Inhibition of NF-kB Activation in Combination with Bcl-2 Expression Allows for Persistence of First-Generation Adenovirus Vectors in the Mouse Liver

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NF-k**B is a key regulator of the innate antiviral immune response, due in part to its transcriptional activation of cytokines and adhesion molecules, which, in turn, function in chemotaxis and activation of inflammatory cells. We reported earlier that viral gene expression in hepatocytes transduced with firstgeneration (E1-deleted) adenoviruses induced NF-**k**B activation, elevation of serum cytokines, and hepatocellular apoptosis during the first days postinfusion. These events did not occur in mice infused with an adenovirus vector deleted for E1, E2, E3, and late gene expression. In the present study, we used an adenovirus expressing an I**k**B**a **supersuppressor (Ad.I**k**BM) and** *bcl-2* **transgenic mice to unravel the role of virus-induced NF-**k**B activation and apoptosis in the clearance of recombinant adenovirus vectors from the liver. The** combined action of I_KBM and Bcl-2 allowed for vector persistence in livers of C57BL/6 \times C3H mice. In the **absence of Bcl-2, I**k**BM expression in mouse livers significantly reduced NF-**k**B activation, cytokine expression, leukocyte infiltration, and the humoral immune response against the transgene product; however, this was not sufficient to prevent the decline of vector DNA in transduced cells. Infusion of Ad.I**k**BM caused extended apoptosis predominantly in periportal liver regions, indicating that NF-**k**B activation may protect transduced hepatocytes from apoptosis induced by adenovirus gene products. To confer vector persistence,** *bcl-2* **transgene expression was required to block virus-induced apoptosis if NF-**k**B protection was inactivated by I**k**BM. Expression of gene products involved in early stages of apoptotic pathways was up-regulated in response to virus infusion in** *bcl-2* **transgenic mice, which may represent a compensatory effect. Our study supports the idea that the suppression of innate defense mechanisms improves vector persistence.**

First-generation recombinant adenoviruses (rAd) deleted for all E1a and E1b genes are widely used for gene transfer in vitro and in vivo (for a review, see reference 17). Systemic application of rAd in mice by tail vein infusion results in predominant transduction of the liver (60). Several studies reported expression of early and late adenovirus proteins in transduced hepatocytes mediated by cellular proteins that can substitute for E1a in its function as transactivator for viral genes (for a review, see reference 28). Expressed viral proteins contribute to toxic effects and elicit an innate and specific immune response directed against the virus or transduced cells (for reviews, see references 7 and 29). In most mouse strains, transgene expression is lost within several weeks after rAd infusion (3, 46). In cases where the decline in transgene expression correlated with the loss of hepatic vector DNA, the etiology of vector clearance was attributed to cytotoxic T-lymphocyte (CTL)-mediated cytolysis of transduced cells involving the Fas (1, 15) and/or perforin/granzyme (65) pathways. In other reports, the humoral immune response against viral and/or transgene products was thought to lead to lysis if transduced cells and/or to interference with the detection of secreted transgene products (33, 45, 46, 57). Alternative factors causing vector clearance include intracellular degradation of vector genomes by innate antiviral mechanisms (initiated, for example, by cytokines such as tumor necrosis factor alpha $[TNF-\alpha]$ and interferons [IFNs] without cell loss or apoptosis induced directly by viral proteins expressed in transduced cells (11, 18, 27, 29, 50).

There have been promising attempts to modulate the antigen-specific host immune response in mice by specific inhibition of costimulatory signals required for B- and T-cell activation (20, 21, 43, 63, 64). Furthermore, newer generations of adenovirus vectors deleted for all viral genes (so-called gutless vectors) that lead to persistent expression of the human α_1 antitrypsin (hAAT) gene at high levels in C57BL/6 mice have been developed (44). Nonetheless, first-generation vectors remain an attractive vehicle for gene therapy of tumors and viral infections due, in part, to relatively easy production of purified virus at high titers and the ability to transduce a variety of cell types in vivo. Studies by Ilan et al. indicate that problems of vector-host interaction may also be addressed by overexpression of adenovirus E3 genes that counteract host defenses (18). In this context, the goal of our study was to understand the role of NF-kB activation and of apoptotic pathways that can be blocked by Bcl-2 in adenovirus clearance from mouse liver.

The NF- κ B proteins with transactivating function represent a heterodimer of p65 (or c-Rel or RelB) with p50 or p52 (for reviews, see references 12 and 52). NF-kB is sequestered in the cytoplasm by tightly bound inhibitory proteins called I_kBs , masking the nuclear localization signal of NF-kB. The most

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important members of this family include $I \kappa B\alpha$, $-\beta$, and $-\varepsilon$, which inhibit different members of the NF- κ B family. Phosphorylation of IkB by IkB kinases results in ubiquitination of IkB, which in turn leads to proteasome-mediated degradation of the inhibitor, allowing NF-kB to enter the nucleus and to function as a transcriptional transactivator for a large variety of genes. IkB kinases can be activated through a number of pathways, including the TNF–TNF- α receptor 1 (TNFR1)–TNFR1associated death domain-containing protein (TRADD)–TRAF2 pathway (12). A variety of viruses, including human immunodeficiency virus, herpes simplex virus, cytomegalovirus, papovaviruses (for reviews, see references 2 and 34), and rAd (8, 29, 31), can activate NF-kB. Viruses have evolved mechanisms to utilize various properties of NF-kB to facilitate their gene expression, replication, and evasion of immune responses. In adenoviruses, NF-kB, for example, transactivates the promoter for the E3 region, which encodes a number of proteins that block TNF- α activity (9). On the other hand, it is known that NF-_KB is a key regulator in the antiviral immune response. This is achieved in part by its potential to coordinately transactivate transcription of genes for inflammatory cytokines, or adhesion molecules, which in turn may induce activation or chemotaxis of immune cells (for a review, see reference 12).

NF-kB also regulates apoptotic pathways (for a review, see reference 51). Depending on the specific cell type, differentiation stage, and duration of NF-kB activation, both a proapoptotic role of NF-kB (14, 30) and a role in protection against apoptosis (4, 5, 59, 62) have been observed.

Bcl-2 and related members of the Bcl-2 family are antiapoptotic proteins (for a review, see reference 24). Bcl-2 is localized in mitochondrial, endoplasmic reticulum, and nuclear membranes. In general, the ratio of death antagonists such as Bcl-2 to agonists (e.g., Bax and Bak) determines whether apoptosis will be suppressed or promoted. Hepatocytes normally lack detectable Bcl-2 expression but produce high levels of Bax and Bak, which sensitizes them to Fas-induced apoptosis (22, 23). Hepatic expression of Bcl-2 in Bcl-2 transgenic mice protected them completely from apoptosis induced by synergistic anti-Fas antibody injection (25, 39). The mechanisms behind the antiapoptotic action of Bcl-2 remain enigmatic. In relation to virus infection, Bcl-2 overexpression in vivo inhibited Semliki Forest virus transcription and early replication and delayed virus-induced apoptosis (42). NF-kB-mediated activation of apoptosis after Sindbis virus infection can be inhibited in specific cell lines by Bcl-2 (30).

Activation of cytokines is a central element of the innate immune response against viruses (34). Cytokines are released or produced after systemic application of rAd in a titer-dependent manner (13). Cytokines can directly affect viral replication and gene expression, can induce apoptosis of infected cells, and/or can stimulate production of leukotrienes or adhesion molecules which are involved in chemotaxis and activation of immune cells (35). Among the cytokines activated after rAd infusion is TNF- α (for a review, see reference 37). In the liver, TNF- α is produced primarily by activated Kupffer cells as a biologically active membrane-bound precursor that is cleaved to produce the mature cytokine released in the serum with systemic effects. Released TNF- α binds to two different receptors on a variety of target cell types, including Kupffer cells and hepatocytes. The two TNFRs p55 and p75 have different intracellular domains involved in activation of different signal transduction pathways leading to cell cycle progression, apoptosis, or differentiation. It is thought that among the inflammatory cytokines, TNF- α plays a dominant role in rAd clearance (11, 29). This conclusion is supported by the observation

that $TNF-\alpha$ knockout mice or Kupffer cell-depleted mice demonstrated less leukocyte infiltration early after infection and a reduced humoral immune response to the virus. The importance of TNF- α in defense against adenoviruses is also suggested by the fact that four of the \sim 25 early adenovirus proteins (E1b-19K, E3-10.4K, and E3-10.4K/14.5K) prevent early TNF- α activity (13). The corresponding genes are removed in E1/E3-deleted first-generation vectors in order to make space for cloning larger inserts.

The duration of transgene expression after gene transfer with first-generation adenovirus varies between different mouse strains (3, 46). In the mouse strain used in this study, reporter gene expression declined to zero by 5 weeks after rAd infusion. The initial goal of this study was to investigate the role of cytokine activation and apoptosis occurring early after rAd infusion in the loss of transgene expression in this mouse strain. Our hypothesis was that virus-induced NF-kB activation was a crucial element in pathways that lead to vector clearance. To dissect the role of NF-kB in activation of cytokines and/or apoptosis, we used an adenovirus vector expressing an $I \kappa B\alpha$ supersuppressor (I_{KBM}). I_{KBM} is not able to respond to activation signals due to mutations in phosphorylation sites and thus remains associated with NF-kB, preventing nuclear translocation. This IkB supersuppressor was used in a number of studies to specifically block NF-kB activation (5, 59, 62). To study the role of Bcl-2-sensitive apoptotic pathways in virus clearance from the mouse liver, we used *bcl-2* transgenic mice $(C57BL/6 \times C3H$ background). We investigated the effects of individual and combined Bcl-2 and IkBM expression on hepatic NF-kB activation, cytokine gene expression, apoptosis, cellular liver infiltrates, vector genome and transgene persistence, and humoral immune response.

MATERIALS AND METHODS

Adenoviruses. Ad/RSVhAAT (Ad.hAAT) (19) contains the Rous sarcoma virus long terminal repeat promoter, the hAAT cDNA (1.4 kb), and the bovine growth hormone polyadenylation signal (bPA). Ad.IkBM contains the 1.0-kb cDNA for the dominant negative $I \kappa B\alpha$ (10) fused at the 5' end to hemagglutinin tag. The IkBM gene was inserted into Ad.PGK between the phosphoglycerokinase (PGK) promoter and the bPA signal (19). The PGK promoter is active in mouse hepatocytes in vivo (19). Ad.Co is a vector that has only the PGK-bPA cassette without any transgene. All adenoviruses were generated by recombination in 293 cells with pJM17 (Microbix, Toronto, Ontario, Canada). pJM17 is an E1-deleted adenovirus type 5 derivative that contains a series of substitutions and deletions that inactivate the expression of the E3-10.4/14.5K and E3-14.7K proteins, which antagonize TNF actions (6). The genes for the E3-11.6K protein and for gp19 are intact in pJM17-derived vectors. Thus, all viruses used in this study are typical first-generation adenoviruses. The plaque titer of all viruses was determined on 293 cells. The presence of replication-competent adenovirus and contamination with endotoxin in virus preparation were excluded by tests described earlier (29). Viruses with a titer of 5×10^{11} PFU/ml were stored at -80° C in 10 mM Tris-Cl (pH 8.0)–1 mM MgCl₂–10% glycerol. IkBM expression in mouse livers after Ad.IkBM gene transfer was confirmed by Western blotting with antibodies specific to the hemagglutinin tag or $I\kappa B\alpha$ (C-21; catalog no. sc-371; Santa Cruz Biotechnology, Santa Cruz, Calif.). IKBM expression was detectable beginning 24 h after Ad.IkBM injection.

Animals. Animal studies were performed in accordance with the institutional guidelines set forth by the University of Washington. All animals were housed in specific-pathogen-free facilities. Human *bcl-2* transgenic (bcl-2tg) mice were provided by Stanley Korsmeyer. Transgenic mice had the *bcl-2* gene under control of the mouse metallothionein promoter. All experimental animals had the same genetic background (C57BL/ $6 \times$ C3H inbred offspring); one group was heterozygous for the *bcl*-2 transgene (bcl-2tg^{+/-}), and the other group was PCR negative for the transgene (bcl-2tg^{-/-}). For breeding, bcl-2tg^{-/-} littermates were mated with bcl-2tg^{+/-} mice because of difficulties in breeding homozygote mice. Mice were screened by PCR with an *bcl-2*-specific primer (5'CTTTGTG GAACTGTACGGCCCCAGCATGCG), an hGX-specific reverse primer (5'G GAGCAGGGACGTCCGGGAGCC) and 500 ng of genomic DNA obtained from mouse tails (PCR conditions, 40 cycles of 45 s at 95°C, 45 s at 55°C, and 1 min at 72°C). For experiments, 5- to 6-week-old bcl-2tg^{-/-} or bcl-2tg^{+/-} mice were used. For induction of *bcl*-2 expression, drinking water with 25 mM ZnSO₄ was given to the mice 4 days before the experiment. *bcl-2* expression after induction was confirmed by Western blotting of liver lysates with Bcl-2-specific antibodies (PharMingen catalog no. 65111A) as described by Pezzella et al. (36). The Bcl-2-specific signal appeared 48 h after provision of the $\rm ZnSO_4$ water and disappeared 5 days after its removal. Adenovirus injection was performed via tail vein infusion with $200 \mu l$ of adenovirus diluted in serum-free Dulbecco modified Eagle medium. Blood samples for analysis were obtained by retro-orbital bleeding. Serum samples for hAAT analysis and hAAT antibody analysis were stored at -20°C. Serum hAAT concentrations were determined by enzyme-linked immunosorbent assay (ELISA) as previously described (19). To obtain liver samples, mice were sacrificed by cervical dislocation.

NF-k**B electrophoretic mobility shift assay (EMSA).** Nuclear extracts were obtained from mouse livers as described previously (29) and stored at -80°C Protein concentrations were measured by the Bradford method. The NF- κ B binding sequence from the class I major histocompatibility complex enhancer element (*H2^k*) was used as a probe. Double-stranded oligonucleotides were end labeled with $[\gamma^{-32}P]$ ATP by using T4 polynucleotide kinase. For each reaction, 10 mg of nuclear protein was incubated with 0.2 ng of labeled oligonucleotide probe for 30 min at room temperature and electrophoresed through 5% polyacrylamide Tris-glycine-EDTA gels. For antibody supershift assays, p50- and p65-specific polyclonal antibodies (Santa Cruz Biotechnology) were used. One microgram of the corresponding antibody was added to the samples after 30 min of incubation with the labeled probe. Gels were dried and exposed to Kodak-AR film for 24 h.

Histological analysis. For histological analysis, liver samples from different lobes were used. For terminal deoxynucleotidyltransferase-mediated dUTP nick end labeling (TUNEL) analysis, liver tissue was frozen in OCT compound (Miles, Inc., Elkhart, Ind.) and cryosectioned in 10-um sections. An In Situ Cell Death Detection kit (Boehringer Mannheim) was used to quantify apoptosis in hepatocytes as specified by the manufacturer. Liver sections were counterstained with hematoxylin and eosin and analyzed by light microscopy at a magnification of \times 60. A higher magnification (\times 190) was used to identify the cellular infiltrate as polymorphonuclear leukocytes. Representative samples were photographed with a Nikon VFM camera. Figure 3 was scanned from color film slides, assembled in Adobe Photoshop, and printed on a dye sublimation printer.

RPA. RNA was isolated from 100 mg of snap-frozen liver tissue and stored in liquid nitrogen, using an RNeasy mini kit (Qiagen). RNA was dissolved on RNase-free water and stored in aliquots at -80° C. The RNase protection assay (RPA) was performed by using PharMingen RPA kits, mAPO-3 Multi-Probe Template Set (45355P) for mouse apoptosis gene expression, and mCK-3b Multi-Probe Template Set for mouse cytokine gene expression according to the protocol provided by the manufacturer. Probes were labeled with $[^{32}P]$ UTP. Quantification of all bands was performed on a model 400S PhosphorImager (Molecular Dynamics, Sunnyvale, Calif.). The signals from specific mRNAs were normalized to signals from housekeeping genes (mouse L32 [mL32] and mouse glyceraldehyde 3-phosphate dehydrogenase [mGAPDH]) run on each lane to adjust for loading differences.

hAAT antibodies. Anti-hAAT antibodies were determined by ELISA as described previously (46). Briefly, ELISA plates coated with anti-hAAT capture monoclonal antibody (MAb) were blocked and then incubated with hAAT protein (calibrator serum 4; Atlantic Antibodies, Stillwater, Minn.) diluted 1:50 in blocking buffer for 2 h at room temperature. For each two sample wells, two additional wells were mock loaded with blocking buffer only to determine whether individual high serum hAAT levels would interfere with the assay. Mouse serum samples, diluted 1:1,000 and 1:10,000, were loaded onto the plate along with a similarly diluted naive serum as a negative control. A murine immunoglobulin G2b (IgG2b) anti-hAAT MAb (178260; Calbiochem, La Jolla, Calif.) serially diluted in blocking buffer from 10⁻² to 10⁻⁶ was also included on each plate as a positive control. Following another 2 h of incubation at room temperature, the plates were incubated with horseradish peroxidase-labeled sheep anti-mouse IgG whole-molecule antibody (A-6782; Sigma, St. Louis, Mo.) for another 2 h. To avoid interference with high endogenous hAAT levels, the values for the control wells (without hAAT calibrator) were subtracted from the sample values.

Southern analysis. For genomic DNA preparation, mouse livers were flushed with 5 ml of phosphate-buffered saline via the portal vein. Genomic DNA was extracted from 100 mg of liver tissue as described earlier (28). DNA concentrations were determined spectrophotometrically. Ten micrograms of genomic DNA was digested with *Bam*HI, run on a 0.8% agarose gel, and electrotransferred to a Hybond N^+ nylon filter (Amersham). The blots were hybridized in rapid hybridization buffer (Amersham) with a [a-32P]dCTP-labeled hAAT probe, using a random priming kit (Gibco BRL).

RESULTS

Inhibition of NF-k**B activation by Ad.I**k**BM.** Recently, we reported that infusion of a first-generation virus (rAd) or a vector deleted for 25 kb, including E1, E2, E3, and late genes, activated NF-kB in mouse livers within minutes, probably as a result of particle-receptor interaction and/or internalization (29). This early NF-kB activation was followed by a second phase of NF-kB activity beginning at day 3 after infusion of rAd but not the deleted vector, suggesting that viral gene

FIG. 1. Hepatic NF-kB activity after adenovirus infusion. bcl-2tg^{-/-} and bcl-2tg^{+/-} mice were injected with 4×10^9 PFU of Ad.Co (without transgene) because the contract matrix of 24.2×10^9 PFU of Ad.Co (without transgene) or 4×10^9 PFU Ad.I_KBM together with an equal dose of Ad.hAAT, and nuclear extracts were analyzed by EMSA at day 3 p.i. Reticulocyte lysate (RL) was used as the marker to determine the position of the p50-p65 complex. Positions of the NF- κ B p50-p65 complex and the p50 homodimers were identified in supershift assays with specific antibodies (30). Each lane represents an individual animal.

expression was responsible for NF-kB activation. The same pattern of NF-kB activation was observed in mice depleted for Kupffer cells, indicating that NF- κ B activation takes place in hepatocytes. A biphasic elevation of serum TNF was observed, with a first peak occurring shortly after rAd infusion as a result of TNF release from Kupffer cells and a second peak that correlated with viral gene expression in hepatocytes (29).

Our initial hypothesis in this study was that cytokines produced in hepatocytes early after rAd transduction are responsible in large part for loss of vector DNA by activation of innate defenses (e.g., apoptosis) and/or by facilitating subsequent antigen-specific clearance mechanisms and that the inhibition of cytokine expression can prolong vector persistence. To study whether virus-induced NF-kB activation is causally linked to expression of cytokine genes, we used IkBM to specifically block the NF-kB activation pathway after rAd infusion. All of the following experiments were performed with four groups of animals. Each bcl-2tg^{+/-} or bcl-2tg^{-/-} mouse was injected with 4×10^9 PFU of Ad.IkBM together with an equal dose of Ad.RSV/hAAT (expressing hAAT as a reporter gene) or 4×10^9 PFU of Ad.Co (rAd without transgene) with 4×10^9 PFU Ad.RSV/hAAT. From earlier experiments, it was known that a dose of 8×10^9 PFU transduces >95% of hepatocytes with 20 to 40 viral genome copies per cell, such that both hAAT and IkBM were coexpressed in host cells (60, 61). Endogenous hepatic Bcl-2 expression was not detectable by Western blotting in bcl-2tg^{+/-} and bcl-2tg^{-/-} mice (data not shown). *bcl-2* transgene expression from the metallothionein promoter in bcl-2tg^{+/-} mice can be induced by adding $ZnSO₄$ to the drinking water (see Materials and Methods).

The activation status of NF-kB was analyzed by EMSA, which reflects the DNA binding activity in nuclear extracts from livers. For the analysis of NF-kB activity, we selected day 3 postinfusion (p.i.) because it corresponded to the second phase of NF-kB activation and could potentially be inhibited by IkBM expressed after Ad.IkBM infusion. NF-kB activation at day 3 p.i. was significantly reduced in nuclear extracts from livers transduced with Ad.IkBM, regardless of the bcl-2tg status (Fig. 1). It cannot be excluded that IkBM expression from

FIG. 2. Concentrations of cytokine RNAs after adenovirus infusion. Adenovirus was administered to mice as described for Fig. 1. At days 4 and 40 p.i., 2.5 µg of total liver RNA was quantified for RNAs specific for the inflammatory cytokines TGF-β1, IFN-γ IL-6, and TNF- α by RPA using the mCK-3b Multi-Probe Template Set (PharMingen) for mouse cytokine gene expression. Noninfused bcl-2tg^{-/-} mice were used as controls. Cytokine mRNA expression in these mice was the same as in naive bcl-2tg^{+/-} animals. Signals from protected fragments were quantified on a PhosphorImager. Represented are the average mRNA intensities normalized to the signals of the housekeeping genes (mL32 and mGAPDH) and expressed as arbitrary units. Results are means \pm standard deviations for at least three animals. (IL-6 mRNA was not detectable by RPA in any of the groups and therefore was not included in the graphic.)

Ad.IkBM was not high enough in all transduced cells to completely block NF-kB activation.

Cytokine expression in relation to NF-k**B activation.** To study the activation of cytokine gene expression in relation to NF-kB status, we decided to quantify cytokine-specific RNA in livers by RPA rather than by measuring serum cytokine levels. The latter cannot identify the organ that produced cytokines and cannot discriminate between de novo expression and cytokine release. We concentrated on the main inflammatory cytokines transforming growth factor β 1 (TGF- β 1), IFN- γ , TNF- α , and interleukin-6 (IL-6). Total liver RNA was isolated at days 4 and 40 after rAd infusion from the four experimental groups described above and analyzed by RPA with a mouse cytokine template set (PharMingen) (Fig. 2). Detectable IL-6 transcription was not found in liver RNA from any of the animals in all four experimental groups. TGF- β 1, IFN- γ , and TNF- α RNA expression was higher by more than a factor of 2 at day 4 after infusion of rAd in the bcl-2tg^{-/-}/Ad.Co group than in naive animals that did not receive adenovirus. Ad.IkBM administration reduced significantly the rAd-induced activation of cytokine gene transcription in bcl-2tg⁺ and bcl-2tg^{-/-} mice. At day 40 after rAd infusion, cytokine expression was not significantly higher than in naive mice. The data suggest that NF-kB was one of the factors responsible for activation of cytokine expression in hepatocytes shortly after rAd infusion.

Histological analysis for liver apoptosis and inflammation. To study the effects of NF-kB and Bcl-2 on hepatic apoptosis and inflammation induced early after rAd transduction, we obtained liver sections from three mice per group at days 1, 2,

5, and 40 p.i. and analyzed them for signs of apoptosis by TUNEL and for neutrophil infiltration. Representative samples from each group obtained at days 1, 2, and 5 p.i. are shown in Fig. 3. Liver histology performed on samples from day 40 p.i. did not differ from those obtained at day 5 p.i. (data not shown). TUNEL signals in about 2 to 3% of hepatocytes were observed as early as 24 h (Fig. 3b) as well as 5 days after Ad.Co infusion into bcl-2tg^{$-/-$} mice (Fig. 3g). In addition to these sparse signals, intense focal hepatocellular apoptosis appeared 24 h after infusion of Ad.I_KBM into bcl-2tg^{-/-} mice, particularly in periportal regions, which theoretically received the highest viral load and probably expressed the highest concentrations of $I \kappa BM$ (Fig. 3c). These focal apoptotic areas were infiltrated with neutrophils by day 2 (Fig. 3d) and resolved by day 5 p.i. (Fig. 3h). Hepatocellular apoptosis was completely absent (Fig. 3f, i, and j) or observed in fewer than 0.1% hepatocytes (Fig. 3e) in bcl-2tg^{+/-} transgenic mice. Isolated TUNELpositive cells in Fig. 3i and j were from nonparenchymal cells or infiltrated inflammatory cells. Cellular infiltrates observed at day 5 p.i. (Fig. 3g and i) were mostly polymorphonuclear neutrophils. Hepatic infiltrates at day 5 p.i. were significantly reduced in livers transduced with Ad.IkBM. Bcl-2 expression in combination with Ad.IkBM transduction resulted in normal liver morphology at day 5 after rAd infusion. Taken together, these data demonstrate that (i) IkBM expression inhibits hepatic leukocyte infiltration after rAd administration, (ii) efficient inhibition of NF-kB by IkBM may be the reason for hepatocellular apoptosis induced by viral proteins, indicating an antiapoptotic role of NF- κ B, and (iii) hepatocellular apoptosis can be blocked by Bcl-2.

FIG. 3. TUNEL analysis of liver sections. Adenovirus was administered to mice as described for Fig. 1. Liver sections were obtained at days 1, 2, 5, and 40 p.i. and analyzed for apoptotic cell death by the TUNEL technique bcl-2tg^{-/-}/Ad.InBM, day 1; (d) bcl-2tg^{-/-}/Ad.InBM, day 2; (e) bcl-2tg^{+/-}/Ad.Co, day 1; (f) bcl-2tg^{+/-}/Ad.InBM, day 1; (g) bcl-2tg^{-/-}/Ad.Co, day 5; (i) bcl-2tg^{-/-/}
Ad.InBM, day 5; (i) bcl-2tg^{+/-}/Ad.Co, day 5;

Expression of gene products involved in apoptotis. Certain chronic apoptotic stimuli may up-regulate expression of death proteins. Moreover, a compensatory reaction in the form of induction of apoptotic gene expression is thought to take place if specific pathways or the balance between death proteins are affected (24, 53). In an attempt to begin to unravel the mechanisms involved in rAd-induced apoptosis, we studied the expression of major components of the cell death pathways triggered by TNF or Fas ligand (FasL). At day 4 after rAd infusion, total liver RNA was quantified by RPA for RNAs specific for murine FasL2 (also called TRAIL), Fas, FLICE (caspase-81), FAF, and Fas-associated death domain-containing protein (FADD) (for the Fas pathway) and TNFR p55 and TRADD (Fig. 4). Transcription of the genes for TNFR p55, FAF, Fas, and FLICE was increased significantly in both groups of bcl-2tg^{+/-} transgenic mice, with the most pronounced activation (threefold) for TNFR p55 expression. To a lesser extent, FAS and FLICE expression was stimulated in bcl-2tg^{-/-} mice that received Ad.I_{KBM}, while TRADD, FasL2, and FADD expression was not significantly increased (Fig. 4A). Induction of gene expression in bcl-2tg^{+/-} mice occurred gradually during the first 4 days after rAd infusion (Fig. 4B). The data suggest that rAd (Ad.Co) infusion did not significantly increase expression of apoptotic genes in bcl- $2tg^{-/-}$ mice, whereas in bcl-2tg^{+/-} mice, expression of certain cell death proteins involved in the early stages of apoptotic pathways was up-regulated.

Effects of *bcl-2* **and I**k**BM on transgene expression and vector DNA persistence.** Based on our hypothesis that inflammation and apoptosis activated by cytokines and/or viral proteins affect vector persistence, we monitored hAAT transgene expression after Ad.RSV/hAAT infusion in relation to Bcl-2 expression and NF-kB inhibition by Ad.IkBM, which can block the above-mentioned mechanisms (Fig. 5). hAAT expression declined to undetectable levels by weeks 5 to 6 in bcl-2tg⁻ and bcl-2tg^{+/-} mice infused with Ad.hAAT in combination with a control virus. Two of three bcl-2tg^{-/-} mice that received Ad.IkBM and Ad.hAAT expressed hAAT for 1 or 70 days. Coinfusion of Ad.I_KBM with Ad.hAAT into bcl-2tg^{+/-} mice resulted in persistent hAAT expression over the analyzed time period of 100 days. Transient induction of Bcl-2 expression throughout the first week after adenovirus infusion was sufficient to confer stable transgene expression.

To understand the reasons for this expression pattern, vector DNA in genomic liver DNA was analyzed by Southern blotting. The analysis was performed on the animals used for hAAT studies represented in Fig. 6 at day 100 p.i. At this time point, hAAT expression was present only in Ad.IkBM-infused $\text{bcl-2tg}^{+/-}$ mice. At day 100 p.i., a high concentration of vector (Ad.hAAT) DNA was detected only in bcl-2tg^{+/-} mice that received Ad.IkBM. Vector DNA in all other groups was significantly reduced but still detectable. Liver DNA from bcl- $2tg^{-/-}$ mice that received Ad.I_{KBM} had a slightly higher amount of vector DNA than did liver DNA from bcl- $2tg^$ animals injected with Ad.Co. The data indicate that inhibition of NF-kB prolonged transgene expression, but the block of apoptosis by coexpression of Bcl-2 was necessary to achieve persistence.

Effects of Bcl-2 and Ik**BM on anti-hAAT antibody production.** The absence of hAAT expression at a time point when vector DNA was still detectable may be related to a block of transgene expression at the level of transcription, translation, or posttranslational processing. Tsui et al. reported that CTLderived cytokines can induce posttranscriptional clearance of hepatitis B virus RNA in infected hepatocytes (58). Furthermore, certain cytokines can inhibit transcription from the Rous

sarcoma virus promoter used for hAAT expression (38). We did not investigate these possibilities but instead concentrated on an observation recently made by our group (45, 46) and others (33), that antibodies to the expressed transgene product (hAAT) reduce the level of detectable serum hAAT in C3H mice. To evaluate this possibility, we determined whether antibodies to hAAT were present in the serum at day 4, 7, 14, and 30 p.i. (Table 1). Anti-hAAT IgG antibody levels in serum samples of bcl-2tg^{-/-} and bcl-2tg^{+/-} mice infused with Ad.IkBM were at least 100 times lower than in those of mice that received the control virus. In most of the Ad.IkBM-infused bcl-2tg^{+/-} mice, anti-hAAT antibodies were undetectable. This indicates (i) that anti-hAAT antibodies interfered with hAAT detection by ELISA and (ii) that inhibition of NF-kB effectively prevented the humoral immune response against the transgene product hAAT.

DISCUSSION

This study addressed the hypothesis that apoptosis of transduced cells is a key mechanism for rAd vector clearance from the mouse liver. We postulated that viral proteins expressed in transduced hepatocytes induce apoptosis by a combination of two major mechanisms: (i) a direct proapoptotic activity inherent to specific adenovirus early or late proteins and (ii) activation of cytokines (e.g., $TNF-\alpha$) that cause apoptosis through direct action or by recruitment of CTLs or NK cells, which trigger apoptotic cell death (Fig. 7). We hypothesized that cytokine induction involves the activation of NF-kB. These hypotheses were based on an earlier study where we demonstrated that viral gene expression in hepatocytes transduced with first-generation adenoviruses induced an innate response characterized by NF-kB activation, elevation of serum cytokines, and hepatocellular apoptosis during the first 3 days p.i. These reactions were absent in mice infused with an adenovirus vector deleted for 25 kb, including the E1, E2, E3, and late genes (29).

In an attempt to delineate the role of virus-induced NF-kB activation and apoptosis in clearance of rAd vectors from the liver, we used IkBM and *bcl-2* transgene expression to block the corresponding mechanisms. Strikingly, the combined action of IkBM and Bcl-2 led to vector persistence in livers of the mouse strain used in this study.

Role of hepatocellular cytokine expression. The down-regulation of cytokine expression by blocking the NF-kB transactivation mechanisms inhibited infiltration and probably activation of immune cells. This is reflected in part by the fact that in mice infused with Ad.IkBM, leukocyte infiltrates in transduced livers were absent and the levels of anti-hAAT antibodies were markedly reduced. Reduced levels of cytokine transcripts detected in total liver RNA at day 4 after infusion of Ad.IkBM may be the result of inhibited cytokine expression in hepatocytes and of subsequently suppressed infiltration of inflammatory cells, which can potentially produce cytokines (20, 64). Our hypothesis that the initial IkBM-mediated block of cytokine activation occurred primarily in hepatocytes and not in nonparenchymal liver cells is supported by the following observations. (i) A number of reports demonstrated the expression of TNF- α , TGF- β 1, and IFN- γ genes in isolated primary hepatocytes or hepatocyte lines after adequate activation (40, 41, 47, 54). (ii) The same pattern of NF-kB activation occurs in mice depleted for Kupffer cells, suggesting that rAd-induced NF-kB activation takes place in hepatocytes (29). (iii) Inhibition of cytokine expression correlates with the blockage of NF-kB activation by IkBM in hepatocytes. Hepatocytes are the cell type that is predominantly transduced after intravenous

FIG. 4. Expression of proapoptotic genes after adenovirus infusion. Adenovirus was administered to mice as described for Fig. 1. (A) At days 4 and 40 p.i., total liver RNA from all four experimental groups was quantified by RPA for RNAs specific for gene products involved in apoptotic pathways. (B) Analysis of gene expression
in bcl-2tg^{+/–} mice at days 1, 2, 3, and 4 after infusi mRNAs were normalized to signals from housekeeping genes (mL32 and mGAPDH) run on each lane to adjust for loading differences and expressed as arbitrary units.
Apoptotic gene expression in the experimental groups is repres at least three animals.

FIG. 5. Serum hAAT levels after Ad.hAAT infusion in relation to *bcl-2* expression and NF- κ B inhibition by I κ BM. A dose of 4 \times 10⁹ PFU of Ad.Co or 4 \times 10⁹ PFU of Ad.IkBM together with an equal dose of Ad.hAAT (first-generation vector) was injected via tail vein into bcl-2tg^{-/-} and bcl-2tg^{+/-} mice. In one group of bcl-2tg^{+/-} animals that received Ad.IkBM (bcl-2tg^{+/-}/Ad.IkBM*), *bcl-2* expression was induced only during the first 7 days after adenovirus infusion by ZnSO₄. All the other animals were under ZnSO₄ for the full duration of the experiment. Serum was analyzed periodically for hAAT by ELISA. Each line represents an individual animal.

rAd infusion in mice. There are no reported data showing that rAd-mediated transgene expression can occur in Kupffer cells or immune cells.

Role of apoptosis induced by viral proteins expressed in transduced hepatocytes. In the absence of cytokine-mediated immune cell recruitment or activation, apoptosis induced by expressed viral proteins may be the central mechanism responsible for the decline in vector DNA in transduced hepatocytes. In general, lytic viruses regulate cell viability, inhibiting apoptosis during the early stages of replication and then promoting apoptosis late in infection so that progeny virions can be released from the cell (for a review, see reference 55). In adenovirus, these processes are regulated mainly by E1a and the antiapoptotic proteins E1B-55K and E1B-19K, which are deleted in rAd vectors. Apoptotic signals observed in mouse livers after infusion of rAd may be causally linked to other adenovirus proapoptotic proteins such as the E4 open reading frame 4 (50), E3-11.6K (56), and pre terminal (pTP) E2 (26) proteins or specific late proteins like penton or fiber (48). Two observations that we made earlier support the etiological role of expressed viral proteins in hepatocellular apoptosis. A small percentage of apoptotic hepatocytes was detected in mouse liver only with adenovirus vectors that expressed early and late viral proteins, not with vectors deleted for viral genes (29). Furthermore, a considerable amount of vector genome is lost during the first 36 h p.i. in both immunocompetent and immunodeficient mice, which may be attributed, in part, to death of infected cells due to virus-induced apoptosis (28). Although apoptosis was observed in only 2 to 3% of all hepatocytes at each time point, it cannot be excluded that an asynchronous turnover of the majority of hepatocytes takes place in the first day(s) after rAd infusion. This hypothesis would be consistent with the DNA replication in \sim 70% of hepatocytes occurring at day 4 p.i. as a result of liver damage and cytokine activation (28, 29, 44).

NF-k**B has a protective role against virus-induced apoptosis in hepatocytes.** Infusion of Ad.IkBM caused extended apoptosis in periportal liver regions that probably received the highest

FIG. 6. Southern blot analysis of adenovirus vector (Adv) DNA in transduced mouse livers. A dose of 4×10^9 PFU of Ad.Co or 4×10^9 PFU of Ad.IkBM together with an equal dose of Ad.hAAT (first-generation vector) was injected into bcl-2tg^{-/-} and bcl-2tg^{+/-} mice. Mice were sacrificed at day 100 p.i. Ten micrograms of *Bam*HI-digested genomic liver DNA was loaded on each lane. Blots were hybridized with a labeled hAAT probe; Exposure was for 4 days. The fragment specific for Ad.hAAT is 2.0 kb. The analysis was performed with animals used for hAAT studies represented in Fig. 5.

viral load and therefore express IkBM at levels sufficient for complete NF-kB inhibition. This finding indicates that NF-kB activation may protect transduced cells from apoptosis induced by adenovirus gene products. This may be why only in combination with Bcl-2 expression did IkBM prevent vector clearance. In general, an antiapoptotic role of NF-kB is supported by the fact that *relA* knockout mice were found to display embryonic lethality due to apoptosis in the liver (4) and by in vitro studies where inhibition of NF-kB activation promoted cell death (5, 59, 62) whereas ectopic NF-kB overexpression protected against apoptosis (51).

Inhibition of apoptosis by Bcl-2. Endogenous Bcl-2 is not expressed at detectable levels in the liver (23, 24). In our studies, *bcl-2* transgene expression in mouse livers prevented virus-induced apoptosis. Bcl-2 protection is most pronounced in the absence of antiapoptotic $NF-\kappa B$ mechanisms (Fig. 7, hypothesis 1). We hypothesize that Bcl-2 can override the virus-induced death signals activated upon the loss of NF-kB activity due to IkBM overexpression. The mechanisms behind the antiapoptotic action of Bcl-2 are still unclear; however, our observation that cytokine expression is slightly reduced in *bcl-2* transgenic mice indicates that Bcl-2 may inhibit the transactivation activity of p65 as suggested previously (14) and neutralize the residual hepatic NF-kB activity left after Ad.IkBM infusion (Fig. 7, hypothesis 2).

Hepatic expression of the *bcl-2* transgene was associated with up-regulated transcription of proapoptotic genes. It appears that all induced gene products are involved in Fas/TNFmediated pathways at or before a step that is blocked by Bcl-2, suggesting that this up-regulation represents a compensatory reaction to a Bcl-2-mediated block (24, 53). More refined analysis with specific inhibitors for Fas, TRAIL (FasL2), caspases, or other components are required to support this hypothesis.

FIG. 7. Hypothetical mechanisms for the roles of NF-kB and Bcl-2 in vector persistence. Expressed adenovirus proteins can induce early apoptosis. (1) N_F - KB has an antiapoptotic role. Bcl-2 blocks virus-induced apoptosis if $NF - \kappa \vec{B}$ protection is inactivated by I_KBM. (2) NF-_KB transactivates cytokine expression, which is a key element in recruitment and activation of immune cells. Hence, NF-kB promotes cytokine/CTL-mediated cell death. Bcl-2 is required to block residual NF-kB activity, which was not inhibited by IkBM.

Mechanism for extinction of transgene expression. We demonstrated that IkBM expression alone was sufficient to inhibit the humoral immune response against the transgene and the infiltration of inflammatory cells into the liver; however, the additional expression of Bcl-2 was required to counteract vector loss in transduced cells. This finding suggests that at least in this mouse strain, a humoral immune response, while affecting the detection of hAAT, was not primarily responsible for vector clearance. Also, loss of vector DNA was a result of at least two mechanisms: (i) the cell-mediated immune response, which is triggered by cytokines and cannot be blocked by Bcl-2 expression alone (Fig. 6, bcl-2tg^{+/-}/Ad.Co) and (ii) early apoptosis induced by viral proteins, which can be prevented by Bcl-2. This conclusion implies that other strategies that can block cytokine activation or action may prolong vector persistence. Notably, a rAd vector that overexpressed all adenovirus E3 proteins suppressed the production of neutralizing antibodies and resulted in transgene persistence (18). The adenovirus E3 region encodes proteins that protect against TNF- and Fas-mediated apoptosis (49). Interestingly, the E3 promoter is transactivated by $NF-\kappa B$ (9), which may be one of the mechanisms for how NF-kB exerts its antiapoptotic function after rAd transduction.

An argument against our hypothesis that early apoptosis induced by cytokines, CTLs, and/or viral proteins, which is

TABLE 1. Anti-hAAT antibody levels in mouse serum after adenovirus infusion*^a*

Mouse/virus	Antibody level			
	Day 4 p.i.	Day 7 p.i.	Day 14 p.i.	Day 30 p.i.
$bcl-2tg^{-/-}/Ad.Co$	$+/+/+$	$+ + +/ + +/ + +$	$+++/++++/++++$	$+++/++++/+++$
bcl-2tg ^{-/-} /Ad.I _K BM	$-/-/-$	$+/-/-$	$+/-$	$+/-$
$bcl-2tg^{+/-}/Ad.Co$	$+/+/+$	$+ +/+ +/+ +$	$+++/++++/++++$	$+++/++++/+++$
bcl-2tg ^{+/-} /Ad.I _K BM	$-/-/-$	$+/-/-$	$+/-/-$	$+/-/-$

^a Virus infusion was done as described in the legend to Fig. 6. Serum obtained at days 4, 7, 14, and 30 p.i. was analyzed for anti-hAAT IgG by a semiquantitative ELISA in comparison to a standard curve of a commercial anti-hAAT MAb. Anti-hAAT antibody levels are expressed for individual animals as not detected (-) or detection similar to that of the standard MAb diluted 1:1,000 $(++)$, 1:100 $(+)$, or 1:10 $(+)$. Results of three separate determinations are shown.

associated with loss of transduced cells, represents a central mechanism for vector clearance is that destruction of $>90\%$ of transduced hepatocytes over a short time should be fatal. However, this argument does not consider that the liver can regenerate quickly in response to damage and cytokine release, particularly if apoptosis is asynchronous. Hepatocellular proliferation in response to rAd was reported by a number of groups (28, 29, 44, 64).

Importance for gene therapy. Our data suggest that transient expression of Bcl-2 at early time points after rAd infusion results in persistent gene expression if the vector is coadministered with Ad.IkBM. If, in addition to transient Bcl-2 expression, a transient, inducible expression of IkBM during a critical period after rAd infusion were sufficient to confer vector persistence, then our study could be the basis for a practicable approach to achieve persistent gene expression for therapeutic purposes. On the contrary, a permanent inactivation of NF-kB will probably have consequences including tumor development and lost protection against infectious agents (for a review, see reference 32).

With respect to permanent Bcl-2 expression, it is notable that an aberrant overexpression of Bcl-2 is often observed in tumors and may be responsible for resistance of tumor cells to apoptotic stimuli such as radiotherapy and chemotherapy (for a review, see reference 16). Nonetheless, our study contributes to improving the understanding of early host antiviral host mechanisms. Further efforts in this direction may include the use of specific compounds for NF-kB inhibition (e.g., oligonucleotide decoys) and blockage of specific apoptotic pathways (e.g., by caspase inhibitors).

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