Pancreatic cancer is one of the most aggressive human malignancies and currently the fourth leading cause of cancer related death in the industrialised world. Despite recent advance in the management of the disease by conventional chemotheraphy, the development of new therapeutic strategies is critically important.

Molecular studies have identified mutation of the KRAS oncogene and overexpression of multiple receptor tyrosine kinases and their cognate ligands as key contributors to the aberrant upregulation of survival and proliferation pathways which give growth advantage to the malignant population.

A key effector of RAS and receptor tyrosine kinases is class I phosphatidylinositol-3-kinase (PI3K). PI3K, a heterodimer which consists of a p110 catalytic subunit and a p85 regulatory subunit, recruits various pleckstrin homology domain containing proteins to the plasma membrane. These include PDK1 and the Ser/Thr protein kinase AKT/PKB, an important downstream target of PI3K/AKT signal transduction at various levels of PI3K, and AAA-AKT, a dominant negative mutant of AKT was employed to interfere with PI3K/AKT signalling at the level of AKT.

Antiproliferative, proapoptotic, and anticancer effects were documented, showing that inhibition of the PI3K pathway in these cell lines suppresses tumour cell growth in vitro and reduces growth in nude mice.

Conclusions: The PI3K/AKT pathway represents a potential therapeutic target for pancreatic cancer, and gene therapy may be one approach to produce selective inhibition.
pcEXV-1 was a gift from Chris Marshall (Institute of Cancer Research, London, UK). The p85β cDNA was generated by polymerase chain reaction. Human PTEN cDNA was purchased from Invitrogen Ltd. It was amplified using the upstream primer 5’-cgg ctc gag gcc acc atg cgg ggt tct cat act cat cat cat cat gg-3’ engineered with a XhoI restriction site and the downstream primer 5’-ccg ctc gag gcc acc atg cgg ggt tct cat act cat cat cat cat gg-3’ engineered with a XhoI restriction site and the downstream primer 5’-ccg ctc gag gcc acc atg cgg ggt tct cat act cat cat cat cat gg-3’. cDNA encoding triple mutant kinase dead AKT (AAA-AKT in which Lys-179 (involved in ATP binding) and Thr-308 and Ser-473 (required for phosphorylation induced activation) have been mutated to Ala) was generated using the Quickchange kit (Stratagene Europe, Paisley, UK) according to the manufacturer’s instructions. For production of adenoviruses (Ad-N17 and Ad-p85β) without the GFP gene, cDNA of N17 and p85β were subcloned into pShuttle-CMV. Recombinant adenoviral constructs containing dominant negative transgenes under the control of the CMV promoter in the E1 region were produced using the Ad-Easy bacterial recombination system (www.coloncancer.org/adeasy.htm) as previously outlined. Adenoviral particles were produced by amplification on 293 cells and subsequent purification by double density CsCl gradient ultracentrifugation. Ad-PTEN was provided by Intron Therapeutics Inc (Houston, Texas, USA). Ad-CMV-null (Ad-null) and Ad-CMV-GFP were purchased from Qbiogene-Alexis Ltd (Bingham, UK). The titre of infectious particles in all preparations of adenoviruses was determined as plaque forming units (PFU) on 293 cells. The titre of infectious particles in preparations of GFP trackable adenoviruses was additionally determined as green fluorescent units. GFP expressing foci were scored by fluorescent microscopy five days post infection.

Adenoviral transduction in vitro
Pancreatic cancer cells were seeded on tissue culture plates (24 well, 6 well, 6 cm diameter or 10 cm diameter) or 8 well chamber slides with DMEM/10% FCS and infected at 37°C for four hours in serum free OptiMEM (Invitrogen Ltd) as soon as they attached. Subsequently OptiMEM was exchanged by DMEM/10% FCS. Examination of GFP expression 48 hours after infection of pancreatic cancer cells with GFP trackable adenoviruses revealed that a ratio of infectious viral particles to cell (multiplicity of infection (MOI)) of 30 in the case of PANC I and 100 in the case of HPAF was required to infect close to 100% of cells (data not shown). These MOIs were used in all in vitro experiments.

Cell viability and apoptosis analysis
Cell viability was assessed by trypan blue exclusion and by MTS (3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxy-methoxyphenyl)-2-(4-sulphophenyl)-2H)-assay performed with a kit (Cell Titer 96 Cell Proliferation Assay; Promega, Southampton, UK). For SubG1 analysis by flow cytometry, 3 x 10⁶ cells were infected. At indicated time points, cells were fixed in ice cold 70% ethanol for at least 30 minutes, washed twice with phosphate/citrate buffer (192 mM Na₂HPO₄, 40 μM citric acid), treated with RNase A, stained with propidium iodide, and analysed using a FACScalibur flow cytometer (Becton Dickinson, Cowley, Oxford, UK). For analysis of annexin V staining, 4 x 10⁶ cells were infected. At indicated time points post infection, cells were harvested and resuspended in HEPES buffer (10 mM HEPES, 140 mM NaCl, 2.5 mM CaCl₂). Annexin V-Alexa647 (Molecular Probes Europe, Leiden, the Netherlands) was added according to the manufacturer’s recommendation followed by incubation in the dark at room temperature for 30 minutes. Propidium iodide was added prior to analysis to allow gating of dead cells.

Protein analysis by western blot
Transfected cells, maintained in the continuous presence of DMEM/10% FCS or starved overnight in serum free medium and stimulated with epidermal growth factor (EGF 50 ng/ml) or insulin (5 μg/ml) for five minutes were lysed on ice in lysis buffer (50 mM HEPES, pH 7.5, 150 mM NaCl, 10% glycerol, 1 mM EDTA, 1% Triton X-100) containing 1 x complete protease inhibitors (Roche Diagnostics GmbH, Mannheim, Germany). To study protein phosphorylation the buffer was additionally supplemented with 1 mM Na₃VO₄, 10 mM NAF, and 10 mM β-glycero phosphate. Lysates were heat denatured at 100°C for 10 minutes before electrophoresis on 10% sodium dodecyl sulphate-polyacrylamide gel electrophoresis gels and transfer onto polyvinylidene difluoride membranes (Millipore UK Ltd, Winchester, UK). Membranes were blocked in 3% bovine serum albumin (BSA) in TBST buffer (10 mM Tris (pH 7.6), 150 mM NaCl, 0.1% Tween 20) and probed with primary antibody in 1.5% BSA in TBST buffer at recommended dilutions. Primary antibodies included pan-ras (Ab-2) (CN Biosciences UK Ltd, Nottingham, UK), phospho-AKT (Ser 473) antibody, AKT antibody, phospho-p44/42 MAP kinase antibody, p44/42 MAP kinase antibody, and phospho-GSK-3 α/β (Ser21/9) (all from New England Biolabs UK Ltd, Hitchin, UK), anti-human PTEN (Cascade Bio Science, Winchester, Massachusetts, USA), antihuman caspase 3 antibody (R&D Systems, Abingdon, UK), anti-GFP mouse monoclonal GFP 3E1 antibody (Research Monoclonal Antibody Service, CR-UK; dilution 1:600), anti-p38 kinase p85β subunit mouse monoclonal p85b4 antibody (Research Monoclonal Antibody Service, CR-UK; dilution 1:500), and actin-C antibody (Santa Cruz Biotechnology Inc, Calne, UK). After probing with an appropriate horseradish peroxidase labelled secondary antibody (Santa Cruz Biotechnology Inc) in 1.5% BSA in TBST for 30 minutes, the signal was detected by the enhanced chemiluminescence system (Amersham Pharmacia Biotech, Bucks, UK) and BioMax MR-1 radiographic film (Kodak, UK).

Immunohistochemistry
Cells were grown and transfected on eight well chamber sides, fixed in 4% phosphate buffered saline (PBS) formaldehyde, and permeabilised with 0.2% Triton X-100 in PBS for 15 minutes. Subsequently, cells were blocked in 1 in 20 diluted goat serum (Dako Ltd, Cambridge, UK) and incubated with anti-human caspase 3 antibody (R&D Systems) at 4°C overnight. Endogenous peroxidase activity was quenched with 0.03% hydrogen peroxide solution. Cells were then incubated with an horseradish peroxidase labelled secondary goat anti-rabbit antibody and finally with DAB+ peroxidase substrate-chromogen solution provided by the Dako EnVision+ System, Peroxidase (DAB) kit (Dako Ltd). Slides were counterstained with haematoxylin.

Analysis of anchorage independent growth
PANC I and HPAF cells were infected for 36 hours before 5 x 10⁶ cells were seeded with 3 ml of DMEM/0.3% agarose onto a base of 3 ml DMEM/0.6% agarose containing 10% FCS in 6 cm diameter dishes. The top agarose layer was kept moist by DMEM. After three weeks, colonies were visualised by adding MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; Sigma-Aldrich Poole, UK) at a concentration of 500 μg/ml. Colonies visible to the eye were scored.

In vivo analyses
In vivo experiments were performed using HPAF cells. Tumour formation by pancreatic cancer cells pretreated with

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therapeutic adenoviruses and the effect of therapeutic adenoviruses on growth of established tumours in vivo were assessed in adult BALB/c nu/nu mice (CR-UK BSU, Clare Hall, South Mimms, UK). Tumour growth was assessed over a four month period following subcutaneous inoculation of 5 × 10^6 cells in 0.1 ml PBS into the right dorsal region of mice. Therapeutic viruses were injected into established tumours when a volume of 15–65 mm³ was reached. A dose of 6 × 10⁸ PFU of the appropriate virus in 60 μl PBS was administered on three consecutive days. On each day of treatment a different direction of needle pass was used. Animals were killed on day 120 or before if they were judged to be terminally sick or tumours reached a critical size of maximally 900 mm³. Tumour volumes were calculated using the equation: tumour volume = longest diameter × shortest diameter² ÷ 6. All control and treatment groups consisted of 10 animals. Analysis of data on tumour growth after intratumoral injection of therapeutic viruses was performed on the median population of animals. The best performer and the worst performer were not included. All procedures were carried out under an appropriate UK Home Office Licence.

Statistics
The unpaired Student’s t test was used for statistical analysis of MTS, trypan blue exclusion, and immunohistochemistry experiments. Annexin V and FACS data were analysed using the χ² test. To analyse if variance exists between average subcutaneous tumour volumes in animals assigned to control and treatment groups before intratumoral injections of therapeutic adenoviruses commenced, the one way ANOVA test was applied. Survival data of in vivo experiments are presented as survival percentages using the Kaplan-Meier method. Analysis of survival data was performed using the log rank test. Results of statistical analyses are expressed as a probability value (p value). Statistical significance was defined as p<0.05. Calculated p values for Student’s t and χ² tests were two tailed.

RESULTS
Transgene expression and dominant negative inhibition of PI3K/AKT signalling in vitro
Western blot analysis of proteins extracted from pancreatic cancer cells after treatment with viruses at equal concentrations showed that RASN17, p85β, PTEN, and AAA-AKT transgenes were efficiently expressed on adenoviral delivery (fig 1A) and that control and therapeutic adenoviruses were similarly infective (fig 1B). The difference in molecular weights between the GFP protein expressed by Ad-GFP control and that expressed by Ad-N17-GFP, Ad-p85β-GFP, Ad-AAA-GFP, Ad-PTEN-GFP, Ad-N17-GFP, Ad-p85β-GFP, or Ad-AAA-GFP, or Ad-p85β at an multiplicity of infection of 30 by western blot analysis of proteins extracted 72 hours post infection was applied. Survival data of in vivo experiments are presented as survival percentages using the Kaplan-Meier method. Analysis of survival data was performed using the log rank test. Results of statistical analyses are expressed as a probability value (p value). Statistical significance was defined as p<0.05. Calculated p values for Student’s t and χ² tests were two tailed.

Figure 1 Transgenes delivered by adenoviral transfer were efficiently expressed in pancreatic cancer cells. (A) Expression of dominant negative inhibitors of the phosphatidylinositol 3-kinase (PI3K)/AKT pathway was assessed in PANC-I cells on infection with Ad-N17-GFP, Ad-AAA-GFP, Ad-PTEN-GFP, or Ad-p85β at an multiplicity of infection of 30 by western blot analysis of proteins extracted 72 hours post infection. (B) Similar infection efficiency of control and therapeutic viruses was determined by immunoblot analysis of GFP in PANC-I cells. Experimental details are as described in (A). β-actin was used as the loading control. Experiments were performed using 30 μg of protein samples. Representative data of two separate experiments are presented.

Figure 2 RASN17, p85β, PTEN, and AAA-AKT were specific inhibitors of intracellular signalling pathways in pancreatic cancer cells. (A) On the second day post infection with control and therapeutic adenoviruses, cells were starved overnight and were then stimulated with epidermal growth factor (EGF). Cells were lysed and 30 μg of total protein were subjected to western blot analysis. (B) Cells were mock infected (lanes 1 and 4) or treated with Ad-GFP control (lanes 2 and 5) or Ad-AAA-GFP (lanes 3 and 6). On the second day post infection, cells were starved and then stimulated with insulin the following morning. Total protein (15 μg) from lysates was subjected to western blot analysis. (C) Experimental details are as described in (A), but 20 μg of total protein from lysates were used. Blots were probed for β-actin as a loading control. Representative data of two separate experiments are presented.
Ad-PTEN-GFP, and Ad-AAA-GFP (fig 1B) was accounted for by the fact that the latter contained the gene of enhanced GFP.19

The dominant negative effect of RASN17, p85β, PTEN, and AAA-AKT has previously been demonstrated in other cell types.20–23 In order to confirm that expression of these dominant negative inhibitors disrupts activation of the PI3K/AKT pathway in pancreatic cancer cells, activation of the pathway, which involves phosphorylation of endogenous AKT on serine 473, was examined on stimulation of starved cells with EGF. As expected, overexpression of dominant negative transgenes RASN17, p85β, and PTEN in PANC I and HPAF cell lines almost completely abolished EGF induced phosphorylation of AKT but did not affect the level of AKT expression (fig 2A). PANC I cells were found to express higher levels of AKT protein than HPAF. This is consistent with previously reported 30-fold amplification of AKT 2 in Panc I cells.24 Unexpectedly, there was low level cross reactivity of the phospho specific antibody with the highly expressed non-phosphorylated dominant negative AAA-Akt (fig 2A). To demonstrate the inhibitory effect of AAA-AKT on activation of endogenous AKT in pancreatic cancer cells, the phosphorylation status of GSK-3-β, which is directly phosphorylated by AKT in response to insulin,24 was examined by western blot analysis. As shown in fig 2B, levels of phosphorylated GSK-3-β generated in response to insulin stimulation were markedly reduced in the presence of overexpressed AAA-AKT.

To investigate if the dominant negative inhibitors have specificity for their proposed cellular targets, their effect on signal transduction through the RAF/MEK/ERK pathway downstream of RAS was also examined. Western blot analysis showed that dominant negative inhibitors p85β, PTEN, and AAA-AKT failed to abolish activation of ERK1 and ERK2. In contrast, RASN17, which inhibits signal transduction upstream of the PI3K/AKT and the RAF/MEK/ERK pathways, inhibited activation of ERK1 and ERK2 (fig 2C). Thus the dominant negative inhibitors employed here function in a specific manner.

Adenovirus mediated transfer of dominant negative inhibitors of the PI3K/AKT pathway decreases the viability of pancreatic cancer cells in vitro

Infection of PANC I and HPAF cells with Ad-N17-GFP, Ad-p85β-GFP, Ad-PTEN-GFP, or Ad-AAA-GFP, but not with Ad-GFP control, induced profound changes in their morphology (fig 3 and data not shown). As these morphological alterations were not seen with Ad-CMV-GFP, which expresses enhanced GFP (data not shown), it was concluded that they were caused by dominant negative inhibition of the PI3K/AKT pathway. We examined the effect of dominant negative expression on the viability of PANC I and HPAF cells by MTS assay as described in (A). Data are presented as percentage OD 490 nm of mock infected cells at 24 hours. They are representative of two separate experiments where n = 3; *p=0.0001.

(B) Viability of PANC I was assessed 72 hours post treatment with therapeutic adenoviruses by MTS assay as described in (A). Data are presented as percentage OD 490 nm of mock infected cells at 24 hours and are representative of two separate experiment where n = 3; *p=0.0002, **p=0.0001, ***p=0.0003, ****p=0.0002.
pathway by p85β, PTEN, or AAA-AKT also significantly decreased the number of viable cells (fig 4B). The data were confirmed by FACS analysis to 120 hours after adenoviral treatment. A significant proportion (p<0.0001) of Ad-N17-GFP and Ad-AAA-GFP infected cells was shown to undergo cell death which was represented by the subdiploid fraction of the cell cycle profile. Approximately 12% of Panc I cells treated with Ad-N17-GFP or Ad-AAA-GFP were detected in subG1 peaks 120 hours post infection (fig 5). Similar results were obtained for HPAF cells (data not shown). From these results we concluded that the PI3K/AKT pathway plays an important role in promoting the growth of pancreatic cancer cells.

Overexpression of dominant negative inhibitors of the PI3K/AKT pathway induces apoptosis in pancreatic cancer cells

We observed a significantly higher subdiploid (subG1) signal in the cell cycle profile of PANC I and HPAF cells treated with dominant negative transgenes compared with mock and control Ad-GFP treated and Ad-PTEN-GFP infected cultures (Ad-N17-GFP and Ad-AAA-GFP). A significant difference was observed between the subG1 signal of control Ad-GFP treated and Ad-N17-GFP or Ad-AAA-GFP infected cultures (Ad-N17-GFP and Ad-AAA; p<0.0001).

Table 1 Dominant negative inhibition of the phosphatidylinositol 3-kinase (PI3K)/AKT pathway induced activation of caspase 3

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Percentage of cells† positive for cleaved caspase 3</th>
<th>Experiment I</th>
<th>Experiment II</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mock infection</td>
<td>5.0 (0.54)</td>
<td>5.7 (0.63)</td>
<td></td>
</tr>
<tr>
<td>Ad-GFP control</td>
<td>7.4 (1.67)</td>
<td>7.4 (0.86)</td>
<td></td>
</tr>
<tr>
<td>Ad-N17-GFP</td>
<td>41.7 (1.81)***</td>
<td>47.4 (1.52)***</td>
<td></td>
</tr>
<tr>
<td>Ad-AAA-GFP</td>
<td>42.7 (1.19)***</td>
<td>44.9 (1.63)***</td>
<td></td>
</tr>
<tr>
<td>Ad-PTEN-GFP</td>
<td>42.5 (1.02)***</td>
<td>40.6 (0.89)***</td>
<td></td>
</tr>
<tr>
<td>Ad-p85β-GFP</td>
<td>46.5 (1.05)***</td>
<td>40.3 (0.79)***</td>
<td></td>
</tr>
</tbody>
</table>

Results are mean (SEM) (n = 4).
†HPAF.
* Determined by immunohistochemistry 144 hours post treatment.
** p<0.0001.

Figure 5 Induction of death in Panc I cells by Ad-N17-GFP and Ad-AAA-GFP. DNA content of cultured cells was examined by propidium iodide staining and FACS analysis 48–120 hours post infection with adenoviruses at an multiplicity of infection of 30. Cell cycle profiles of mock infected Panc I and Ad-GFP, Ad-N17-GFP, and Ad-AAA-GFP infected Panc I cultures at 120 hours are presented. The subG1 signals at time points 48–120 hours are summarised at the bottom. A significant difference was observed between the subG1 signal of control Ad-GFP treated and Ad-N17-GFP or Ad-AAA-GFP infected cultures (Ad-N17-GFP and Ad-AAA; p<0.0001).

Figure 6 Expression of dominant negative inhibitors in pancreatic cancer cells induced activation of caspase 3. Expression of the active fragment of cleaved caspase 3 was assessed in PANC I cells on infection with therapeutic adenoviruses by western blot analysis of proteins extracted 120 hours post infection. Total protein (20 μg) from lysates was used. Blots were probed for β-actin as a loading control. Representative data of two separate experiments are presented.
Ad-GFP infected cells, which indicates that inhibition of the PI3K/AKT pathway induces cell death in pancreatic cancer cells (data not shown). To investigate if the dominant negative inhibitors induce apoptosis, pancreatic cancer cells treated with therapeutic adenoviruses were analysed for proteolytically processed caspase 3 by immunohistochemistry and western blotting. A significant difference in the number of apoptotic cells compared with Ad-GFP control infected cells was detected by immunohistochemistry (Ad-N17-GFP, Ad-p85β-GFP, Ad-PTEN-GFP, Ad-AAA-GFP; p<0.0001). Between 40% and 48% of treated cells stained positive when an antibody specific for cleaved caspase 3 was applied after infection with therapeutic adenoviruses (table 1). Activated caspase 3 was also detected in protein lysates from PANC I and HPAF cells treated with Ad-N17-GFP, Ad-p85β-GFP, Ad-PTEN-GFP, or Ad-AAA-GFP but not in cell lysates from mock infected and Ad-GFP treated cells (fig 6 and data not shown). Finally, analysis of externalised phosphatidylserine using the phosphatidylserine binding protein annexin V revealed induction of apoptosis by Ad-N17-GFP, Ad-p85β-GFP, Ad-PTEN-GFP, and Ad-AAA-GFP in PANC I and HPAF cells 96 hours and 120 hours post infection, which was significant (table 2). Collectively, these findings indicate that dominant negative inhibition of the PI3K/AKT pathway induces substantial cell death and suggest that pancreatic cancer cells die predominantly by apoptosis.

Inhibition of the PI3K/AKT pathway using RASN17, p85β, PTEN, and AAA-AKT abolishes anchorage independent growth of pancreatic cancer cells

To investigate if dominant negative inhibition of the PI3K/AKT pathway suppresses anchorage independent growth, we analysed the formation of colonies in soft agar. Colony formation was almost completely inhibited in both cell lines on expression of RASN17, p85β, PTEN, or AAA-AKT (table 3). Not only was the number of colonies substantially decreased so also was their size.

While infection with Ad-GFP control did not have a significant antitumour effect in short term viability and death assays with adherent pancreatic cancer cells, treatment of PANC I and HPAF cells with this virus reduced the number of colonies formed in soft agar by approximately 40% (table 3). Ad-GFP also had a beneficial therapeutic effect in an intraperitoneal in vivo mouse model with HPAF cells. In this experiment, Ad-null, which lacks the gene for GFP, had no effect in mice compared with mock treatment (data not shown). Thus GFP seems to have a non-specific antitumour effect, which is not observed in short term viability and death assays.

Table 2: Dominant negative inhibitors of the phosphatidylinositol 3-kinase (PI3K)/AKT pathway induced apoptosis in pancreatic cancer cells†

<table>
<thead>
<tr>
<th>Treatment</th>
<th>PANC I</th>
<th>HPAF</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mock infection</td>
<td>2.2</td>
<td>4.6</td>
</tr>
<tr>
<td>Ad-GFP</td>
<td>3.8</td>
<td>3.7</td>
</tr>
<tr>
<td>Ad-N17-GFP</td>
<td>45.6**</td>
<td>29.7***</td>
</tr>
<tr>
<td>Ad-AAA-GFP</td>
<td>41.2**</td>
<td>28.1***</td>
</tr>
<tr>
<td>Ad-PTEN-GFP</td>
<td>36.4***</td>
<td>27.5***</td>
</tr>
<tr>
<td>Ad-p85β-GFP</td>
<td>37***</td>
<td>23.1***</td>
</tr>
</tbody>
</table>

Representative data of two separate experiments.
†As analysed by annexin V/propidium iodide staining 120 hours post treatment.
‡Annexin V positive.
§10 000 cells analysed per sample.
**p<0.0001.

Table 3: Dominant negative inhibition of phosphatidylinositol 3-kinase (PI3K)/AKT signalling suppressed anchorage independent growth of pancreatic cancer cells†

<table>
<thead>
<tr>
<th>Treatment</th>
<th>PANC I</th>
<th>HPAF</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mock infection</td>
<td>100</td>
<td>100</td>
</tr>
<tr>
<td>Ad-Null control</td>
<td>111</td>
<td>86</td>
</tr>
<tr>
<td>Ad-GFP control</td>
<td>65</td>
<td>63</td>
</tr>
<tr>
<td>Ad-N17-GFP</td>
<td>0.11</td>
<td>0</td>
</tr>
<tr>
<td>Ad-AAA-GFP</td>
<td>0.23</td>
<td>0</td>
</tr>
<tr>
<td>Ad-PTEN-GFP</td>
<td>5.87</td>
<td>0.76</td>
</tr>
<tr>
<td>Ad-p85β-GFP</td>
<td>0.35</td>
<td>0.15</td>
</tr>
</tbody>
</table>

†As analysed by the formation of colonies in agarose medium.
‡Results are mean (n = 2); two representative experiments of three separate experiments are presented.

Dominant negative inhibition of the PI3K/AKT pathway shows therapeutic potential in vivo

We constructed adenoviruses which express dominant negative transgenes but lack an additional expression cassette for GFP. Efficient transgene expression in pancreatic cancer cells on infection with Ad-N17, Ad-p85β, or Ad-PTEN was verified by western blot analysis (data not shown).

The effect of dominant negative inhibition of the PI3K/AKT pathway on the ability of pancreatic cancer cells to establish tumours was studied in nu/nu mice subcutaneously injected with HPAF cells after pre-infection with Ad-N17, Ad-p85β, or Ad-PTEN. Animals injected with pretreated HPAF cells did not form tumours. In contrast, tumours developed and grew rapidly in control groups of animals injected with mock or Ad-null infected HPAF cells and all mice had to be killed by days 58 and 60, respectively. In contrast, animals injected with Ad-N17, Ad-p85β, or Ad-PTEN infected HPAF cells were tumour free and survived until the experiment was terminated (fig 7A, B).

We also assessed the effect of RASN17, p85β, and PTEN on growth of established tumours formed by HPAF cells in a subcutaneous nu/nu mouse model. Animals with subcutaneous tumours of HPAF cells were assigned to control and treatment groups with no significant difference between average tumour volumes at the start of adenoviral treatment (p = 0.98) (data not shown). The rates of tumour growth in Ad-N17, Ad-p85β, and Ad-PTEN treated groups were decreased compared with mock infected and Ad-null treated control groups (fig 7C). Animals injected with Ad-N17 or Ad-p85β survived significantly longer than Ad-null treated animals (Ad-N17, p = 0.0028; Ad-p85β, p = 0.0047) (fig 7D). However, no significant difference between survival curves of Ad-PTEN and Ad-null control treated mice was observed (p = 0.598) (fig 7D). Ad-N17 and Ad-p85β performed significantly better in prolonging the survival of animals than Ad-PTEN (p = 0.0076; p = 0.026) while no significant difference between Ad-N17 and Ad-p85β was found (p = 0.617). Collectively, these data show that RASN17, p85β, and PTEN are equally effective in suppressing the establishment of tumour deposits while dominant negative RASN17 and p85β but not PTEN are effective therapeutic genes for the treatment of established pancreatic tumours.

DISCUSSION

We found that the PI3K/AKT pathway plays an important role in maintaining the neoplastic phenotype of pancreatic cancer cells which harbour mutations in the KRAS oncogene,
express high levels of a number of growth factor receptors, and produce the cognate ligands of these receptors. Inhibition of the pathway suppressed tumour cell growth in vitro and in vivo and significantly improved the survival of nude mice bearing pancreatic tumour xenografts. These findings are in agreement with a recent study of inhibition of PI3 kinases with the chemical inhibitors wortmannin and LY294002 which induced apoptosis in a panel of pancreatic cancer cell lines, including the two cell lines employed in our investigations (PANC 1 and HPAF). We showed that inhibition of AKT using AAA-AKT was as effective as inhibition of PI3K by p85β or PTEN in inducing apoptosis. Furthermore, dominant negative AKT, p85β, and PTEN were equally potent in suppressing the growth of pancreatic cancer cells in soft agarose.

Treatment of pancreatic cancer cells with inhibitory transgenes of the PI3K/AKT pathway almost completely suppressed anchorage independent growth of pancreatic cancer cells in soft agarose and inhibited the formation of tumours deposits by these cells in nude mice. These data support the contention that the PI3K/AKT pathway plays an important role in protecting pancreatic cancer cells from apoptosis in the absence of attachment to a solid support and might explain their highly metastatic phenotype.

Both PI3K/AKT and RAF/MEK/ERK pathways are activated by RAS, which is downstream of growth factor receptor signalling. In addition, the activity of these pathways is regulated by cell-cell contact and integrin receptors which mediate interaction with the extracellular matrix. Thus

![Image of Figure 7]

**Figure 7** Dominant negative inhibition of phosphatidylinositol 3-kinase (PI3K) signalling suppressed pancreatic tumour growth in vivo. (A, B) HPAF cells were pre-infected with control and therapeutic viruses for 48 hours before being injected into right dorsal flanks of nu/nu mice. (A) Data (mean (SEM)) represent average tumour volumes (n = 10). They are presented until the first animal in each group was sacrificed. (B) Kaplan-Meier survival curves (Ad-N17, Ad-p85β, Ad-PTEN; p < 0.0001). (C, D) Control and therapeutic viruses were administered by intratumoral injection of subcutaneous tumours of nu/nu mice on three consecutive days. (C) Data (mean (SEM)) represent average tumour volumes (n = 8). Tumour volumes are shown until the first animal was sacrificed in each group. (D) Kaplan-Meier survival curves.
shown to be required for malignant transformation of pancreatic cancer cells.25-26 The dominant negative RAS mutant RASN17 is a potent inhibitor of wild-type RAS but less efficiently abolishes the function of constitutively active RAS mutants. It was previously shown that transformation of RIE-1 cells with KRAS caused upregulation of an EGF receptor dependent signalling loop which requires the function of wild-type endogenous RAS. This activity was required for transformation as treatment of cells with an EGF receptor specific inhibitor reversed their transformed phenotype.27 It has also been postulated that RASN17 may have inhibitory effects on the ERK MAP kinase pathway downstream of RAS.28

The finding that the PI3K/AKT pathway plays a pivotal role in the pathogenesis of pancreatic cancer encourages the preclinical development of modalities for efficient and safe delivery of therapeutic genes such as p85β to pancreatic tumours in patients. The potential toxicity of this transgene when administered in humans remains to be tested, but in view of the much lower intrinsic levels of death signals in normal cells compared with tumour cells and the greater selectivity of p85β compared with existing chemical inhibitors of PI3K signalling, we suggest that it might be relatively well tolerated. As survival signalling by PI3K might contribute to the inherent drug resistance of pancreatic cancers,29 gene therapy involving dominant negative PI3K inhibition may be particularly useful in combination with existing chemotherapeutic regimens in the treatment of pancreatic cancer.

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Authors’ affiliations

V Stoll, G Vassaux, N R Lemoine, Cancer Research UK Molecular Oncology Unit, Faculty of Medicine, Imperial College London, London, UK

V Calleja, Cancer Research UK Molecular Oncology Unit, Faculty of Medicine, Imperial College London, London, UK, and Signal Transduction Laboratory, Cancer Research UK London Research Institute, London, UK

J Downward, Signal Transduction Laboratory, Cancer Research UK London Research Institute, London UK

Conflict of interest: None declared.

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Dominant negative inhibitors of signalling through the phosphoinositol 3-kinase pathway for gene therapy of pancreatic cancer

V Stoll, V Calleja, G Vassaux, J Downward and N R Lemoine

doi: 10.1136/gut.2004.046706

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