ADENOVIRAL GENE THERAPY FOR NON-SMALL CELL LUNG CANCER

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Academic Dissertation

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To Meri-Vuokko and Petteri

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LIST OF ORIGINAL PUBLICATIONS

This thesis is based on the following original publications, which are referred to in the text by their Roman numerals.

- I Särkioja M, Kanerva A, Salo J, Kangasniemi L, Eriksson M, Raki M, Ranki T, Hakkarainen T, Hemminki A. Noninvasive imaging for evaluation of the systemic delivery of capsid-modified adenoviruses in an orthotopic model of advanced lung cancer. Cancer 2006; 107:1578-88.
- II Hakkarainen T, Särkioja M, Lehenkari P, Miettinen S Ylikomi T, Suuronen R, Desmond RA, Kanerva A, Hemminki A. Human mesenchymal stem cells lack tumor tropism but enhance the antitumor activity of oncolytic adenoviruses in orthotopic lung and breast tumors. Hum Gene Ther 2007; 18: 627-41.
- III Särkioja M, Hakkarainen T, Eriksson M, Ristimäki A, Desmond RA, Kanerva A, Hemminki A. The cyclo-oxygenase 2 promoter is induced in nontarget cells following adenovirus infection, but an AU-rich 3' untranslated region destabilization element can increase specificity. J Gene Med 2008; 10: 744-53.
- IV Särkioja M, Pesonen S, Raki M, Hakkarainen T, Salo J, Ahonen MT, Kanerva A, Hemminki A. Changing the adenovirus fiber for retaining gene delivery efficacy in the presence of neutralizing antibodies. Gene Therapy 2008; 15: 921–29.

ABBREVIATIONS

5-FU 5-fluorouracil

AAV adeno associated virus

ACE angiotensing-converting enzyme

Ad3 adenovirus serotype 3 Ad5 adenovirus serotype 5 BAd3 bovine Ad type 3

CAR coxsackie-adenovirus receptor

CD cytosine deaminase
CDK cyclin dependent kinase
CEA carcinoembryogenic antigen

CMV cytomegalovirus
Cox-2 cyclooxygenase 2
CXCR4 chemokine receptor
DC dendritic cells

EGF epidermal growth factor

Ep-CAM epithelial cellular adhesion molecule

Fab antibody fragment FGF fibroblast growth factor

FLAG Asparagin, Tyrosine, Lysine, Asparagin, Asparagin, Asparagin,

Asparagin Lysine

GCV Ganciclovir GM growth media

HD vector helper-dependent vector hmSC human mesenchymal stem cell heparan sulfate proteoglycan

HSV-tk herpes simplex virus type I thymidine kinase hTERT human telomerase reverse transcriptase

i.ha intra hepatic artery
i.p intraperitoneal
i.t intratumoral
i.v intravenous
IFN interferon

INK4 inhibitors of cyclin-dependent kinase CDK4

kb kilobase
kD kiloDalton
KS kaposi carcoma
LacZ β-galactocidase
luc firefly luciferase

MAP mitomycin C+doxorubicin + cisplatin
MAPK mitogen-activated protein kinase
Mdr-1 multiple drug resistance 1

MSC mesenchymal stem cell

MTS 3-(4, 5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)- 2-(4-

sulfophenyl-2H tetrazolium,inner salt

Nab neutralizing antibody NF-κB nuclear factor-κB

NGR asparagine-glycine-arginine
NRP human lung surfactant protein D
NSCLC Non-small cell lung cancer

orf open reading frame PEG polyethylene glycol Pfu plaque forming unit

pK polylysin PKR protein kinase R

PLGA polyethylene glycol-cationic lipid and poly (lactic-glycolic) acid

PSA prostate-specific antigen

Rb retinoblastoma

RGD-4C arginine-glycine-aspartatic acid

sCAR soluble form of CAR

SCCA2 squamous cell carcinoma antigen

SCCHN squamous cell carcinoma of the head and neck

SCLC small cell lung cancer

SIGYLPLP Seriini-Isoleucine-Glycine-Tyrosine-Leucine-Proline-Leucine-Proline

 $\begin{array}{lll} SLP1 & secretory leukoprotease inhibitor 1 \\ TAG-72 & tumor associated glycoprotein 72 \\ TCID_{50} & tissue culture infective dose_{50} \\ TK/GCV & thymidine kinase ganciclovir \\ TSP & tumor/tissue specific promoter \\ \end{array}$

TTS dual promoter with hTERT and human surfactant protein A1 promoters.

uPA urokinase plasminogen activator

uPAR urokinase plasminogen activator receptor

UTR untranslated region
VA RNA virus-associated RNA

vp viral particles

ABSTRACT

Adenoviral gene therapy is an experimental approach to cancer refractory to standard cancer therapies. Adenoviruses can be utilized as vectors to deliver therapeutic transgenes into cancer cells, while gene therapy with oncolytic adenoviruses exploits the lytic potential of viruses to kill tumor cells. Although adenoviruses demonstrate several advantages over other vectors - such as the unparalleled transduction efficacy and natural tropism to a wide range of tissues - the gene transfer efficacy to cancer cells has been limited, consequently restricting the therapeutic effect. There are, however, several approaches to circumvent this problem.

We utilized different modified adenoviruses to obtain information on adenovirus tropism towards non-small cell lung cancer (NSCLC) cells. To enhance therapeutic outcome, oncolytic adenoviruses were evaluated. Further, to enhance gene delivery to tumors, we used mesenchymal stem cells (MSCs) as carriers. To improve adenovirus specificity, we investigated whether widely used cyclooxygenase 2 (Cox-2) promoter is induced by adenovirus infection in nontarget cells and whether selectivity can be retained by the 3'untranslated region (UTR) AU-rich elements. In addition, we investigated whether switching adenovirus fiber can retain gene delivery in the presence of neutralizing antibodies.

Our results show that adenoviruses, whose capsids were modified with arginine-glycine-aspartatic acid (RGD-4C), the serotype 3 knob, or polylysins displayed enhanced gene transfer into NSCLC cell lines and fresh clinical specimens from patients. The therapeutic efficacy was further improved by using respective oncolytic adenoviruses with isogenic 24bp deletion in the E1A gene. Cox-2 promoter was also shown to be induced in normal and tumor cells following adenovirus infection, but utilization of 3'UTR elements can increase the tumor specificity of the promoter. Further, the results suggested that use of MSCs could enhance the bioavailability and delivery of adenoviruses into human tumors, although cells had no tumor tropism *per se*. Finally, we demonstrated that changing

adenovirus fiber can allow virus to escape from existing neutralizing antibodies when delivered systemically.

In conclusion, these results reveal that adenovirus gene transfer and specificity can be increased by using modified adenoviruses and MSCs as carriers, and fiber modifications simultaneously decrease the effect of neutralizing antibodies. This promising data suggest that these approaches could translate into clinical testing in patients with NSCLC refractory to current modalities.

1 Introduction

Lung cancer is one of the most common cancers and the leading cause of cancer-related mortality worldwide. It can be divided into two types: small-cell lung cancer (SCLC) and non-small cell lung cancer (NSCLC). Non-small cell lung cancer can be subdivided further into squamous cell carcinoma, adenocarcinoma, and large-cell carcinoma and it accounts for approximately 80% of all lung cancers. A paucity of effective treatments exists for NSCLC patients. Treatments include chemotherapy and surgery, but in most cases the cancer is at advanced stage when it is detected and these treatments are not curative. Consequently, there is a need for new therapies (Ramalingam and Belani 2008). A promising treatment alternative for many types of cancers refractory to current modalities is adenoviral gene therapy. Adenoviruses are excellent vectors for gene therapy since they can infect many types of cells, can be grown to high titers, do not integrate into host genome, and, above all, they have low pathogenicity for humans (Glasgow et al. 2004). Importantly, there are encouraging results in recent randomized trials concerning glioma, prostate, hepatocellular, and head and neck cancers (Immonen et al. 2004; Li et al. 2007; Shirakawa et al. 2007). However, three potential limitations may have an impact on the efficacy of adenoviruses: 1) Adenoviruses do not transduce efficiently clinically relevant cancer cells, since they exhibit variable expression of adenovirus primary receptor coxsackie adenovirus receptor (CAR), 2) Adenoviruses do not have a natural preference to replicate only in tumor cells, 3) Replication may also be inefficient in large tumor masses, which are very complex in structure, and 4) Adenoviruses have a strong immunogenicity that especially affects systemic re-administration of viruses.

To circumvent some of these problems, adenoviruses can be retargeted to cancer cells by improving their abilities to attach to heterologous receptors other than CAR, which can enhance gene transfer to the desired tissues (Bauerschmitz *et al.* 2002). Further, adenovirus replication can be controlled at transcriptional level by specifically engineered deletions in viral genes or tissue-specific promoters controlling crucial replication regulations, such as E1A, resulting in viruses that replicate only in cells where promoters are expressed

(Saukkonen and Hemminki 2004). Replication can also be controlled via posttranscriptional regulation with elements affecting replication at the mRNA level (Ahmed *et al.* 2003; Gahery-Segard *et al.* 1998). To avoid neutralizing antibodies (NAbs) raised by adenoviruses upon re-exposure, different serotypes of adenoviruses can be used, (Hashimoto *et al.* 2005), alternatively, even small changes in the Ad5 fiber knob can allow the virus to escape from pre-existing capsid-specific NAb (Gahery-Segard *et al.* 1998; Rahman *et al.* 2001). Further, using carriers such as mesenchymal stem cells (MSCs), which are not recognized by NAbs, can enhance gene transfer and antitumor efficacy (Studeny *et al.* 2004).

We used capsid-modified adenoviruses to determine the adenovirus tropism towards NSCLC. In addition, we tested whether adenoviruses could be further targeted by regulating gene expression posttranscriptionally. Also, we present some unpublished data suggesting that although gene expression can be beneficially modulated with post-transcriptional control elements in the context of plasmids, the effect was not powerful enough to be useful in the context of recombinant adenoviruses. We also studied the possibility of utilizing MSCs as carriers for oncolytic adenoviruses, thereby combining MSCs with the anti-tumor effect provided by oncolytic adenoviruses. Finally, we investigated whether the changing adenovirus fiber can retain gene transfer efficiency in the presence of adenovirus neutralizing antibodies.

2 Lung cancer

Lung cancer is the second (among men) and fourth (among women) most common cancer in Finland, with 2140 new cases in 2006 (Finnish Cancer Registry; www. cancerregistry.fi). The most common histological type of NSCLC is adenocarcinoma, followed by squamous-cell carcinoma and large-cell carcinoma. The prognosis of NSCLC is poor because the disease is often at an advanced stage when detected. The overall 5-year survival rate is only 8% for men and 12% for women.

Treatment of NSCLC remains a challenge. Surgical resection at on early stage represents the only treatment associated with a high likelihood of attaining 5-year survival. However,

since the NSCLC is often detected at a later stage, combination therapies with radio- and chemotherapy are used.

2.1 Molecular mechanism of NSCLC

Development of cancer, including NSCLC, involves accumulation of multiple molecular abnormalities over a long period of time. These include chromosomal abnormalities, inactivation of tumor suppressor genes, activation of oncogenes, expression of hormone receptors and, production of growth factors. The most common genetic alterations involve mutations in the p53/p14ARF, p16/retinoblastoma (Rb) and in K-ras pathways. These mutations involving tumor suppressor genes may be a rate-limiting event in the development of cancer (Brambilla *et al.* 1999).

2.1.1 Tumor suppressors

p53 is a tumor suppressor gene that is mutated in 2/3 human lung cancers, and it is commonly seen in squamous cell carcinomas. In normal cells, when DNA damage occurs, p53 can halt the cell cycle at the G1 checkpoint, preventing the cycle from proceeding to the S phase until the DNA is repaired or it can induce the cell to apoptosis. In cancer cells, by contrast, when p53 is mutated, the control is lost. The p16/Rb pathway regulates entry and progression through the cell cycle. Disruption in one or more factors in this pathway can lead to tumor genesis. p16 and pRb proteins are critical members of the Rb pathway, and they are mutated in 80% of NSCLC (Belinsky et al. 1998; Tanaka et al. 1998). In normal cells, when cell cycle protein p16 is needed, it functions as an inhibitor of the CDK4/6 cyclin-D complex. This leads to dephosphorylation of protein Rb, which in turn cannot release transcription factor E2F that binds to DNA to stimulate the synthesis of proteins necessary for cell division. No transcription of the genes necessary for the cell cycle to progress from G1 to S phase, therefore occurs. If CDK4/6-cyclin D complex is not under inhibition, it increases phosphorylation of the retinoblastoma protein (pRb), which can then release E2F and transcription will occur and the cell cycle continues from G1 to S phase. In tumor cells with defects in p16 or Rb function or a CDK4/6 cyclin D complex with increased activity, E2F is constitutively active (Jakubczak et al. 2003) (Figure 1.).

a. Normal cell

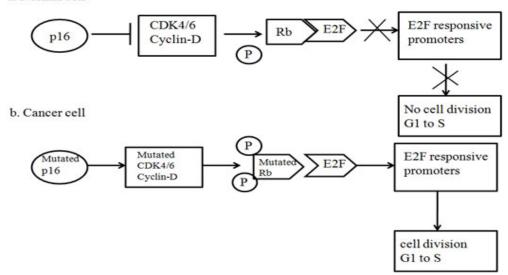


Figure 1. p16/Rb pathway. a) In normal cells, when p16 is needed, it inhibits CDK4/6 Cyclin-D complex that dephosphorylates protein Rb which prevents Rb to interact with E2F and the cell cycle is arrested to G1. **b).** In cancer cells, the E2F transcription factor is constantly active and defects in p16, CDK4/6 Cyclin-D complex or pRb, don't affect on cell division.

2.1.2 Oncogenes

Oncogenes are typically mutated genes that have an important role in cell signaling pathways. They encode proteins that control cell proliferation, apoptosis, or both. They can be activated by structural alterations resulting from a mutation or a gene fusion (Konopka *et al.* 1985; Tsujimoto *et al.* 1985). The most important abnormalities detected are mutations involving the *ras* family of oncogenes. The *ras* oncogene family has three members: H-*ras*, K-*ras*, and N-*ras*. These genes encode a protein on the inner surface of the cell membrane with GTPase activity and may be involved in signal transduction. K-*ras* mutations are most commonly seen in 30% in adenocarcinomas of the lung (Westra *et al.* 1993).

3 Cancer gene therapy

Cancer is a complex disease requiring combinations of several therapeutic modalities to treat it effectively. Gene therapy was originally devised to treat disease by replacing a lost or defective gene. Now it has broadened to encompass a variety of approaches of introducing genetic materials into cells for therapeutic purposes. These strategies can be roughly categorized into two main categories: immunologic and molecular (Heo DS 2002). One of the frequently used genetic immunotherapy strategy involves the transfer of the genes of the immune-stimulant molecules such as cytokines such as *interleukin 12* (IL12) gene to boost T-cell mediated immune response against cancer (Shi *et al.* 2002; Barajas *et al.* 2001). Other approaches includes the manipulation of antigen presenting cells such as dentritic cells to enable them of more active tumor antigen presentation in the tumor tissue (Vollmer *et al.* 2002; Vulink *et al.* 2008) and direct genetic vaccination by the antigenencoding genes (Hanke *et al.* 2002; Grim *et al.* 2000).

Molecular approaches involves suicide conversion gene therapy, use of anti-oncogenes to inhibit oncogene activity, utilization of tumor suppressor genes, anti-angiogenesis gene therapy, transfer the drug resistance genes to overcome the dose-limiting toxicity of traditional chemotherapy, the use of anti-viral vectors to deliver anticancer reagents to the tumor tissue and also adenovirus gene therapy that will be discussed in the chapters 4-6. One of the most investigated suicide gene/prodrug systems is the herpes simplex virus thymidine kinase (HSV-TK)/ganciclovir (GCV) system where non-tocix prodrug GCV is converted to the toxic form. The efficiency of the system has been demonstrated successfully in patients with malignant glioma in a clinical trial where the mean survival of patients was prolonged by 81% when treated with TK/GCV therapy after surgery (Immonen *et al.* 2004).

Anti-oncogenes are mostly targeted against *bcl-2*, *c-myc* genes and *K ras* gene family. Treatments with anti-oncogenes have resulted in reduction in tumor growth both *in vitro* and *in vivo* with melanoma and colon cancer (Hu *et al.* 2001; Duggan *et al.* 2001; Chana *et al.* 2002; Tokunaga *et al.* 2000).

Successful transfection of the functional wt p53 into cancer cells has showed therapeutic outcomes in patients. When combined with radiation, no viable tumor was observed in

63% of patients after three month of the completion of the therapy (Swisher *et al.*2003). The main targets for anti-angiogenesis cancer therapy includes inhibiting angiogenic inducers (namely, vascular endothelial growth factor and angiopoietin) or introducing angiogenic inhibitors such as angiostatin, endostatin, Interleukin-12, and p53 (Chen *et al.* 2001).

One of the main problems with tradiotional chemotherapy is the toxicity to normal cells. To protect cells and to enable the use of larger drug doses, drug resistance genes such as multiple drug resistance gene-1 (*Mdr-1*), can be transferred nto the hematopoietic progenitors (Licht and Peschel 2002; Koc *et al.* 1999).

To improve carriers capacity to deliver anticancer reagents, as well as synthetic oligonucleotides, antibodies and RNA, in addition to therapeutic genes, nonviral vectors have been developed. Mostly used carriers include liposomes and polymers that can be packed with the DNA of desired anticancer agent. Molecules are delivered into cells by endocytosis (Wheeler *et al.* 1996; Sudimak *et al.* 2000; Qian *et al.* 2002).

4 Adenoviral cancer gene therapy

Adenoviruses are probably the most studied viral gene transfer vectors, but other viruses are used as well. Numerous cancer gene therapy studies have been conducted with retroviruses (Cole *et al.* 2005; Thanarajasingam *et al.* 2007), lentiviruses, vaccinia viruses, adeno-associated virus (AAV), and herpes simplex virus to mention but a few (Kirn *et al.* 2007; Mueller and Flotte 2008; Pellinen *et al.* 2004; Tyler *et al.* 2008).

4.1 Adenoviruses

Adenoviruses are nonenveloped DNA viruses. They have a characteristic morphology with an icosahedral capsid, which consists of three major proteins: hexon (II), penton base (III), and a knobbed fiber (IV), along with minor proteins: VI, VIII, IX, IIIa, and IVa2 (Stewart *et al.* 1993). Highly basic protein VII, small peptide *mu*, and a terminal protein (TP) are associated with DNA. Protein V provides a structural link to the capsid via protein VI (Figure 2). Thus far, at least 51 serotypes of human adenovirus have been identified and

classified in to six distinct subgroups, A-F (De Jong *et al.* 1999). Adenovirus serotypes 1-7 cause the most common infections in humans (Arstila 2004). In the context of gene therapy, the most studied adenovirus serotype 5 (Ad5) belongs to subgroup C.

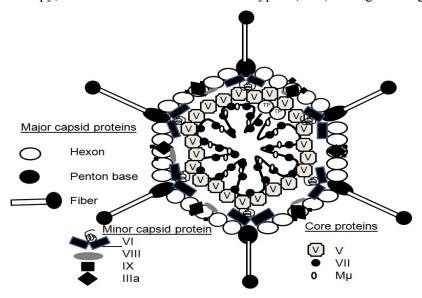


Figure 2. Structure of adenovirus 5 virion. [modified from the figure in (Russell 2000)].

The adenovirus infection cycle can be divided into two phases, early and late, occurring before and after virus DNA replication. The first phase covers the entry of the virus into the host cell (Figure 3), followed by the passage of the virus genome into the nucleus and selective transcription and translation of early viral genes. Early genes are needed for the transcription and translation of late genes, which lead to the assembly of structural proteins and maturation of the infectious virus. Entry of the virus into the host cell is mediated through specific receptors on the target cell. The virus binds to a receptor via the knob portion of the fiber. The receptor for adenoviruses belonging to subgroup A-F, but excluding subgroup B, is known to be CAR, at least *in vitro*. (Roelvink *et al.* 1998). This traditionally accepted mechanism of cell entry is now known to be more complex at least in vivo after systemic delivery and it is not even known in humans. More intense research

is required to reveal virus-host interactions in terms of viral components, blood cell and serum interactions of adenoviral vectors and mechanisms behind the adenovirus tropism (Baker 2007a and b; Nicklin et al. 2005; Waddington et al. 2008). Some adenovirus serotypes seem to have additional binding specificities, suggesting that other receptors besides CAR may be needed for the entry (Segerman et al. 2000). After the interaction of the virus with the receptor, the entry proceeds via clathrin-mediated endocytosis. The key step is the interaction of the penton base arginine-glycine-aspartatic acid (RGD) motif with host cell $\alpha_v \beta$ integrins, leading to endosome formation and internalization. After internalization, adenovirus DNA is transported to the nucleus, which is followed by early gene protein synthesis. The early gene region consists of four genes: E1, E2, E3, and E4. E1 gene products can be subdivided further into E1A and E1B. E1A proteins modulate cellular metabolism in a way that makes cell more susceptible to virus replication. They interfere with, for instance, the cell division processes and the regulation of NF-κB, p53, and pRb proteins. E1B functions cooperatively with E1A by inhibiting apoptosis. E2 is also subdivided into two separate transcription regions; E2A and E2B. Their gene products (including DNA polymerase) are required for DNA replication. E3 gene products are, on the other hand, responsible for defending against the host immune system, enhancing cell lysis, and releasing of virus progeny. E4 gene products mainly facilitate virus RNA metabolism, promote virus DNA replication, and prevent host protein synthesis. Transcription of late genes leads to production of the virus structural components and the encapsidation and maturation of virus particles in the nucleus (Russell 2000).

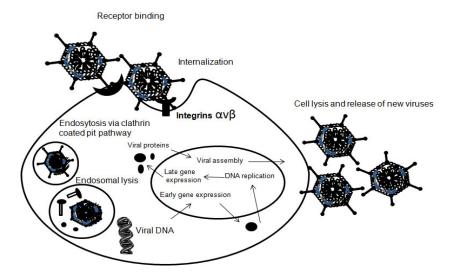


Figure 3. Schematic illustration of the adenovirus 5 infection pathway. Cell entry is initiated by binding of the fiber knob to its primary receptor CAR, which is followed by clathrin-mediated endocytosis. Interaction of the penton base RGD motif with host cell integrins $(\alpha v\beta)$ is needed for the process to proceed. After endosomal lysis, viral DNA is transported to the nucleus, where viral genes are expressed.

Adenoviruses might be useful gene transfer vehicles for treatment of various diseases (Bainbridge *et al.* 2008; Hedman *et al.* 2003; Immonen *et al.* 2004; Li *et al.* 2007; Shirakawa *et al.* 2007). However, clinical trials have shown that uncontrolled, high-dose gene transfer and expression could have side effects (Raper *et al.* 2003). Viruses should therefore be targeted to the preferred tissue on a transductional or transcriptional level. The former implies restriction of viral entry to target cells, whereas the latter involves selective gene expression in the cells of interest.

4.2 Transductional targeting

Adenoviruses display unparalleled transduction efficacy and natural tropism to a wide range of epithelial tissues. Receptor recognition could be one of the most important factors involved in cell tropism. Increased CAR expression appears to have a growth inhibitory effect on some cancer cell lines, while loss of CAR expression correlates with tumor

progression and advanced disease (Okegawa *et al.* 2001; Bauerschmitz *et al.* 2002; Miller *et al.* 1998a; Cripe *et al.* 2001; Dmitriev *et al.* 1998). In addition, CAR has been suggested to play a role in cell adhesion, and its expression may be cell-cycle dependent (Cohen *et al.* 2001). As efficient gene transfer is one of the key issues for successful gene therapy, low expression level of CAR on tumor cells causes a major challenge (Okegawa *et al.* 2001; Seidman *et al.* 2001). To circumvent deficiency of CAR, viruses can be retargeted by improving their abilities to attach to heterologous receptors. There are two distinct approaches to retarget adenoviruses transductionally: adapter molecule-based targeting and targeting via genetic manipulation of the adenovirus capsid (Figure 4).

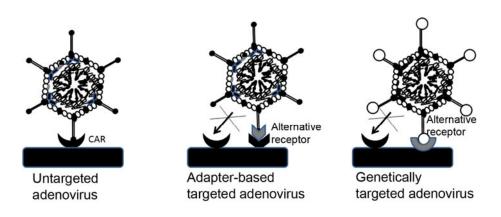


Figure 4. Transductional targeting of adenoviruses. Transductional targeting can be used to target adenovirus to alternative cellular receptors instead of its primary receptor CAR. Targeting can be based on use of adapter molecules or genetically modified fibers that allow the virus to bind alternative receptors.

4.2.1 Adapter molecule-based targeting

The idea behind adapter molecule-based targeting is to use adapter molecules that crosslink the adenovirus vector to alternative cell surface receptors, bypassing the native CAR-based tropism (Figure 4). The majority of currently used adapter molecules are bispecific, which both ablates the native CAR binding and formats a novel tropism to cellular receptors, highly expressed on cancer cells. Various adapter molecules from

bispecific antibodies, chemical conjugates between antibody fragment (Fab) and cell selective ligand (ex. folate), Fab antibody and peptide ligand conjugates to recombinant fusion proteins that incorporate Fabs and peptide ligands have been used successfully. The first *in vitro* study of adenovirus targeting via adapter molecule was done by Douglas *et al* (Douglas *et al*. 1996). They targeted to the folate receptor, overexpressed on many malignant cells with anti-knob neutralizing Fab-folate conjugate. A similar targeting system was used with FGF receptor-positive Kaposi's sarcoma (KS) cells *in vitro*, where basic fibroblast growth factor (FGF2) was fused to anti-knob Fab (Goldman *et al*. 1997). Other Fab-ligand conjugates have been targeted to epithelial cellular adhesion molecule (Ep-CAM) (Haisma *et al*. 1999), tumor-associated glycoprotein (Tag-72) (Kelly *et al*. 2000), epidermal growth factor (EGF) receptor (Miller *et al*. 1998b), and CD 40 cell marker (Tillman *et al*. 1999), with promising results.

An interesting alternative method to the chemical conjugates was developed utilizing a truncated soluble form of CAR (sCAR) fused to EGF (Dmitriev *et al.* 2000). Approaches resulted in increased reporter gene expression in many cancer cell lines overexpressing either EGF receptor or CD40 *in vitro*. When targeted to the EGF receptor, up to a 9-fold increase in gene transfer was seen compared with nontargeted adenovirus.

The above approaches have also been tested *in vivo*. A bispecific antibody was used to target adenovirus to the angiotensin-converting enzyme (ACE) expressed in the pulmonary capillary endothelium. Systemic delivery of adenovirus resulted in a 20-fold increase in the lungs compared with the untargeted control. Moreover, adenovirus gene transfer to liver decreased 83% (Reynolds *et al.* 2000). Another unique lung-targeting approach was developed by Everts *et al*, where sCAR was fused with with a single-chain anti-CEA antibody (MFE-23). sCAR-MFE targeted adenovirus decreased liver gene expression by > 90 % and increased transduction to the CEA-positive epithelial tumor cells after systemic delivery (Everts *et al.* 2005; Li *et al.* 2007).

4.2.2 Genetic targeting

The advantage of the adapter-based targeting approach is the wide range of ligands that can be incorporated into fusion molecules, but there are also several drawbacks. The structure itself is complex, which results in creation of a heterogeneous population of

virions, and the stability of the complexes in the human circulation is not very well characterized. This might be disadvantageous in human trials. Therefore, one-component systems where adenoviruses capsid proteins are genetically manipulated have been created. Overall, there are three basic ways to modify adenovirus fiber to redirect its tropism: 1) so-called fiber pseudotyping, 2) ligand incorporation into the fiber knob, and 3) 'de-knobbing' of the fiber coupled with ligand addition.

Understanding of adenovirus infection biology indicates that genetic modifications are usually located in the fiber, the primary capsid determinant of adenoviral tropism. Many clinically relevant cancer cell types are refractory to adenovirus infection due to variable or aberrant expression of the primary receptor CAR (Bauerschmitz et al. 2002; Cripe et al. 2001; Hemminki et al. 2002; Miller et al. 1998a; Okegawa et al. 2000; Seidman et al. 2001). Genetic replacement of fiber knob was first accomplished by Krasnykh et al (Krasnykh et al. 1996). Adenovirus serotype 5 knob was replaced by the knob from serotype 3, which uses a different receptor than Ad5 in cell entry. Ad3 belongs to subgroup B and primary receptors for adenoviruses belonging to the group have recently been suggested, including CD46, a complement regulatory protein expressed on the surface of nucleated human cells (Liszewski et al. 1991), and CD80 and CD86, costimulatory molecules present on mature dendritic cells and B lymphocytes and involved in stimulating T-lymphocyte activation (Short et al. 2004). The receptor for adenovirus 3 is still unknown, although there are suggestions that it could be on one of the above receptors or some coreceptor or combination of receptors (Sirena et al. 2004; Tuve et al. 2006a; Sirena et al. 2004). Ad5/3 chimera has displayed enhanced CAR independent transduction on variety of neoplastic cells (Krasnykh et al. 1996; Davidoff et al. 1999; Kanerva et al. 2002b). Promising results with Ad5/3 chimeras have encouraged many researchers to evaluate other serotype chimeras. Ad5/35, Ad5/11, and Ad5/7 have shown enhanced transduction refractory to the CAR in many types of cells with clinical importance and reduced localization to the liver (Shayakhmetov et al. 2000; Stecher et al. 2001; Gall et al. 1996; Mizuguchi & Hayakawa 2002; Schoggins et al. 2005).

The fiber knob region is the major attachment site for the primary receptor at least *in vitro*. Consequently, modification of the knob region has become a major research focus for targeting gene delivery. Studies have exploited two locations within the knob, C-terminus

and HI loop, which can be manipulated without losing function of the fiber. Since the Cterminus is solvent-exposed, Wickham et al. added a polylysin tail to mediate adenovirus binding through heparan sulfate proteoglycans (HSPGs), yielding promising results (Wickham et al. 1997; Kangasniemi et al. 2006; Ranki et al. 2007a). $\alpha_v \beta$ integrin-binding arginine-lysine-aspartic acid (RGD) motifs have also successfully been incorporated into the C-terminus (Wickham et al. 1997). Another promising knob location, the HI-loop, which is exposed outside and can tolerate rather large (up to 100 amino acids) peptide insertions, is located between β-strands H and I in the knob monomer. FLAG peptide was introduced to the site first by Krasnykh et al, and other retargeting motifs in the HI loop have been evaluated. RGD-4C peptide in the HI loop is widely used with enhanced infectivity to wide range of tumor cells in vitro and in vivo (Cripe et al. 2001; Dmitriev et al. 1998; Hemminki et al. 2001; Kanerva et al. 2002b; Kangasniemi et al. 2006; Ranki et al. 2007a). Other peptide that has provided cancer selective targeting when inserted into the HI loop includes asparagine-glycine-arginine (NGR) (Mizuguchi et al. 2001). Wu et al, combined a polylysin tail in the C-terminus and a RGD-4C motif in the HI loop, and enhanced gene transfer was achieved with several CAR-deficient cell lines, as well as with human pancreatic islet cells (Wu et al. 2002). Some studies have also used both genetic fiber modification and an adapter-based approach where incorporating proteins with immunoglobulin binding domains into the C-terminus or the HI loop resulted in enhanced gene transduction in cells containing the Fc region of Ig (Henning et al. 2005; Korokhov et al. 2003; Volpers et al. 2003).

Since the ligand incorporation methods described above do not necessarily ablate CAR binding, a native fiber has been replaced by a knobless fiber and simultaneous adding of targeting ligand to the fiber has resulted in a more targeted adenovirus (Falgout and Ketner 1988; Von Seggern *et al.* 1999). However, the major barrier to this approach is that the knob region contains a stabilization element for the fiber element. To circumvent the problem, external trimerization elements, such as the bacteriophage T4 fibritin or the neck region of human lung surfactant protein D (NRP), have been used (Hedley *et al.* 2006; Krasnykh *et al.* 2001; Magnusson *et al.* 2001). Li *et al.* described a novel trimerization element within the adenovirus fiber protein containing an N-terminus tail and shaft repeats

that was able to form stable virions and serve as a platform for the generation of tissue-specific adenovirus vectors (Li *et al.* 2006).

The need for targeting adenoviruses has lead to searches for other capsid proteins that could be manipulated such as hexon, penton, and pIX. Due to the location of the hexon protein, it is an attractive site for incorporating targeting ligands. Hexon hypervariable regions have a loop that protrudes outside the capsid and Vigne *et al.* exploited the possibility to attach an RGD motif to this loop. Notably, the stability of the virion was unaffected and enhanced gene delivery to non CAR cells was achieved (Vigne *et al.* 1999). The RGD motif in the penton base was replaced with a peptide that targeted adenoviruses to the metastatic melanomas and glioblastomas (Wickham *et al.* 1995). Recently, pIX was found to be a versatile capsid location suitable for both targeting and imaging motifs (Dmitriev *et al.* 2002; Le *et al.* 2004).

4.3 Transcriptionally targeted adenoviruses

To achieve more tumor cell-selective adenoviruses, gene expression can be regulated transcriptionally. Regulation is usually mediated by tumor-specific promoters. Transcriptional targeting of adenovirus transgene expression or replication restricts the transcription unit expression to the cells where the promoter being used is active. A number of tumor, or tissue-specific promoters have been applied in cancer gene therapy (Glasgow *et al.* 2004; Saukkonen & Hemminki 2004). Table 1 summarizes some of these.

Table 1. Selected tumor-specific promoters used in adenoviral gene therapy.

Promoter	Transgene(s)	Disease application	Reference
CEA	LacZ	Gastric cancer	(Brand et al. 1998;
	HSVtk	Lung cancer	Osaki <i>et al</i> . 1994).
SLPI	HSVtk	Ovarian cancer	(Barker et al.
			2003a).
Cox-2	HSVtk	Gastric cancer	(Yamamoto et al.
	Luciferase	Pancreatic cancer	2001).
	HSVtk		(Wesseling et al.
	HSVtk	Ovarian cancer	2001).
			(Casado et al. 2001).
	Luciferase	Lung cancer	(Sarkioja <i>et al</i> .
			2008).
Midkine (MK)	Luciferase	Pancreatic cancer	(Wesseling et al.
	HSVtk		2001).
hTERT	E1A_IRES_E1B	Lung cancer	(Hashimoto et al.
	GFP/TRAIL	Breast cancer	2008).
			(Lin et al. 2002).
E2F	E1A	Lung cancer	(Tsukuda <i>et al</i> .
		Ovarian cancer	2002).
CXCR4	E1A	Lung cancer	(Zhu et al. 2007b).
	E1A	Breast cancer	(Rocconi et al.
	E1A	Renal cancer	2007).
			(Haviv et al. 2004).
SCCA2	Luciferase	Lung cancer	(Oshikiri <i>et al</i> .
			2006).
TTS		Lung cancer	(Fukazawa et al.
			2004).

Abbreviations used: CEA, carcinoembronic antigen; SLP1, secretory leukoprotease inhibitor; Cox-2, cyclooxygenase 2; hTERT, human telomerase reverse transcriptase; CXCR4, chemokine receptor; SCCA2, squamous cell carcinoma antigen; TTS, dual promoter with hTERT and human surfactant protein A1 promoters.

Carcinoembryonic antigen (CEA) promoter was one of the first tissue-specific promoters explored. CEA is expressed in many types of cancers. Herpes simplex virus thymidine kinase (HSVtk) expression driven by the CEA promoter left CEA-negative cell lines resistant to ganciclovir (GCV) prodrug therapy, while CEA-positive A549 cells were 1000-fold more sensitive (Osaki et al. 1994). Further, Hashimoto et al. used human telomerase reverse transcriptase promoter (hTERT)-driven adenovirus E1A and E1B genes connected to the internal ribosome entry site (IRES) to target lung cancer cells in vitro and in vivo, which led to a 50% reduction in cell viability in normal cells versus lung cancer cells (Hashimoto et al. 2008). In addition, E2F, CXCR4, SCCA2, and TTS have been used to target adenoviruses to lung cancer tissue, with promising results. E2F transcription factors are critical regulators of cell growth and are often overexpressed in cancer cells because of frequent aberrations in the pRb/E2F/p16(INK4a) pathway. As a result, a majority of tumor cells exist in a high proliferative state with E2F constantly active, making it a good candidate promoter for targeting adenovirus (Tsukuda et al. 2002). Chemokine receptor CXCR4 is highly expressed in lung cancer and CXCR4 promoterdriven RGD-modified adenovirus showed a 10-fold increase in replication in lung cancer tissue slices, in contrast to a 10-fold decrease in the human liver slices (Spano et al. 2004; Zhu et al. 2007b).

In addition to tissue-specific promoters, adenoviral transgene expression or replication can also be controlled at a posttranscriptional level. Eukaryotic mRNAs contain 3'-untranslated regions (UTRs), the AU- rich elements of which are involved in posttranscriptional control of gene expression. These elements are frequently present in 3-UTRs of many proto-oncogene and cytokine mRNAs (Dixon *et al.* 2000; Kruys *et al.* 1989). The usual physiological function of the elements is to rapidly downregulate mRNA transcripts when the threat (e.g. infection) has passed. Ahmed *et al.* used CMV promoter-driven adenovirus, where the E1A was under the regulation of the 3'UTR element from the cyclooxygenase 2

(Cox-2) gene, allowing RAS/P-(MAPK)-specific stabilization on the mRNA (Ahmed *et al.* 2003).

Tumor-specific promoters and other regulatory elements have the potential to increase specificity and decrease toxicity of adenoviral gene therapy. However, adenoviruses targeted only with promoters or regulators are dependent on CAR for entry. Combining transductional and transcriptional targeting is an effective way to achieve a synergistic effect on gene transfer (Barker *et al.* 2003b; Reynolds *et al.* 2001).

5 Oncolytic adenoviruses

Although nonreplicating first-generation adenoviruses have provided high *in vitro* and *in vivo* transduction rates and good safety data, clinical trials have suggested that the single-agent antitumor effect may not be sufficient for all treatment approaches. Advanced human solid tumor masses are larger and more complex than xenograft tumors, which restrict sufficient access of virions to tumor cells. Necrotic, hyperbaric, hypoxic, and stromal areas can for instance, be problematic. To overcome this problem, oncolytic adenoviruses have been developed. Their antitumor effect is based on oncolysis caused by replication *per se*, whereas normal tissues are spared due to low replication (Figure 5). Hypothetically, the replication would continue until all cancer cells are lysed, and released viruses would even disseminate into the bloodstream and infect distant metastasis.

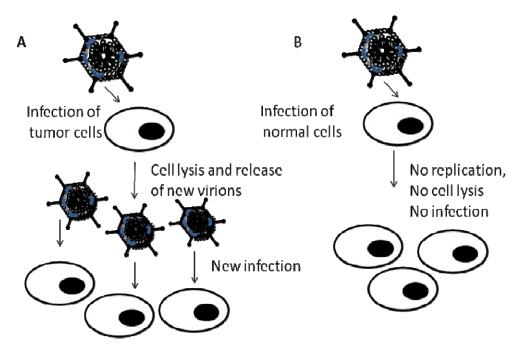


Figure 5. Oncolytic adenoviruses. A) Infection of tumor cells results in viral replication, cell lysis, and release of new viruses. B) Normal cells are spared due to lack of replication.

Adenovirus early 1 (E1) gene is responsible for replication and most development of oncolytic adenoviruses have focused on the genetic engineering of E1 genes. There are two approaches that have been used to restrict viral replication to target cells and to spare normal tissue. The first published oncolytic adenovirus, d11520, later called ONYX-015 has two mutations in the E1B gene coding the E1B-55kD protein and is very similar to the H101, the first licensed oncolytic agent (Bischoff *et al.* 1996; Xia *et al.* 2004b). Adenovirus E1B-55kD protein binds to tumor suppressor p53 to inactivate it for induction of S-phase-like state, which is required for viral replication. Further, it is responsible for forming a different complex with E4orf6 that shuts off host mRNA nuclear export and host protein synthesis. It also facilitates the nuclear localization of transcription factor YB1 to activate the E2 late promoter (Holm *et al.* 2002). Tumor cells often have an inactivated p53-p14ARF pathway, and thus, viruses should only replicate in these cells. However some tumor cells seem to fail to support E1B-55kD deleted adenoviral replication.

Recently O'Shea *et al* suggested that it may be due to failure in late mRNA export that prevents host protein shutoff (O'Shea *et al*. 2005).

Another type of Ad contains a 24 bp deletion in constant region 2 (CR2) of the E1A gene. This domain is responsible for binding the Rb tumor suppressor/cell cycle regulator protein, thereby allowing adenovirus to induce an S-phase-like state. As a result, viruses with the Δ 24-E1A deletion have a reduced ability to overcome the G1-S checkpoint and replicate efficiently only in cells where this interaction is not necessary, e.g. in tumor cells defective in the Rb-p16 pathway. (Heise et al. 2000b; Fueyo et al. 2000; Suzuki et al. 2001).

A third type of deletion used to confer cancer selectivity to adenovirus is the deletion of virus-associated (VA) RNA genes. These small type I and II RNAs inactivate the protein kinase R (PKR). PKR plays a major role in the interferon (IFN) antiviral cell defense, and by binding to PKR, VA-RNAs counteract this IFN response to adenovirus, and conversely, adenoviruses defective in VA-RNAs are blocked by IFN via protein translation inhibition. However, many cancer cells present a truncated IFN pathway, and adenoviruses with VA RNAs deleted retain replication in these cells, but not so in normal cells (Cascallo *et al.* 2006).

In addition, viral replication can be restricted to target cells by using tumor-specific promoters (TSPs). TSPs are usually added to control *EIA*, but alternatively, other genes such as *E4* can also be regulated. Various promoters have been used and some selected ones are presented in Table 1.

5.1 Oncolytic adenoviruses combined with targeting moieties

To improve adenovirus-based replication in the context of clinically applicable cancer gene therapy, it is essential to increase targeting to tumor cells and decrease targeting to the liver due to possible toxicity. Adenovirus 5-based oncolytic viruses may fall victims to variable CAR expression, and a number of strategies exists to improve Ad5 based viral tropism. Previously described transductional targeting methods have been combined with oncolytic viruses with good results. Serotype chimeric Ad5/3-Δ24 has shown increased therapeutic efficacy at least in ovarian, kidney, breast, and gastric cancers (Kanerva *et al.* 2003; Guse *et al.* 2007b; Kangasniemi *et al.* 2006; Ranki *et al.* 2007a). Ad5-Δ24RGD, which has RGD

modification in the HI loop also has shown efficient cell killing in numerous cancer cells, including ovarian cancer, glioma and osteosarcoma (Lamfers *et al.* 2002; Suzuki *et al.* 2001; Witlox *et al.* 2004). Further, Ranki *et al.* added seven polylysins to the C-terminus of the Ad5 Δ 24 creating Ad5.pK7- Δ 24, which displayed enhanced oncolysis of breast cancer cells *in vitro* and *in vivo* (Ranki *et al.* 2007).

Recent studies have utilized double, or even triple-targeted adenoviruses to achieve more specificity. Combining a transductional targeting moiety with a tumor-specific promoter has resulted in a pancreatic and ovarian cancer specific adenovirus featuring Ad3 knob and Cox-2 promoter driving E1A with a 24 bp deletion in the CR2 (Bauerschmitz *et al.* 2006; Ramirez *et al.* 2008). Guse *et al.* used many double- and triple-targeted adenoviruses with Ad3 knob, RGD, and pK7 motifs combined with Cox-2 promoter driving CR1 and/or CR2 deletions with promising results in ovarian and renal cancer xenografts (Guse *et al.* 2007a).

5.2 Armed oncolytic adenoviruses

Despite successful targeting and specific replication of oncolytic adenoviruses in the tumor tissue, clinical trials with early generation adenoviruses have shown that complete eradication of solid tumor masses rarely occurs. This might be due to intratumoral complexities such as stromal matrix and other extracellular barriers within the tumor tissue (Cheng et al. 2007). A useful approach for improving the potency of replicating agents is to arm viruses with therapeutic transgenes, e.g. genes encoding prodrug-converting enzymes (Hermiston and Kuhn 2002). Common approaches include HSV-TK and Escherichia coli cytosine deaminase (CD), which convert nontoxic prodrugs [ganciclovir (GCV) and 5-fluorocytosine, respectively] into cytotoxic metabolites. The activated drugs can spread into surrounding cells creating a so called bystander effect. Reports combining a TK/GCV system with replicating adenoviruses have been rather controversial. Some studies have demonstrated that GCV enhances oncolytic potency of adenoviruses expressing HSV-tk (Nanda et al. 2001; Raki et al. 2007; Wildner et al. 1999), in contrast to others where oncolytic potential was not improved possibly due to the inhibition of viral replication by GCV(Hakkarainen et al. 2006; Lambright et al. 2001; Morris & Wildner 2000; Wildner & Morris 2000). Double suicide therapy with both HSV-

tk and CD genes has proven to be rather efficient, especially when combined with radiotherapy (Freytag *et al.* 1998; Rogulski *et al.* 2000; Wu *et al.* 2005).

5.3 Oncolytic adenoviruses in combination with conventional therapies

Combining adenoviral gene therapy with traditional therapies is a powerful tool to achieve enhanced or even synergistic cell killing and antitumor efficacy. The mechanism of the additive or synergistic effect of combined therapy is not yet thoroughly known, although a few hypotheses have been tendered. Firstly, expression of adenoviral E1A protein may sensitize cells to chemotherapy or radiotherapy-mediated cell killing (Duque et al. 1998; Sanchez-Prieto et al. 1996). Secondly, chemotherapeutic agents may enhance the level of virus replication. Finally, chemotherapeutic agents may affect the receptor status of the target cells or each agent may work independently within the tumor tissue. There are a number of studies where oncolytic adenoviruses have been combined with chemo- or radiotherapy, and some of them have even proceeded to clinical testing. ONYX-015 was the first virus to undergo clinical trials combined with chemotherapy. ONYX-015 was combined with 5-fluorouracil (5-FU) and cisplatin to treat squamous cell carcinoma of the head and neck carcinoma (Xia et al. 2004a; Khuri et al. 2000a). ONYX-015 was also combined with gemcitabine (pancreatic cancer) (Hecht et al. 2003), mitomycin C+doxorubicin + cisplatin (MAP) (advanced sarcoma) (Galanis et al. 2005a), and carboplatin + taxol (lung metastasis) (Nemunaitis et al. 2001), among other but a few studies. Chemo- and radiotherapies were combined with HSV-tk and CD-expressing adenovirus to treat prostate cancer with good safety data (Freytag et al. 2003b); these treatments may provide a potential long-term benefit for the patients (Freytag et al. 2007).

5.4 Cells as carriers of oncolytic adenoviruses

Although infectivity and specificity of adenoviruses can be enhanced, obstacles remain that may hinder the bioavailability of viruses, especially after intravenous administration. Despite retargeting, the majority of systemically administered adenoviruses are taken up by Kupffer cells in the liver, decreasing the amount of available virus (Alemany *et al.* 2000). In addition, adaptive and innate immune responses may eliminate therapeutic

viruses (Bessis *et al.* 2004). Further, tumors are known to be very heterogeneous in structure, containing hypoxic, necrotic, and stromal areas, which may hinder spreading of the adenovirus within the tumor. A potential solution to solve these problems is the use of cells as viral carriers (Sanz *et al.* 2005; Raykov and Rommelaere 2008; Fritz and Jorgensen 2008). This might have advantages over systemic viral administration, as attenuated neutralization and improved viral targeting could result in smaller viral doses administered to patients, which may in turn reduce the systemic side effects. However, the use of cells as agents for systemic virus delivery depends on 1) their ability to be transduced with adenovirus, 2) their ability to allow the replication and release of oncolytic adenoviruses and 3) their ability to home specifically to tumor. There a several cell types that have been used as carriers of viruses suchs as antigen-specific T-cells (Cole *et al.* 2005; Yotnda *et al.* 2004), macrophages (Griffiths *et al.* 2000), endothelial progenitor cells (Jevremovic *et al.* 2004), non antigen specific T-cells (Ong *et al.* 2007) and mesenchymal stem cells (MSCs)(Dvorak 1986; Studeny *et al.* 2004a).

MSCs are precursor cells that originate either from e.g. bone marrow or fat tissue, and they can differentiate into adipocytes, chondrocytes, osteoblasts, myoblasts, and tenocytes (Prockop 1997). One function of the MSCs is to repair injured tissue, and it is also hypothesized that the tumor environment resembles injured tissue, and therefore, circulating MSCs would home to tumors. (Dvorak 1986; Studeny *et al.* 2002; Studeny *et al.* 2004a). In some studies adenoviruses have been targeted to tumor tissue using MSCs as carriers. Nakamura *et al.* targeted adenoviral vector coding human interleukin-2 within MSCs with an augmented antitumor effect and prolonged survival of glioma-bearing rats (Nakamura *et al.* 2004). Stoff-Khalili *et al.* used hMSCs to target oncolytic adenovirus in to lung metastases of breast cancer which reduced the growth of lung metastases *in vivo* (Stoff-Khalili *et al.* 2007a).

6 Clinical trials with oncolytic adenoviruses

Cancer is a complex disease that often eludes successful treatment due to its propensity to evolve or adapt in the face of current therapeutic regimes. Viral oncolytic therapy has been

under investigation as a novel anticancer strategy and the first cancer trial with wild type adenovirus was done as early as 1950s. Since the late 1990s oncolytic adenoviruses have been utilized to treat phase I-III cancer patients. The first oncolytic adenovirus used in a clinical trial was ONYX-015, which has a 55 kb deletion in the *E1b* gene, and thus, has tumor selectivity to cancer cells with mutant p53. The virus established a clinical proof-of-concept for oncolytic virotherapy in a first phase I trial in which the virus was directly injected into head and neck tumors with 14% regression rates (Ganly *et al.* 2000; Nemunaitis *et al.* 2000). When ONYX-015 was combined with cisplatin and 5-FU in the phase II trial, approximately 65% of patients showed tumor regression (Khuri *et al.* 2000b; Nemunaitis *et al.* 2000). A phase III trial was done in China with a closely related virus H101 and investigators reported 79% response rate in combination-treated patients versus 40% in controls without virus treatment (p<0.001) (Xia *et al.* 2004b) and later the virus got a official selling permit in China.

Another oncolytic adenovirus used in clinical trials is CV706. CV706 is a prostate-specific antigen (PSA)-selective, replication-competent adenovirus that showed in the phase I study good safety data and intraprostatic replication (DeWeese *et al.* 2001). In addition, another prostate specific, intravenously administered virus cg 7870 has been used in phase I study to treat hormone refractory metastatic prostate cancer with promising safety data (Small *et al.* 2006). Overall, trials with onvolytic adenoviruses have reported good safety data, but only marginal efficacy. Selected oncolytic adenovirus trials are summarized in Table 2. For a more detailed listing of gene therapy trials, see

http://www.abedia.com/wiley/index.html.

Table 2. Clinical trials with oncolytic adenoviruses.

Virus/treatment	Genetic modification	Phase	Route	Max.dose	Disease	Beneficial/ total	Reference
ONYX-015	E1b-55 kD deletion	I	i.t.	1x10 ¹¹ pfu	SCCHN	2/22	(Ganly <i>et al.</i> 2000).
ONYX-015	E1b-55 kD deletion	I	i.t.	1x10 ¹¹ pfu	Pancreatic cancer	0/23	(Mulvihill et al. 2001).

	•	•		•	-		
ONYX-015	E1b-55 kD deletion	I	i.v.	2x10 ¹³ vp	Cancer metastatic to lung	0/10	(Nemunaitis et al. 2001).
ONYX-015	E1b-55 kD deletion	I	i.p.	1x10 ¹¹ pfu/d x 5d	Ovarian cancer	0/16	(Vasey et al. 2002).
ONYX-015+ 5-FU+ leucovorin	E1b-55 kD deletion	I	i.ha.	2x10 ¹² vp	Colorectal cancer metastatic to liver	1/11	(Reid <i>et al.</i> 2001).
ONYX-015	E1b-55 kD deletion	I	i.t.	1x10 ¹⁰ pfu	Glioma	3/24	(Chiocca et al. 2004).
ONYX-015+ etanercept	E1b-55 kD deletion	I	i.v.	1x10 ¹² pfu	Advanced ca.	0/9	(Nemunaitis et al. 2007).
CV 706	PSA promoter- enhancer contr. E1A	I	i.t.	1x10 ¹³ vp	Prostate cancer	5/20	(DeWeese et al. 2001).
Ad5- CD/TKrep + GCV/ 5-FU + radiation	E1b-55 kD deletion + TK/CD	I	i.t.	1x10 ¹² vp	Prostate cancer	15/15	(Freytag et al. 2003a).
ONYX-015 + 5-FU	E1b-55 kD deletion	I-II	i.t. i.ha. i.v.	3x10 ¹¹ pfu	HCC and colorectal cancer metastatic to liver	3/16	(Habib <i>et al.</i> 2001).
ONYX-015	E1b-55 kD deletion	II	i.t.	2x10 ¹¹ vp X 10d	SCCHN	5/40	(Nemunaitis et al. 2001).

		•	-				
ONYX-015 + cisplatin and 5-FU	E1b-55 kD deletion	II	i.t.	1x10 ¹⁰ vp/d X 5d	SCCHN	19/37	(Khuri <i>et al</i> . 2000a).
ONYX-015 + Gemcitabine	E1b-55 kD deletion	I-II	i.t.	2x10 ¹¹ vp/wk 8 cycles	Pancreatic cancer	2/21	(Hecht et al. 2003).
ONYX-015	E1b-55 kD deletion	II	i.v.	2x10 ¹² vp/ 2 weeks	Metastatic colorectal cancer	0/18	(Hamid <i>et al.</i> 2003).
CG7870	Rat probasin controlling <i>E1A</i>	I-II	i.t.	1x10 ¹³ vp	Prostate cancer	10/20	DeWeese 2003
H101 + cisplatin/ adriamycin + 5-FU	E1b-55 kD deletion	III	i.t.	1.5x10 ¹² vp/d X 5d	SCCHN	71/160	(Xia et al. 2004b).

ONYX-015 +	<i>E1b</i> -55 kD	1-II	i.t.	5x10 ¹⁰ pfu	Sarcoma	1/6	(Galanis et
MAP	deletion						al. 2005b).
chemotherapy							

Abbreviations used: i.t., intratumoral; i.v., intravenous; i.p., intraperitoneal; i.ha, intrahepatic artery, 5-FU, 5-fluorouracil; MAP, mitomycin C+doxorubicin + cisplatin; SCCHN, squamous cell carcinoma of the head and neck; vp, viral particles; pfu, plaque-forming units; HSV-tk, herpes simplex virus thymidine kinase; CD, cytosine deaminase

7 Adenoviruses and immune response

Effective readministration of adenovirus might be important factor in achieving full potential of adenoviral gene therapy for treating many diseases. More than 50 different

serotypes of adenoviruses are known to infect humans. Due to the ubiquitous nature of adenoviruses, a majority of the human population has been exposed to them (also to Ad5), leading to the development of adenovirus-specific neutralizing antibodies (NAbs). A high neutralizing antibody titer might not be a limiting factor in local treatment (Cichon *et al.* 2001; Hodges *et al.* 2005), but it can decrease the bioavailability of systemically administered adenoviruses. NAbs can be induced against all adenoviral capsid proteins, such as hexon, penton and fiber and there may even be synergy between them (Gahery-Segard *et al.* 1998; Rahman *et al.* 2001). Strategies that are being examined to circumvent immune response to adenovirus vectors include immunosuppression, immunomodulation, serotype switching, use of targeted adenovirus vectors, microencapsulation of adenovirus vectors, use of helper-dependent (HD) adenovirus vectors, and development of nonhuman adenovirus vectors.

7.1 Manipulation of adenoviruses to circumvent immune response

Following intravenous inoculation, adenoviruses are generally taken up by Kupffer cells in the liver, and by macrophages in the spleen leading to rapid induction of an innate immune response. The innate immune response is activated following recognition of molecular patterns on the adenovirus capsid by receptors on macrophages and dendritic cells (DCs), resulting in activation of multiple signaling pathways, such as mitogen-activated protein kinase (MAPK) and nuclear factor (NF)-κB pathways, that augment expression of several proinflammatory cytokines and chemokines (Bruder and Kovesdi 1997; Lieber et al. 1997; Muruve et al. 1999). The use of such immunosuppressive agents as cyclosporine and cyclophosphamide to block antibodies raised against adenovirus enhances the duration of transgene expression (Smith et al. 1996). Since macrophages in the liver activate the immune response, there are approaches where depletion of macrophages and dendritic cells has resulted in increased transgene expression (Ranki et al. 2007b; Schiedner et al. 2003). Further, alteration of the immunodominant epitopes of adenovirus capsid can also help to evade immune response. Covalent attachment of polymers, such as polyethylene glycol (PEG), to the adenovirus capsid has been shown to inhibit antibody-mediated virus neutralization but there is evidence that it also decreases tumor transduction of the virus (Croyle et al. 2002; Croyle et al. 2005).

Adenoviral serotypes are classically determined based on neutralization assay and it is therefore logical that different serotype can overcome NAb. In some studies, serotype switching in vector construction has helped to overcome immune response (Kass-Eisler et al. 1996; Mastrangeli et al. 1996). Especially the use of subgroup B adenoviruses, such as Ad3, Ad11 and Ad35 has been promising since these are not dependent on CAR-mediated internalization and can evade pre-existing Ad5 immunity (Barouch et al. 2004; Stone et al. 2005; Vogels et al. 2003; Kanerva et al. 2002b). The adenoviral fiber knob has been shown to induce DC activation and maturation (Molinier-Frenkel et al. 2003), and therefore, fiber knob modifications to incorporate cellular ligands with novel cell-binding capacity might confer targeting and be one way to decrease vector immunogenicity (Nanda et al. 2005) Adenoviral microencapsulation with, for instance, polyethylene glycol-cationic lipid and poly (lactic-glycolic) acid (PLGA) copolymer has been demonstrated to guard adenovirus vectors from neutralizing antibodies and to be capable of effective transduction and gene expression on target cells (Beer et al. 1998; Chillon et al. 1998; Steel et al. 2004). Lastly, utilization of helper dependent (HD) adenoviruses has resulted in controversial results in evading immune response. HD adenoviral vectors lack all coding sequences of the adenovirus genome, except the packaging sequence and inverted terminal repeats, have been shown to elicit minimal cell-mediated immune response (Morsy et al. 1998) but led to acute toxicity accompanied by activation of the innate immune response after systemic administration in baboons (Brunetti-Pierri et al. 2004).

To extend the range of Ad vectors that can be used to evade human adenovirus neutralizing immune response, a number of nonhuman adenoviruses, such as bovine Ad type 3 (BAd3), canine Ad type 2, porcine Ad type 3, ovine Ad ,and chimpanzee Ad have been exploited for gene transfer, without hindrance from cross-neutralizing effects (Bangari & Mittal 2004; Farina *et al.* 2001; Hemminki *et al.* 2003; Hofmann *et al.* 1999; Mittal *et al.* 1995).

Aims of the study

- 1. To determine the best receptor binding motif for NSCLC cell lines and fresh NSCLC primary tissue samples *in vitro* and *in vivo* and to evaluate gene transfer and antitumor efficacy of oncolytic adenoviruses through non-CAR receptors. In addition, the biodistribution, and the oncolytic potency of adenoviruses were evaluated *in vivo* with noninvasive imaging. (I).
- 2. To examine the capacity of capsid-modified adenoviruses to infect and replicate in MSCs. Further, biodistribution, tumor-homing ability, and tumor-killing efficacy of systemically delivered, virus-loaded MSCs in orthotopic lung and breast cancer tumor models were evaluated (II).
- **3.** To investigate induction of the Cox-2 promoter both in normal and tumor cells after adenovirus infection and to explore the utility of AU-rich elements for regaining promoter selectivity (III).
- **4.** To evaluate whether changing the adenovirus fiber knob could allow the virus to overcome pre-existing adenoviral NAbs in the context of low or high NAb titer, in mice with advanced orthotopic NCSLC, and in human clinical samples of both cancerous and nonmalignant lung tissue **(IV).**

Materials and methods

Detailed description of the methodologies used can be found in the original publications.

1 Cell lines, human mesenchymal stem cells (MSCs), and primary NSCLC (I-IV)

Table 3. Human cell lines used in studies I-IV.

Transformed embryonic ATCC (Manassas, I,III VA) NCI-H460 Lung large-cell ATCC I HTB-177 carcinoma NCI-H661 Lung large-cell ATCC I HTB-183 carcinoma LNM35/EGFP Lung large-cell Takashi Takahashi I,II, III, IV carcinoma expressing enhanced green Institute, Japan) fluorescent protein SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I carcinoma (University of	Cell line	Description	Source	Study
NCI-H460 Lung large-cell ATCC I HTB-177 carcinoma NCI-H661 Lung large-cell ATCC I HTB-183 carcinoma LNM35/EGFP Lung large-cell Takashi Takahashi I,II, III, IV carcinoma expressing enhanced green Institute, Japan) fluorescent protein SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I		•	· ·	I,III
HTB-177 carcinoma NCI-H661 Lung large-cell ATCC I HTB-183 carcinoma LNM35/EGFP Lung large-cell Takashi Takahashi I,II, III, IV carcinoma expressing (Honda Research enhanced green Institute, Japan) fluorescent protein SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I	ATCC CRL-1573	kidney cells	VA)	
NCI-H661 Lung large-cell ATCC I HTB-183 carcinoma LNM35/EGFP Lung large-cell Takashi Takahashi I,II, III, IV carcinoma expressing (Honda Research enhanced green Institute, Japan) fluorescent protein SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I	NCI-H460	Lung large-cell	ATCC	I
HTB-183 LNM35/EGFP Lung large-cell carcinoma expressing enhanced green fluorescent protein SW900 Lung squamous cell HTB-59 Carcinoma Lung squamous cell Kazuya Kondo I,II, III, IV Carcinoma I,II, III, IV Carcinoma ATCC I Kazuya Kondo I	HTB-177	carcinoma		
LNM35/EGFP Lung large-cell carcinoma expressing enhanced green fluorescent protein SW900 Lung squamous cell HTB-59 Lung squamous cell Kazuya Kondo I,II, III, IV Carcinoma I,II, III, IV Carcinoma III, III, IV ATCC I Kazuya Kondo I	NCI-H661	Lung large-cell	ATCC	I
carcinoma expressing (Honda Research enhanced green Institute, Japan) fluorescent protein SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I	HTB-183	carcinoma		
enhanced green Institute, Japan) fluorescent protein SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I	LNM35/EGFP	Lung large-cell	Takashi Takahashi	I,II, III, IV
fluorescent protein SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I		carcinoma expressing	(Honda Research	
SW900 Lung squamous cell ATCC I HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I		enhanced green	Institute, Japan)	
HTB-59 carcinoma Ma44-3 Lung squamous cell Kazuya Kondo I		fluorescent protein		
Ma44-3 Lung squamous cell Kazuya Kondo I	SW900	Lung squamous cell	ATCC	I
	HTB-59	carcinoma		
carcinoma (University of	Ma44-3	Lung squamous cell	Kazuya Kondo	I
(011, 1101) 02		carcinoma	(University of	
Tokushima, Japan),			Tokushima, Japan),	
NCI-H520 Lung squamous cell ATCC I	NCI-H520	Lung squamous cell	ATCC	I
HTB-182 carcinoma	HTB-182	carcinoma		
Calu-3 Lung adenocarcinoma ATCC I	Calu-3	Lung adenocarcinoma	ATCC	I
HTB-55	HTB-55			
NCI-H23 Lung adenocarcinoma ATCC I	NCI-H23	Lung adenocarcinoma	ATCC	I
CRL-5800	CRL-5800			

A549	Lung adenocarcinoma	ATCC	I,III,IV
CCL-185			
201T	Lung adenocarcinoma	Jay D. Hunt (Louisiana State University, New Orleans, LA),	Ι
M4A4-LM3	Breast cancer	S.Goodison (University of Florida, FL)	II
Hs173we CRL- 7834	Human fibroblast	Gerd Bauerschmitz, (University of Duesseldorf, Germany)	III

Fresh NSCLC primary tissue samples were collected after signed informed consent from patients diagnosed with advanced lung cancer at the Helsinki University Central Hospital. The study was approved by the hospital Ethics Committee (I). Bone marrow-derived human mesenchymal stem cells were isolated as described previously (Leskelä *et al.* 2003). Adipose stem cells were obtained from human subcutaneous and intraperitoneal adipose tissues (II).

2 Adenoviral vectors and replicating adenoviruses (I-IV)

Table 4. Replication deficient and oncolytic adenoviruses used in this study

Virus	E1A	Transgene	Capsid	Target	Source	Study
			modification	receptor		
Adluc1	Deleted	Luciferase	-	CAR	(Dmitriev	I,II,
					et al.	IV
					1998).	
A5/3luc1	Deleted	Luciferase	Serotype 3	Unknown	(Kanerva	I, IV
			knob	and CD46	et al.	
					2002a).	
Ad5lucRGD	Deleted	Luciferase	RGD motif	$\alpha_{\rm v} \beta$	(Dmitriev	I,II,
			in the HI	integrins	et al.	IV

			loop	and CAR	1998).	
Ad5(GL)	Deleted	Luciferase	-	CAR	(Wu et al.	I, II,
		and GFP			2002).	IV
Ad5.pK7(GL)	Deleted	Luciferase	7 polylysins	Heparan	(Wu et al.	I, II,
		and GFP	in the C-	sulfates	2002).	IV
			terminus	and CAR		
Ad5.RGD.pK7(GL)	Deleted	Luciferase	RGD motif	$\alpha_v \beta$	(Wu et al.	I, II
		and GFP	in the HI	integrins,	2002).	
			loop, seven	Heparan		
			polylysins in	sulfates,		
			the C-	CAR		
			terminus			
Ad5.LacZ	Deleted	β-	-	CAR	(Sarkar et	I
		galactosidase			al. 2005).	
Ad5.pK7LacZ	Deleted	β -	seven	Heparan	(Sarkar et	II
		galactosidase	polylysins in	sulfates,	al. 2005).	
			the C-	CAR		
			terminus			
Ad5F/K21	Deleted	β-	21 polylysins	Heparan	(Sarkar et	I
		galactosidase	in the C-	sulfates,	al. 2005).	
			terminus	CAR		
Ad5Cox2Lluc	Deleted	Cox-2	-	CAR	This study	
		promoter			unpublised	
		luciferase			data	
Ad5Cox2LlucUL	Deleted	Cox-2	-	CAR	This study	
		promoter			unpublised	
		Luciferase,			data	
		3'UTR				
Ad300wt	Intact	-	-	CAR	ATCC	I,II,
(Ad5 wild-type)						III
Ad5Δ24E3+	24-bp	-	-	CAR	(Suzuki et	I, II,
	deletion				al. 2001).	III
	in CR2					

Ad5/3 -Δ24	24-bp	-	Serotype 3	Unknown,	(Kanerva	I, II,
	deletion		knob	CD46	et al.	III
					2003).	
Ad5- Δ24RGD	24-bp	-	RGD motif	$\alpha_{\rm v} \beta$	(Suzuki et	I, II
	deletion		in the HI	integrins,	al. 2001).	
			loop			
Ad5.pK7-Δ24	24-bp	-	seven	Heparan	(Ranki et	I, II
	deletion		polylysins in	sulfates,	al. 2007).	
			the C-			
			terminus			

2.1 Construction of Cox-2 and 3'UTR element-expressing plasmids (III)

In the plasmids pShuttleCox-2Lluc and pShuttleCox-2Lluc + 3'-UTR, the Cox-2L promoter (-1492/+59) was amplified by PCR from full-length Cox-2 containing pGL3/Cox-2 using forward primer 5'-AAAAGATCTGAGGTACCTGGT- 3'to place a *BgIII* site at the 5'-end of Cox-2L and reverse primer 5'-

AAAAAGCTTCGCTGAGGA-3' to place a *HindIII* site at the 3'-end. A 603-bp long 3'-UTR fragment from Cox-2 was amplified from pZeo/Luc3 + 3'-UTR using forward primer 5'-AAAATCTAGAAAGTCTAATGATC-3' and reverse 5'-

AAAATCTAGAAACTTTAAG-3' to place *XbaI* at both ends. The Cox-2L promoter and 3'-UTR were cloned into luciferase-containing plasmid pGL3basic. The final constructs: pShuttleCox-2Lluc and pShuttleCox-2Lluc + 3'-UTR were created by digesting pGL3Cox-2Lluc and pGL3Cox-2Lluc + 3'-UTR with *KpnI* and *SalI* and subcloning into pShuttle. pShuttleCMVluc and pShuttleCMVluc + 3'-UTR were also created: the luciferase gene and 3'-UTR were digested with *KpnI* and *SalI* from pGL3 basic and inserted into pShuttleCMV. All plasmids were purified by agarose gel electrophoresis.

Cox-2L 1491 bp	Luciferase gene 1656bp	3'UTR 603bp	polyA	
Cox-2L 1491 bp	Luciferase gene 1656bp	polyA		
CMV 592 bp	Luciferase gene 1656bp	3'UTR 603bp pol		
CMV 592 bp	Luciferase gene 1656bp	polyA		

Figure 6. Schematic presentation of the plasmids constructed in this study: pShuttleCox-2Lluc + 3'-UTR, pShuttleCox-2Lluc, pShuttleCMVluc + 3'-UTR and

pShuttleCMVluc.

Table 5. The plasmids used in this study

Plasmid	Description	Source	Used	
			in	
pGL3basic	pGL3 cloning vector	Promega, Madison, WI	III	
pGL3/Cox-2	pGL3 vector containing full-length	Ari Ristimäki,	III	
	Cox-2	University of Helsinki,		
		Finland		
pZeo/Luc3 + 3'-UTR	pZeo/Luc3 vector containing full-	Ari Ristimäki	III	
	length 3'UTR			
pShuttleCMV	pShuttle vector containing CMV	Stratagene, La Jolla, CA	III	
	promoter			
pShuttleCox-2Lluc	pShuttle vector containing Cox-2L	This study	III	
	promoter driving luciferese			
pShuttleCox-2Lluc +	pShuttle vector containing Cox-2L	This study	III	
3'-UTR	promoter driving luciferese and			
	3'UTR element			
pShuttleCMVluc	pShuttle vector containing CMV	This study	III	
	promoter driving luciferase			
pShuttleCMVluc +	pShuttle vector containing CMV	This study	III	
3'-UTR	promoter driving luciferase and			
	3'UTR element			
pAdEasyI	E1/E3 deleted Ad5 genome	This study unpublished		
		data		

2.2 Construction of non replicating adenoviruses

Previously described plasmids pShuttleCox-2luc and pShuttleCox-2luc+3'UTR were digested with *PmeI* and homologous recombination was done with AdEasy I plasmid. Resulting plasmid was further digested with *PacI* and transfected with 293 cells.

2.3 *In vitro* transfection of plasmids (III)

 1.5×10^5 cells were seeded into 12-well plates 17–20 h prior to transfection; 1 µg of each plasmid were transfected with Fugene 6 reagent (Roche, Basel, Switzerland) in a ratio of 3:2 (Fugene 6/plasmid) following the manufacturer's instructions.

2.4 High titer production of viruses (I-IV)

Replication-deficient and oncolytic adenoviruses were propagated in 293 and A549 cells, respectively. Viruses were purified by standard cesium chloride gradient techniques, and quality control was performed by PCR. Viral particles (vp) were determined by spectrophotometry, infectious units are expressed as the median tissue culture infective dose₅₀ (TCID₅₀) (Svensson *et al.*1999).

3 In vitro studies

3.1 Adenovirus-mediated gene transfer assays (I-III)

Cells and tissue samples were infected with Ad5luc1, Ad5(GL), Ad5/3luc1, Ad5lucRGD, Ad5.pK7(GL), or Ad5.RGD.pK7(GL) for 30min and 1 hour respectively. Luciferase activity was determined 24 h postinfection with a luciferase assay system (Promega) according to the manufacturer's protocol. Values are presented relative to syngeneic control viruses [Ad5luc1 and Ad5(GL)], which were assigned a value of 1. For immunohistochemistry, NCI-H460 cells were infected in chamber slides (Nunc, Naperville, IL). The next day, the cells were fixed with 10% formalin and permeabilized with 0.1% Tween 20/phosphate-buffered saline (PBS). Blocking was done with 1% bovine

serum albumin (BSA)/PBS for 30 min at room temperature. Antihexon was used as a primary antibody at a concentration of 1.5 mg/ml (LabFrontier, Seoul, Korea). Texas Redlabeled secondary antibody (AbCAM, Cambridge, UK) was used at a concentration of 2 mg/ml. Cells were stained with Hoechst dye (Oriola, Espoo, Finland), positive cells were counted, and fluorescence images obtained (Olympus Ix81, Biosystems, Munich, Germany).

3.2 Determination of receptor expression by flow cytometry (II)

For flow cytometric analysis, cells (2×10^5) were incubated for 20 min at 4°C with the following primary antibodies at a 1:200 dilution: anti-CAR (clone RmcB; Upstate Cell Signaling Solutions/ Millipore, Lake Placid, NY), anti-HSPG (clone F58-10E4; Seikagaku, Falmouth, MA), anti- $\alpha_v\beta_3$ integrin (clone LM609; Chemicon International/Millipore, Temecula, CA), anti- $\alpha_v\beta_5$ integrin (clone P1F6; Chemicon International/Millipore), or anti-CD46 (clone E4.3; BD Biosciences, San Jose, CA). Cells were washed with fluorescence-activated cell-sorting (FACS) buffer (phosphate buffered salin, PBS containing 2% serum) and incubated with a 1:200 dilution of phycoerythrin-labeled secondary antibody (goat anti-mouse polyclonal antibody; BD Biosciences) for 30 min at 4°C. Cells were washed with FACS buffer and analyzed by flow cytometry (FACSCalibur; BD Biosciences) to determine receptor expression levels. Unstained cells were used as a negative control.

3.3 Cytotoxicity assay (I, II)

The 3-(4, 5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl) 2-(4-sulfophenyl-2H-tetrazolium, inner salt ["MTS"] cell killing assay was performed, and viability was measured using the CellTiter 96 AQ_{ueous} One Solution Cell Proliferation Assay (MTS assay; Promega), when at least one tested virus showed a cytotoxic effect in the two highest dilutions. In the crystal violet assay, cells were stained with crystal violet when the cytotoxic effect was as above.

3.4 Semiquantitative polymerase chain reaction (III)

Transfected Hs173We cells were incubated with Proteinase K (Qiagen, Valencia, CA, USA) at 37°C for 30 min. Proteinase K was then inactivated by 20 min of incubation at

95°C. Amplification (30, 40, and 60 cycles, annealing at 50°C) of a 77-bp DNA fragment was carried out by using primers that were positioned at the left end of the adenovirus genome, between the left inverted terminal repeat and the packaging signal. Upstream: 5'-GCGACGGATGTGGCAAAAGT-3'(150-171bp) and downstream: 5'-CCTAAAACCGCGCGAAAA-3' (211-228bp).

3.5 Assessment of neutralizing antibody (NAb) titers (IV)

A549 cells were seeded on 96-well plates and cultured overnight. Serum collected from the immunized ICR mice was incubated at 56°C for 60 min to inactivate the complement and then a 4-fold dilution series of serum was prepared in growth media without serum. Ad5(GL) (100 vp/cell) was mixed with diluted serum samples and incubated at room temperature on a rocker for 30 min, and the mix was added to cells in quadruplicates. Two hours later, the infection media was replaced by fresh growth media. After 48 h, luciferase expression was measured by FluorStar Optima luminometer (BMG LabTech, Offenburg, Germany) according to the manufacturer's instructions. NAb titer was determined as the lowest degree of dilution that blocked gene transfer by more than 50%.

3.6 Effect of NAbs on gene transfer (IV)

Human NSCLC and normal lung tissue samples were homogenized and infected with Ad5(GL), Ad5/3luc1, and Ad5lucRGD (1000 vp / cell). Viruses were preincubated with (0, 1:512, and 1:4 dilutions) inactivated serum from Ad5(GL)-immunized mice at room temperature for 30 min before infection of tissue explants. A luciferase assay was performed in quadruplicate 24 h later by the using Luciferase Assay System (Promega).

4 In vivo studies

4.1 Mice and mice models used (I-IV)

Female NMRI nude mice and immunocompetent ICR mice aged 3-4 weeks were obtained from Taconic (Ejby, Denmark) and quarantined for 2 weeks. All animal experiments were

approved by the Experimental Animal Committee of the University of Helsinki and the Provincial Government of Southern Finland.

Subcutaneous models were created as follows: 1×10^6 293 and 5×10^5 A549 cells were injected into the flanks of mice. Tumors were allowed to develop for 30 days (293) or 14 days (A549) to reach 5 mm in diameter. An orthotopic model of locally advanced lung cancer was created by injecting $2x10^6$ cells (LNM35/EGFP) into the left lung of NMRI nude mice in a total volume of 200 ml using a syringe with a 27G needle. The orthotopic breast cancer model used in these studies has been described in more detail by Ranki *et al.* (Ranki *et al.* 2007b). Briefly, 2×10^6 M4A4-LM3 cells were injected into both the right and the left second lowest mammary fat pads. Tumors were allowed to develop until they reached a diameter of approximately 4 mm. Before injections, the mice were anesthetized with medetomidine (Domitor; Orion Pharma, Espoo, Finland) and ketamine (Ketalar; Pfizer, New York, NY) at 1:2.

4.2 Biodistribution assay (I, IV)

After tumors had developed (I) or 1 month after immunization (IV) $3x10^{10}$ vp were injected through the tail vein, and 48 h later, organs were harvested and frozen at -80°C. Organs were homogenized in Cell Culture Lysis Buffer (Promega), and luciferase expression was measured by using FluoStar Optima luminometer.

4.3 Therapeutic assay (I, II)

Advanced intrapulmonary disease was allowed to develop for 5 days and mice were then divided randomly into 5 groups (n=12 mice per group) (**I**) and (n=6-10 mice per group) (**II**). Viruses were injected into the tail vein (2x 10¹⁰ vp/mouse). Mice were treated on days 5, 12, and 19 (**I**). Viruses injected along with MSC groups are as follows: either MSCs (7x10⁵ cells/mouse, n=5), virus [1.4x 10⁸ vp of Ad5.pK7(GL) ,7 x10⁷ vp of Ad5.pK7- Δ24, n=10], MSCs infected with nonreplicating adenovirus [Ad5.pK7(GL) at 200 vp/cell, Ad5pK7LacZ at 100 vp/cell, 7x10⁵ cells/mouse, n=6] or MSCs infected with oncolytic adenoviruses [Ad5.pK7(GL) at 200 vp/cell, Ad5.pK7-Δ24 at 100 vp/cell, 7x10⁵ cells/mouse, n=10] 7, 11, and 15 days after cancer cell inoculation. In the breast cancer model, tumors were measured in two dimensions with calibers and tumor volumes were

calculated according to the following formula: length x width 2 x 0.5. The health of the mice was monitored daily, and mice were killed according to humane endpoint guidelines.

4.4 In vivo transfection of plasmids and infection of viruses (III)

Transfection was accomplished by using *in vivo* JetPei cationic polymer transfection reagent (Poly transfection, Illkirch, France) following the manufacturer's protocol. Twenty four hours after transfection, 5×10^9 vp/tumor of Ad300wt in 100 μ l was injected intratumorally. Tumors were collected and luciferase expression was analyzed as described previously in the biodistribution assay.

4.5 Noninvasive *in vivo* imaging (I-III)

Imaging was done by using the IVIS 100 Imaging System (Xenogen, Alameda, CA). Fluorescence emission was quantitated by determination of regions of interest, which were normalized to reference regions in the same image, but without tumor. When luciferase was imaged, luciferin substrate (4.5 mg/mouse) was injected intraperitoneally before bioluminescence imaging.

4.6 Induction of NAbs (IV)

ICR mice were immunized with viruses subcutaneously. Injection of $3x10^8$ vp occured either once on day 0 or thrice on days 0, 3, and 6. Control animals were mock-immunized with serum-free media. On day 27, blood was collected from the hindlimb artery and allowed to clot, and serum was then separated by centrifugation and stored at -20°C for further experiments.

4.7 Effect of NAbs on gene transfer (IV)

Orthotopic tumors of advanced lung cancer were allowed to develop for 9 days before mice were injected with either viruses or serum at 1:512 or 1:4. Serum was pooled from ICR mice immunized once or thrice with Ad5(GL). Tumors were harvested 48 h later and luciferase assay was performed by FluorStar Optima. The data are protein-normalized (to

account for possible differences in tumor size) and presented relative to gene transfer obtained with no NAb.

5 Statistics (I-IV)

Pairwise comparisons were performed with Student's t test, one-way analysis of variance (ANOVA) with Bonferroni's post-hoc test, or ANOVA followed by Dunnett's pairwise multiple comparison t-test. Survival data were analyzed with a log-rank test using SPSS software (version 11.5; SPSS Inc., Chicago, IL) or PROC LIFETEST (SAS version 9.1; SAS Institute, Cary, NC), using a Weibull distribution. Analysis of whether MSCs could improve the efficacy of oncolytic virus was performed with χ^2 test (II). For comparison of tumor size, data were analyzed with a repeated measures growth model with PROC MIXED (SAS version 9.1). The effects of treatment group, time, and the interaction of treatment group and time were evaluated by F- tests (II). Analysis of TCID⁵⁰ data was performed with an unpaired t- test with Welch's correction (II). The results from Cox-2 induction $in\ vivo\ (III)$ were analysed with Multtest, version 9.1 (SAS Institute Inc., Cary, NC, USA). For all analyses, p<0.05 was deemed significant.

Results and discussion

1 Modified adenoviruses show enhanced gene transfer to NSCLC cell lines, MSCs, and primary lung cancer tissues *in vitro* (I, II)

One of the major concerns in adenoviral gene therapy is effective targeting since the primary receptor CAR exhibits variable expression on cancer cells (Bauerschmitz et al. 2002; Kanerva et al. 2002b). We hypothesized that by using modified adenoviruses, we could enhance gene transfer to NSCLC cell lines, bone marrow, or fat tissue-derived human mesenchymal stem cells and lung cancer tissue samples fresh from patients. Ten NSCLC cell lines (Figure 1 in Study I) were infected with Ad5luc1, Ad5/3luc1, Ad5lucRGD, Ad5(GL), Ad5. pK7(GL), Ad5.pK7.RGD(GL), Ad5LacZ, and Ad5F/K21 replication-deficient adenoviruses (Table 4). On the basis of our results, $\alpha_v \beta$ integrins and HSPG targeted viruses enhanced transduction up to 270- and 100-fold respectively in NSCLC cell lines. To examine whether the increased gene transfer was due to enhanced infection of the cells, we stained Ad5luc1-, and Ad5lucRGD-infected cells with antihexon antibody, which confirmed the hypothesis. In addition, we investigated gene transfer in primary NSCLC samples fresh from patients without any passaging, since geno- and phenotypes of clinical tumor specimens better resemble human tumor tissue. The results of four adenocarcinoma and two squamous cell carcinoma samples were in line with those of the NSCLC cell lines. The HSPG targeted Ad5pK7(GL) virus had up to 200-fold better gene transfer compared to the unmodified control virus (p < 0.001) (Figure 2 in Study I). Both unmodified and genetically engineered MSCs have been widely used for various therapeutic purposes preclinically, and a number of studies have suggested an immuneprivileged nature for MSCs and their ability to avoid allogeneic rejection in human and animal models, broadening their potential for therapeutic use (Aggarwal and Pittenger 2005; Chamberlain et al. 2004; Le Blanc et al. 2004). Antitumor activity of retrovirus vector-transduced MSC, producing IL-12, has been tested in a mouse melanoma model(Elzaouk et al. 2006), and MSC, transduced by a lentivirus vector to express human

OPG (osteoprotegerin) has been evaluated for the treatment of osteolysis in a myeloma bone disease model (Rabin *et al.* 2007). Modified MSCs loaded with adenoviruses have been successfully used preclinically as treatment alternatives for glioma, melanoma, and ovarian and breast cancers (Komarova *et al.* 2006; Stoff-Khalili *et al.* 2007b; Stoff-Khalili *et al.* 2007b; Studeny *et al.* 2002; Studeny *et al.* 2004b; Sonabend *et al.* 2008). Unfortunately, the infectivity of MSCs with serotype Ad5 viruses has been limited due to low expression of CAR (Conget & Minguell 2000; Mizuguchi *et al.* 2005; Studeny *et al.* 2002).

Based on our results, $\alpha_v \beta$ integrin- or HSPG-targeted adenoviruses increased gene transfer 1100- and 11000-fold compared with the untargeted virus. To further confirm the data, we studied the expression levels of target receptors on MSCs by flow cytometry. Interestingly, the most expressed receptor was CD46, a proposed receptor of Ad3, although only slightly enhanced transduction of Ad3-targeted virus was seen (Figure 1 in Study II). These results are in accordance with data suggesting that CD46 is not the only receptor of Ad3 (Gaggar *et al.* 2003; Tuve *et al.* 2006b).

2 Biodistribution of capsid-modified adenoviruses (I)

The orthotopic murine model of advanced NSCLC created in this study may resemble the actual disease to some degree since it is very aggressive. The disease spreads from the primary tumor (located in the left lung with this model), to the mediastinal lymph nodes and later to the peritoneum. Replication-deficient viruses Ad5luc1, Ad5/3luc1, Ad5lucRGD, Ad5(GL), Ad5. pK7(GL), and Ad5.pK7.RGD(GL) were used to examine biodistribution after systemic delivery (Figure 4 in Study I). Although systemic delivery of viruses might be useful for treatment of disseminated diseases, there are many hurdles that a therapeutic vector faces in the bloodstream, such as the innate and adaptive immune response, potential adsorption to erythrocytes and thrombocytes and clearance by specialized cells in the liver and spleen (Baker *et al.* 2007; Mohr *et al.* 2008). In our study, no statistical differences were observed between viruses. Although Ad5lucRGD and Ad5. pK7(GL) viruses showed moderately high gene delivery to the

lungs, kidneys, peritoneum, and ovaries, transduction to the tumor was also seen with the same viruses. The route of administration clearly also affects biodistribution since when administered systemically, adenovirus is exposed to many factors in the blood, such as coagulation factors IX and X and complement component C4-binding protein, which might determine the biodistribution of the virus (Shayakhmetov *et al.* 2005).

3 Capsid-modified adenoviruses display improved cell killing efficacy in NSCLC in vitro and in vivo (I)

Even though efficacy has been seen following local administration, nonreplicating adenoviral agents have not yet demonstrated systemic therapeutic efficacy in humans. To overcome this problem, oncolytic adenoviruses have been developed. An advantage gained by using replication-competent oncolytic viruses is the potential for local amplification within the tumor, following initial transduction of even a small number of cells. In theory, this oncolytic process continues as long as target cells for the virus persist. This process when rigorously controlled in the oncolytic adenovirus context could be of enormous therapeutic benefit. Oncolytic potency of replicating agents is mostly determined by the infectivity of the viruses (Douglas *et al.* 2001; Hemminki *et al.* 2001). Unsurprisingly, targeted oncolytic adenoviruses have shown impressive preclinical efficacy (Kanerva *et al.* 2003; Ranki *et al.* 2007; Suzuki *et al.* 2001).

We evaluated the potency of oncolytic adenoviruses harboring the same capsid modifications with a cell viability assay *in vitro* (Figure 3 in Study **I**). In contrast to previous results with replication-deficient viruses, Ad5/3-Δ24 (Table 4) showed the best cell killing efficacy in all cell lines tested. The finding could be explained by the structure of RGD-4C- or pK7-containing virions, which may lead to less rapid packaging (in comparison to gene delivery), perhaps due to the "sticky" nature of these moieties. *In vivo*, with the same panel of viruses in an orthotopic model of advanced lung cancer, the median survival times times were 19, 22, 23, 26, 26, and 26 days for no virus, Ad5luc1, Ad5/3-Δ24, Ad5-Δ24E3, Ad5.pK7-Δ24, and Ad5-Δ24RGD, respectively (Figure 5a in Study **I**). The therapeutic efficacy of Ad5-Δ24E3, Ad5.pK7-Δ24, and Ad5-Δ24RGD was

significantly enhanced compared with the untreated group (p<0.01). One mouse from the Ad5- Δ 24RGD-treated group survived until the end of the experiment (93 days). Ad5/3- Δ 24 virus did not enhance survival of the mice, which might be partly explained by the biodistribution data. This finding is in accordance with data obtained from an orthotopic breast cancer model (Ranki *et al.* 2006). The route of administration clearly has an effect on the infectivity and oncolytic potency of Ad5/3- Δ 24 since after intraperitoneal administration of Ad5/3- Δ 24 in an orthotopic ovarian cancer model, survival was enhanced dramatically (Kanerva *et al.* 2003).

4 MSCs target lungs in vivo (II)

To use MSCs as carriers for adenoviruses, they should support the replication of oncolytic adenoviruses. To study replication and subsequent cytolysis, we infected in vitro passaged MSCs with a panel of oncolytic adenoviruses containing a 24 bp deletion in the E1A region and cell viability was analyzed by MTS assay (Figure 2 in Study II). The oncolysis correlated with the gene transfer of the viruses with the same targeting moieties (Figure 1 in Study II). HSPG- and $\alpha_v \beta$ integrin-targeted oncolytic viruses showed significantly higher cytolysis rates in both adipose tissue- and bone marrow-derived cells. Ad5/3- Δ 24 virus also enhanced cell killing significantly compared with the unmodified virus (All p < 0.001). The data revealed that cytolysis correlates with gene transfer. Previous finding of Komarova et al. showed that MSCs enable the replication of 5/3 targeted adenovirus with intact E1 (Komarova et al. 2006). Our viruses have a 24 bp deletion in E1A, allowing them to replicate in cells with a defective p16/Rb pathway. However, in vitro passaged MSCs feature phosphorylation of the Rb protein, which would be expected to allow replication of Δ 24-type viruses, explaining the replication seen here (Heise et al. 2000a). We found no differences between the ability of adipose tissue- and bone marrow-derived cells to support the replication and allow release of active virions.

Since MSCs have been shown to target tumors and act as a shield to pre-existing antibodies (Aggarwal & Pittenger 2005; Le Blanc *et al.* 2004; Mohr *et al.* 2008), these engineered stem cells might serve as therapeutic Trojan horses, therefore enhancing the

therapeutic efficacy *in vivo* compared with the virus alone. Recently, Gutova *et al.* provided evidence that urokinase plasminogen activator (uPA) and its receptor uPAR, involved in chemotaxis and cell guidance during normal development, are upregulated in invasive tumors and might augment neural and mesenchymal stem cell tropism (Gutova *et al.* 2008).

In this study, we tested whether bone marrow-derived MSC could support viral replication and release in the context of orthotopic lung tumors. Mice received a single dose of intravenous oncolytic adenovirus (Ad5.pK7-Δ24) or MSCs infected either with replicationdeficient or oncolytic adenovirus. Luciferase-expressing virus was included in each group, which allowed us to use the noninvasive bioimaging system to follow the biodistribution of the virus (Figures 3 and 4 in Study II). Mice were imaged 15 min, and 3, 7, and 10 days after treatment. Initially, the luciferase signal was detected from the lungs of all MSCtreated mice, but not in the virus-alone group because of a lack of sufficient time for luciferase production. On the third day, the oncolytic virus-loaded MSCs displayed a strong luciferase signal from the liver, suggesting that viruses had escaped from the MSCs and transduced the liver. A 60 time weaker luciferase signal was captured from the livers of mice treated with virus only. Replication-deficient virus with MSCs showed weak luciferase expression in the lungs. On the next imaging days, a weak luciferase signal was detected from the liver of both groups of MSC-treated mice, but not mice treated with virus alone (Figure 3 in Study II). In addition, the kinetic studies revealed that viral release from MSC carriers was relatively rapid because 24 h after MSC injection (i.e. 72 h after infection) some of the released virus was already detectable in the liver. Further, Dillabeled MSCs were injected intravenously into tumor-free, lung tumor- or breast tumorbearing mice to see more clearly whether the luciferase signal was from the lungs or from the tumor. A strong fluorescence signal was obtained symmetrically from both lungs, suggesting accumulation preferentially in the lungs instead of specificity for tumors. No MSCs were found in breast tumors. In conclusion, we were able to demonstrate that MSCs loaded with oncolytic viruses homed to lungs of mice bearing orthotopic tumors. In contrast to previous reports, no tumor tropism was seen (Komarova et al. 2006; Studeny et al. 2004b).

5 Oncolytic adenoviruses and MSCs loaded with oncolytic adenoviruses show therapeutic efficacy in an orthotopic murine model of advanced lung and breast cancers (II)

Therapeutic efficacy of MSCs packed with oncolytic adenoviruses was investigated in orthotopic models of advanced lung and breast cancers (Figure 6 in Study II). Mice were treated with virus only (Ad5.pK7- Δ 24),MSC only, MSC + oncolytic adenovirus (Ad5.pK7- Δ 24), MSC + replication-deficient adenovirus (Ad5pK7LacZ) and median survival times were 23, 25, 27, and 28 days, respectively, for lung tumor-bearing mice, and survival was significantly enhanced (p=0.0031) with MSC loaded with oncolytic adenovirus. In an advanced breast cancer model the MSCs with oncolytic adenoviruses inhibited tumor growth significantly (p<0.001) compared with control mice. These results suggest that despite MSCs lacking a specific tumor-targeting capacity, they were able to release the virus for subsequent transduction of tumors, and enhanced therapeutic efficacy was seen.

6 In vivo monitoring of oncolytic adenoviruses (I, II)

Imaging has become an indispensable tool in cancer research, clinical trials, and medical practice. Imaging systems can be grouped by the energy used to derive visual information (X-rays, positrons, photons, or sound waves) (Groot-Wassink *et al.* 2004), the spatial resolution attained (macroscopic, mesoscopic, or microscopic), or the type of information obtained (anatomical, physiological, cellular, or molecular). In the context of adenoviral gene therapy, longitudinal monitoring of tumor burden and viral spreading has advantages over survival alone.

In this study, fluorescence and bioluminescence imaging were used to follow up tumor burden and the spreading of adenoviruses *in vivo*. Overall, imaging data correlated well with date achieved from biodistribution and survival experiments. The fluorescence imaging used in Study I (Figure 6) even revealed individual variation in mice treated with oncolytic adenoviruses. Mice treated with Ad5-Δ24RGD were grouped according to

response. Some mice had progressive disease, the disease stabilized in others, and others had dramatic responses. In addition, imaging allowed us to track MSCs loaded with the adenoviruses more precisely in mice (Figures 3-5 in this Study II)

7 Cox-2 and CMV promoter are induced following adenovirus infection, but the Cox-2 3'UTR element can retain specificity *in vitro* and *in vivo* (III)

Achievement of oncolytic replication in tumor cells while avoiding nonspecific replication in normal tissues remains a major challenge in oncolytic gene therapy. In this regard, the elucidation of the critical pathways involved in the adenoviral infectious cycle has provided the basis for multiple levels of control to achieve the goal of selective replication. Tumor-specific promoters are at the foundation of transcriptional targeting, and Cox-2 promoter has been among the more popular candidates (Bauerschmitz et al. 2006; Casado et al. 2001; Hirschowitz et al. 2002; Liang et al. 2004; Nettelbeck et al. 2003; Yamamoto et al. 2001; Yamamoto et al. 2003; Kanerva et al. 2005). However, the physiological function of Cox-2 as an enzyme expressed upon inflammation, infection and injury as well as in tumors is very well established (Kujubu et al. 1991; Smith 1992; Turini & DuBois 2002). The Ras/MAPK group, one of the common pathways to control the cell cycle, is mutated in most cancers. Cox-2 expression is both upregulated transcriptionally and downregulated posttranscriptionally by the RAS/p38/MAPK pathway (Lasa et al. 2000). Downregulation is in partly through AU-rich elements within the 3'-UTR, and their physiological function is to rapidly downregulate Cox- 2 mediated transcripts, when the threat (e.g. infection) has passed. Naturally, cancer cells have learned to take advantage of this mechanism and Cox-2 is stabilized through defected Ras pathway in tumor cells (Cok & Morrison 2001; Dean et al. 1999; Dixon et al. 2000). In this study, we hypothesized that Cox-2 promoter is induced by adenoviral infection, but the specificity could be regained by utilization of 3'UTR elements. First, nonmalignant 293 cells were transfected with pShuttleCox2Lluc plasmid featuring Cox-2L promoter driving luciferase, and the cells were then infected with a wild-type Ad5, a selectively

oncolytic adenovirus Ad5- Δ 24E3, or an E1-deleted adenovirus (Ad5LacZ) (Figure 2 A-C in Study III). Infection with a wild-type adenovirus or a conditionally replicating virus resulted in significant Cox-2 induction (p<0.05 and p<0.01, respectively). Replication-deficient adenovirus also induced Cox-2 in these E1-containing cells, which allows replication of E1-deleted viruses (p<0.05). Further, we sought to investigate whether this Cox-2 induction is a cell type-specific phenomenon, and human fibroblasts were transfected and infected as previously, except that Ad5/3- Δ 24 virus was used instead of Ad300wt. Induction of the promoter seemed to be related to replication or E1 proteins since Cox-2 was induced by replication-competent adenoviruses but not by replication-deficient Ad5LacZ virus (Figure 3 in Study III). In addition, Ad5/3- Δ 24 virus, which replicates more efficiently in most cell types, also (Bauerschmitz *et al.* 2006; Kanerva *et al.* 2003). induced higher Cox-2 levels (Figures 3 and 4 in Study III).

When the same cell lines were transfected with a plasmid containing 3'UTR element after the luciferase gene (pShuttleCox2Lluc+3'UTR) and infected with the same viruses, the 3'UTR element was able to abrogate induced Cox-2 expression in nonmalignant cells, but had little impact on malignant cells (Figures 2–4 in Study III). Interestingly, our control plasmids with CMV promoter with or without 3'UTR showed that CMV promoter is also induced by adenoviral infection to a similar degree as the Cox-2 promoter in all cell lines, and the UTR element also had a similar activity on the CMV promoter (Figures 2-4, panels D-F).

The induction of Cox-2 and CMV promoters was also studied *in vivo*. Subcutaneous 293 or A549 tumors were allowed to develop 30 and 14 days, respectively, and were then transfected with plasmids followed by intratumoral injection of wild-type adenovirus the next day. To investigate the induction, bioluminescence was imaged 15 min and 24 h after infection. Interestingly, there was a 4.3-fold induction in Cox-2 expression only 15 min after infection and this persisted to the next imaging. 3'UTR did not have time to affect expression during the first round of imaging, but 24 h later luciferase expressed by the Cox-2–3'UTR complex in infected cells decreased from 55 039 to 51 716, whereas in uninfected cells the luciferase continued to increase as expected (from 4339 to 14 519). The destabilization signal therefore appeared to work as predicted. However, there were variations typical of *in vivo* luciferase imaging and the complexities associated with *in vivo*

plasmid transfection followed by intratumoral infection, and therefore, no statistically significant differences between groups could be seen. After the second round of imaging, the tumors were excised and the amount of luciferase protein was analyzed. The results were in line with imaging data. Adenovirus infection induced expression of the Cox-2 and CMV promoters in both nonmalignant and malignant cells, and the 3'UTR element had an effect on the induction in 293 cells (Figures 5 and 6 in Study III).

In contrast to the promising data achieved from Cox-2 and 3'UTR expressing plasmids, the effect of the 3'UTR element was not powerful enough to downregulate the expression Cox-2 promoter driven luciferase in the context of replication deficient adenoviruses in human fibroblasts (Figure 5, unpublished data).

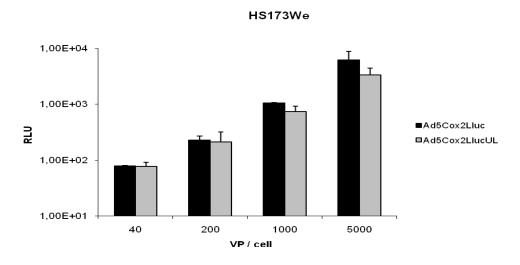


Figure 5. Replication deficient adenovirus expressing 603bp long 3'UTR element has a minor impact on downregulation the expression of luciferase. Human fibroblasts were infected with Ad5Cox2Lluc and Ad5Cox2LlucUL with, 40, 200, 1000 and 5000 vp/cell. Luciferase was imaged 24h after infection.

8 Switching the adenoviral fiber retains viral gene transfer in the presence of NAbs (IV)

The use of adenoviral vectors in gene therapy has been hampered by significant problems with attendant host immune responses that limit their safety and efficacy *in vivo* (Wilson 1996). Adenoviral vectors can effectively elicit the innate immune response immediately after infection, leading to the secretion of proinflammatory cytokines and chemokines in mice, humans, and nonhuman primates and that might damage healthy tissue (Raper *et al.* 2003; Schnell *et al.* 2001; Zhang *et al.* 2001). Problems particularly may arise after systemic delivery or upon readministration of adenoviruses. Recently, Zhu *et al.* described adenoviruses to elicit an innate immune response through the TLR9-MyD88 pathway in plasmacytoid DCs (pDCs) and a TLR-independent pathway in non-pDCs, leading to production of mainly type I IFNs (Zhu *et al.* 2007a).

We investigated whether the changes in the adenoviral fiber could allow the virus to escape from NAbs. To test the gene transfer of Ad5 in the presence of NAbs, immunocompetent mice were immunized with capsid-modified replication-deficient adenoviruses (see Table 6), and the inhibitory effect on Ad5(GL) gene transfer efficacy was analyzed. We used a single immunization to model a low level of NAb and a triple immunization to simulate a high NAb titer. Triple immunization with Ad5(GL) resulted in complete blocking of *in vitro* gene transfer by Ad5(GL). By contrast, after triple immunization with Ad5/3luc1 or Ad5pK7(GL), Ad5(GL) was not blocked. Anti-Ad5lucRGD NAbs marginally inhibited Ad5(GL) gene transfer (Table 2 in Study IV). These are consistent with previous results showing an ability of RGD or 5/3 modifications of the Ad5 fiber to avoid pre-existing anti-Ad5 NAb present in human serum or ascites (Hemminki et al. 2001; Kanerva et al. 2003). Immunized mice were injected intravenously with Ad5(GL), and organs were harvested for luciferase measurement (Table 6). A single immunization with Ad5(GL) reduced subsequent gene transfer with Ad5(GL) to the liver (p<0.001), heart (p<0.001), spleen (p<0.05), pancreas (p<0.05), and kidney (p<0.01), in contrast to a single immunization with the other viruses, which did not significantly inhibit gene transfer of Ad5(GL) in most organs. However, triple immunization did produce a

stronger antibody response, and NAbs to capsid-modified adenoviruses started to reduce gene transfer of Ad5(GL). Liver transduction was decreased more than 1000-fold with Ad5(GL) compared with a single immunization with the same virus (Figures 2 and 3 in Study **IV**). Although a degree of cross-blocking was seen for all viruses, Ad5/3luc was affected the least, probably due to the complete knob-swap in the fiber, suggesting that anti-fiber knob NAbs might be most relevant.

Table 6. Induction of neutralizing antibodies and determination of their effect on gene transfer in immune-competent mice.

Group	I	II	Ш	IV	V	VI	VII	VIII	IX	X
DAY										
0 (s.c.)	Ad5(GL)	Ad5(GL)	Ad5/3luc1	Ad5/3luc1	Ad5lucRG D	Ad5lucRG D	Ad5pK7(GL)	Ad5pK7(GL)	mock	mock
3 (s.c.)		Ad5(GL)		Ad5/3luc1		Ad5lucRG D	,	Ad5pK7(GL)		mock
6 (s.c.)		Ad5(GL)		Ad5/3luc1		Ad5lucRG D		Ad5pK7(GL)		mock
27	Nab	NAb	NAb	NAb	NAb	NAb	NAb	NAb	NAb	NAb
28 (i.v.)	Ad5(GL)	Ad5(GL)	Ad5(GL)	Ad5(GL)	Ad5(GL)	Ad5(GL)	Ad5(GL)	Ad5(GL)	Ad5(GL)	Ad5(GL)
29	organs &	organs & serum	organs	organs	organs	organs	organs	organs	organs	organs

Abbreviations: i.v., intravenous injection (3 x 10^{10} vp); NAb, determination of neutralizing antibody titer from serum; s.c., subcutaneous immunization (3 x 10^8 vp); vp, viral particle

The effect on gene transfer in the presence NAbs was also evaluated *in vivo*. Mice bearing orthotopic tumors of advanced lung cancer were injected intravenously with Ad5(GL), Ad5/3luc1, Ad5lucRGD, or Ad5pK7(GL) or with the same viruses preincubated with serum collected from Ad5(GL)-immunized mice. A low titer of anti-Ad5 NAb did not block intravenous Ad5(GL), but a high titer almost completely prevented gene transfer. Regardless of anti-Ad5 NAb titer, effective gene delivery was achieved with Ad5lucRGD, Ad5/3luc1, and Ad5pK7(GL) (Figure 4 in Study **IV**).

Two human squamous cell carcinomas, one adenocarcinoma, and one non-malignant lung tissue specimen were obtained fresh from patients and infected with viruses preincubated with serum collected from Ad5(GL)-immunized mice. Ad5(GL) was completely blocked by a high NAb titer, whereas Ad5/3luc1 and Ad5pK7(GL) continued to allow gene transfer. Both Ad5/3luc1 and Ad5pK7(GL) showed enhanced gene delivery compared with the Ad5(GL) in the presence of low or high NAb. Intriguingly, anti-Ad5 NAb blocked gene transfer by capsid-modified viruses to normal human lung explants much more than to tumor explants (Figures 5 and 6 in Study IV). This might be explained by the fiber knob-independent mechanisms proposed to be important in systemic biodistribution of mice and primates (Bayo-Puxan *et al.* 2006; Smith *et al.* 2003a; Smith *et al.* 2003b). In conclusion, our data suggest that relative escape from the NAbs is achieved after systemic delivery using capsid-modified adenoviruses.

Summary and conclusions

The overall aim of this study was to obtain information on adenovirus tropism towards NSCLC cells and to enhance therapeutic outcome with oncolytic adenoviruses. To further improve the specificity, we sought to determine whether the Cox-2 promoter is induced by adenovirus infection and whether selectivity could be regained by 3'UTR elements. Finally, we tested whether switching the adenovirus fiber can retain gene delivery in the presence of NAbs.

HSPG- and $\alpha_v\beta$ integrin-targeted viruses displayed enhanced gene transfer to NSCLC cells *in vitro* and *in vivo*. Respective oncolytic viruses showed effective cell killing and this translated into a therapeutic benefit in a highly aggressive orthotopic model of large-cell NSCLC. Combination therapies with radio- or chemotherapy and arming oncolytic viruses with immunostimulatory or cytotoxic transgenes could lead to an even better therapeutic outcome.

Most importantly, the data laid the foundation for clinical testing of these viruses in patients with NSCLC refractory to current treatments. The use of cells as oncolytic virus delivery vehicles offers a number of potential advantages. One lies in the tumor-homing capacity of certain cells. A second advantage is that carrier cells can be engineered to express a set of transgenes exceeding the cloning capacity of the viral vector considered, thereby acting as local virus factories allowing high amplification of the inoculated virus. Finally, oncolytic viruses harbored by carrier cells remain at least partly protected from inactivation by the host until released at their destination.

MSCs possess multilineage properties making them useful for a number of potential therapeutic applications, and they have suggested targeting many different cancers. Infectivity of MSCs could be enhanced by using capsid-modified adenoviruses. In addition, MSCs were able to support the replication and viral release of oncolytic adenoviruses harboring a 24-bp deletion in the E1A region both *in vitro* and *in vivo*. Further, MSCs loaded with oncolytic adenoviruses prolonged the survival of mice with

advanced lung cancer and showed enhanced antitumor efficacy with breast cancer tumors after systemic delivery. These data suggest that MSCs could improve the bioavailability of adenoviruses in the context of advanced and aggressive disease.

Viruses are known to induce Cox-2 (not previously known for adenovirus), but the mechanisms behind this phenomenon are poorly understood. It is unclear whether Cox-2 induction occurs due to a protective antiviral response initiated by human cells, or whether viruses can also actively promote Cox-2 induction for their own purposes. Our results suggest that both Cox-2 and CMV promoters are induced by adenoviral infection, but selectivity could be rescued by the 3'UTR element of the Cox-2. It would be interesting to study the activity of these elements in the context of oncolytic adenoviruses, and further, to determine whether other tumor specific promoters could be controlled by these elements. Another approach to enhance the efficacy of adenoviral gene therapy is to reduce the effect of Nabs. Our data proposed that minor changes in adenovirus fiber can allow the virus to escape from pre-existing Nabs, and therefore, more effective systemic readministration could be achieved. It might be even more beneficial to use viruses from different serotypes, or viruses pseudotyped with the entire fiber (instead of just the knob) or hexon might yield further advances. It would also be interesting to study MSCs in the context of NAbs.

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