

Substitution of Adenovirus Serotype 3 Hexon onto a Serotype 5 Oncolytic Adenovirus Reduces Factor X Binding, Decreases Liver Tropism, and Improves Antitumor Efficacy

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Abstract

Following intravascular delivery, an important route of administration for many clinical applications, the liver is the predominant site of adenovirus serotype 5 (Ad5) sequestration, thereby posing a risk of toxicity. In this regard, it has recently been shown that the Ad5 capsid binds to the blood coagulation factor X (FX) via the Ad5 hexon protein. This interaction mediates the majority of Ad5 liver transduction. Patient FX levels can be diminished by the administration of warfarin, a vitamin K inhibitor in the liver that decreases FX production; however, warfarin is a potent anticoagulant and can have a number of undesired side effects. Therefore, genetic modification of the virus to ablate FX binding is the preferred approach. Modifications of the hexon protein, specifically within the hypervariable 5 (HVR5) and 7 (HVR7) regions, have produced Ad5 vectors that show minimal liver sequestration. Our laboratory has pioneered adenovirus hexon modifications, including insertion of peptide ligands into the hypervariable regions and substitution of the adenovirus hexon with hexon proteins from alternate serotypes. Substitution of the adenovirus serotype 3 (Ad3) hexon protein onto the Ad5 capsid has been further characterized with regard to its interaction with FX and incorporated into an infectivity-enhanced conditionally replicative adenovirus (CRAd). *In vitro* evaluation of these hexon-modified vectors showed decreased binding to FX and decreased cell transduction via FX-mediated pathways. Furthermore, *in vivo* biodistribution studies in mice exhibited a decrease in liver sequestration. With the use of xenograft tumor models, the antitumor efficacy of the hexon-modified CRAds was enhanced over nonmodified controls. *Mol Cancer Ther*; 9(9); 2536–44. ©2010 AACR.

Introduction

Adenovirus (Ad)-based vectors represent the most commonly used vectors in gene therapy clinical trials (1), with the majority of these vectors based on serotype 5 (Ad5). Furthermore, cancer represents the disease most commonly addressed. To achieve an effective adenovirus-based cancer therapy, systemic administration (i.e., i.v. administration) is the ideal mode of delivery. However, effective i.v. administration of adenovirus-based vectors has been hindered by the innate liver tropism of Ad5 and concerns of hepatotoxicity. Liver tropism also

results in a sequestration of the administered adenoviral vector away from the desired tumor targets and a rapid clearance, thereby necessitating higher doses of administered vector and resulting in poor antitumor efficacy (2).

Key components of this process and how to overcome the liver tropism of Ad5 have recently been elucidated and may involve multiple redundant and synergistic mechanisms (3). In this regard, the Ad5 capsid interacts with multiple vitamin K–dependent coagulation factors (4). Specifically, coagulation factor X (FX) binds to the Ad5 capsid protein hexon and mediates the majority of the Ad5 liver tropism, presumably via heparin sulfate proteoglycan–mediated pathways (5–7). Through crystal structure analysis and mutagenesis studies, it has been discovered that hypervariable regions (HVR) 5 and 7 are critical for the interaction with FX (8). Multiple vectors have been published that involve mutation of one or more of these HVRs, either by point mutation or peptide insertion. These hexon modifications have been incorporated into nonreplicative vectors, and some of the modifications are complex and may not be easily translated to other vectors. Our group initially pioneered Ad5 hexon modifications with substitution of the adenovirus serotype 3 (subspecies B, Ad3) hexon onto the Ad5 capsid (9) and peptide insertions into the various HVRs (10).

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Pretreatment with the drug warfarin, which inhibits vitamin K-dependent coagulation factor production, can deplete circulating FX levels and decrease the liver tropism of i.v. administered Ad5. Pretreatment with warfarin prior to the administration of an oncolytic adenovirus did increase the therapeutic window of the oncolytic adenovirus and resulted in an enhanced antitumor efficacy in two separate studies (11, 12). However, warfarin has numerous side effects, multiple drug interactions, and a narrow therapeutic index, and can place the patient in a hypocoagulable state that may present other problems. Therefore, incorporation of a genetic modification of the hexon protein is the preferred approach, eliminating the need for anticoagulation therapy. One of the initial hexon modifications showing FX binding has been advanced to a replicative oncolytic adenovirus and has shown improved antitumor efficacies compared with its unmodified control (13).

Ablating Ad5 liver tropism is only a portion of improving adenovirus tumor targeting. In this context, Ad5-based vectors selectively transduce target cells via the adenovirus fiber protein binding to the primary cellular receptor, the coxsackie and adenovirus receptor (CAR). However, CAR is poorly expressed on many tumor targets, thereby limiting transduction (14). One means of improving tumor target transduction is by enhancing the infectivity of the vector with substitution of the knob domain of the fiber protein with that of another adenovirus serotype whose primary receptor has a better tumor cell expression profile. We had previously shown that substitution of the Ad3 knob domain improves tumor cell transduction for a variety of cancer types and improves antitumor efficacy of oncolytic adenoviruses (15, 16).

We hypothesized that ablating Ad5 capsid binding to FX via a genetic modification could be incorporated into an infectivity-enhanced oncolytic adenovirus to reduce liver tropism and increase the antitumor efficacy.

Materials and Methods

Cell lines

HEK-293, A549, HepG2, and Skov3.ip1 cells were purchased from the American Type Culture Collection (July 2009; no further authentication was done). All cells were grown in DMEM/F12 50:50 supplemented with 10% fetal bovine serum, 1% penicillin/streptomycin, and 1% L-glutamine, cultured in a humidified incubator at 37°C with 5% CO₂.

Animals

All animal care was done under protocols approved by the Institutional Animal Care and Use Committee. Five-week-old female C57BL/6 mice and 4-week-old female athymic, nude mice were purchased from Frederick Cancer Research and housed in University of Alabama at Birmingham Animal Resources Program facilities.

Reagents, antibodies

Purified human factors X and XI, and antifactor X monoclonal mouse antibodies were purchased from Haematologic Technologies. Antiadenovirus capsid polyclonal antibody was a gift from Dr. Joanne Douglas (University of Alabama at Birmingham, Birmingham, AL). Goat anti-mouse antibodies conjugated to horseradish peroxidase (HRP) and goat anti-rabbit antibodies conjugated to HRP were purchased from Dako, Inc. CM5 bio-core sensor chips were purchased from GE Healthcare.

Adenoviral vectors

Wild-type Ad48 was purchased from the American Type Culture Collection. Construction of nonreplicative Ad5, Ad5HVR5-6H, and Ad5H3 encoding a green fluorescent protein (GFP)-luciferase (luc) expression cassette under the cytomegalovirus (CMV) promoter in the E1 coding region were previously published (9, 10). Nonreplicative Ad5HBAP encoding a red fluorescent protein (RFP) expression cassette under the CMV promoter in the E1 coding region was provided as a gift by Dr. Michael Barry (Mayo Clinic, Rochester, MN; ref. 17). To generate the nonreplicative Ad5HBAP encoding the isogenic GFP-luc cassette as the above vectors, the Ad5 hexon sequence with insertion of the BAP peptide was cloned by PCR using the virus provided by Dr. Barry as a template and using the primers F: TAACAAGTTTGAACCCACG and R: CATGGGATCCACCTCAAAGTC. The PCR product was digested with *Dra*III and *Bam*HI and was inserted into the previously published hexon shuttle vector pHS (9) to generate HBAP/pHS. Homologous recombination in *Escherichia coli* strain BJ5183 (Stratagene) with the *Swa*I-digested Ad5 backbone containing the E1 CMV GFP-Luc cassette and hexon deletion was used to generate the final nonreplicative Ad5HBAP vector encoding GFP-Luc in E1. Restriction digests and sequencing confirmed the appropriate sequences. Replicative vectors were constructed on an adenovirus backbone with the Δ24 E1 mutation and deletion of E3 with replacement of the E3 protein ADP and with either Ad5 fiber or with substitution of the Ad3 fiber knob domain onto the Ad5 fiber. Hexon modifications were introduced by digesting the nonreplicative backbones with *Sfi*I (unique *Sfi*I enzyme sites flank the hexon protein coding region) and then insertion into the replicative backbones by digestion with *Sfi*I and subsequent ligation. Adenoviral vector genomes were then digested with *Pac*I and transfected into HEK-293 cells. Nonreplicative viruses were upscaled via HEK-293 cells whereas, replicative viruses were upscaled with A549 cells after initial transfection with HEK-293 cells. Virus was purified by CsCl double ultracentrifugation and dialysis into the appropriate buffer.

Slot blot analysis

Polyvinyl pyrrolidone (PVDF) membrane (Bio-Rad) was preincubated with 100% methanol and then rinsed with TBS. Virus particles (vp; 10¹⁰ vp/sample) were

then exposed to the PVDF membrane using a slot blot chamber. The PVDF membrane was washed five times with TBS with Tween-20 and then incubated with TBS plus 1% nonfat dry milk blocking reagent (Bio-Rad) for 30 minutes at room temperature. Purified human FX and all antibodies were diluted in TBS plus 1% nonfat dry milk blocking reagent. The membrane was incubated with either purified human FX (8 $\mu\text{g}/\text{mL}$) or anti-adenovirus polyclonal rabbit antibody (1:1,000 dilution) for 1 hour at room temperature. The membranes were washed five times with TBS with Tween-20 between all incubations and incubated with TBS plus nonfat dry milk blocking reagent for 30 minutes at room temperature prior to the next antibody incubation. The FX-incubated samples were incubated with anti-FX monoclonal mouse antibodies (Haematologic Technologies) followed by anti-mouse goat-HRP antibodies (DAKO, Inc.). Antiadenovirus incubated samples were incubated with anti-rabbit goat-HRP (DAKO, Inc.). Following a final wash cycle (5 times with TBS with Tween-20), both PVDF membranes were developed with diaminobenzidine (Sigma-Aldrich).

Surface plasmon resonance analysis

Purified human FX and FXI proteins were covalently immobilized to CM5 biocore sensor chips by amine coupling. Viruses (10^{10} virus particles) were diluted in 10 mmol/L HEPES (pH 7.4), 150 mmol/L NaCl, and 5 mmol/L CaCl_2 . Samples were processed at 37°C . Samples were analyzed using a Biocore 2000 (GE Healthcare). Virus samples were passed over the coagulation factor coupled chips at $30\ \mu\text{L}/\text{minute}$. Chip surfaces were regenerated between virus samples by injection of 10 mmol/L HEPES (pH 7.4), 150 mmol/L NaCl, 5 mmol/L CaCl_2 , 3 mmol/L EDTA, and 0.005% Tween-20. All virus samples were analyzed in duplicate.

In vitro factor X studies

A549, HepG2, and Skov3.ip1 cells were seeded in 24-well plates at a density of 5×10^4 cells/well and allowed to grow 24 hours as described above. Viruses were diluted to quantities sufficient for 10 vp/cell and 100 vp/cell in a volume of 200 $\mu\text{L}/\text{well}$. Selected virus samples were then incubated for 30 minutes at room temperature with FX at 4 $\mu\text{g}/\text{mL}$, 8 $\mu\text{g}/\text{mL}$, or 16 $\mu\text{g}/\text{mL}$. Virus was then added to wells and infection allowed for 2 hours at 37°C . After 2 hours, virus and media were removed and fresh growth media were added to each well. Cells were allowed to grow for 48 hours and then analyzed for luciferase expression via Promega luciferase assay kit (Promega). Protein concentrations were calculated with Bio-Rad direct protein assay kit (Bio-Rad). All samples were done in triplicate.

Crystal violet assay

Skov3.ip1 cells were seeded in 12-well plates at a density of 2.5×10^4 cells/well and allowed to grow for 24 hours as described above. Viruses were diluted to quantities sufficient for a range of 0.01 to 1,000 vp/cell in 200 $\mu\text{L}/\text{well}$ of 2% infection

media. Viruses were allowed to infect cells for 2 hours, after which virus and media were removed and cells were grown in 2% growth media for 10 days. After 10 days, media were removed and cells were washed with PBS then fixed with 10% formaldehyde. Remaining cells were stained with crystal violet solution 10 minutes and rinsed with deionized water.

In vivo biodistribution studies

For biodistribution studies, one group (3–5 mice/group) of 5-week-old C57BL/6 mice were pretreated with warfarin (5 mg/kg) dissolved in peanut oil administered s.c. 3 days and 1 day prior to vector injections. The mice were then injected with 10^{11} vp/mouse via tail vein injections. After 48 hours the mice were sacrificed and the organs were harvested. Organs were lysed with cell culture lysis reagent (Promega) and mechanically homogenized with glass beads. Cell lysate was analyzed for luciferase expression with Promega luciferase assay kit, and protein concentrations were obtained with Bio-Rad direct protein assay kit. For tumor studies, skov3.ip1 cells were grown to near confluency as described above. Cells were harvested and 1×10^6 cells/mouse were injected into the flank of 4-week-old athymic nude mice. Tumors were allowed to establish over 4 to 6 weeks. Mice in the warfarin pretreatment group were dosed with warfarin (5 mg/kg) at 3 days and 1 day prior to vector injections. Mice were injected with 10^{11} vp/mouse via tail vein injections. After 48 hours the mice were sacrificed and the organs and tumors were harvested. DNA was isolated using Qiagen DNeasy Blood and Tissue Kit (Qiagen). Quantitative PCR was done on DNA samples using primers specific for luciferase transgene and mouse or human β -actin.

In vivo antitumor efficacy studies

Skov3.ip1 cells were grown to near confluency as described above. The cells were harvested, and 1×10^6 cells/mouse were injected into the flank of 4-week-old athymic nude mice. Tumors were allowed to establish over 4 to 6 weeks, and the mice placed into experimental groups with tumors ranging from 0.3 to 0.5 cm in diameter. Mice in the warfarin pretreatment group were dosed with warfarin (5 mg/kg) at 3 days and 1 day prior to vector injections. Mice were injected with 10^{11} vp/mouse via tail vein injections. Tumors were measured with calipers at days 0, 3, 7, 14, 21, and 28 by measuring the average diameter and using the formula $\text{length} \times \text{width}^2 \times 0.5$ (18). Five mice from each experimental group were sacrificed at each time point beyond day 0, and tumor and liver samples were harvested for viral DNA analysis. All remaining mice were sacrificed at 28 days, and liver and tumor samples were harvested. DNA was isolated from liver and tumor samples using Qiagen DNeasy Blood and Tissue Kit (Qiagen). Quantitative PCR was done on DNA samples using primers specific for Ad E4 coding region and mouse or human β -actin.

Statistical analysis

All data are expressed as means. *In vitro* experiments were done in triplicate. *In vivo* experiments were done

Table 1. Characteristics of adenoviral vectors used

Virus	Hexon modification	Hexon amino acid sequence	Fiber (knob domain; F)	E1 transgene (for nonreplicative)
Ad5	None (wild-type Ad5)	...F ₂₆₅ FSTTEATAGNGDNLTPKV ₂₈₄ ...	Ad5 or Ad3 (F5/3)	GFP-Luc
Ad5HVR5-6H	Ad5 capsid with 6-histidine insertion into HVR5 of hexon protein	...F ₂₆₅ FSTTLGSHHHHHHGLSLTPKV ₂₈₄ ...	Ad5	GFP-Luc
Ad5H3	Ad5 capsid with substitution for Ad3 hexon	First 55 aa of Ad5 hexon, then Ad3 hexon aa sequence, then last 45 aa of Ad5 hexon	Ad5 or Ad3 (F5/3)	GFP-Luc
Ad5HBAP	Ad5 capsid with biotin acceptor protein (BAP) inserted into HVR5 of hexon protein	...F ₂₆₅ FSGST ...82 aa BAP sequence ...GTPKV ₂₈₄ ...	Ad5 or Ad3 (F5/3)	GFP-Luc
Ad48	None (wild-type Ad48 capsid)	Wild-type Ad48	Ad48	None

with at least four animals per group. Statistical analysis was carried out using Student's *t*-test, or for multiple comparisons, ANOVA and Tukey's pairwise comparison using SAS software. Statistical significance was set at $P < 0.05$.

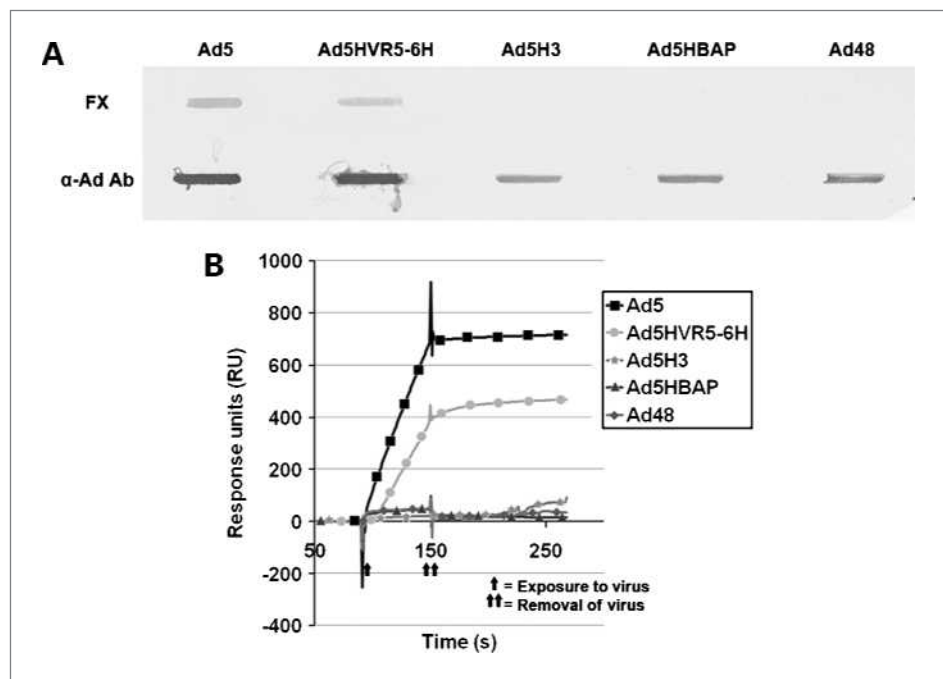
Results

Modification of the Ad5 capsid hexon protein ablates adenovirus binding to human coagulation factor X

To further investigate the interaction of FX with the adenovirus capsid we examined several adenovirus mutants that either had 6-histidine (6-His) peptide insertions

into the Ad5 hexon protein or substitution of the Ad5 hexon protein with that of another adenovirus serotype, Ad3 (Ad5H3). Characteristics of the vectors used in this study are presented in Table 1. Slot blot analysis (Fig. 1A) showed binding of FX to wild-type Ad5 hexon containing virus (Ad5), as well as to an Ad5 virus with a 6-His peptide inserted into HVR5 (Ad5HVR5-6H). No detectable binding was detected between an Ad5-based capsid with the Ad3 hexon-substituted capsid (Ad5H3) or with two of the previously published viruses that do not bind FX, Ad5HBAP, and Ad48. These FX-adenovirus interactions were further confirmed by surface plasmon resonance (SPR) analysis (Fig. 1B). Purified human factors X

Figure 1. Substitution of Ad3 hexon onto Ad5 capsid ablates binding to FX. A, slot blot assay. Viruses bound to PVDF membrane were incubated with FX or with polyclonal antiadenovirus capsid antibody. Ab, antibody. B, SPR analysis of FX binding to hexon variants. Purified human FX was covalently coupled to a streptavidin biosensor chip. Adenoviral vectors were then exposed to the chip and binding interactions were analyzed. B, overlay sensograms representative of two replicates at 37°C (additional replicates were done at 25°C showing similar results; data not shown).



and XI were covalently bound to biocore sensor chips, and viruses were exposed to each protein with differential binding measured. Wild-type Ad5 and Ad5HVR5-6H both showed specific binding to FX, whereas no specific binding was observed for Ad5H3, Ad5HBAP, or Ad48. Additional adenovirus mutants were evaluated with these initial assays but failed to show any significant decrease in binding to FX compared with wild-type Ad5 (data not shown).

Hexon modifications decrease factor X-mediated uptake *in vitro*

To investigate whether the biochemical binding FX to our adenovirus mutants observed in the slot blot and SPR analysis translated to differences in cell transduction we examined the effect of preincubation of adenoviral vectors with FX prior to *in vitro* cell infection. Adenoviral vectors were preincubated with increasing concentrations of purified human FX and then allowed to infect either A549 (high CAR expression), HepG2 (liver representative, moderate CAR expression), or SKOV3 (low CAR expression) cells. Luciferase transgene expression was measured and normalized for transgene expression of adenoviral vectors without preincubation with FX (Fig. 2). Ad5 vectors with wild-type hexon (Ad5) had an increase in transgene expression with increasing FX concentrations as would be expected, and this effect was most notable in the SKOV3 cells secondary to their low CAR expression resulting in a low level of Ad5 transduction without the FX. Ad5HVR5-6H exhibited some

increased transduction with FX preincubation but to a lesser extent than Ad5. Ad5H3 and Ad5HBAP showed little to no change in transduction with preincubation with FX, but Ad5H3 did exhibit some increase in transduction at the highest levels of FX (16 $\mu\text{g}/\text{mL}$), whereas physiological levels were around 8 $\mu\text{g}/\text{mL}$.

Modification of adenovirus hexon decreases liver transduction in mice

The biodistribution of our panel of vectors was investigated in C57BL/6 mice (Fig. 3A). The mice were injected with adenoviral vectors via tail vein injections, and at 36 hours organs were harvested and luciferase transgene expression was assayed. Ad5H3 and Ad5HBAP exhibited approximately 100-fold decrease in liver transduction, comparable with Ad5 in warfarin pretreated mice (Ad5 + warf). Ad5HVR5-6H had a slight decrease in liver transduction, but to a lesser extent than Ad5H3 and Ad5HBAP. There were no other significant variations in the biodistribution of the other vectors with the exception of an increase in the Ad5HVR5-6H vector to the spleen and a decrease in the Ad5 plus warfarin to the spleen.

Substitution of Ad3 hexon onto Ad5 capsid results in improved tumor to liver transgene ratio

Flank tumors using skov3.ip1 cells were established in athymic nude (*nu/nu*) mice. Following establishment of tumors adenoviral vectors were injected via tail vein injections. Luciferase transgene transcript copy numbers in liver and tumors were measured by quantitative PCR

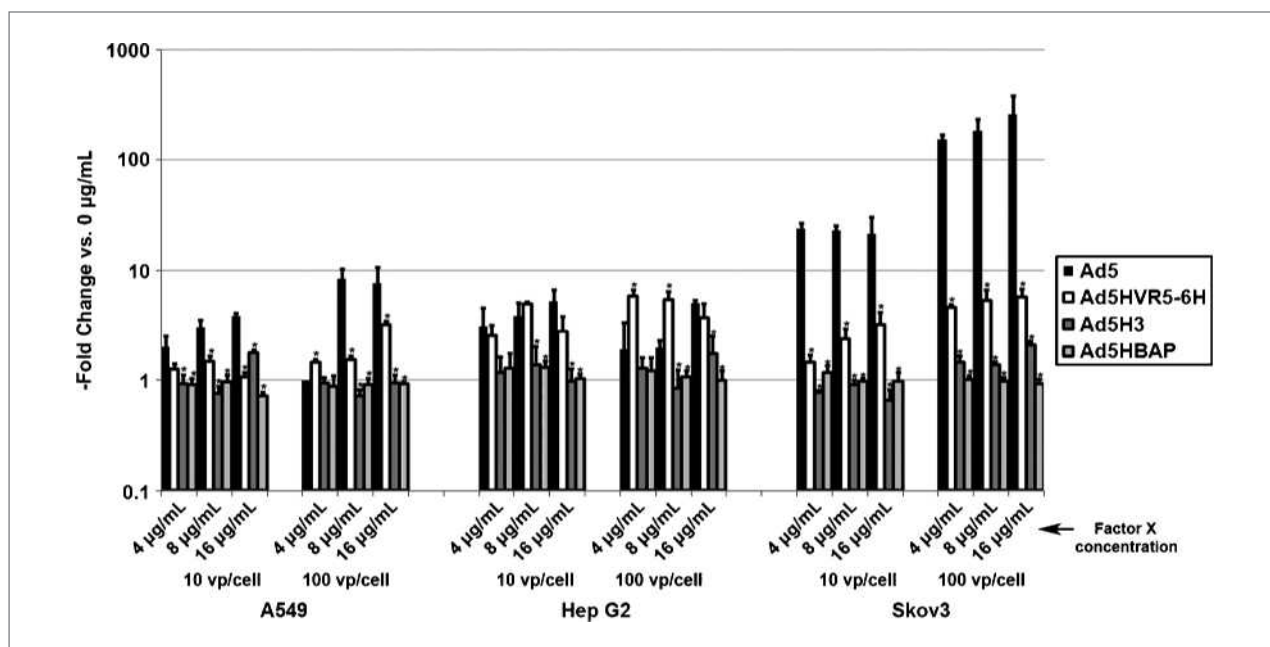


Figure 2. Hexon modifications decrease FX-mediated uptake *in vitro*. Adenoviral vectors were preincubated with increasing concentrations of FX and then allowed to infect either A549 (high CAR expression), HepG2 (liver representative, moderate CAR expression), or SKOV3 (low CAR expression) cells. Luciferase transgene expression was measured at 36 hours and normalized for protein and divided over vectors without preincubation with FX. Ad5 vectors with wild-type hexon (Ad5) had an increase in transgene expression with increasing FX concentrations. *, $P < 0.05$ compared with Ad5.

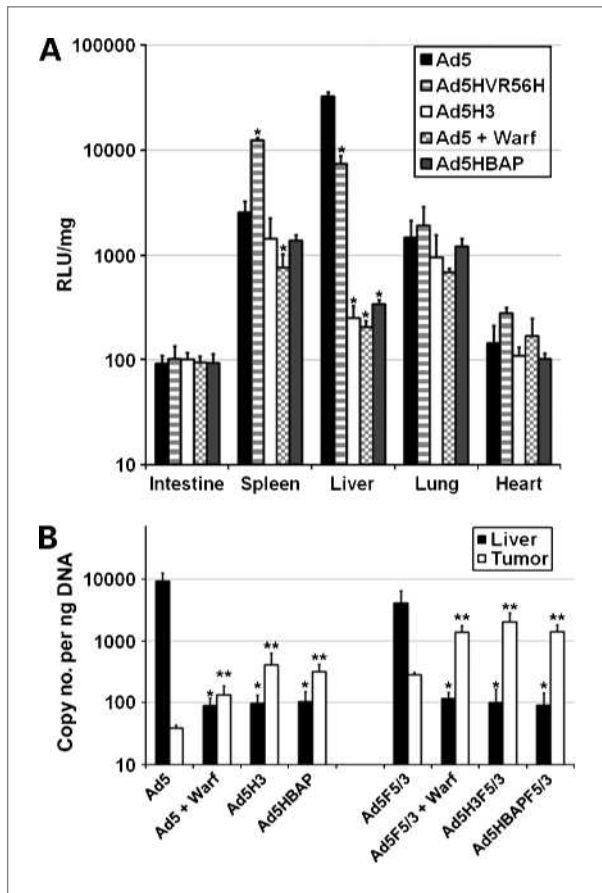


Figure 3. Substitution of Ad3 hexon onto Ad5 capsid results in improved biodistribution in mice. **A**, modification of Ad5 hexon decreases liver transduction in mice. C57BL/6 mice were injected with adenoviral vectors and at 36 hours organs were harvested and luciferase transgene expression was assayed. RLU, relative light unit. *, $P < 0.05$. **B**, substitution of Ad3 hexon onto Ad5 capsid results in improved tumor to liver transgene ratio. Athymic nude mice with skov3.ip1 flank tumors underwent tail vein injections with adenoviral vectors. Luciferase transgene transcript copy numbers in liver and tumors were measured by quantitative PCR and normalized for genomic β -actin DNA. Furthermore, comparison was made between infectivity-enhanced vectors with substitution of Ad3 fiber knob domain (right grouping) and vectors with Ad5 fiber knob domain (left grouping). *, $P < 0.05$ compared with liver control; **, $P < 0.05$ compared with tumor control.

and normalized for genomic β -actin DNA (Fig. 3B). Two panels of vectors were investigated: (a) vectors with wild-type Ad5 fiber protein (left grouping of Fig. 3B, Ad5, Ad5 + warf, Ad5H3, Ad5HBAP), and (b) infectivity-enhanced vectors with substitution of Ad3 fiber knob domain (right grouping of Fig. 3B, Ad5/3, Ad5/3 + warf, Ad5H3F5/3, Ad5HBAPF5/3). Vectors with wild-type Ad5 hexon proteins (Ad5 and Ad5F5/3) without warfarin showed the highest levels of transcript copy numbers in the liver with a 100-fold decrease in luciferase transcript copy numbers when pretreated with warfarin or with hexon modification (Ad5H3 and Ad5H3F5/3, and Ad5HBAP and Ad5HBAPF5/3). The infectivity-enhanced

vectors (right grouping of Fig. 3B) that had a hexon modification or were administered to animals pretreated with warfarin resulted in a 10-fold increase in luciferase transcript copy numbers in tumors compared with vectors with wild-type Ad5 fiber knob domains (left grouping of Fig. 3B).

Incorporation of hexon modification into conditionally replicative adenovirus context ablates factor X binding but does not impair cell killing

The Ad3 hexon substitution was translated to a conditionally replicative adenoviral (CRAd) vector. The backbone of the vector contained the δ 24 mutation (24 bp deletion) in the E1A protein, thereby conferring replication to pRb-deficient cells only and a deletion of the E3 region with preservation of the E3 protein ADP. Two series of CRAds were constructed: (a) one set with wild-type Ad5 hexon or with Ad3 hexon, and (b) one set with the wild-type Ad5 fiber or with the Ad3 fiber knob domain in place of the Ad5 fiber knob domain (Table 1). SPR analysis of human FX binding to these CRAd vectors was done to ensure the ablation of FX binding (Fig. 4A). The same procedure done earlier was used to evaluate the CRAd vectors and it further confirmed that substitution of Ad3 hexon ablated binding to FX. To ensure the substitution of the Ad3 hexon was compatible with cell killing, a crystal violet assay was done to investigate the cytolytic effect of the CRAds (Fig. 4B). There was no impairment of cell killing by substitution of the Ad3 hexon onto the CRAd capsids (Fig. 4Bii and Biii, right samples) versus wild-type Ad5 hexon (Fig. 4Bii and Biii, left samples). Skov3 cells, which are low in CAR expression, were used to further illustrate the infectivity enhancement seen with substitution of the Ad3 fiber knob domain onto CRAd vectors (Fig. 4Biii), which resulted in approximately 100- to 1,000-fold increases in cell killing compared with CRAds with Ad5 fiber knob (Fig. 4Bii).

Adenovirus hexon modification by substitution of the Ad3 hexon onto Ad5 capsid improves antitumor efficacy after systemic vector administration

To evaluate the efficacy of these infectivity-enhanced CRAds *in vivo* we used a s.c. flank tumor xenograft model and administered the CRAds via tail vein injection. We established Skov3.ip1 flank xenograft tumors in athymic nude mice. The vectors (10^{11} vp) were administered i.v. by tail vein injection (day 0), with one group of animals being pretreated with warfarin at the time of injection as a control. The mice injected with the Ad3 hexon-containing vectors (Ad5 Δ 24E1H3F5/3; Fig. 5A, light gray diamonds) had significantly smaller flank tumors through 28 days postinjection compared with placebo-treated animals (Fig. 5A, black squares). The Ad5 hexon-containing vector (Ad5 Δ 24E1F5/3; Fig. 5A, dark gray triangles) showed a mild decrease in tumor size immediately after injection of vector but had negligible size differences at each time point through 28 days compared with placebo (PBS)-treated animals. Mice that

were pretreated with warfarin (Ad5 Δ 24E1F5/3 + warfarin; Fig. 5A, gray circles) at the time of injection showed a decrease in tumor size for early time points, but this effect was only temporary as the warfarin wore off and FX levels were replenished.

The tumor sizes correlated with viral replication as measured by E4 copy numbers in liver and tumor specimens (Fig. 5B). The wild-type Ad5 hexon-containing vector (Ad5 Δ 24E1F5/3) had the highest levels of virus present in the liver at day 3 with very low E4 copy numbers in the tumor specimens. The warfarin-treated mice and Ad3 hexon-substituted vectors showed an improvement in replication of virus in tumor compared with liver. Viral replication in the tumors of the warfarin-pretreated mice was only significantly measurable up to day 7, whereas CRAd replication persisted with the Ad3 hexon-substituted vectors until between 14 and 21 days. It should also be noted that persistent virus in the liver would not be expected as human adenovirus does not

replicate in mice tissues. These data further show that effective i.v. administration of adenoviral vectors can be achieved by appropriate ablation of adenovirus binding to FX.

Discussion

Substitution of the Ad3 hexon onto the Ad5 capsid confers ablation of binding to FX and thereby decreases liver tropism and enhances the antitumor therapeutic window. Herein, we have shown that these hexon-substituted vectors do not bind FX, which translates into decreased liver transduction and improved tumor transduction. Furthermore, by ablating FX binding and decreasing liver sequestration the effects of infectivity enhancement are improved, as shown by improved tumor transduction with the Ad3 fiber knob modified vectors in conjunction with hexon modification.

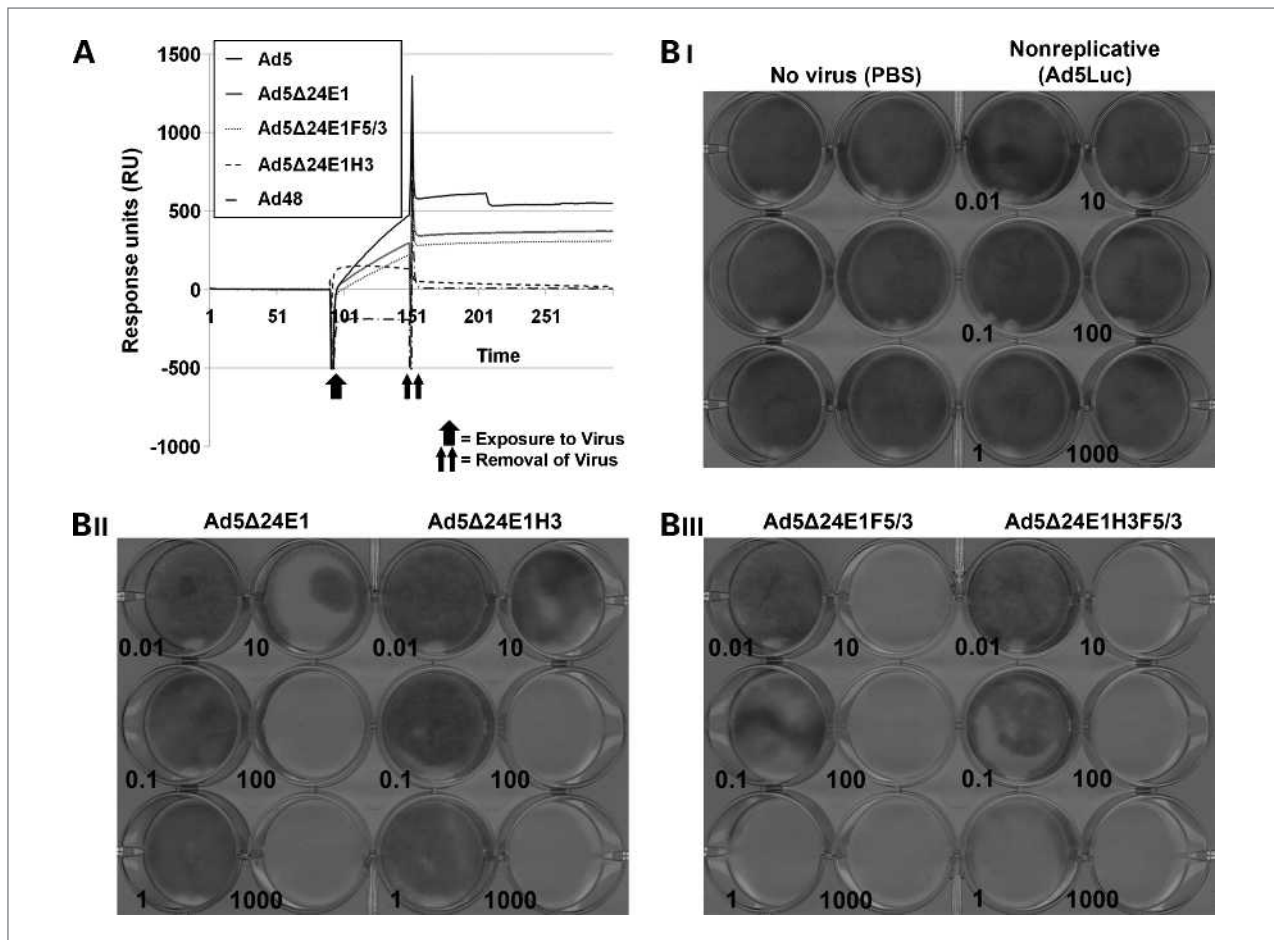


Figure 4. Incorporation of hexon modification into CRAd context ablates FX binding but does not impair cell killing. **A**, SPR analysis of purified human FX binding to CRAd vectors. Purified human FX was covalently coupled to a streptavidin biosensor chip. Adenoviral vectors were then exposed to the chip and binding interactions were analyzed. **A**, overlay sensograms representative of two replicates done at 37°C. **B**, crystal violet assay showing no impairment of cell killing by substitution of Ad3 hexon onto CRAd capsid. Cytocidal effect of vectors on skov3.ip1 cells was assayed using crystal violet dye assay at 10 days postinfection (vp/cell ratio in bottom left corner of sample). Images presented are representative of three replicates done for each vector and dose.

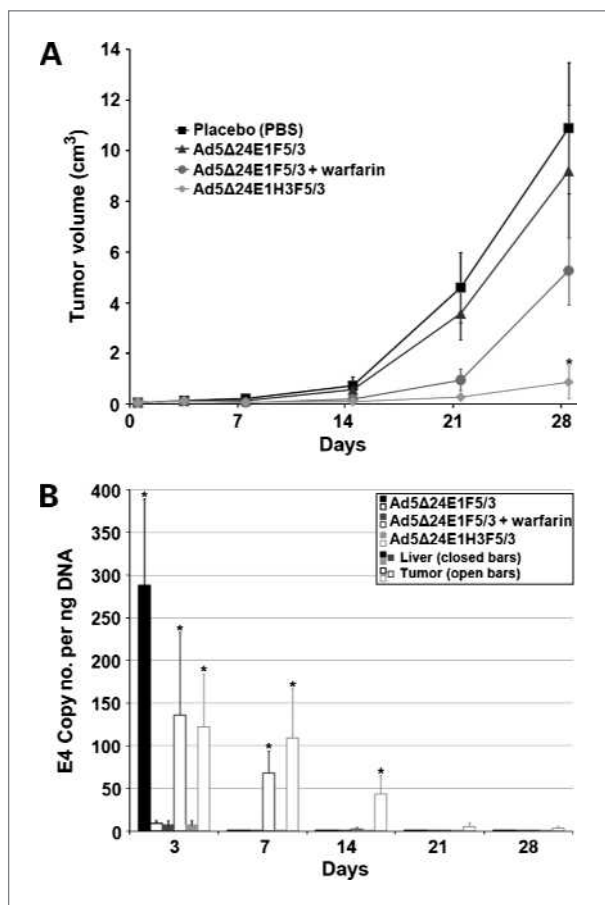


Figure 5. Hexon modification results in effective systemic targeting of oncolytic adenovirus. Athymic nude mice with skov3.ip1 s.c. flank tumors were treated with the panel of infectivity enhanced CRAds. A, tumor size was followed up to 28 days. Ad5Δ24E1F5/3, wild-type Ad5 hexon-containing vector (dark gray triangle); Ad5Δ24E1F5/3 + warfarin, wild-type Ad5 hexon-containing vector pretreated with warfarin at days -3 and -1 (gray circles); Ad5Δ24E1H3F5/3, Ad3 hexon-substituted vectors (light gray diamonds). B, viral replication was measured by E4 copy number in liver and tumor specimens. Closed bars, liver; open bars, tumor. *, $P < 0.05$ compared with placebo.

It was surprising that the Ad5HVR5-6H vector bound to FX and did not exhibit any decreased liver transduction based on previously published findings on the role of HVR5 in binding to FX. After further investigation, however, it is likely that the lack of ablation of binding was due to a combination of factors. Some of these factors are that enough of the residues in HVR5 responsible for binding to FX were preserved (8), HVR7 remained intact, and the peptide insertion was insufficient to confer a conformational change in the hexon protein. Previously published HVR5 mutants did not preserve the threonine amino acid at position 269 (which was preserved in our vectors) and included insertions of at least 8 amino acids (6–8).

The treatment of animals with warfarin during the administration of adenoviral vectors has been previously

shown to result in improved antitumor efficacies (11, 12). The lack of a lasting effect of the warfarin-treated animals with the CRAd agent containing the wild-type Ad5 hexon is likely because the progeny virus contains the wild-type Ad5 hexon and can bind to FX, which is repleted after dosing with warfarin wears off and therefore the circulating FX would help clear the progeny virus. In our experiments therapeutic doses of warfarin were only administered immediately prior to the initial adenovirus injection.

In this study we substituted the Ad3 hexon onto the Ad5 capsid. Therefore, it remains to be clearly determined if the ablation to binding FX of this vector results from an intrinsic lack of binding by the Ad3 hexon or if it is the conformational change from incorporation of this foreign hexon into the Ad5 capsid that ablates any binding to FX. Although previous experiences have noted a decrease in viral titer production with this modification, we were able to generate adequate viral titers (1×10^{12} vp/mL) for our experimental purposes with comparable 50% tissue culture infective doses (data not shown).

Overcoming the obstacles to safe and effective systemic administration of adenoviral vectors is essential in advancing an effective anticancer adenovirus-based therapy. Several key components of achieving this have been recently elucidated, notably the liver tropism mediated by Ad5 hexon binding to coagulation FX. There are now several published adenovirus hexon mutants that show ablation of FX binding with resultant decreased liver tropism. Which of these mutants is the “best” will depend on further comparison analysis and the result may differ for the particular application. Beyond its role in binding to FX, hexon serves as one of the major capsid proteins and modification of it may lead to vector instability. However, modifications may also serve multiple purposes beyond ablating FX binding, such as retargeting through peptide insertions or changing the immunogenicity of the vector.

Three major approaches will likely be advanced for FX-ablating hexon modifications. Two strategies retain the majority of the Ad5 hexon: (a) minimal point mutations of the required residues in HVR5 and HVR7 to minimally affect the vector capsid yet ablate FX binding, and (b) insertion of peptides into HVR5 and/or HVR7 that ablate FX binding and allow further benefits such as retargeting, epitope display, or avoidance of immune recognition. The third (c) entails hexon pseudotyping with adenovirus serotypes that do not bind FX. The first approach has been successfully achieved by substitution of three residues (two in HVR5 and one in HVR7) with corresponding residues from Ad26 (8). The second approach has been initiated by some groups, but the full potential of such has not been completely shown. The third approach involves substitution of the majority of the hexon protein with that of an alternate serotype that ablates FX binding and potentially provides additional benefits such as decreased immunogenicity for repeat vector administration.

The findings presented herein offer another strategy for adenovirus hexon modification for ablation of FX binding and decreasing liver tropism. Furthermore, the synergistic effect of combining infectivity enhancement strategies with decreased liver sequestration provide promise that effective targeting and therapeutic effects following i.v. administration can be achieved. To apply adenoviral vectors for several systemic disease targets, including metastatic cancer, will necessitate effective i.v. administration. Several obstacles will have to be overcome to achieve this end; these include decreasing liver toxicity and sequestration of injected vector, improving target cell transduction, and decreasing vector-immune system interactions. Adenovirus pseudotyping of the hexon and fiber proteins with alternate serotypes, as

presented here, may offer the opportunity to overcome several of these obstacles.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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