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E2F1 and p53 Are Dispensable, whereas p21 warren Cooperates with Rb to Restrict Endoreduplication and Apoptosis during Skeletal Myogenesis

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We describe temporal and genetic analyses of partially rescued Rb mutant fetuses, mgRb:Rb-/-, that survive to birth and reveal specific defects in skeletal muscle differentiation. We show that in the absence of Rb, these fetuses exhibit increased apoptosis, *bona fide* endoreduplication, and incomplete differentiation throughout terminal myogenesis. These defects were further augmented in composite mutant fetuses, mgRb:Rb-/-:p21-/-, lacking both Rb and the cyclin-dependent kinase inhibitor p21^{Waf1/Cip1}. Although E2F1 and p53 mediate ectopic DNA synthesis and cell death in several tissues in Rb mutant embryos, both endoreduplication and apoptosis persisted in mgRb:Rb-/-:E2F1-/- and mgRb:Rb-/-:p53-/- compound mutant muscles. Thus, combined inactivation of Rb and p21^{Waf1/Cip1} augments endoreduplication and apoptosis, whereas E2F1 and p53 are dispensable during aberrant myogenesis in Rb-deficient fetuses. © 2000 Academic Press

Key Words: Rb; p21; E2F1; p53; cell cycle; apoptosis; endoreduplication; myogenesis.

INTRODUCTION

The retinoblastoma tumor suppressor, Rb, is a transcriptional cofactor whose phosphorylation and activity oscillate during the cell cycle under control of cyclin-dependent kinases (Cdks) (Sherr and Roberts, 1999). The Cdks and their associated G1 cyclins are subject to noncatalytic inhibition by specific cyclin-dependent kinase inhibitors (CKIs; e.g., p16) and universal CKIs (e.g., p21, p27, p57) (Sherr and Roberts, 1995). During most of the G1 phase of the cell cycle, Rb suppresses the expression of genes required for progression into, and execution of, the DNA replication phase by forming specific complexes with certain transcription factors, such as E2F1 (Nevins *et al.*, 1997;

² To whom correspondence should be addressed at the Departments of Medicine and Medical Biophysics, University of Toronto, Toronto General Research Institute–University Health Network, 67 College Street, Room 407, Toronto, Ontario M5G 2M1, Canada. Fax: (416) 340-3453. E-mail: eldad.zacksenhaus@utoronto.ca. thereby linking mutations in Rb to p53-dependent apoptosis, at least in some tissues (Bates *et al.*, 1998; Pomerantz *et al.*, 1998). Indeed, Rb and p53 are often inactivated in human cancer, leading to deregulation of the cell cycle and survival of the transformed state (Gottlieb and Oren, 1998). Mice lacking functional Rb die at midgestation and exhibit ectopic DNA synthesis, apoptosis, and incomplete differentiation during neurogenesis, erythropoiesis, and lens development (Clarke *et al.*, 1992; Jacks *et al.*, 1992; Lee *et al.*, 1992, 1994; Macleod, 1999; Mulligan and Jacks, 1998). This spectrum of defects is consistent with the restricted pattern of expression of Rb during embryogenesis

in the nervous system, lens, liver, and skeletal muscles

(Jiang et al., 1997). Studies on the role of Rb during terminal

Dyson, 1998). Following its phosphorylation prior to a restriction point late in G1, Rb dissociates from its partners and the progression into S phase is allowed. In addition to S

phase, E2F1 also induces apoptosis when overexpressed in

vitro (DeGregori et al., 1997; Wu and Levine, 1994). Free

E2F1 can transcriptionally activate p19^{ARF}, which stabilizes

p53 by debilitating the function of its regulator MDM2,

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FIG. 1. Temporal analysis of aberrant myogenesis in Rb mutant embryos. (A–F) H&E staining of intercostal muscles from Rb mutant fetuses and control littermates (original magnification100×). Large nuclei, indicated by arrowheads, are evident as early as E15.5 and accumulate during late fetal development. (G, H) PCNA staining of back muscles (100×). Mutant muscles contain PCNA-positive nuclei within myotubes. (I–K) TUNEL analysis of neck muscles reveals cell death throughout terminal myogenesis (40×). (L) RT-PCR analysis of MCK gene expression in E14.5–E18.5 mgRb:Rb–/– fetuses and wild-type littermates with β -actin as internal control. (M–P, R–U) *In situ* hybridization analysis of mgRb:Rb–/– mutant fetuses and control littermates, using ³⁵S-labeled MCK or myogenin antisense riboprobes. Shown are bright-field images in which silver grains appear as black dots. (Q) Expression of MCK in E17.5 mgRb:Rb–/– :p21–/– double-mutant fetus. Abbreviations: H, heart; D, diaphragm; L, lung; IM, intercostal muscles; T, tongue.

myogenesis, which occurs after Rb-/- embryos die, were made possible in fetuses partially rescued to birth by a Rb minigene (Zacksenhaus *et al.*, 1996). At embryonic day (E) 17.5, these mutant fetuses express the transgenic Rb protein in the brain but not in muscles or other tissues (Zacksenhaus *et al.*, 1996). The partially rescued mgRb: Rb-/- fetuses exhibit severe skeletal muscle defects, including increased cell death and shorter myotubes, ectopic DNA synthesis, the appearance of giant nuclei in residual myotubes, and expression of early skeletal muscle structural genes (myosin heavy chain and cardiac actin) but not late muscle markers (MCK and MRF4).

Inactivation of both Rb and E2F1 results in extension of life span by several days and inhibition of apoptosis in lens and central nervous system (CNS) but to a lesser extent in the peripheral nervous system (PNS) (Tsai *et al.*, 1998). Apoptosis in the lens and CNS, but not the PNS, is also mediated by a p53-dependent pathway (Macleod *et al.*, 1996). Accordingly, mutations in p53 increase the spectrum of tumors in Rb+/-:p53-/- double mutants relative to Rb+/- heterozygote mice (Williams *et al.*, 1994a). In addition to apoptosis, p53 can also induce cell cycle arrest by direct transcriptional activation of the Cdk inhibitor p21^{Waf1/Cip1}. p53-dependent activation of p21 following γ -irradiation leads to cell cycle arrest by inhibiting phosphorylation of Rb (Brugarolas *et al.*, 1999). Notably, γ -irradiation or overexpression of p21 *in vitro* induces cell cycle arrest in Rb-positive cells but endoreduplication in cells lacking Rb (Niculescu *et al.*, 1998). p21 (as well p16) can also suppress apoptosis during myoblast differentiation *in vitro* through Rb or other pathways (e.g., AKT; Wang and Walsh, 1996; Fujio *et al.*, 1999).

Since components of the Rb pathway exhibit tissuespecific effects, the analysis of the Rb pathway in particular tissues is important in understanding normal development as well as neoplastic transformation of specific cell lineages. To determine the role of E2F1, p53, and p21^{Waf1/Cip1} in the aberrant myogenesis in mgRb:Rb-/- embryos, we performed a temporal and genetic analysis on compound mutant mice lacking Rb and E2F1, p53, or p21^{Waf1/Cip1}. Our results show that ectopic DNA synthesis and apoptosis in Rb-deficient muscles are mediated by a pathway, yet to be defined, which is independent of E2F1 or p53. Inactivation of both Rb and p21^{Waf1/Cip1} leads to increased endoreduplication and apoptosis, indicating that these two negative regulators cooperate to facilitate cell cycle exit during terminal myogenesis.

MATERIALS AND METHODS

Intercrosses and Genotyping

The mgRb:Rb+/- mice and p53-/-, E2F1-/-, and p21-/knockout mice were described previously (Brugarolas et al., 1995; Field et al., 1996; Jacks et al., 1994; Zacksenhaus et al., 1996). For timed pregnancy, the morning of vaginal plug observation was considered E0.5. In the experiments described below, we analyzed three to six double-mutant embryos, mgRb:Rb-/- mutants, and control littermates in every assay. All mutant embryos were genotyped in duplicates using DNA extracted separately from limb and tail biopsies. Tissues were treated overnight with proteinase K (400 μ g/ml) at 55°C. The DNA was extracted once with phenol: chloroform (1:1) by shaking for 1 h in a mixer (Fisher; Model 5432) followed by centrifugation, isopropanol precipitation, one wash in 70% ethanol, and resuspension in 100 μ l TE buffer (10 mM Tris \cdot Cl, pH 8.0, 1 mM EDTA). PCRs included DNA 2 µl, MgCl₂ (25 mM) 3 μ l, PCR buffer (10× 100 mM Tris · HCl, pH 8.3, 500 mM KCl) 5 µl, dNTP (10 mM) 1 µl, gelatin (2 mg/ml) 2.5 µl, primers (200 ng/ μ l) 1 μ l \times 2, in 50- μ l reactions. PCR conditions included 30 cycles of 1 min each at 94°C (denaturation), 55-60°C (annealing), and 72°C (extension). Primers for mgRb, Rb, E2F1, and p21 were as described (Brugarolas et al., 1995; Field et al., 1996; Zacksenhaus et al., 1996). The primers for p53 were p53-1 (5'-GTA TCT GGA AGA CAG GCA GAC) and p53x7 (5'-GAT GGT GGT ATA CTC AGA GCC), for wild-type allele, and neo-2 (5'-TCC TCG TGC TTT ACG GTA TCG) and p53x7 for mutant allele.

Histology and Immunohistochemistry

Embryos were embedded in 4% paraformaldehyde in PBS, dehydrated, paraffin embedded, and cut at 4–5 μ m for hematoxylin and eosin (H&E) staining, fluorescence *in situ* hybridization (FISH), and image cytometry and 8 μ m for TUNEL, immunohistochemistry, and *in situ* hybridization. Immunohistochemical analysis of myosin heavy chain (Fast) and PCNA was performed with monoclonal antibodies (Sigma) at 1:300 dilution followed by biotin-labeled anti-mouse and ABC kit from DAKO.

RNA in Situ Hybridization and Reverse Transcription-PCR

In situ hybridization analysis for Rb, MCK, myogenin, p21, and other markers was performed as described (Jiang *et al.*, 1997). All probes were used at 2×10^5 cpm/ μ l with the exception of cardiac-actin and MCK probes, which were used at 5×10^4 cpm/ μ l. The slides were coated with emulsion and exposed for 10–20 days. RT-PCR analysis (25 cycles) of MCK was performed with forward primer EZ211 (5'-CAG ACA AGC ATA AGA CCG) at nt 379 and reverse primer EZ212 (5'-TTG TCG TTG TGC CAG ATG) at nt 737, yielding a fragment of 359 bp.

Apoptosis Analysis

Sections were deparaffinated and hydrated by two changes of xylenes, 5 min, followed by 100, 96, 90, and 80% ethanol and ddH₂O for 3 min each. For nuclear condensation analysis, sections were treated with Hoechst 33258 (Sigma; 0.5 mg/ml in water) for 30 min, washed, and observed under fluorescence microscope. For TUNEL analysis, we used the following modified procedure (Gavrieli *et al.*, 1992). For nuclei stripping, slides were immersed in $1 \times$ PBS for 5 min and then incubated for 2 min with 5 μ g/ml proteinase K in PBS. The sections were rinsed 1×1 min and 2×5 min in PBS and then 2×5 min in ddH₂O. For H₂O₂ treatment, sections were covered with 3% H₂O₂ for 15 min and then rinsed with ddH₂O twice for 2 min and once for 5 min. For equilibration, the sections were covered with $1 \times$ TdT buffer with Na-cacodylate (supplied with the TdT enzyme; Boehringer Mannheim) and BSA 0.25 μ g/ μ l using Parafilm as coverslips for 30 min. For the reaction, the equilibration buffer was removed and replaced by the same buffer containing 10 µM Bio-16-dUTP (Boehringer Mannheim) and 0.25 U/µl TdT (Boehringer Mannheim) with the same coverslips in humid atmosphere for 60 min at 37°C. For termination of the reaction, the slides were washed twice for 15 min in prewarmed 4imesSSC buffer at 37°C. The slides were further washed 2×5 min in PBS and 2×5 min in PBS-T (PBS plus 0.1% Triton X-100) at room temperature. ABC reagent (DAKO) was prepared according to the manufacturer's instructions in PBS-T and added for 30 min. Slides were washed 2 imes 5 min in PBS-T and 2 imes 5 min in PBS. One milliliter of DAB (3',3'-diaminobenzadine) solution was made by mixing 50 μ l of DAB (2.5 mg/ml in 0.1 M NaPO₄, pH 7.6), 23 μ l of 1% NiCl2, 1 μ l of H₂O₂ in NaPO₄, pH 7.6. Slides were treated with DAB for 10 min and washed extensively with tap water. Counterstaining with methyl green (0.5% w/v in 0.1 M Na-acetate, pH 4.0) was done for 30 s, followed by several washes with tap water, 2 imes30 s in 100% butanol and 2×3 min in xylenes. The slides were mounted with Permount (Fisher).

Bone Staining

Embryos were fixed in 95% ethanol. Following genotype analysis by PCR, the skin and viscera were carefully removed and the skeleton was incubated in 1 vol of acetic acid, 4 vol of 95% ethanol, and 7.5 mg Alcian blue (Sigma) per 50 ml to stain ossification centers (Lufkin *et al.*, 1992). After 24 h, the solution was removed, and the skeleton was washed in 95% ethanol for 1 h and then with 2% (original protocol) or 1% (modified protocol) KOH for 24 h. Staining of cartilage in Alizarin red (Sigma) in 1% KOH was done overnight followed by clearing in 20% glycerol, 1% KOH for a week with daily changes. Skeletons were finally stored in 50% glycerol, 50% ethanol.

Image Cytometry

Sections of 5 μ m were dewaxed, rehydrated, and then stained by a standard Feulgen technique using the Quantitative DNA Staining Kit (Becton-Dickinson Cellular Imaging System, San Jose, CA; Cat. No. 54100140). Image cytometry analysis was performed with the CAS 200, a video-based interactive image cytometer (Becton-Dickinson) (Bacus and Bacus, 1994; Berezowsky et al., 1995). After instrument calibration using CAS control slides provided by the manufacturer, microscopic fields of muscle tissue were randomly selected. Within each field, all the isolated nuclei were picked manually. In order to get the largest number of nuclei from each field, manually drawn boundaries were necessary in many cases to assist the software in picking up single nuclei. Ten to twenty fields were selected and more than 300 nuclei for each section were analyzed in total. The images of selected nuclei were digitized and the optical densities and images recorded. The integrated optical density of each Feulgen-stained nucleus was considered proportional to the amount of DNA per nucleus. The optical density was converted into the weight of DNA, and a histogram of all processed nuclei, as well as a scatter plot, was generated by the software. For a better comparison, the processed nuclei for each sample were subclassified into five groups based on their DNA content: class 1 (less than or equal to 12 pg, diploid to tetraploid), class 2 (between 12 and 24 pg), class 3 (between 24 and 48 pg), class 4 (between 48 and 96 pg), and class 5 (more than 96 pg).

Fluorescence in Situ Hybridization (FISH)

Genotype analysis of mouse Y chromosome by PCR with primers specific to the *Sry* gene was performed with forward primer TDY1, 5'-GAC TGG TGA CAA TTG TCT AG, and reverse primer TDY2, 5'-TAA AAT GCC ACT CCT CTG TG, with male samples generating a product of 292 bp (Gubbay *et al.*, 1990). The mouse Y chromosome paint probe (biotin-labeled) (Breneman *et al.*, 1993) was purchased from Cedarlane Laboratories (Cat. No. 1187-YMB-02). All other reagents were from Oncor (tissue kit, S1337-TC; detection kit biotin–FITC, S1333-BF; and hybridization kit, S1340-Kit). The procedures were performed as recommended by the manufacturers with minor modifications (available upon request). The slides were finally mounted with 9 μ l of DAPI (2.5 ng/ml) and 3 μ l of PI (0.1 μ g/ μ l) in anti-fade and observed under fluorescence microscope (Axioscope II; Zeiss).

RESULTS

Temporal Analysis of Aberrant Myogenesis in Rb Mutant Fetuses

We previously described the generation and analysis of partially rescued Rb mutant fetuses, in which the neurogenic defect in Rb-/- embryos is specifically rescued by a Rb minigene, mgRb (Zacksenhaus et al., 1996). Recent analysis of Rb promoter-lacZ transgenic mice (Jiang et al., 2000) indicates that the same Rb promoter used to generate the mgRb minigene can direct transgene expression exclusively to the developing nervous system but not to other tissues, such as liver and skeletal muscles, where endogenous Rb is normally expressed (Jiang et al., 1997). The mgRb:Rb-/- fetuses can therefore be viewed as null for Rb throughout myogenesis. Our previous analysis identified major skeletal muscle defects in E17.5-E18.5 mgRb:Rb-/fetuses. To address the temporal consequences of Rb loss during terminal myogenesis, mgRb:Rb-/- embryos were analyzed from E13.5 to E18.5 (Fig. 1). H&E staining revealed that large nuclei accumulated within myotubes; there were very few large nuclei at E13.5-E14.5 but they were clearly detected by E15.5 (Figs. 1A and 1B); and by E18.5 more than 20% of the myotube nuclei appeared abnormally enlarged (Fig. 1F). The formation of large nuclei was accompanied by ectopic DNA synthesis within the myotube as detected by staining with antibodies specific to the proliferating cell nuclear antigen PCNA (Figs. 1G and 1H, also see below). Thus, in the absence of Rb, myotubes are unable to permanently withdraw from the cell cycle and instead reenter S phase and accumulate large nuclei. Concomitant with this abnormal cell cycle exit, there was a marked increase in cell death throughout terminal myogenesis (Figs. 1I-1K).

The cellular abnormalities described above were accompanied by specific reduction in expression of the late muscle marker, muscle creatine kinase (MCK) (Fig. 1L), but not early muscle markers such as myosin heavy chain (MHC), myogenin, and cardiac actin (see below). Some low levels of MCK transcripts were, however, detectable by RT-PCR analysis at E16.5–E18.5 even in mutant fetuses (Fig. 1L). To determine whether the reduction in MCK gene expression affected all or a subset of myotubes, we performed in situ hybridization analysis on sagittal sections of mutant and control littermates. Expression of MCK was uniformly reduced in mgRb:Rb-/- mutant embryos at E13.5-E15.5 (Figs. 1O and 1P and data not shown), but progressively elevated and readily detectable in E16.5-E18.5 fetuses (Figs. 1R-1U). In contrast, expression of cardiac actin (not shown) and myogenin (Figs. 1M and 1N) was unaffected as determined by in situ hybridization on adjacent sections. The results indicate that Rb is required for the proper transcriptional activation of MCK during myogenesis but despite the excessive apoptosis and ectopic DNA synthesis, a low level of kinase expression was gradually attained during late fetal development even in the absence of Rb.



FIG. 2. mgRb:Rb-/- myotubes undergo endoreduplication and become polyploid. (A–C) Images of E18.5 muscle sections of control (A) and Rb mutant fetuses (B, C) stained by the Feulgen technique. Note the disorganized appearance and abnormally large nuclei in mutant muscles (arrows). (D–F) Image cytometry analysis of E18.5 control (D) and Rb mutant muscles from trunk (E) and tongue (F). The Rb mutant muscles contain nuclei with DNA mass of 2N to >32N. (G) Scatter-plot analysis of E18.5 mgRb-/- mutant fetuses revealing direct relationship between large nuclei and DNA mass. (H–M) Fluorescence *in situ* hybridization analysis with Y chromosome painting of myotube nuclei of E18.5 control and mutant fetuses. Fetuses were genotyped with primers for the mouse Y-chromosome-specific *Sry* gene and analyzed as described under Materials and Methods. (H) A control female fetus showing no positive signal. (I) A wild-type male fetus showing single focus of staining per nuclei along the myotubes. (J) DAPI staining of the FISH image shown in (K). (K–M) Three examples of mgRb:Rb-/- male mutant fetuses with large nuclei (demarcated) containing multiple Y chromosomes.

Nuclei in mgRb:Rb-/- Myotubes Undergo Endoreduplication and Become Polyploid

The giant nuclei in the Rb-deficient myotubes could reflect nonspecific DNA repair/synthesis or polyploidy due to genuine endoreduplication. To address the nature of these nuclei, we first stained histology sections with Feulgen reagent, which provides a quantitative, linear staining in proportion to the amount of DNA in the nuclei (see Materials and Methods). Microscopic observations readily detected large nuclei in muscle areas but not in other tissues in mutant fetuses (Figs. 2A–2C). Image cytometry analysis revealed that nuclei from normal E18.5 skeletal muscles displayed a DNA content of 2N–4N (Fig. 2D). This DNA distribution represented postmitotic nuclei in myotubes and mitotic nuclei from myoblasts and other cell lineages that overlapped the muscle fibers. In contrast, a fraction of the nuclei from mgRb:Rb-/- muscles had a DNA content of more then 4N (Figs. 2E and 2F). Remarkably, some nuclei contained more than 32N chromosomes, presumably due to more than five successive DNA replications without intervening mitosis. A scatter-plot analysis comparing DNA mass and nuclear size revealed a linear relationship (Fig. 2G), thus establishing that the nuclei with excess DNA are the giant nuclei.

We next used FISH to determine whether the excess DNA in the large nuclei in mgRb:Rb-/- fetuses was organized into multiple, distinct chromosomes. Embryos were sexed by PCR with primers for the mouse



FIG. 2—Continued

Y-chromosome-specific *Sry* gene (Gubbay *et al.*, 1990, see Materials and Methods). Tissue sections were then hybridized with a biotin-labeled Y chromosome paint and treated with DAPI plus propidium iodide (PI) to stain the Y chromosomes and nuclei, respectively. Sections from female fetuses produced no signal (Fig. 2H), whereas normal male fetuses revealed a regular pattern of nuclei along the myotube with a single focus of staining per nucleus (Fig. 2I). In contrast, large nuclei from mgRb:Rb-/- male fetuses contained several foci of staining, indicative of multiple Y chromosomes (Figs. 2J-2M). Thus, in the absence of Rb, nuclei in residual myotubes are unable to permanently withdraw from the cell cycle and instead undergo endoreduplication and become polyploid.

Increased Endoreduplication and Apoptosis in Composite Mutant Fetuses Lacking both Rb and p21^{Waft//Cip1}

The observed endored uplication in mgRb:Rb-/- muscles could be mediated by a G2 block exerted by p21^{Waf1/Cip1}, as ectopic expression of this CKI leads to endored uplication in Rb-deficient cells but not Rb-positive cells *in vitro* (Niculescu *et al.*, 1998). Furthermore, p21^{Waf1/Cip1} was readily detected by *in situ* hybridization in skeletal muscles of normal fetuses and might be even slightly elevated in mgRb:Rb-/- mutant littermates (Figs. 3A-3B). To determine whether endored uplication in mgRb: Rb-/- fetuses was mediated by p21^{Waf1/Cip1}, we generated and analyzed mgRb:Rb-/-:p21-/- compound mutant fetuses. Some mgRb:Rb-/-:p21-/- fetuses harvested at E18.5 appeared unwell with patches of hemorrhage, indicating that loss of both genes had an adverse effect on development. We therefore recovered mgRb:Rb-/-: p21-/- mutant fetuses at E16.5-E17.5 (frequency 7/117 = 5.9%), at stages when the compound mutants appeared viable and grossly indistinguishable from mgRb:Rb-/- single-mutant littermates. Additional compound mutant fetuses were generated by mating mutant mice homozygous for p21^{Waf1/Cip1} (i.e., mgRb:Rb+/-:p21-/-).

PCNA staining of the mgRb:Rb-/-:p21-/- doublemutant muscles readily detected large positive nuclei within myotubes, indicating that endoreduplication occurred even in the absence of p21^{Waf1/Cip1}(Figs. 3E and 3F). To quantify the effect of p21 on endoreduplication, we performed image cytometry analysis on single and compound mutant muscles (Fig. 3G). On average, there were over twice as many nuclei with 32*N* chromosomes (class 4) and nine times more nuclei with >32*N* chromosomes (class 5) in mgRb:Rb-/-:p21-/- tongue muscles compared with mgRb:Rb-/- littermates (Fig. 3G).

Immunohistochemical analysis with antibody specific to MHC revealed that myotubes in E16.5 mgRb:Rb-/-: p21-/- fetuses were generally shorter and more disorganized than in single-mutant fetuses (Figs. 4A-4C). Expression of MCK was also reduced in the compound mutant fetuses (Fig. 1Q). Some large nuclei in the mgRb:Rb-/-: p21-/- embryos appeared to collapse (Figs. 4C, 4F, 4I, and 4L, red and yellow arrowheads; Fig. 3F, arrow) and may represent retraction of the myotubes, apoptosis, or mitotic

catastrophe. Nuclear staining with the fluorescent dye Hoechst 33258 revealed multiple nuclei with chromatin condensation in mgRb:Rb-/- and mgRb:Rb-/-:p21-/muscles but very few in wild-type fetuses (Figs. 4G-4I). Both apoptosis and mitotic catastrophe involve nuclear and chromatin condensation but the former is distinguished by DNA degradation, the hallmark of programmed cell death (Chan et al., 1999, and references therein). In situ analysis of DNA degradation (TUNEL) revealed that some, but not all, of the collapsed nuclei were TUNEL-positive (Fig. 4L), thus precluding unequivocal definition of these collapsed nuclei. There appeared to be more cell death in mgRb: Rb-/-:p21-/- muscles than in mgRb:Rb-/- mutants. To quantify cell death, apoptotic nuclei were scored in several areas from three single mutants and three compound mutants at E16.5 (Fig. 4J). A moderate but significant increase in the frequency of apoptotic nuclei was found in muscles of mgRb:Rb-/-:p21-/- double-mutant relative to mgRb: Rb-/- single-mutant fetuses (t test 0.005; t critical = 2.776). On average there were 60% more apoptotic nuclei in muscle areas of Rb-deficient fetuses when p21 was absent. Thus, the combined loss of Rb and $p21^{Waf1/Cip1}$ during myogenesis leads to relatively more ectopic DNA synthesis, which is accompanied by increased cell death.

Enhanced Skeletal Defects in mgRb:Rb-/-:p21-/-Compound Mutant Fetuses

The mgRb:Rb-/-:p21-/- double-mutant fetuses exhibited abnormal posture and hunchback that appeared more severe than in mgRb:Rb-/- fetuses. To test the skeletons of mgRb:Rb-/-:p21-/- fetuses, we performed Alizarin red and Alcian blue staining to reveal bones and cartilage, respectively. The skeletons of mgRb:Rb-/-:p21-/- mutant embryos were fragile and readily dissociated under standard conditions, but were successfully obtained under lower concentrations of KOH (1% instead of 2% in the standard protocol—see Methods and Materials).

The limb bones in the mgRb:Rb-/- and mgRb:Rb-/-: p21-/- mutant embryos were moderately shorter compared with those of control littermates (Figs. 5B vs 5A). This defect cannot be explained merely as a delay in development because the timing of ossification of the metacarpal and phalangeal bones of limbs (Figs. 5A and 5B), a faithful indicator of developmental stages, was comparable in the mutant and wild-type animals. The humerus process (deltoid tuberosity) was significantly less developed in mutant fetuses and in some mutants was completely missing (Figs. 5A-5D, arrows). At higher magnification, the bones (e.g., scapula, Figs. 5E and 5F) of the mgRb:Rb-/-:p21-/double-mutant fetuses appeared more perforated and the boundaries between the ossification center and the cartilage were not as distinct as in wild-type littermates. In wild-type and mgRb:Rb-/- mutant fetuses the ribs were connected to the sternum at an angle, whereas in the double-mutant fetuses the ribs joined the sternum perpendicularly (Figs.

5G–5J). The upper region of the xiphoid process at the end of the sternum was abnormal in both mgRb:Rb–/– and mgRb:Rb–/–:p21–/– mutant embryos (Figs. 5H and 5J, arrows). In the double-mutant fetuses there was also abnormal ossification of the sternum (Fig. 5J, upper arrow). The abnormalities in the deltoid tuberosity and the sternum likely represent indirect consequences of the muscle deficiency in Rb-deficient fetuses, as similar bone defects are also observed in myogenin-deficient mice (Hasty *et al.,* 1993). Thus, the increased skeleton abnormalities in mgRb: Rb–/–:p21–/– double mutants compared with mgRb: Rb–/– single mutants further support the conclusion that inactivation of p21^{WafI/CIp1} further augments the muscle defects in Rb-deficient fetuses.

Endoreduplication and Apoptosis in Skeletal Muscles of Rb-Deficient Fetuses Are Independent of E2F1 and p53

In the lens and CNS but not PNS of Rb mutant embryos, both ectopic DNA synthesis and apoptosis are mediated by E2F1- and p53-dependent pathways (Macleod *et al.*, 1996; Tsai *et al.*, 1998). To test the roles of E2F1 and p53 in endoreduplication and apoptosis in Rb-deficient muscles, we generated composite mgRb:Rb-/-:p53-/- and mgRb: Rb-/-:E2F1-/- fetuses. The mgRb:Rb-/-:E2F1-/- mutant fetuses were readily obtained (6/74 = 8.1%), while mgRb:Rb-/-:p53-/- embryos were recovered at a lower frequency (6/137 = 4.4%), in agreement with the lethal effect of p53 loss on some female embryos (Sah *et al.*, 1995).

The double-mutant fetuses exhibited a hunchback appearance and only a subtle response to stimulation, similar to mgRb:Rb-/- single-mutant fetuses (Zacksenhaus et al., 1996), suggesting that the absence of E2F1 or p53 did not further rescue the mutant embryos or the muscle defects. Accordingly, histology and PCNA-staining analyses revealed no obvious differences in the morphology of the skeletal muscles or the frequency of polyploid nuclei between the single- and the double-mutant fetuses (data not shown). Indeed, image cytometry analysis indicated that the loss of neither E2F1 nor p53 had any measurable effect on aberrant entry of Rb mutant muscles into an endoreduplicating cycle (Fig. 6A). TUNEL analysis during terminal myogenesis in compound mgRb:Rb-/-:p53-/- and mgRb: Rb-/-:E2F1-/- mutant fetuses revealed that apoptosis was also not affected by the absence of E2F1 or p53 (Figs. 6B-6G). In contrast, apoptosis in the lens of mgRb: Rb-/-:p53-/- fetuses was reduced over 10-fold (Figs. 6H-6I), in accord with previous reports (Morgenbesser et al., 1994; Tsai et al., 1998). We conclude that in contrast to the lens and CNS but similar to the PNS, Rb-deficient skeletal muscles undergo ectopic DNA synthesis and apoptosis through a pathway, yet to be defined, which is independent of E2F1 and p53.

DISCUSSION

Control of Myogenesis by Rb

In the absence of Rb, initial myoblast fusion and formation of multinucleated myotubes are near normal until E15.5-16.5. Thereupon, myotubes become relatively shorter presumably due to the accumulative effects of the excessive apoptosis and endoreduplication that occur during terminal myogenesis in the Rb-deficient fetuses. Expression of MCK but not several other muscle markers was significantly reduced in E13.5-E15.5 embryos but gradually elevated thereafter (Fig. 1). Thus, Rb is required for permanent withdrawal from the cell cycle, otherwise apoptosis and endoreduplication ensue, and for transcriptional activation of a subset of muscle genes (Fig. 7A). Although the expression of MCK never reaches wild-type levels, the gradual increase in MCK gene expression in Rb mutant fetuses from E13.5 to E18.5, despite the ectopic DNA synthesis, excessive apoptosis, and muscle degeneration, is somehow puzzling. Other factors, perhaps p107 and p130, may partially compensate for Rb loss. Alternatively, Rb may be required to coordinate several aspects of cell cycle exit and terminal differentiation but many Rb-/- cells may survive the initial crisis and continue to differentiate. albeit less effectively, even in the absence of Rb. The latter scenario is consistent with the observations that Rb-/cells can contribute efficiently to most adult tissues in chimeric mice (Maandag et al., 1994; Williams et al., 1994b).

Rb was shown to interact directly with myogenic factors such as MyoD and myogenin (Gu et al., 1993) and the adipogenic factor C/EBP (Chen et al., 1996) and to positively participate in transcriptional activation. These interactions are, however, difficult to reproduce (Zhang et al., 1999b) and the mechanisms by which Rb cooperates with these factors are ill defined. Whereas a complex containing Rb, E2F1, and an E2F-recognition site can readily be demonstrated experimentally, analogous complexes containing Rb, MyoD (or C/EBP), and the appropriate DNA binding sites have not been observed (Chen et al., 1996; Gu et al., 1993). Recently, a stable protein-DNA complex, containing Rb, the bHLH protein tal-1, and other factors, that controls erythroblast gene expression has been documented, directly implicating Rb in regulating not only cell cycle exit but also differentiation in a physiological context (Vitelli et al., 2000). Rb was also shown to bind Pax3, an earlier muscle factor that controls MyoD expression in body muscles (Wiggan et al., 1998). Analysis of Rb-/- fibroblasts engineered to express myogenic factors indicates that transcriptional activation by MEF2C is also affected (Novitch et al., 1999). However, there are clearly some differences between fibroblasts expressing myogenic factors and myoblasts. For example, MyoD:Rb-/- fibroblasts do not fuse well and fail to express both MCK and MHC (Gu et al., 1993; Novitch et al., 1996), whereas Rb-/- myoblasts do form myotubes and exhibit near-normal expression of MHC in vitro

(Schneider *et al.*, 1994) and *in vivo* (Figs. 4A and 4B). Other pathways may impinge upon MCK gene expression in the absence of Rb. It has recently been shown that Rb-deficient fibroblasts have elevated levels of activated Ras and that suppression of Ras activity in these fibroblasts restores their ability to induce MyoD-responsive genes (Lee *et al.*, 1999). However, so far, we were unable to detect elevated GTP-loaded RAS in mgRb:Rb-/- muscles (Y. Liu and E. Zacksenhaus, unpublished data). Our observation that Rb controls the normal expression profile of MCK *in vivo* should encourage additional analysis to explore the mechanisms of this regulation in Rb mutant mice and myoblasts.

Cooperative Effects of Rb and p21^{Waf1/Cip1} during Myogenesis

We demonstrated herein that nuclei in Rb-deficient myotubes undergo bona fide endoreduplication leading to the accumulation of DNA up to 32N chromosomes and more (Fig. 2). Endoreduplication in human is rare and occurs only in certain specialized lineages such as megakaryocytes (Zimmet et al., 1997). More often, endoreduplication is associated with genomic instability, loss of cell cycle control, and neoplastic transformation. Although the loss of Rb normally leads to ectopic DNA synthesis or apoptosis, there might be scenarios in which the transition through G2 is inhibited in Rb mutant cells, leading to endoreduplication and genomic instability. Such a scenario was demonstrated in Rb mutant cells overexpressing p21^{Waf1/Cip1} or irradiated to induce p53 and endogenous p21^{Waf1/Cip1} (Niculescu et al., 1998). In both cases endoreduplication occurred only in cells lacking Rb. Rb was also implicated in limiting rereplication and aneuploidy in human cells treated with microtubule inhibitors (Khan and Wahl, 1998). Thus, the endoreduplication observed in Rb mutant muscles in vivo represents a pathological process that may occur during the progression of cancer.

Although p21^{Waf1/Cip1} can induce endoreduplication in Rb mutant cells in vitro, our genetic analysis revealed that despite its high expression in mgRb:Rb-/- myotubes, inactivation of this CKI does not prevent endoreduplication in mgRb:Rb-/-:p21-/- muscles. In fact, mgRb:Rb-/-: p21-/- mutant embryos exhibited increased polyploidy and apoptosis, reduced MCK gene expression and muscle mass, and more severe bone defects (Figs. 3-5 and 7A). The elevated cell death in mgRb:Rb-/-:p21-/- muscles may be triggered by the increased cell proliferation that occurs when both Rb and p21 are inactivated (Fig. 3) and/or may reflect an anti-apoptotic effect of p21. The enhanced cell proliferation despite the increased cell death in mgRb: Rb-/-:p21-/- mutant muscles is consistent with the accelerated tumorigenicity observed in Rb+/-:p21-/mutant mice (Brugarolas et al., 1998). Embryos lacking both p21 and p57 demonstrate muscle defects very similar to those in Rb mutant fetuses, suggesting that Rb may be a major target of these CKIs (Zhang et al., 1999). However,



FIG. 3. Compound mutant fetuses lacking both Rb and $p21^{Wart1/Clp1}$ exhibit increased endoreduplication. (A, B) RNA *in situ* hybridization of p21 in a E17.5 mgRb:Rb+/- control embryo (A) and a mgRb:Rb-/- mutant littermate (B). Abbreviations: B, back muscles; IM, intercostal muscles; D, diaphragm; T, tongue. (C-F) Compound mutant fetuses, defective in both Rb and p21, exhibit increased PCNA-positive nuclei within myotubes. Arrow in F indicates a "collapsed" nucleus. (G) Quantitative image cytometry analysis. Class 1 corresponds to nuclei with 2N-4N chromosomes and classes 2 to 5 represent polyploid nuclei (see Materials and Methods). The mgRb:Rb-/-:p21-/- double-mutant fetuses exhibited increased numbers of 32N and >32N nuclei compared to mgRb:Rb-/- single mutants at E16.5. The increased endoreduplication is most evident in the tongue (T). Numbers indicate the percentage of nuclei in the respective class.

the enhanced muscle phenotype in mgRb:Rb-/-:p21-/- mutant embryos suggests that p21 likely regulates other factors in addition to Rb. Thus, p21^{Waf1/Cip1} may modulate

the activity of cyclin E/Cdk2 (Brugarolas *et al.*, 1998; Mal *et al.*, 2000) or cyclin-independent Cdk2 activity (Gil-Gomez *et al.*, 1998), the Rb-related proteins p107 or p130, myogenic





FIG. 6. Endoreduplication and apoptosis in skeletal muscles of Rb-deficient fetuses are independent of E2F1 and p53. (A) Image cytometry analysis of E18.5 mgRb:Rb+/- control, mgRb:Rb-/- single-mutant, and mgRb:Rb-/-:E2F1-/- and mgRb:Rb-/-:p53-/- compound-mutant fetuses. The absence of E2F1 or p53 does not significantly affect endoreduplication. (B-F) TUNEL analysis of muscle sections from mgRb:Rb+/- control, mgRb:Rb-/- mutant, and mgRb:Rb-/-:E2F1-/- and mgRb:Rb-/-:p53-/- compound-mutant fetuses in control fetuses exhibit no apoptotic activity. The arrow indicates apoptotic nuclei in skin epithelia. Arrow in (C) indicates a rare, large apoptotic cell in Rb mutant fetuses. (G-I) TUNEL analysis of mgRb:Rb-/- and mgRb:Rb-/- ip53-/- fetuses demonstrating suppression of apoptosis in the absence of p53 in lens but not muscles of Rb-deficient fetuses. The image of skeletal muscles in (G) was taken from an area adjacent to the lens of the mgRb:Rb-/-:p53-/- mutant embryo shown in (I).

FIG. 4. Compound mutant fetuses lacking both Rb and p21^{Waf1/Cip1} exhibit reduced muscle mass and increased apoptosis. (A–C) Immunohistochemical staining of E16.5 Rb mutant fetuses with antibodies reactive to myosin heavy chain (fast). The myotubes in mgRb:Rb-/-:p21-/- mutant muscles are shorter, are less organized, and contain many collapsed nuclei (red arrowheads) (original magnification, 10×). (D–F) H&E staining demonstrating giant (black arrows) and collapsed (red arrowheads) nuclei in E16.5 mgRb: Rb-/-:p21-/- mutant muscles (40×). (G–I) Fluorescence microscopy of muscle sections stained with Hoechst 33258. Purple arrows point to condensed nuclei; yellow arrows indicate large collapsed nuclei. (J–L) TUNEL analysis showing increased apoptosis in E16.5 mgRb: Rb-/-:p21-/- mutant muscles compared with mgRb:Rb-/- littermates. Both TUNEL-positive (yellow arrowheads) and TUNEL-negative (red arrowhead) collapsed nuclei are observed in mgRb:Rb-/-:p21-/- mutant (L, top and bottom). Purple arrows indicate apoptotic nuclei of regular size. (J) The number of TUNEL-positive nuclei represents the average from three different areas counted at 40× magnification from three independent mgRb:Rb-/- and mgRb:Rb-/-:p21-/- fetuses.

FIG. 5. Compound mutant fetuses lacking both Rb and p21^{Waft/Clp1} exhibit more severe skeletal defects. Skeletons were stained with Alizarin red (ossification centers) and Alcian blue (cartilage). (A, B) Limbs of mgRb:Rb+/-:p21+/- and mgRb:Rb-/-:p21-/- compound-mutant fetuses at E17.5. The mutant limb is smaller, but the ossification centers in the metacarpal and phalangeal bones indicate a similar developmental stage. (C, D) Higher magnification of the humerus demonstrating lack of deltoid tuberosity in mutant fetuses (arrows). (E, F) Higher magnification of the scapula of E16.5 mgRb:Rb-/-:p21-/- mutant and control littermate fetuses. The mutant scapula appears more brittle/perforated and lacks defined boundaries between bone and cartilage. (G, H) Sternum of mgRb:Rb-/- mutant and control littermate at E16.5. (I, J) Sternum of mgRb:Rb-/-:p21-/- mutant and control littermate at E16.5. In both pairs, the xiphoid process at the end of the sternum is abnormally developed in the mutant fetuses (arrows). Note the abnormal ossification of the sternum in the double-mutant fetuses. The reduced concentrations of KOH used to obtain intact skeletons of the more fragile mgRb:Rb-/-:p21-/- double-mutant fetuses yielded incomplete clearing of the skeleton preparations in A-D and I-J.



FIG. 7. Distinct pathways downstream of Rb in muscles and other tissues. The genetic analysis described herein establishes the roles of the Cdk inhibitors p21^{Waf1/Cip1}, E2F1, and p53 in aberrant myogenesis in Rb mutant fetuses. (A) Inactivation of Rb results in apoptosis; endoreduplication; reduced expression of MCK and MRF4, but not MHC; and reduced muscle mass and shorter myotubes. These defects are augmented by the loss of p21^{Waf1/Cip1} but not by loss of E2F1 or p53. (B) At least two pathways exist downstream of Rb. In lens and CNS, E2F1 and p53 mediate ectopic DNA synthesis and apoptosis. In skeletal muscles and PNS, inappropriate DNA synthesis and cell death occur through a pathway(s), yet to be defined, which is independent of E2F1 and p53.

factors (Zhang *et al.*, 1999a), or survival factors such as AKT (Fujio *et al.*, 1999).

E2F1- and p53-Dependent and Independent Apoptotic Pathways Downstream of Rb

We demonstrated herein that both endoreduplication and apoptosis in Rb mutant muscles are independent of E2F1 and p53 (Fig. 6). In accord with our results, recent analysis has demonstrated that myoblasts and other cell types can undergo p53-independent apoptosis in vitro (Cerone et al., 2000; Fimia et al., 1998; Agah et al., 1997). These and other studies (Macleod et al., 1996; Tsai et al., 1998) indicate that there are at least two pathways that control cell cycle progression and apoptosis in Rb-deficient mice (Fig. 7B). In the CNS and lens, ectopic DNA synthesis and programmed cell death are mediated through E2F1- and p53-dependent pathways. In the PNS, loss of E2F1 reduces to some degree inappropriate entry into S phase, but there is still substantial cell proliferation and apoptosis in the Rb-E2F1 double mutants (Tsai et al., 1998). Thus, in the PNS and skeletal muscles, the E2F1/p53 pathways do not operate or are redundant. Interestingly, we have recently found that Apaf1 is required for apoptosis in the CNS and lens but less so in PNS and skeletal muscles (Z. Guo, T. Mak, and E. Zacksenhaus, unpublished data). The elucidation of this E2F1/ p53/Apaf1-independent pathway(s) is important as it may be disrupted during the progression of tumors in which the p53 pathway is intact. Curiously, however, p53 is often inactivated in rhabdomyosarcoma, a pediatric cancer of skeletal muscles (Diller et al., 1995), indicating that p53 may play an important role in muscle physiology at least in some contexts that are presumably different from the cell death observed during myogenesis in Rb-deficient mice.

Rb and **Bone Development**

Most of the bone defects observed in the mgRb:Rb-/fetuses, which are further augmented in mgRb:Rb-/-: p21 - / - compound mutants, likely reflect an indirect consequence of muscle degeneration, as they are also observed in myogenin knockout mice. The relatively shortened limbs and brittle/pierced appearance of the bones in mgRb: Rb - /-: p21 - /- fetuses (Fig. 5) may, however, represent an autonomous role for Rb in bone development. A requirement for Rb in bone development would be in accord with the increased risk for osteosarcoma in individuals with germ-line mutations in Rb (Weichselbaum et al., 1988). Notably, ectopic expression of Rb in the Rb mutant osteosarcoma cell line Saos-2 induces differentiation (Sellers et al., 1998), indicating that Rb can be a limiting factor in osteoblast differentiation. Mutant mice devoid of both p107 and p130 display increased chondrocyte density, delayed endochondral bone formation, and short limbs (Cobrinik et al., 1996). In this regard, it will be of interest to test mgRb:Rb-/-:p107-/- and mgRb:Rb-/-:p130-/- mutant fetuses for synergistic effects of Rb and its relatives on development of the bone as well as skeletal muscles and other tissues.

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