expression of β -casein (D, white arrows) in the same lobules. (E) Relative apoptosis was determined by counting TUNEL-positive nuclei in multiple random areas under 400× magnification. (F and G) RT-PCR and Western blot analyses of milk gene expression in transgenic and wild-type littermate females at day 6 of first involution. The level of milk genes was slightly higher in the transgenic females. (H and M) Enlarged lobuloalveolar-like structures in multiparous retired MMTV-Rb Δ K11#29 transgenic breeders but not in control littermates. (H–K) H-E staining (H and I, 10×; J and K, 40×). (L and M) Gomori trichrome staining for collagen (L, 126×; M, 63×). ECM is present in transgenic (black arrow) but not in wild-type females (white arrow). (N–P) Western blot analyses of the indicated proteins in multiparous retired breeders (N and P) and 4-mo-old nulliparous females (O).

arous transgenic mice (Fig. 8 O). Taken together, the results presented in Fig. 8 suggest that $Rb\Delta K$ suppress mammary epithelial cell death at least in part by modulating the level or activity of E2F1, IGFR-1R α , and PKB/AKT.

Discussion

We report that targeted expression of phosphorylation-resistant constitutively active Rb proteins in the mammary gland leads to ductal growth suppression, precocious differentiation, extended survival of the mammary epithelium, and paradoxically to mammary tumors. The growth suppression is moderate, affects ductal but not alveolar development, and represents the only phenotype that is consistent with the role of Rb as a tumor suppressor. The precocious differentiation and mammary tumors were less expected and are discussed below.

Mechanisms of precocious differentiation in MMTV-Rb∆K transgenic mice

Although the role of Rb in cell proliferation and apoptosis is well established, it is not clear whether Rb affects terminal differentiation directly through interaction with differentiation factors or indirectly by allowing proper cell cycle exit and cell survival (Dyson, 1998; Macleod, 1999; Harbour and Dean, 2000b). In this study, we show for the first time that active Rb can induce cellular and molecular differentiation in vivo. Precocious differentiation of the mammary



ectopic cell proliferation and apoptosis that perturb the differentiation program. Alternatively, Rb may be required for the onset of terminal differentiation via direct or indirect stimulation of differentiation factors (e.g., glucocorticoid receptor, C/EBP, MyoD, etc.). (B) As shown in this study, activation of Rb in the mammary gland results in suppression of cell proliferation, precocious differentiation, extended survival of the mammary epithelium, and mammary tumors. The precocious differentiation may reflect the ability of activate Rb to extend the survival of differentiated cells and/or to directly promote differentiation. We propose that inhibition of apoptosis by active Rb can lead to the accumulation of oncogenic alterations and neoplastic transformation in susceptible tissues such as the mammary epithelium.

gland has also been described in transgenic mice overexpressing activated β -catenine (Imbert et al., 2001), stromolysin (Sympson et al., 1994), dominant negative TGF β R (Gorska et al., 1998), E-cadherin (Delmas et al., 1999), and in mice lacking P-cadherin (Radice et al., 1997).

The coinduction of Rb and β -casein at mid-pregnancy (Fig. 2; unpublished data) suggests that Rb may be involved directly in transcriptional activation of milk genes (Fig. 9 B). This model is consistent with numerous examples, demonstrating a required role for Rb in terminal differentiation of certain cell types and transcriptional activation by MyoD (Gu et al., 1993; Sellers et al., 1998; Puri et al., 2001), C/EBP (Doppler et al., 1995), the glucocorticoid receptor (Stocklin et al., 1996), and other factors. From a cancer treatment point of view, the possibility that activation of Rb can induce differentiation is attractive, since it may allow the development of drugs that would propel breast tumor cells not only to exit the cell cycle but also to terminally differentiate. However, although the lack of Rb can be a limiting factor for differentiation, activation of the Rb pathway may not be sufficient to force terminal differentiation. Indeed, we have not been able so far to detect any effect of overexpressing Rb Δ Ks on the β-casein promoter in transient reporter assays in Rb-positive mammary epithelial HC11 cells (unpublished data).

In an alternative model, the Rb Δ K transgene may induce precocious development by extending the life span of differentiated mammary epithelial cells with each estrous cycle, hence leading to the accumulation of differentiated mammary epithelial cells and milk secreting lobuloalveoli. This "survival model" is consistent with other phenotypes observed in our transgenic mice, most notably the development of mammary tumors in the MMTV-Rb Δ K and WAP-Rb Δ p34 mice. We note that the two models are not mutually exclusive, and activated Rb may promote both the onset and survival of the differentiation state (Fig. 9 B).

Mechanisms of mammary adenocarcinoma in MMTV/WAP-Rb Δ K transgenic mice

Paradoxically, we found that transgene expression of constitutively active Rb in the mammary gland induces mammary adenocarcinoma. We propose that these tumors may be induced in three steps. First, growth suppression by $Rb\Delta K$ may exert a selection pressure in favor of transforming mutations that accelerate cell proliferation. Second, the suppression of cell death by the Rb Δ K transgenes may promote the survival and accumulation of transformed cells that would otherwise die by apoptosis. Third, notwithstanding the limited number of tumors analyzed so far (Fig. 7 O), our results suggest that downregulation of transgene expression may permit rapid clonal expansion of transformed cells and the development of full-blown mammary adenocarcinomas. Thus, in accordance with the "survival model" described above suppression of apoptosis by active Rb may extend the life span of mammary epithelial cells that have partly differentiated during the estrous cycle, leading to precocious differentiation, or have acquired transforming mutations, resulting in mammary adenocarcinoma (Fig. 9 B). There is a reciprocal relationship between the phenotypes of Rb knockout mice and E2F1 transgenic mice on the one hand (Fig. 9 A) and the Rb Δ K transgenic mice and E2F1-null mice on the other hand (Fig. 9 B) (Field et al., 1996; Yamasaki et al., 1996; Pierce et al., 1998, 1999). This symmetry is consistent with the antagonistic effects of Rb and E2F1 on cell cycle progression and apoptosis so that both activation and inactivation of these factors induce cancer in susceptible tissues (Fig. 9).

We note that transgenic expression of the survival factor Bcl2 in the mammary gland is not sufficient to induce mammary adenocarcinoma, although it accelerates tumorigenicity in double transgenic mice expressing Bcl2 and Myc (Jager et al., 1997). Similarly, a BCl2 transgene induces hyperplastic splenic follicles, which do not progress to B cell lymphoma in transgenic mice (McDonnell et al., 1989). In contrast, expression of BCl2 in the T cell lineage induces peripheral T cell lymphoma (Linette et al., 1995). Activation of PKB/AKT in the mammary gland also delays involution and promotes mammary adenocarcinomas in conjunction with polyomavirus middle T antigen but does not by itself induce tumors (Hutchinson et al., 2001). The moderate phenotypes of the MMTV-BCl2 and -PKB/AKT transgenic mice may reflect the expression levels of the transgenes relative to the endogenous proteins, or the differential sensitivity of different tissues to perturbations in distinct apoptotic and survival pathways.

Although unlikely, it is formally possible that by virtue of the mutations in the Cdk phosphoacceptor sites the Rb Δ K alleles have acquired activities that are not associated normally with wild-type Rb. We note that numerous other MMTV transgenic mice expressing dominant-active alleles have been created, for example, neu/erbB2 (Siegel et al., 1999), B-catenin (Imbert et al., 2001), TGFB (Pierce et al., 1993), and PKB/AKT (Hutchinson et al., 2001). The Rb Δ K alleles have been used in diverse transcriptional and cellular contexts in vitro and in vivo and invariably demonstrated the same albeit superior functions as native unphosphorylated Rb (Hamel et al., 1992; Geng et al., 1996; Knudsen et al., 1998; Sellers et al., 1998; Brown et al., 1999; Harbour et al., 1999). Most importantly, using Rb Δ K11 as bait in exhaustive yeast twohybrid interactive screens of cDNA libraries from mammary and prostate glands we have identified multiple known and novel Rb binding factors, all of which could also bind wildtype Rb (but not nonfunctional Rb $\Delta 22$ mutant; unpublished data]). Thus, there is no evidence to our knowledge that supports the notion that the Rb Δ K alleles are neomorphs, though we cannot categorically rule out this possibility.

In addition to the inhibition of apoptosis, it is conceivable that constitutive expression of Rb may promote tumorigenicity by inducing DNA damage. Rb can modulate the progression through S and G2 (Karantza et al., 1993; Knudsen et al., 1998; Lukas et al., 1999; Knudsen et al., 2000), and unscheduled activation of Rb may interfere with DNA synthesis and/or DNA repair processes. Moreover, in response to DNA damage ATM phosphorylates and stabilizes E2F1, thereby inducing apoptosis through p53 and p73 (Irwin et al., 2000; Lin et al., 2001). Thus, constitutive activation of Rb may result in the accumulation of damaged DNA and the acquisition of transforming mutations. If this model is correct, the hyperplastic nodules in MMTV/WAP-Rb Δ K transgenic mice should exhibit uncharacteristically high chromosomal anomalies, a possibility that should be tested in future studies. Activation of endogenous Rb by Cdk inhibitors might induce similar chromosomal aberrations and elicit tumors as in our transgenic mice.

The results presented in Fig. 8 suggest that inhibition of apoptosis by active Rb involve an E2F1, IGFR1 α , and PKB/ AKT pathway. Recent analysis of anchorage-independent epithelial cell survival in vitro has indicated that the Rb– E2F1 complex regulates PKB/AKT activity via IGF-1 and suppresses rather than induces cell survival (Yu et al., 2001a). These observations establish a link between Rb, E2F1, and PKB/AKT, and conceivably this same pathway may be used by Rb Δ K to prolong cell survival via PKB in the mammary epithelium (Scheid and Woodgett, 2001). Future analysis of MMTV-Rb Δ K:E2F1^{-/-} composite mice will directly establish the role of E2F1 in the development of neoplasia in our transgenic mice.

In contrast to the MMTV/WAP-Rb Δ K transgenic mice, cyclin D1–deficient mice do not succumb to mammary tumors but are rather protected from a subset of oncogenic pathways in the mammary gland (Sicinski et al., 1995; Yu et al., 2001b). However, cyclin D1 has other targets in addition to Rb that might inhibit tumor progression in cyclin D1^{-/-} mice. For example, cyclin D1 binds and stimulates the estrogen receptor by acting as a bridging factor between the ER and steroid receptor coactivators (Neuman et al., 1997; Zwijsen et al., 1997, 1998). In addition, cyclin D1 induces the phosphorylation of all members of the Rb protein family, including p107 and p130, which are coexpressed during mammary gland development (our unpublished observations), and together may mediate a strong inhibition of proliferation in the cyclin D1^{-/-} mammary epithelium. Thus, the activation of Rb cannot be equated with the loss of cyclin D1. Furthermore, the adverse effect of the Rb pathway may be unraveled only when activation is transient; constitutive activation of Rb in cyclin D1-null mice may not allow mammary epithelial cells that acquired transforming mutations to proliferate and progress into full-blown mammary adenocarcinomas.

Implications for cancer therapy

Since deregulated proliferation is the hallmark of tumor development, antiproliferative agents have become a major therapeutic approach to cancer treatment. However, as stated recently (Evan and Vousden, 2001) many growth deregulatory mutations have pleiotropic and tissue-specific effects on cell proliferation and apoptosis, and the therapeutic consequences of antiproliferative agents are thus difficult to predict.

Our results illustrate this potential problem. Although activation of Rb suppresses epithelial cell proliferation in pubescent females, the Rb Δ K transgenic mice eventually develop breast tumors. This scenario may also occur with antiproliferative drugs in the course of cancer therapy. Systemic activation of Rb by therapeutical drugs such as Cdk inhibitors may extend the survival of cells destined to die by apoptosis, leading to the accumulation of oncogenic alterations, clonal expansion, and ultimately cancer. Several Cdk inhibitors (e.g., flavopiridol) that target the Rb pathway are currently in preclinical and clinical development (Bange et al., 2001). Although such drugs have beneficial inhibitory effect on tumor growth, they have also been documented to suppress apoptosis through Rb (Osuga et al., 2000). Our results suggest that such drugs should be examined cautiously in long-term assays in vivo to determine their systemic effects on tissues with high turnover such as the mammary epithelium.

Combined, the coupled effects of Rb on cell proliferation and apoptosis appear to render this tumor suppressor oncogenic in certain tissues both when mutated and activated. This inherent caveat may limit the use of certain Cdk inhibitors, which restrict cancer cell proliferation by activating Rb, to short term treatments. Alternatively, inasmuch as that cancer progression often involves the inactivation of both Rb and p53 long-term cancer therapy may require combinatorial drugs that would induce the Rb pathway but also antagonize its survival effect.

Materials and methods

Plasmid construction and verification

pECE-HA-Rb Δ **K11.** pECE-HA-Rb Δ p34 is a mouse Rb cDNA with an HA tag at the NH₂ terminus and eight mutations in the indicated Cdk sites (a gift from Dr. P. Hamel, University of Toronto) (Hamel et al., 1992; Fig. 1 A).

The plasmid was transformed into dcm⁻ bacteria to allow digestion with Stul. The Cdk sites, S773, T814, and T819, in exon 23 of RbAp34 were mutated to alanine by PCR-directed mutagenesis using Pfu polymerase. Complex PCR included a forward primer (EZ52, 5'-GCACCCACA-AAAATGGCTCCGAGATCTAGAATCTTGG) and a reverse primer (EZ53, 5'-CGGAGCCATTTTTGTGGGTGCTGGCAGACCTTCTG). The flanking primers were as follows: a 5' primer that overlapped the Stul site at the junction of exons 22 and 23 (EZ51, 5'-TAATAGGCCTCCTACCTTGGCAC-CAATACCTCACATTCC) and a 3' reversed primer that overlapped the Saul site downstream of the Rb stop codon. The \sim 450-bp PCR product was digested with Stul and Saul and cloned into pECE-HA-RbDp34, partially digested with Stul, and further linearized with Saul. The PCR region in the resulting pECE-HA-RbAK11 plasmid was sequenced on both strands to confirm the site directed. The ability of RbAK11 to bind large T, localize to the nucleus, induce the flat cell phenotype in Saos-2 cells, and suppress E2F activation was confirmed by transfecting pECE-HA-RbΔK11 DNA into COS and Saos2 cells using standard methods (Zacksenhaus et al., 1993, 1996a)

pBS-MMTV-HA-Rb∆p34 and pBS-MMTV-HA-Rb∆K11. Briefly, the linker in pBluescript was replaced by a double strand linker containing SacI, Notl, Xhol, Notl, and KpnI sites, yielding pBS-EZ15/16. A 2.684-kb Xhol fragment containing the MMTV-LTR and rabbit β-globin cassette was transferred from mmtv-KCR plasmid (Matsui et al., 1990) provided by Dr. Robert Coffey (Vanderbilt University, Nashville, TN) and subcloned into pBS-EZ15/16 (pBS-MMTV). The Rb∆K alleles were subcloned into the unique EcoRI site in pBS-MMTV (Fig. 1B).

WAP-HA-RbΔp34. pUC-mWAP, containing a 6.75 kb of the mouse WAP gene (Campbell et al., 1984) cloned into a unique EcoRI, was purchased from American Type Culture Collection. A linker Kpnl, HindIII, Sall was used to replace a 1.5 kb between the first and third exons of the WAP gene, yielding pUC-mWAP(K, H, S). The pECE-HA-RbΔp34 plasmid was modified so that it contained a 5' HindIII and 3' Sall site, flanking the RbΔp34 coding cDNA. The HA-RbΔp34 was excised as a 2.7-kb HindIII-Sall fragment and subcloned into pUC-mWAP(K, H, S), partially digested with HindIII and further linearized with Sall. The junction fragments at the 5' and 3' ends in WAP-HA-RbΔp34 were confirmed by sequencing.

pBS-MMTV-HA-Rb Δ p34 and pBS-MMTV-HA-Rb Δ K11 were analyzed in vitro to confirm the expression of the Rb alleles, interaction with large T, and nuclear localization using antibodies specific to both Rb and HA. To induce the MMTV-LTR promoter, COS cells were transfected with pBS-MMTV-HA-Rb Δ p34 or pBS-MMTV-HA-Rb Δ K11 together with an expression plasmid for the glucocorticoid receptor (Giguere et al., 1986; Zacksenhaus et al., 1996a), and the cells were cultured in the presence of dexamethasone for 36 h before analysis.

Generation, genotyping, and analysis of transgenic mice

Microinjection of 2 ng/ml DNA into pronuclei of C57BL/6×SJL or FVB mice and generation of chimeric mice were done in the transgenic mouse facilities at the Hospital for Sick Children and Ontario Cancer Institute. Transgenic mice were initially identified by Southern blot hybridization and later genotyped by PCR. Primers used for PCR of MMTV-RbΔks were as follows: forward primer EZ99 from exon 23 of Rb (5'-ATTTCAGAAG-GTCTGCCAAC) and reverse primer EZ38 from the β-globin cassette (5'-GAATAAGGAATGGACAGCAGGG); WAP-RbΔp34 (EZ99 and EZ72, 5'-CATGTCATGACACAGTCGAC). Transgenic females and control wild-type littermates were caged together. For timed pregnancy, the morning of vaginal plug observation was considered as P0.5.

Southern, Northern, and Western blots, and RT-PCR analysis

Genotyping by Southern blot and PCR analyses was performed on tail DNA as described (Zacksenhaus et al., 1996b). To detect MMTV-RbAKs, DNA was digested with Sstl, Southern blotted, and hybridized with a ³²PdCTP SstI-EcoRI 1.0-kb Rb cDNA fragment from the plasmid. WAP-RbAp34 transgene was detected after digestion of genomic DNA with Pvull and hybridization with a ³²P-labeled KpnI-Pvull 1.9-kb probe from WAP-Rbdp34. Total RNA was isolated from mammary glands or other tissues with Trizol Reagent (GIBCO BRL). For Northern blots, PolyA+ RNA samples (2 µg) were resolved on 1% agarose formaldehyde gels, transferred to nylon membranes, and hybridized to a ³²P-labeled mouse Rb Kpnl-Sstl 1.9 kb DNA probe. Protein lysates from mammary gland tissues were prepared as described (Yao et al., 1999). Anti-HA monoclonal antibody was used directly from hybridoma supernatant. Anti-Rb G3-245 antibody was from Pharmingen (14001A). E2F1 (C-20), IGF1Ra (sc-712), cyclin D1 (sc-450), BCI-XI (sc-7195), and p21 (sc-397) were from Santa Cruz Biotechnology, Inc. Total AKT (#9272) and phospho-AKT (#9271) polyclonal antibodies were from NEB. Anti-keratin 18 was obtained from Sigma-Aldrich. The β -casein monoclonal antibodies were a gift from Dr. Charlotte Kaetzel (University of Kentucky, Lexington, KY) (Kaetzel and Ray, 1984).

MMTV-Rb Δ K transcripts were detected using reverse transcribed cDNA (0.3–0.4 μ g) as a template with forward primer EZ99 and reverse primer EZ73 (5'-TAGCCAGAAGTCAGAT-GCTC). WAP-Rb Δ p34 was detected with EZ99 and EZ72. Primers for WAP were forward EZ178 (5'-TAGCAG-CAGATTGAAAGCATTATG) and reverse EZ179 (5'-GACACCGGTACCAT-GCGTTG), for β -casein were forward EZ182 (5'-TCCTGGATCCAGC-CTATTGCTCAAC) and reverse EZ183 (5'-AGTCCATGGGTCGAATTC), for β -actin were forward EZ70 (5'-CTTCTACAATGAGGTGGGTG) and reverse EZ71 (5'-TGGCATAGAGGTCTTTACGG), and for keratin 18 were forward EZ245 (5'-TGGTCTAGCAGATTGAGG) and reverse EZ246 (5'-CTGGT-GTCATGAGTCGGG).

Whole-mount analysis

Right inguinal mammary glands (#4) were resected and flatten fixed in 4% paraformaldehyde, treated sequentially in 100% ethanol, 100% acetone, 100% ethanol, and 95% ethanol, and stained with iron hematoxylin. The glands were destained with 50% acidified ethanol followed by ascending ethanol, Xylenes, and finally stored in methyl salicylate.

Histology, in situ hybridization, and TUNEL

Left inguinal mammary glands (#4) were fixed in 4% paraformaldehyde, treated with $1 \times PBS$, saline, 50% ethanol, and 70% ethanol (in saline), paraffin embedded, and sections were stained with hematoxylin and eosin (H-E) or Gomori-trichrome. In situ hybridization and TUNEL were performed exactly as described (liang et al., 1997, 2000).

Analysis of mammary adenocarcinomas

Mammary glands with visible palpable tumors were resected, fixed in 4% paraformaldehyde, and stained with H-E. Pathology analysis was performed by Dr. Colin McKerlie (Sunnybrook Hospital, Toronto, Canada). Some mammary tumors were cultured in α -MEM medium supplemented with 10% FCS, 10 ng/ml EGF, and 5 µg/ml insulin.

Immunohistochemistry

IHC analysis was performed with a DAKO streptABComplex/HRP kit. PCNA monoclonal antibody (P8825; Sigma-Aldrich) was used at 1:300 dilution. β -casein monoclonal antibody was diluted 1:800. Rabbit antikeratin wide spectrum screening polyclonal antibody (DAKOZ0622) was diluted 1:200. Rb (C-15) polyclonal antibodies from Santa Cruz Biotechnology, Inc. was diluted 1:50. HA monoclonal antibodies from tissue culture supernatant (at 1:20–1:200) or from Boehringer (1583816) (at 4 µg/ml) were used as in Chang et al. (1995).

We thank Dr. Paul Hamel for provision of pECE-HA-Rb Δ p34, Drs. Robert Coffey and Sean Egan for mmtv-KCR, Dr. Charlotte Kaetzel for the R β C1 anti– β -casein antibody, Andrew Ho for advice on Gomori-trichrome staining, Dr. Colin McKerlie for pathology analysis, and Drs. Jim Fata and Brenda Gallie for critical comments on the article.

E. Zacksenhaus is a recipient of a scholarship award from the Cancer Research Society Inc./Canadian Institute of Health Research. This study was supported by the Canadian Breast Cancer Research Initiative (08469) and the U.S. Army Breast Cancer Research Program (DAMD17-97-1-7321).

Submitted: 15 June 2001 Revised: 15 November 2001 Accepted: 15 November 2001

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