ABSTRACT

The employment of conditionally replicative adenoviruses (CRAds) constitutes a promising alternative for cancer treatment; however, in the case of esophageal adenocarcinoma (EAC) the lack of an appropriate tumor-specific promoter and relative resistance to adenovirus infection have hampered the construction of CRAds with clinically applicable specificity and efficacy. By combining transcriptional targeting with infectivity enhancement for CRAds, we generated novel cyclooxygenase-2 (Cox-2) promoter-controlled replicative viral agents for the treatment of EAC. We used infectivity enhancement based on incorporation of an RGD-4C motif into the HI loop of the adenoviral (Ad) fiber knob domain as well as replacement of the Ad5 knob with the Ad3 knob. The Cox-2 promoter was highly active in EAC, whereas showing no significant activity in Cox-2-negative cell lines and primary cells isolated from normal mouse esophagus and stomach. Evaluation of infectivity-enhanced vectors revealed that the transduction and virus-cell binding ability of Ad5/Ad3-chimeric Cox-2 promoter-driven CRAds are selective and potent agents for the treatment of EAC.

INTRODUCTION

The incidence of esophageal adenocarcinoma (EAC) has increased rapidly since the mid-70s and is now the fastest expanding type of cancer in Western countries (1–5). The highest rate of increase was nearly 10% per year in the United States (6). Although the reason for this rapid rise is unclear, it is well known that 10% of patients with gastroesophageal reflux disease eventually develop Barrett’s esophagus, a metaplasia of the lower esophageal epithelium, resulting with a 1% annual risk of developing adenocarcinoma (7, 8). Improvements in the diagnosis of EAC at earlier stages as well as the treatment of late-stage disease remain insignificant, whereas the overall 5-year survival rate is <10% (9, 10). In this regard, novel oncolytic virus approaches using the most advanced therapeutic vectors may offer tent agents for the treatment of EAC.

Ad5/Ad3-chimeric Cox-2 promoter-driven CRAds are selective and potent agents for the treatment of EAC. This article must therefore be hereby marked in accordance with 18 U.S.C. Section 1734 solely to indicate this fact.

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Adenoviruses for Esophageal Adenocarcinoma Treatment

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The use of conditionally replicative adenoviruses (CRAds) for the treatment of esophageal adenocarcinoma (EAC) is hampered by the lack of an appropriate tumor-specific promoter and relative resistance to adenovirus infection. To address these limitations, novel vectors were developed that incorporate an RGD-4C motif into the HI loop of the adenoviral fiber knob domain and replace the Ad5 fiber knob with the Ad3 fiber knob. The Cox-2 promoter, which is highly active in EAC, was used to drive CRAd replication. These vectors were found to be highly selective in EAC cells, with minimal activity in normal cell lines and primary cells isolated from normal mouse esophagus and stomach. The use of Cox-2 promoter-controlled CRAds offers a promising alternative for the treatment of EAC.
mediated by the binding of the RGD-modified Ad fiber protein to integrins, which are overexpressed on the surface of target cancer cells (51, 52). Another strategy to achieve CAR-independent cell entry includes fiber-knob chimerism where the fiber knob of Ad5 is replaced with that of another Ad type (53, 54). Notably, the Ad type 3 (Ad3) receptor, the identity of which has not yet been established (55), has been shown recently to be a better target for Ad vectors designed to treat ovarian cancer, squamous cell carcinoma of the head and neck, and renal cancer (56–59).

In this study, we applied these strategies of Ad tropism modification to derive novel Cox-2 promoter-based CRAds as therapeutic agents for EAC. Importantly, we demonstrate that the oncolytic potency of the Cox-2 CRAds transcriptionally targeted to EAC can be augmented significantly by directing them to the Ad3 receptor. This modality of combining transcriptional targeting and Ad5/Ad3 modification to increase the therapeutic index provides a powerful and promising approach to develop oncolytic viruses for EAC.

**MATERIALS AND METHODS**

**Cell Lines.** OE19 and OE33 human EAC lines were obtained from the European Collection of Cell Culture (CARM Centre for Applied Microbiology and Research, Wiltshire, United Kingdom) and were grown in DMEM (Mediatech, Herndon, VA). TE7 (human EAC; Institute of Development, Aging and Cancer, Tohoku University, Sendai, Japan), A549 (Cox-2 positive lung cancer cell line; American Type Culture Collection (ATCC), Manassas, VA), and BT474 (Cox-2 negative breast cancer cell line; ATCC) cells were cultivated in RPMI 1640 (Mediatech). The medium for BT474 was supplemented with bovine insulin (0.01 mg/ml; Life Technologies, Inc., Rockville, MD). Human ovarian adenocarcinoma cell line SKOV3.ipi (Ad3 receptor-positive control, kindly provided by Dr. Janet Price, M.D. Anderson Cancer Center, Houston, TX) was propagated in DMEM/F12, 50:50 (Mediatech). Chinese hamster ovary cells (Ad3 receptor-negative control) were grown in HAM’s F-12 medium. Human hepatoma cell line HepG2 (CAR positive control; ATCC), transformed human embryonic retina cell line 911 (a kind gift from Dr. Alex J. van der Eb, Leiden University, Leiden, the Netherlands), and 293 cells (ATCC) were maintained in DMEM. Each medium was also supplemented with 10% fetal bovine serum (FBS; HyClone, Logan, UT), 2 mm l-glutamine and penicillin (100 IU/ml), and streptomycin (100 mg/ml). Human umbilical vein endothelial cells (HUVEC) were obtained from ATCC and cultured in EGM-2 medium (Cambrex Biosciences, Walkersville, MD). All of the cells were incubated at 37°C in a humidified environment with 5% CO2.

**Adenoviral Vectors.** Replication-incompetent adenoviral vectors (AdCox2L, AdCox2MLuc, and AdCMVLuc) encoding the firefly luciferase (Luc) reporter gene were constructed as described previously (30). Ad5/3F (sense) 5' CATTAGGAATTCGAACAGGC3' and Ad5/3F (antisense) 5' CTACATTAATGACCACTGCGAAGC3' were used as nonselective replicative control vectors.

**Validation of Viral Constructs.** Several critical vector components of the CRad genomes were confirmed by PCR. Viral DNA was isolated from purified virions using a Blood DNA kit (Qiagen), and aliquots corresponding to 101 genomes were analyzed by PCR. Thermal cycling conditions were: initial denaturation (5 min at 95°C), followed by 25 cycles of amplification (30 s at 95°C, 30 s at 55°C, 1 min at 72°C), and final extension (10 min at 72°C). The regions detected and expected band size were as follows: wild-type Ad5 (485 bp), E1a (338 bp), Cox-2 promoter (405 bp), forward direction CRAd (392 bp), Ad5 fiber knob region (wild-type fiber: 247 bp and RGD fiber: 274 bp), and Ad5/Ad3 fiber knob region (293 bp). For the Ad5/Ad3 fiber knob region, the following primers were used: Ad5/3F (sense) 5' CCTCTTATTGGTGACACCAAGGC3' and Ad5/3F (antisense) 5'C ATCATAAATTGGGACAGATTGTTG3'. Other primer sequences were already reported (31).

**Analysis of Cox-2 RNA Levels.** Total cellular RNA was extracted from semiconfluent cell cultures in 10-cm dishes using the RNeasy mini RNA extraction kit (Qiagen). Cox-2 and glyceraldehyde-3-phosphate dehydrogenase RNAs were detected in a semiquantitative manner using the Gene Amp Gold RNA PCR Core kit (Perkin-Elmer, Branchburg, NJ) as described previously (31).

**Flow Cytometry.** Cultured cells were trypsinized, washed with PBS, centrifuged, and resuspended in ice-cold PBS supplemented with 1% BSA and 0.1% NaN3. Cells (106) were incubated for 1 h at 4°C with mouse antibodies RMcB to CAR (1:1000; produced using hybridoma purchased from ATCC), P1F6 to ov integrin (1:1000), P1F6 to ov integrin (1:1000), P4G11 to β1 integrin (1:50), or HA5 to ε5β1 integrin (1:100; Chemicon, Temecula, CA). The cells were then washed twice with PBS/BSA/NaN3 buffer and incubated with secondary FITC-labeled goat anti-mouse IgG serum (1:50; Sigma). After another PBS/BSA/NaN3 rinse, 2.5 μg/ml propidium iodide (Sigma) was added to sort out dead cells from the sample. The cells (105) were analyzed immediately by flow cytometry at the University of Alabama at Birmingham FACS Core Facility.

**Promoter and Gene Delivery Analysis with Luciferase Expression Ads.** EAC cells grown in 24-well plates were infected with Ad vectors at a multiplicity of infection of 50 pfu/cell. The infection medium was replaced.
with the appropriate growth medium 2 h later. Two days after infection, the cells were lysed with cell culture lysis buffer (Promega, Madison, WI), and Lue activity was determined with the Luciferase Assay System (Promega). Experiments were performed in triplicate and standardized with protein concentration quantitated by the DC protein assay (Bio-Rad, Hercules, CA).

The Cox-2 promoter selectivity was also analyzed in mouse esophageal and gastric primary cells. The esophagus and stomach of a Balb/C mouse were aseptically isolated and cut by scissors into small pieces, followed by wash with PBS twice. The tissues were dispersed with Liver Digestion Medium (Life Technologies, Inc.) for 3 h at 37°C, rocking every 15 min. After adding 25% volume of FBS to stop digestion, the cells were centrifuged and resuspended in 20% FBS DMEM/F12 containing 1 μM dexamethasone. The cells were plated and infected as described above.

**In Vitro Analysis of Cytoidal Effect by Crystal Violet Staining.** To analyze virus-mediated cell killing, 25,000 cells/well were plated in 12-well plates and infected with the viruses in 200 μl of growth medium containing 5% FBS at an multiplicity of infection of either 0.01 or 0.1 vp/cell. After 3 h, 1 ml of the growth medium was added. Two days later, the infection medium was replaced with 1% FBS medium. After 12 days (18 days for TE7 cell line) of cultivation, the cells were fixed with 10% buffered formalin for 10 min and stained with 1% crystal violet in 70% ethanol for 20 min, followed by washing with water and drying.

**Virus Binding Assay.** To analyze virus-cell binding, 50,000 cells/well cultivated in 24-well plates were infected with 2000 vp/cell in 100 μl of DMEM and incubated for 1 h at 4°C. At this temperature, virus internalization would not occur. After washing with PBS three times, cells were scrapped from the plates and processed with a QIAamp Blood Mini kit (Qiagen). The isolated DNA was analyzed by real-time PCR analysis to determine the Ad DNA copy number with E4-specific primers at the University of Alabama at Birmingham Gene Therapy Center Correlative Laboratories as described before (31). For the standard curve, E4 template DNA with known copy numbers (10^6, 10^5, 10^4, and 10^3) was used. All of the PCR reactions were carried out using the Light Cycler System (Roche Molecular Biochemicals, Indianapolis, IN) as recommended by the manufacturer. Thermal cycling conditions were: initial denaturation for 10 min at 95°C, and 40 cycles of 15 s at 95°C and 1 min at 60°C. Data were analyzed with the Light Cycler software. The primers were designed to detect a 68 bp-long sequence within E4 region: forward primer (5′GGAGTGGCCGCGAGACA3′, nucleotides 34007–34027) and reverse primer (5′ACTAGTCGGGCTTCTAC3′, nucleotides 34074–34056) and 6-carboxyfluorescein labeled probe (6-carboxyfluorescein-TGGCATGACACTAC-FBS medium. On the next day, 100

the number of living cells was determined by a colorimetric method using the Cell Titer 96 Aqueous Non-Radioactive Cell Proliferation Assay (Promega) as described by the manufacturer. Absorbance was measured at a wavelength of 490 nm in an E-Max spectrophotometer (Molecular Device Corporation, Sunnyvale, CA), and standard curves were generated by analyzing the known number of live cells. On the basis of this curve, the number of living cells was calculated.

**Viral Progeny Production Assay.** To determine production of virus progeny, 250,000 cells seeded in 12-well plates were infected with Ad vectors at a multiplicity of infection of 1 vp/cell in 100 μl of 5% FBS medium. On the next day, 100 μl of the growth medium containing 1% FBS was added. The cells were incubated for 12 days (18 days for TE7), and the number of surviving cells was analyzed by a colorimetric method using the Cell Titer 96 Aqueous Non-Radioactive Cell Proliferation Assay (Promega) as described by the manufacturer. Absorbance was measured at a wavelength of 490 nm in an E-Max spectrophotometer (Molecular Device Corporation, Sunnyvale, CA), and standard curves were generated by analyzing the known number of live cells. On the basis of this curve, the number of living cells was calculated for the experimental groups using the SOFTmax computer software (Molecular Device Corporation). All of the experiments were repeated in triplicate.

**In Vivo Antitumor Effect in an EAC Xenograft Model.** Female ncr/nu nude mice (Frederick Cancer Research, Frederick, MD; 6–8 weeks of age) were used to establish EAC xenografts. OE19 cells (2.4 × 10^6 per injection site) were inoculated into the flanks of mice. When the nodules reached a size of 8–10 mm in maximum diameter, a single virus dose (10^9 vp in 100 μl PBS) of CRAds or control viruses was injected intratumorally. The condition of the mice was monitored daily, and the tumor diameter was measured twice a week with calipers. The tumor volume was calculated using the formula: tumor volume = width^2 × length/2. In accordance with institutional approved animal experimental protocols, mice were euthanized 16 or 21 days after viral injection due to the overgrowth of tumors in the control group. All of the animals received humane care based on the guidelines set by the American Veterinary Association. All of the experimental protocols involving live animals were reviewed and approved by the Institutional Animal Care and Use Committee of the University of Alabama at Birmingham.

**Statistical Methods.** Statistical analysis of CRAd efficacy in vitro and in vivo was performed with a two-tailed t test. Data are expressed as a mean ± SD of at least three sets of results. Results were considered statistically significant when P < 0.05.

**RESULTS**

**Confirmation of CRAd Structure.** All of the CRAd vector structures were confirmed by PCR analysis of the viral DNA. None of the vectors was contaminated with nonselective replication-competent Ad resulting from spontaneous homologous recombination during vector propagation in E1-transcomplementing cell lines (Fig. 2A). The CRAds possessed both the E1a gene and the Cox-2 promoter sequence as part of their E1 expression cassettes (Fig. 2, B and C). A 392-bp fragment corresponding to the forward direction cassette was amplified from CRAdCox2F, RGDCRAdCox2F, and 5/3CRAdCox2F with a sense primer recognizing the left inverted terminal repeat sequence and an antisense primer recognizing the Cox-2 promoter sequence (Fig. 2D). In the analysis of the fiber region, vectors with wild-type Ad5 fiber (CRAdCox2F and CRAdCox2R) produced a 247-bp signal, whereas vectors with the RGD-modified fiber (RGDCRAdCox2F and RGDCRAdCox2R) yielded a 274-bp band (Fig. 2E), the increased size of which represents the 9 amino acid insertion in the HI loop. Ad5/Ad3-chimeric CRAds (5/3CRAdCox2F and 5/3CRAdCox3R) amplified with 5/3F primers showed a 293-bp signal, which confirms the presence of the Ad3 knob in the fiber (Fig. 2F). These data validate the structure and stability of the CRAd constructs.

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**Fig. 2.** Confirmation of conditionally replicative adenoviruses (CRAd) structure by PCR. The structure of the viral DNA was analyzed using sets of primers corresponding to several important regions of the virus. A, detection of wild-type adenovirus (Ad)5 (485 bp). Primers recognizing the wild-type Ad5 left-end sequence did not amplify any fragments from any of the six cycloxygenase (Cox)–2 CRAds. B, detection of the E1a sequence (338 bp). All of the CRAd vector structures contain the E1a sequence because they possess Cox-2 promoter-driven E1 expression cassettes. C, detection of the Cox-2 promoter sequence (405 bp). All of the CRAd vector structures contain the Cox-2 promoter sequence to drive E1 expression. D, direction of the E1 expression cassette. E, Ad5 fiber structure. Primers designed to distinguish the presence of an RGD motif in the HI loop of the Ad fiber knob domain amplified a shorter 247-bp fragment from CRAdCox2F and CRAdCox2R. The same primers amplified a longer 274-bp fragment containing a 9 amino acid insertion from RGDCRAdCox2F and RGDCRAdCox2R. For all gels, lane 1, CRAdCox2F; lane 2, CRAdCox2R; lane 3, RGDCRAdCox2F; lane 4, RGDCRAdCox2R; lane 5, 5/3CRAdCox2F; and lane 6, 5/3CRAdCox2R.
The Cox-2 promoter is strong and selective for EAC. To analyze the activity of the Cox-2 promoter in Ads, two Luc expression vectors with two different lengths of the Cox-2 5' upstream control region (Cox2M and Cox2L) were tested in OE19, OE33, and TE7 EAC cells, Cox-2 negative HUVEC and BT474 cells, and primary cells isolated from normal mouse esophagus and stomach (Fig. 4). The Cox2M promoter showed significantly higher activity in EAC cells compared with the Cox2L promoter. In all three of the EAC cell lines, Cox2M and Cox2L showed high transgene expression levels relative to CMV promoter-driven Luc expression, whereas showing no significant activity in HUVEC, BT474, and primary mouse esophageal and gastric cells.

Expression of CAR and Integrins αβ3, αβ5, αβ1, and α5β1 on Human EAC Cell Lines. To establish the biological basis for the relative resistance of EAC to Ad5 infection, we evaluated the level of CAR and integrins on the cell surface using antibodies specific for CAR and integrins. We found that EAC cell lines were clearly negative for CAR expression (Fig. 5). Flow cytometry also demonstrated that EAC cells express low levels of αβ3 and αβ5 integrins (Fig. 5), whereas αβ1 and α5β1 integrins were present on the cell surface in rather large amounts (Table 1).

Analysis of Transduction, Binding, and Oncolytic Efficiency of RGD- and 5/3-Modified Vectors in EAC. To determine whether incorporation of an RGD-4C motif into the HI loop of the Ad fiber knob domain or replacement of the Ad5 knob with the Ad3 knob would enhance the infectivity of Ad vectors in EAC cells, three identical replication-incompetent CMV promoter-driven Luc expression vectors with wild-type (Ad5Luc1), RGD-modified (Ad5RGDLuc1), and Ad5/Ad3-chimeric (Ad53Luc1) fibers were used (Fig. 6). OE19, OE33, and TE7 EAC cell lines demonstrated significantly higher levels of transgene expression with the RGD-modified vector (2.7-5.7-fold higher than that of Ad5Luc1) and even greater levels with the Ad5/Ad3 chimera (5.4-87.2-fold higher transgene expression in comparison with Ad5Luc1 and 2.0-15.2-fold higher relative to Ad5RGDLuc1). The infectivity enhancements of RGD-modified and Ad5/Ad3 vectors in each cell line were consistent regardless of the viral dose during infection (50 and 500 vpcell; data not shown).

To evaluate virus-cell binding, the cells were incubated with non-selective wild-type Ad5 vectors, including Ad5WT (unmodified fiber), RGDWT (RGD-modified fiber), and Ad5MG553 (Ad5/Ad3-chimeric fiber) for 1 h at 4°C. The isolated total cellular DNA was analyzed by real-time PCR to determine the bound adenoviral DNA copy number.
modified Ads outperformed unmodified Ads, whereas chimeric vectors
cassette (CRAdCox2F, RGDCRAdCox2F, and 5/3CRAdCox2F) exhib-
ting intact (Fig. 9). Whereas the Cox-2-negative control cell line and HUVEC remained
staining showed oncolysis after infection with all six of the CRAds,
cycles of viral replication. In all three of the EAC cell lines, crystal-violet
thelial cells with a low titer (0.1 vp/cell) of each virus to allow multiple
lines, Cox-2-negative cell line BT474, and human umbilical vein endo-
selective oncolytic effect of Cox-2 CRAds, we infected EAC cells
CRAds with a Cox-2 promoter-driven E1 expression cassette. To assess
replicative virus by constructing RGD-modified and Ad5/Ad3-chimeric
tional targeting and Ad tropism modification were combined into a
CRAds
in Vitro
Table 1 Expression of αv, β1, and α5β1 integrins on surface of EAC

<table>
<thead>
<tr>
<th>Integrin</th>
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<th>OE33</th>
<th>TE7</th>
<th>293*</th>
<th>A549b</th>
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<td>34.79</td>
<td>36.03</td>
<td>74.54</td>
<td>84.44</td>
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<tr>
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<td>60.34</td>
<td>55.43</td>
<td>28.87</td>
<td>69.36</td>
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<tr>
<td>α5β1</td>
<td>64.92</td>
<td>10.24</td>
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* αv and α5β1 integrins positive control.
* b αvβ1 and β1 integrins positive control.

In all three of the EAC cell lines, the numbers of viral DNA copies with AdMG553 were significantly higher than that of AdWt and the RGD counterpart (Fig. 7), correlating with the gene transfer data. The RGD-modified vector showed enhanced binding in OE19 and TE7 cells, whereas no binding improvement was observed in OE33 cells.

Furthermore, we analyzed the oncolytic potency of the above nonselective wild-type vectors. We infected OE19, OE33, and TE7 cells with 0.01 vp/cell of each virus and stained the fixed surviving cells with crystal violet 12 days after infection (Fig. 8). In all of the EAC cell lines, infectivity-enhanced viruses demonstrated a stronger cytidal effect compared with unmodified AdWt. The enhancement in cell killing effect was much more evident in cells infected with the Ad5/Ad3 chimera.

**Increased Oncolytic Efficiency of Infectivity-Enhanced Cox-2 CRAds in Vitro.** The established concept of Cox-2 promoter transcriptional targeting and Ad tropism modification were combined into a replicative virus by constructing RGD-modified and Ad5/Ad3-chimeric CRAds with a Cox-2 promoter-driven E1 expression cassette. To assess the selective oncolytic effect of Cox-2 CRAds, we infected EAC cell lines, Cox-2-negative cell line BT474, and human umbilical vein endothelial cells with a low titer (0.1 vp/cell) of each virus to allow multiple cycles of viral replication. In all three of the EAC cell lines, crystal-violet staining showed oncolysis after infection with all six of the CRAds, whereas the Cox-2-negative control cell line and HUVEC remained intact (Fig. 9A). CRAds with the left to right direction E1 expression cassette (CRAdCox2F, RGDCRAdCox2F, and 5/3CRAdCox2F) exhibited stronger oncolytic effect than CRAds with the reverse direction cassette (CRAdCox2R, RGDCRAdCox2R, and 5/3CRAdCox2R). RGD-modified Ads outperformed unmodified Ads, whereas chimeric vectors performed even better. The cytotoxicity of Ad5/Ad3 CRAds was similar to that of the replicative control viruses containing the wild-type early genes (Ad5Wt, RGDWt, and AdMG553); however, these Ad5/Ad3 Cox-2 CRAds maintained their replication specificity because they did not affect BT474 and HUVEC. Nonreplacive controls (Ad5SLuc1, RGDAd5Luc1, and 5/3Ad5Luc1) did not cause any oncolysis.

The replication and cytidal effects of CRAds were additionally confirmed with a quantitative cell viability assay (Fig. 9B). In EAC cell lines, the quantitative cytotoxicity assay showed oncolysis with all of the Cox-2-based CRAds, whereas no cytidal effect was observed in BT474 cells and HUVEC. In OE19, OE33, and TE7 cell lines, the percentages of surviving cells after infection with RGDCRAdCox2F compared with uninfectected cells were 64.7%, 50.6%, and 81.4% (P = 0.0369, 0.0019, and 0.0338, respectively), whereas viability percentages after treatment with RGDCRAdCox2R were 77.5%, 70.4%, and 90.9% (P = 0.0231, 0.0045, and 1.746, respectively). When Ad5/Ad3 CRAds were used for infection, only 26.0%, 18.5%, and 25.1% (5/3CRAdCox2F; P = 0.0001, 0.0002) and 20.8%, 37.6%, and 25.2% (5/3CRAdCox2R; P = 0.0006, 0.0002, and 0.0003) of cells remained alive, respectively. In comparison with unmodified CRAds, RGD CRAds showed unremarkable increase in killing in two of three EAC cell lines (OE19 and OE33). In contrast, Ad5/Ad3 vectors demonstrated significantly improved oncolysis in all three of the cell lines, and this effect was superior to that of RGD-modified CRAds.

**Increased Viral Progeny Production and Selective Replication of Infectivity-Enhanced CRAds in Vitro.** We next determined whether the increased cytotoxic effect of infectivity-modified CRAds correlated with increased Ad replication and progeny production. We infected OE19 (EAC cells) and BT474 (Cox-2 negative cells) with 1 vp/cell of each virus. Twelve days after infection, the cells and media were harvested, and the viral titer was determined by a plaque assay on 911 cells (Fig. 10). In OE19, RGDCRAdCox2F and RGDCRAdCox2R produced progeny titers of 1.27 × 10⁸ and 2.42 × 10⁹ pfu/ml, which were 3.0 and 5.7 times higher than those of their unmodified counterparts CRAdCox2F and CRAdCox2R, respectively. The final titers of the Ad5/Ad3 CRAds (5/3CRAdCox2F and 5/3CRAdCox2R) were 2.27 × 10⁹ and 2.53 × 10⁹ pfu/ml, respectively, which were 53.3 and 59.4 times higher relative to fiber-unmodified CRAds, and 17.9 and 10.48 times higher in comparison with RGD CRAds. This experiment also demonstrated that 5/3CRAdCox2F replicated in OE19 with an efficiency similar to that of the wild-type replicative control. In the BT474 Cox-2-negative cell

![Table 1 Expression of αv, β1, and α5β1 integrins on surface of EAC](image)

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* αv and α5β1 integrins positive control.
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![Fig. 6. Transduction efficiency in EAC cells with infectivity-enhanced Ad vectors.](image)

Human EAC cell lines (OE19, OE33, and TE7) and four control cell lines (A549 integrin positive), HepG2 (CAR positive), SKOV3.ip.1 (Ad3 receptor positive), and Chinese hamster ovary (CHO; integrin positive, CAR and Ad3 receptor negative) were infected with the Ad5/Ad3 chimera.

![Fig. 7. Binding of infectivity-enhanced adenovirus (Ad) vectors to EAC cells.](image)

Human EAC cell lines (OE19, OE33, and TE7) and two control cell lines [SKOV3.ip.1 (Ad3 receptor positive) and Chinese hamster ovary (CHO; integrin positive, CAR and Ad receptor negative)] were infected with AdWt, RGDWt, and AdMG553 for 1 h at 4°C. Isolated total DNA was analyzed by real-time PCR to determine the Ad DNA copy number. Whereas the RGD-modified vector displayed slightly enhanced binding in two of three EAC cells (OE19 and TE7), DNA copy number for Ad5/Ad3-chimeric AdMG553 was significantly higher than that of unmodified AdWt and the RGD counterpart in all three of the EAC cell lines: P < 0.05 (*), P < 0.005 (**), P < 0.001 (**), Bars, ±SD from triplicate experiments.

![Fig. 8. Effect of infectivity-modified CRAds on the survival of each EAC cell line.](image)

In OE19, OE33, and TE7 cell lines, the percentages of surviving cells after infection with RGDCRAdCox2F compared with uninfectected cells were 64.7%, 50.6%, and 81.4% (P = 0.0369, 0.0019, and 0.0338, respectively), whereas viability percentages after treatment with RGDCRAdCox2R were 77.5%, 70.4%, and 90.9% (P = 0.0231, 0.0045, and 1.746, respectively). When Ad5/Ad3 CRAds were used for infection, only 26.0%, 18.5%, and 25.1% (5/3CRAdCox2F; P = 0.0001, 0.0002) and 20.8%, 37.6%, and 25.2% (5/3CRAdCox2R; P = 0.0006, 0.0002, and 0.0003) of cells remained alive, respectively. In comparison with unmodified CRAds, RGD CRAds showed unremarkable increase in killing in two of three EAC cell lines (OE19 and OE33). In contrast, Ad5/Ad3 vectors demonstrated significantly improved oncolysis in all three of the cell lines, and this effect was superior to that of RGD-modified CRAds.
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enhanced viruses demonstrated stronger oncolytic effect than unmodified AdWt. The increase in cell killing effect was superior with Ad5/Ad3-chimeric AdMG553.

line, the replication of Cox-2 CRAds was 200–450-fold less than in the
EAC OE19 cell line, whereas nonselective wild-type viruses
(AdWt, RGDWt, and AdMG553) generated higher titers of virus
progeny in BT474 than in EAC cells.

Therapeutic Efficacy of Infectivity-Enhanced CRAds in Vivo. In vivo analysis of antitumor efficacy of CRAds was performed using
an OE19 s.c. xenograft model in nude mice. Established tumors were
treated with a single intratumoral injection of 10^{10} vp of each virus, and the tumor size was monitored. On day 16 after injection, the
CRAdCox2F group (relative tumor volume 5.83 ± 0.89; n = 6) and the
CRAdCox2R group (5.96 ± 1.85; n = 6) showed tumor growth
suppression, but the effect was not statistically significant in compari-
tion to the untreated group (9.37 ± 6.68; n = 5) and the group that
received a nonreplicative Luc expression vector (7.80 ± 3.02; n = 7;
Fig. 11). Although RGDCRAdCox2F (day 16; 5.49 ± 2.06; n = 7) and
RGDCRAdCox2R (day 16; 5.75 ± 2.72; n = 7) showed slightly
stronger oncolytic effect than unmodified CRAds, no statistically
significant difference was found between these groups and the non-
replicative Ad groups. In contrast, the sizes of the tumors treated with
chimeric 5/3CRAdCox2F (day 16; 3.11 ± 2.12; n = 8) and
5/3CRAdCox2R (day 16; 4.30 ± 1.39; n = 6) were significantly
reduced compared with Ad5Luc1, unmodified CRAds, and their RGD
counterparts. On day 21, both Ad5/Ad3 CRAds yielded therapeutic
effects that were superior to other CRAds [relative tumor volumes:
5/3CRAdCox2F (2.52 ± 0.82; n = 8) versus CRAdCox2F
(7.26 ± 1.05; n = 5; P < 0.00001) and RGDCRAdCox2F
(6.22 ± 3.1; n = 7; P < 0.01); and 5/3CRAdCox2R (5.17 ± 2.3;
versus CRAdCox2R (7.23 ± 4.08; n = 5; P > 0.05) and
RGDCRAdCox2R (6.03 ± 1.85; n = 4; P > 0.05)]. Notably in this
experiment, 5/3CRAdCox2F was even more effective than the wild-
type replicative control AdMG553, which also contains the Ad5/Ad3
surface modification (4.35 ± 1.85; n = 7; P < 0.05). Thus, the in vivo
data indicate that whereas the unmodified Cox-2 CRAds and the RGD
counterparts demonstrated minimal in vivo therapeutic effect, the

Fig. 8. Oncolytic potency of tropism modified vectors in EAC. Human EAC cell lines
OE19, OE33, and TE7) were infected with 0.01 vp/cell of AdWt, RGDWt, or AdMG553.
Twelve days later, the cells were fixed and stained with crystal violet. Data are repre-
sentative samples from at least three experiments performed on each cell line. Infectivity-
enhanced viruses demonstrated stronger oncolytic effect than unmodified AdWt. The increase in cell killing effect was superior with Ad5/Ad3-chimeric AdMG553.

Fig. 9. In vitro cytocidal effect of infectivity-
enhanced cyclooxygenase (Cox)-2 conditionally replic-
itive adenoviruses (CRAds) in EAC. A, human
EAC cell lines (OE19, OE33, and TE7), Cox-2-
negative cell line BT474 and human umbilical vein
endothelial cells (HUVEC) were infected with
CRAds at a multiplicity of infection of 0.1 vp/cell.
Adenoviral cytotoxicity was analyzed by crystal vi-
olet staining. CRAds with the forward direction E1
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olet staining. CRAds with the forward direction E1
conditions (CRAdCox2F) exhibited stronger oncolytic effect than
RGDCRAdCox2R (6.03 ± 1.85; n = 4; P > 0.05). Notably in this
experiment, 5/3CRAdCox2F was even more effective than the wild-
type replicative control AdMG553, which also contains the Ad5/Ad3
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Adenoviral cytotoxicity was analyzed by crystal vi-
olet staining. CRAds with the forward direction E1
cassette (F) exhibited stronger oncolytic effect than
reverse-direction CRAds (R). RGD-modified CRAds
outperformed unmodified vector, whereas Ad5/Ad3
CRAds performed even better. Data are representa-
tive samples from at least three experiments per-
formed on each cell line. B, in parallel, cell viability
was determined with a colorimetric cell proliferation
assay. The results are shown as percentages of living
cells remaining relative to uninfected cells. All of the
Cox-2-based CRAds demonstrated oncolytic killing
in all three of the EAC cell lines, whereas BT474 and
HUVEC remained intact. In comparison to unmod-
ified CRAds, RGD CRAds demonstrated unremark-
ably higher oncoly in two of three EAC cell lines
(OE19 and OE33). In contrast, Ad5/Ad3 CRAds exhibited significantly improved oncoly in all
three cell lines: P < 0.05 (**), P < 0.005 (**),
P < 0.001 (***). Bars, ±SD from four experiments.
was 200-fold higher than in BT474. In OE19 cells, the replication of Cox-2 CRAds produced significantly higher titers of infectious particles compared with unmodified CRAds. OE19 s.c. xenografts in nude mice were treated with a single intratumoral injection of $10^{10}$ vp of each virus. The tumor size was measured and shown as relative volume compared with day 0. Mice from the untreated group and the groups treated with nonreplicative control viruses had to be sacrificed on day 16 due to excessive tumor size. Compared with the nonreplicative control on day 16, both Ad5/Ad3 CRAds ($5/3$CRAdCox2F and $5/3$CRAdCox2R) significantly suppressed the size of established tumors whereas the unmodified CRAd showed significant antitumor effect. On day 21 the chimeric $5/3$Cox2CRAdF demonstrated even stronger oncolytic effect than the wild-type replicative control AdMG553.

**DISCUSSION**

Advanced EAC is a highly aggressive disease with a poor long-term prognosis (3, 5, 8–10). Curative treatment consists mostly of surgery, which is not feasible for all patients; distant metastases, local irresectability, and poor condition of the patient allow only 20–30% to be potential candidates for surgical resection (63, 64). Therefore, new therapeutic approaches such as oncolytic viruses are needed to provide an effective treatment for irresectable EAC. Because EAC is easily accessible with the guidance of an endoscope, we developed our CRAd agents with the intention of applying direct intratumoral injection. However, even in the case of direct injection, ectopic transgene expression and liver toxicity remain major problems. In the setting of EAC, blood of the lower esophagus is collected into the portal vein, which flows directly to the liver. Therefore, mitigation of liver toxicity is extremely important. In this study, we applied CRAds based on Cox-2 promoter selectivity and enhanced their infectivity to develop novel therapeutic agents for EAC.

Tumor-specific promoters are actively being explored (24–32). However, many promoters, which show the desired properties in plasmid vectors, may become promiscuous in the context of a viral genome (23, 30). In our previous studies, the Cox-2 promoter was determined to be a promising tumor-specific promoter for many gastrointestinal cancers and demonstrated high levels of transgene expression in pancreatic, gastric, and colon cancers in vitro and in vivo, whereas maintaining low activity in normal tissues, including the liver (28–31). In this study, the utility of the Cox-2 promoter for EAC was initially determined by using two Luc expression vectors with two different lengths of the Cox-2 5' upstream control region (AdCox2LLuc and AdCox2MLuc). Both vectors showed high transgene expression levels comparable with the CMV promoter-driven vector in all of the EAC cell lines, whereas showing tumor specificity as indicated by minimal activity in HUVEC, the Cox-2-negative BT474 breast cancer cell line, and primary cells isolated from normal mouse esophagus and stomach (Fig. 4). These results were in agreement with the reverse transcription-PCR analysis, which indicated high Cox-2 mRNA transcription levels in EAC cells (Fig. 3). These findings clearly demonstrate the high activity and selectivity of the Cox-2 promoter in the tested EAC cells as well as their fidelity in an Ad backbone. Because the 1491-bp Cox2L promoter showed lower background expression in the liver compared with the 942-bp Cox2M promoter (30), we selected the Cox2L promoter for CRAd construction.

Viral replication and spread are the key factors of oncolytic efficacy. The most crucial parameter of efficient viral spread is the infection ability of the viral progeny, because transduction of surrounding cancer cells is required for amplification and oncolysis throughout the tumor (12, 52). In this regard, the natural CAR deficiency in various tumors (65) not only limits the initial infection event but also affects the potential therapeutic effect of CRAds afforded by viral replication downstream (66). We and others have reported that the efficacy of CRAds may be enhanced by incorporating a RGD-4C motif into the HI loop of the Ad fiber knob domain, which would allow the virus to bind to integrins overexpressed frequently on target cancer cells (31, 52). The native tropism of Ad vectors may also be altered by substituting the Ad5 knob with knobs from different Ad serotypes (53, 54). In particular, recent studies demonstrated that vectors displaying the Ad5 shaft/Ad3 knob were superior to Ad5

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**Fig. 11. Superiority of adenovirus (Ad5/Ad3-chimeric cyclooxygenase (Cox)-2 conditionally replicative adeno-viruses (CRAd) in an in vivo human esophageal adenocarcinoma model.** OE19 s.c. xenografts in nude mice were treated with a single intratumoral injection of $10^{10}$ vp of each virus. The tumor size was measured and shown as relative volume compared with day 0. Mice from the untreated group and the groups treated with nonreplicative control viruses had to be sacrificed on day 16 due to excessive tumor size. Compared with the nonreplicative control on day 16, both Ad5/Ad3 CRAds ($5/3$CRAdCox2F and $5/3$CRAdCox2R) significantly suppressed the size of established tumors whereas the unmodified CRAd showed significantly stronger antitumor effect than the unmodified, RGD-modified, and reverse-direction counterparts: $P < 0.05$ (asterisk), $P < 0.005$ (double asterisk), $P < 0.001$ (triple asterisk). Bars, ±SD from triplicate experiments.

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vectors in infection of ovarian cancer, squamous cell carcinoma of the head and neck, and renal cancer (56–59).

Before analyzing whether RGD-modified or Ad5/Ad3-chimeric vectors would enhance Ad transduction efficiency in EAC, we first explored the biological basis of EAC refractoriness to Ad infection by evaluating human EAC cell lines for CAR and integrin expression. We found that the profound paucity of the primary Ad receptor in OE19, OE33, and TE7 cells is the major factor limiting transduction efficiency of Ad5 vectors in these cells (Fig. 5). Interestingly, the integrins αvβ3 and αvβ5, which have been described previously as predicting factors for RGD-modified virus infectivity in many gastrointestinal cancers (67), were found to be limited on the surface of EAC cells as well. On the other hand, αv, β1, and αβ1 integrins, also known to bind to the RGD motif of fbronectin (68–70), were highly expressed on EAC cells (Table 1). The presence of these receptors provides the rationale for the RGD-retargeting strategy based on the Arg-Gly-Asp motif for specific recognition of integrins. In fact, a subsequent gene transfer assay, comparing RGD-modified vectors to unmodified ones, demonstrated higher levels of transgene expression after infection with RGD vectors (Fig. 6).

The identification of the Ad3 cell surface receptor is still under investigation (55). Although it is possible to analyze the Ad3 receptor expression by flow cytometry using the Ad3 knob protein (59), a binding analysis performed with viral particles should represent functional binding during infection better than an assay using only the fiber knob of the virus. Therefore, we developed a novel approach to analyze the effect of infectivity enhancement in the context of virus-cell binding. The data obtained in this experiment demonstrated that whereas an RGD vector yielded remarkable improvements in binding, the number of bound Ad5/Ad3-chimeric virus was significantly higher than an unmodified vector and an RGD version in all three of the EAC cell lines (Fig. 7). These data suggest that the Ad3 receptor is highly expressed on human EAC cells and may be a better target for infecting EAC cells. Importantly, these binding data correlated with results from the Luc gene transfer assay (Fig. 6) and oncolytic potency experiment using the infectivity-enhanced vectors (Fig. 8).

Because promising results were obtained with both RGD and Ad5/Ad3 targeting strategies, we combined Cox-2 promoter transcriptional selectivity with infectivity enhancement by constructing RGD-modified and Ad5/Ad3-chimeric Cox-2-based CRAds. The Cox-2 promoter was incorporated into the E1 expression cassettes of these CRAds in both forward (F) and reverse (R) directions (Fig. 1). During vector amplification, the Cox-2 CRAds propagated consistently regardless of the presence of fiber modifications, and the yield was comparable with the E1-deleted vectors. The resulting viruses maintained their correct genomic structures and were free from nonselectively replicative recombinants (Fig. 2), indicating their feasibility for clinical grade vector production.

For any therapeutic agent to be clinically useful, both safety and efficacy are required. Therefore, to characterize the safety and selectivity of Cox-2 CRAds we used assays to compare their killing, replication, and progeny production potential in human EAC cells versus control Cox-2-negative cells BT474 and/or HUVECs. In vitro cytotoxicity studies revealed that the Cox-2 CRAds replicated and caused oncolysis in all three of the EAC cell lines but not in BT474 and HUVEC (Fig. 9), thus confirming the dependence of their replication on the Cox-2 promoter activity. The CRAds with the left to right direction E1 expression cassette (F) showed stronger cytotoxic effect than those with the reverse direction (R) version. Overall, the cytotoxicity of infectivity-enhanced CRAds was stronger than that of unmodified vectors. As predicted by the binding assay, the Ad5/Ad3-chimeric CRAds exhibited significantly improved oncolysis similar in degree to the wild-type control in all three of the EAC cell lines compared with the unmodified and RGD-modified vectors. We suspect that low expression of αvβ3 and/or αvβ5 integrins (known to be the secondary receptors for adenoviral internalization; Ref. 50) on EAC cells may be the reason why RGD-enhanced CRAds have been not so effective in killing EAC. To additionally characterize the oncolytic property of our CRAds, we performed a virus progeny production assay. The results indicate that in human EAC OE19 cells, the replication of Cox-2 CRAds was 200–450-fold greater than in BT474 Cox-2-negative cells, whereas nonselective wild-type replicative viruses generated similar high titers of virus progeny in both OE19 cells and in BT474 cells (Fig. 10). On the basis of the cytotoxicity and progeny production experiments, the Cox-2 CRAds maintained replication control specificity in vitro. Of note, our data seem to show a correlation between increased cytopathic effect of infectivity-enhanced CRAds and their replication and progeny production ability. Specifically, in OE19 EAC cells the RGD Cox-2 CRAds (F and R) produced 3.0 and 5.7 times higher pfu titers than their unmodified counterparts, whereas the final titers of the Ad5/Ad3 CRAds were 53.3- and 59.9-fold higher, respectively (Fig. 10). Altogether, these results clearly confirm that infectivity-enhanced Cox-2 CRAds were able to specifically replicate in and kill EAC cells in vitro and that the chimeric CRAds with Ad5 shaft/Ad3 knob were superior in transduction and replication in EAC cells.

To assess whether Cox-2 CRAds can cause oncolysis in cancer cells in vivo, the therapeutic effect of unmodified, RGD-modified, and Ad5/Ad3 CRAds was analyzed and compared in an EAC xenograft model. To allow multiple cycles of replication along with destruction of tumor cells to occur, we treated OE19 xenografts with a single s.c. injection of each virus. In vivo data revealed that on day 16, both Ad5/Ad3 CRAds (F and R) significantly suppressed the tumor growth of established OE19 xenografts, whereas their RGD counterparts showed only slightly stronger oncolysis compare to unmodified vectors, which exhibited remarkable therapeutic abilities (Fig. 11; day 16). Impressively, on day 21 the chimeric Cox-2 CRAd with the forward direction E1 cassette (5/3CRAdCox2F) statistically outperformed both unmodified CRAdCox2F and RGD-modified RGDCRAdCox2F, reflecting achievement of better infection with the Ad5/Ad3 fiber. Furthermore, the anti-tumor effect of the forward-direction 5/3CRAdCox2F was greater than that of the reverse-direction 5/3CRAdCox2R or wild-type replicative control virus AdMG553 (Fig. 11; day 21). The evident superiority in cytopathic effect of 5/3CRAdCox2F in vivo suggests the therapeutic utility of this agent for the treatment of EAC.

In summary, we have shown the feasibility of the Cox-2 promoter and the applicability of RGD- and Ad5/Ad3-based fiber modification for EAC. We have combined transcriptional targeting with infectivity enhancement for CRAds and demonstrated that Ad5/Ad3-chimeric Cox-2 CRAds are effective and selective therapeutic agents for EAC. Thus, these novel CRAds have great clinical potential for the treatment of EAC.

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Infectivity-Enhanced Cyclooxygenase-2-Based Conditionally Replicative Adenoviruses for Esophageal Adenocarcinoma Treatment

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