

Placenta Growth Factor, PLGF, Influences the Motility of Lung Cancer Cells, the Role of Rho Associated Kinase, Rock1

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ABSTRACT

Placenta growth factor (PIGF) is a member of the VEGF family and has been implicated in the aggressive capacity of solid tumours, partly via its impact on angiogenesis. The present study determined the direct biological function of endogenous PIGF in lung cancer cells. From the human non-small cell lung cancer cell line A549 which expressed good level of PIGF, we created sublines within which PIGF expression was knockdown by way of anti-PIGF ribozyme transgenes. Remarkable reductions of both PIGF mRNA and protein by the ribozyme transgenes were revealed in A549 transfectants (A549^{ΔPIGF}) using RT-PCR and Western blotting respectively. A549^{ΔPIGF} cells exhibited significantly reduced migration and adhesion compared with the wild-type (A549^{WT}) and the empty plasmid control (A549^{ΔEF/His}) cells. Immunocytochemistry and Western blotting further revealed that the expression of ROCK1, Rho associated kinase, was also reduced in A549^{ΔPIGF} cells, in comparison with wild-type and control cells. In addition, A549^{ΔPIGF} cells lost its response to a ROCK inhibitor, which otherwise strongly inhibited the motility of A549^{WT} and A549^{ΔEF/His} cells. These data indicate that PIGF directly regulates the motility of human lung cancer cells and that this regulation is critically dependent on ROCK-1. The study further indicates that PIGF is a potential therapeutic target in lung cancer. *J. Cell. Biochem.* 105: 313–320, 2008. © 2008 Wiley-Liss, Inc.

KEY WORDS: PLACENTAL GROWTH FACTOR; MOTILITY; LUNG CANCER; ROCK

Lung cancer is one of the most prevalent cancers globally. It is the top cancer killer in both men and women [Parkin et al., 2001]. Over the last 30 years lung cancer has become one of the leading cause of death in China and indeed a number of other countries, a subject area currently under intense investigation scientifically and clinically [Parkin et al., 1999]. Most of the patients are already at an advanced stage of the disease at the time of diagnosis, an important contributing factor to the very poor prognosis for these patients [Ettinger and Kris, 2001]. Up to 40% of the patients will have detectable distant disease (metastasis) at presentation [Parker et al., 1997]. Metastasis, the leading cause of death for patients with solid tumours is the outcome of a complex biological phenomenon. Cancer cells would have already escaped from the normal growth control, departed from the primary site, degraded and invaded the basement matrix and extracellular matrix and travelled to the destination organ, before a metastatic tumour is

seen. Cellular motility and invasiveness of cancer cells as well as angiogenesis are amongst critical factors in the metastatic process.

Placental growth factor (PIGF), one member of the VEGF family, was first cloned from placenta [Maglione et al., 1991] and has been shown to cooperate with VEGF in the formation of vasculature [Khaliq et al., 1996; Ahmed et al., 2000]. PIGF may activate the VEGF receptor, Flt1, resulting the transphosphorylation of Flk1 and amplifying VEGF derived angiogenesis, in which VEGF/PIGF heterodimer can activate VEGF receptors, forming Flk1/Flt1 heterodimers and angiogenesis process [Autiero et al., 2003]. PIGF has been implicated in the progression of a number of solid tumours. This involvement was initially believed to be the results of induction of angiogenesis by PIGF. However, in recent years, evidence began to indicate that PIGF may directly act on cancer cells. PIGF protein level is significantly higher in many tumours compared with non-tumour foci, such as breast cancer, lung cancer, colorectal cancer

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and gastric cancer and is correlated with patient's survival [Chen et al., 2004; Parr et al., 2005; Wei et al., 2005; Zhang et al., 2005]. In patients with renal cancer, the plasma PIGF levels were significantly higher than non-cancer people, and plasma PIGF was correlate with patients' survival [Matsumoto et al., 2003]. While the correlation between PIGF and disease progression was thought to be the result of PIGF induced angiogenesis in cancer, recent in vitro and in vivo studies have shown that exogenous PIGF have a direct stimulatory effects on cancer cells, such as the motility and invasion of breast cancer cell [Taylor and Goldenberg, 2007]. The in vitro study has shown over-expression of PIGF-2 in cancer cell lines, such as lung cancer cell A549, colon cancer cell HCT116 and glioblastoma U87-MG. Overexpression of PIGF in in vivo models can inhibit tumour growth by depleting VEGF monodimers [Xu et al., 2006]. The later has provided tentative evidence that PIGF may have a dual action in cancer, direct effect on endothelial cells (induction of angiogenesis) and direct effect on cancer cells (inducing motility and invasion). However, the spectrum that PIGF has a direct impact on cancer and the mechanism(s) by which PIGF acts on cancer is yet to be investigated. We have recently reported that, in clinical lung cancer, PIGF was positively stained in lung cancer cells using immunohistochemistry. We further demonstrated that PIGF mRNA was significantly higher in advanced stage lung tumours and was correlated with poor prognosis in the patients [Zhang et al., 2005]. In the present study, we went on to investigate if PIGF had a direct effect on human lung cancer cells and the possible mechanism by which PIGF may influence the biological behaviours of lung cancer cells.

MATERIALS AND METHODS

CELL LINES AND MATERIALS

Human lung cancer cell line A549 (American Type Culture Collection, Manassas, VA), were maintained in Dulbecco's Modified Eagle's medium (DMEM) supplemented with 10% fetal calf serum (FCS) and antibiotics. Goat anti-human PIGF, mouse anti-human ROCK1, mouse anti-human GAPDH antibodies and the ROCK inhibitor (Y-27632) were obtained from Santa Cruz Biotechnology, Inc., Santa Cruz, CA. Matrigel (reconstituted basement membrane) was purchased from Collaborative Research Product (Bedford, MA). Other kits and reagents were obtained from Sigma-Aldrich, Poole, UK, unless otherwise stated.

CONSTRUCTION OF RIBOZYME TRANSGENE TARGETING HUMAN PLGF

Anti-human PIGF hammerhead ribozymes were designed based on the secondary structure of the gene generated using the Zuker RNA mFold program [Zuker, 2003]. Hammerhead ribozymes that specifically target a GTC and a UUC site of human PIGF (GenBank NM_173074) were generated using touch-down PCR (primers as listed in Table I). The ribozymes were then cloned into the pEF6/v5-His vector (Invitrogen), using ampicillin and blasticidin as selection markers for prokaryotic and mammalian cells, respectively. After verification, purified ribozyme transgenes and empty plasmids were transfected into A549 cells individually using an Easyject Plus

TABLE I. PCR Primer Sequences

PIGF	Forward	CCCTGGGGTCTCCTCCTTC
	Reverse	TGCGGCGATGAGAATCTGC
Ribozyme for hPIGF	Forward	CTGCAGCAAGGGAACAGCCTCAT CTGATGAGTCCGTGAGGA
	Reverse	ACTAGTCTGAGAAGATGCCGGTT TCGTCTCACGGACT
β -actin	Forward	ATGATATCGCCCGCTCG
	Reverse	CGCTCGGTGAGGATCTCA

electroporator (EquiBio, Kent, UK). After up to 2 weeks of selection with Blasticidin (5 μ g/ml), stable transfectants were verified for the success in knocking down the PIGF transcript. Successful strains were grown to sufficient number and used in current study. The above procedure generated the following transfected cells from A549: empty vector control: A549^{pEF/His}, PIGF knockdown cells, A549 ^{Δ PIGF} and the wild-type: A549^{WT}.

EXTRACTION RNA AND RT-PCR

RNA was obtained using Total RNA Isolation Reagent (ABgeneTM). 0.25 μ g RNA was used to produce the first strand cDNA using a reverse transcription kit (Sigma, Poole, Dorset, UK). PCR was undertaken using a REDTaqTM ReadyMix PCR reaction mix (Sigma-Aldrich, Inc.). Cycling conditions were 94°C for 5 min, followed by 40 cycles of 94°C for 30 s, 55°C for 30 s and 72°C for 40 s. This was followed by a final extension of 10 min at 72°C. The quality of cDNA was verified by 580 bp β -actin PCR with the anneal temperature of 55°C. The annealing temperature of PIGF was 58°C. Products were separated by 2% agarose gel and visualised under UV light after staining with ethidium bromide.

IMMUNOPRECIPATAION, SDS-PAGE AND WESTERN BLOTTING

Cells were pelleted and extracted in HCMF buffer (160 mM NaCl, 0.6 mM Na₂HPO₄, 0.1% w/v glucose and 0.01 M Hepes, pH 7.4) containing 0.5% Triton X-100, 2 mM CaCl₂, 100 μ g/ml phenylmethylsulfonyl fluoride, 1 μ g/ml leupeptin and 1 μ g/ml aprotinin for 30 min. Protein concentrations were measured using fluorescamine and quantified by using a multifluorescanner (Denly, Sussex, UK). Equal amounts of protein from each cell sample were separated on a 12% (for PIGF) or 8% (for ROCK1) polyacrylamide gel. For immunoprecipitation experiments, both cell lysate and condition media were collected. Equal amount of cell lysate and condition medium were incubated with anti-PIGF IgG (4 μ g/ml) to precipitate PIGF. The immune complexes subsequently formed were then precipitated by the incubation with protein A/G-agarose conjugate for 1 h and subsequent centrifugation at 13,000g for 15 min. After washing the immunoprecipitates with the lysis buffer three times, the samples were boiled in SDS-PAGE sample buffer for 5 min prior to separation using SDS-PAGE. The secondary peroxidase-conjugated secondary antibodies were used at a final dilution 1:1,000. Protein band signal was visualized with the SupersignalTM West Dura system (Pierce Biotechnology, Inc., Rockford, IL), and images obtained using an UVITech imager (UVITech, Inc., Cambridge, England, UK).

IN VITRO CELL GROWTH ASSAY

This was based on a previously reported method [Jiang et al., 2005]. Cells were plated into 96-well plates at 2,000 cells/well followed by a period of incubation. Respective plates were fixed in 10% formaldehyde on the day of plating and daily for the subsequent 5 days. 0.5% crystal violet (w/v) was used to stain cells. Following washing, the stained crystal violet was dissolved with 10% (v/v) acetic acid and the absorbance was determined at a wavelength of 540 nm using an ELx800 spectrophotometer. Absorbance represents the cell number.

CELL-MATRIX ADHESION ASSAY

This was based on a method we previously reported [Jiang et al., 1995a,b]. Ninety six-well plates were coated with 5 μ g of matrigel and allowed to dry. Following rehydration, 40,000 cells were added to each well. After 40 min of incubation non-adherent cells were washed off using BSS buffer. The remaining cells were fixed with 4% formalin and stained with 0.5% crystal violet. The number of adherent cells was then counted under microscopy.

IN VITRO MOTILITY ASSAY USING CYTODEX-2 BEADS (PHARMACIA, PISCATAWAY, NJ)

We followed the protocol described by Rosen et al. [1990]. Cells (1×10^6) were incubated with 100 μ l Cytodex-2 beads in 10 ml DMEM medium overnight to coat the cytodex-2 beads. After the medium was aspirated and the beads were washed with DMEM, they were aliquoted into wells of 24-well plate (100 μ l/well). After 4 h of incubation, the beads were washed off. The cells that migrated onto the bottom of each well were fixed with 4% formalin for 5 min and were stained with 0.5% crystal violet. The cells were counted under a microscope. The experiment was performed three times.

ELECTRIC CELL-SUBSTRATE IMPEDANCE SENSING (ECIS) BASED MOTILITY ASSAY

The ECIS 1600R model instrument and 8W10 arrays (Applied Biophysics, Inc., NJ) were used in the study [Giaever and Keese, 1991; Keese et al., 2004]. Following treating the array surface with a Cysteine solution, the arrays were incubated with complete medium for 1 h. The same number of lung cancer cells, A549^{pEF/His}, A549 ^{Δ PIGF}, or A549^{wt} (300,000 per well) were added to each well. After reaching confluence, medium or ROCK inhibitor (10 μ M) was added to the respective well. The cells were then immediately subject to wounding using the integrated elevated field module in the instrument in the 1600R model (5v, 30 s for each well). The changes of cellular impedance were immediately recorded after wounding (4,000 Hz). The data was analysed using the ECIS RbA modelling software, supplied by the manufacturer.

IMMUNOCYTOCHEMICAL STAINING FOR ROCK1 PROTEIN

Cells were fixed with 4% formaldehyde and then permeabilized with 0.1% Triton X-100 for 5 min in Tris buffer saline. After blocking with horse serum (10%) for 60 min, the cells were probed with anti-

ROCK1 antibody for 1 h, followed by extensive washing. Horseradish peroxidase-conjugated anti-mouse antibody was then added for 1 h and visualized using the Vectastain[®] ABC system. Slides were mounted with Sterilyte.

STATISTICAL ANALYSIS

Minitab[®], version 14 was used for analysis. Normally and Non-normally distributed data were assessed using the two sample *t*-test and Mann-Whitney test respectively.

RESULTS

KNOCKDOWN OF PLGF EXPRESSION BY RIBOZYME TRANSGENES

Following selection, we obtained strains of transfected cells. The expression of PIGF was determined in these transfected cells. As shown by RT-PCR analysis, PIGF mRNA expression was almost completely eliminated from A549 cells by anti-PIGF ribozyme transgene (A549 ^{Δ PIGF}), a clear contrast to the level of expression in wild-type (A549^{WT}) cells and in A549 empty plasmid (A549^{pEF/His}) cells (Fig. 1). Knock-down of PIGF mRNA was also reflected by a reduction in its protein level: a decrease in the PIGF protein level was seen in both condition medium and cell lysate from A549 ribozyme transgene (A549 ^{Δ PIGF}) cells (Fig. 2).

LOSS OF ENDOGENOUS PLGF RESULTED IN A REDUCTION OF MOTILITY AND CELL-MATRIX ADHESION IN A549 CELLS

We first tested the effect of loss of PIGF (A549 ^{Δ PIGF}) on cell-matrix adhesion of the lung cancer cells. A549 ^{Δ PIGF} cells exhibited a significant reduction of cell-matrix adhesion: the number of adherent cells for the A549 ^{Δ PIGF} cells being 13.3 ± 3.95 , $P < 0.05$ compared with both A549^{WT} (25.6 ± 7.90) and A549^{pEF/His} (28.1 ± 6.90) (Fig. 3B).

In the cell carrier based cell motility assay, we found that the motility reduced significantly in A549 cell with the loss of PIGF. The number of migrating A549 ^{Δ PIGF} cells was 23 ± 8 compared with 57.5 ± 13.70 for A549^{WT} cells and 52.12 ± 15.0 for A549 ^{Δ control} cells ($P < 0.05$) (Fig. 3A).

To further investigate the effect of knock-down PIGF on cell motility, ECIS system was utilised to determine the migration of

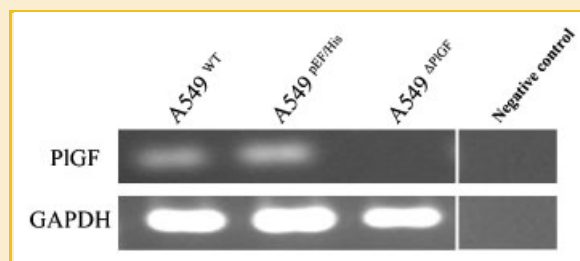


Fig. 1. The effects of ribozyme transgenes on PIGF expression using RT-PCR. mRNA of PIGF was eliminated in A549 ^{Δ PIGF} cells compared with the wild-type (A549^{WT}) and empty plasmid control (A549^{pEF/His}) cells.

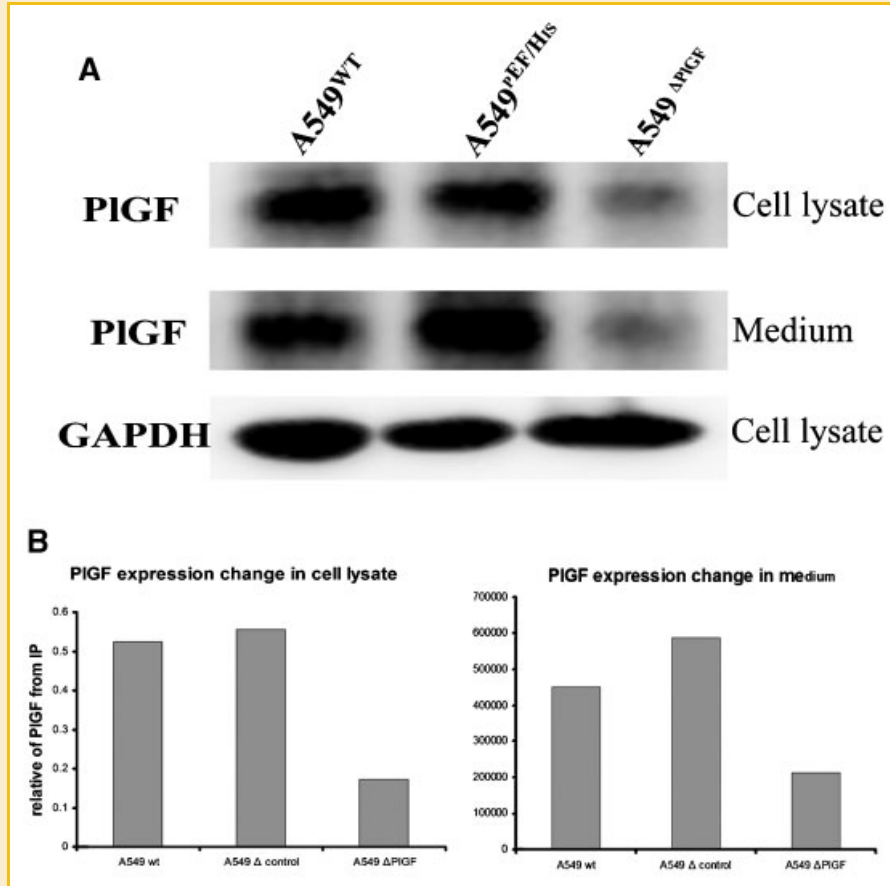


Fig. 2. Knockdown of PIGF by the ribozyme transgene at protein level using immunoprecipitation and western blot analysis. A: PIGF proteins in cell lysates and conditioned medium as shown by immunoprecipitation and Western blotting. Both the cytosolic and secreted PIGF were reduced by the ribozyme transgene. GAPDH was also determined from the original cell lysate and used as an internal reference. B: The bands of PIGF were quantified and the volume of bands were normalised against corresponding GAPDH bands.

A549 cells. The migrating capacity was remarkably reduced in A549^{ΔPIGF} cells after loss of PIGF, compared with A549^{WT} and A549^{P^{EF}/His} cells (Fig. 5B).

The knockdown of PIGF expression had no effect on the growth of lung cancer cells as shown in Figure 3C.

INFLUENCE ON ROCK1 EXPRESSION IN A549 CELLS BY KNOCK-DOWN OF PLGF

Rho kinase (ROCK) belongs to a family of serine/threonine kinases, and is the Rho GTPase effector. ROCK is involved in regulation of cell adhesion and migration. It has been demonstrated that VEGF/PIGF can induce cell migration resulting Rho GTPase activation in leukaemia [Casalou et al., 2007]. In order to examine the relationship between endogenous PIGF and ROCK family, the expression of ROCK1 was examined in A549 cells with and without the transgenes. Surprisingly, we observed a decrease in the level of ROCK1 protein in A549^{ΔPIGF} cells. This was reproduced in both Western blot (Fig. 4A,B) and immunocytochemistry (Fig. 4C) analyses.

KNOCK-DOWN OF PLGF IN LUNG CANCER CELLS RENDERED CELLS LESS RESPONSIVE TO ROCK INHIBITOR

In order to test the role of ROCK1 in PIGF induced motility change in A549 cell, we used a small molecule ROCK inhibitor, Y27632. Both cytodex-2 beads motility assay and ECIS assay were employed. The motility was inhibited by exposure to ROCK inhibitor in both A549^{WT} and A549^{P^{EF}/His} cells, the number of migrating cells being 43.75 ± 12.08 without ROCK inhibitor and 28.33 ± 4.830 with the inhibitor for A549^{WT} cells, and 41.92 ± 10.040 and 27.5 ± 5.37 respectively for A549^{P^{EF}/His} cells (Fig. 5A). Interestingly, A549^{ΔPIGF} lost its response to the ROCK inhibitor in that the motility indices in both ROCK inhibitor treated and non-treated were virtually the same: 17.42 ± 4.89 versus 17.75 ± 6.34 .

In ECIS based analysis, ROCK inhibitor was able to inhibit the migration of A549^{WT} and A549^{P^{EF}/His} cells in comparison with the respective controls, but to less extent compared to the A549^{ΔPIGF} cells (Fig. 5B). Thus similar to the observations seen in the carrier bead motility assay, A549^{ΔPIGF} cells failed to respond to the ROCK inhibitor.

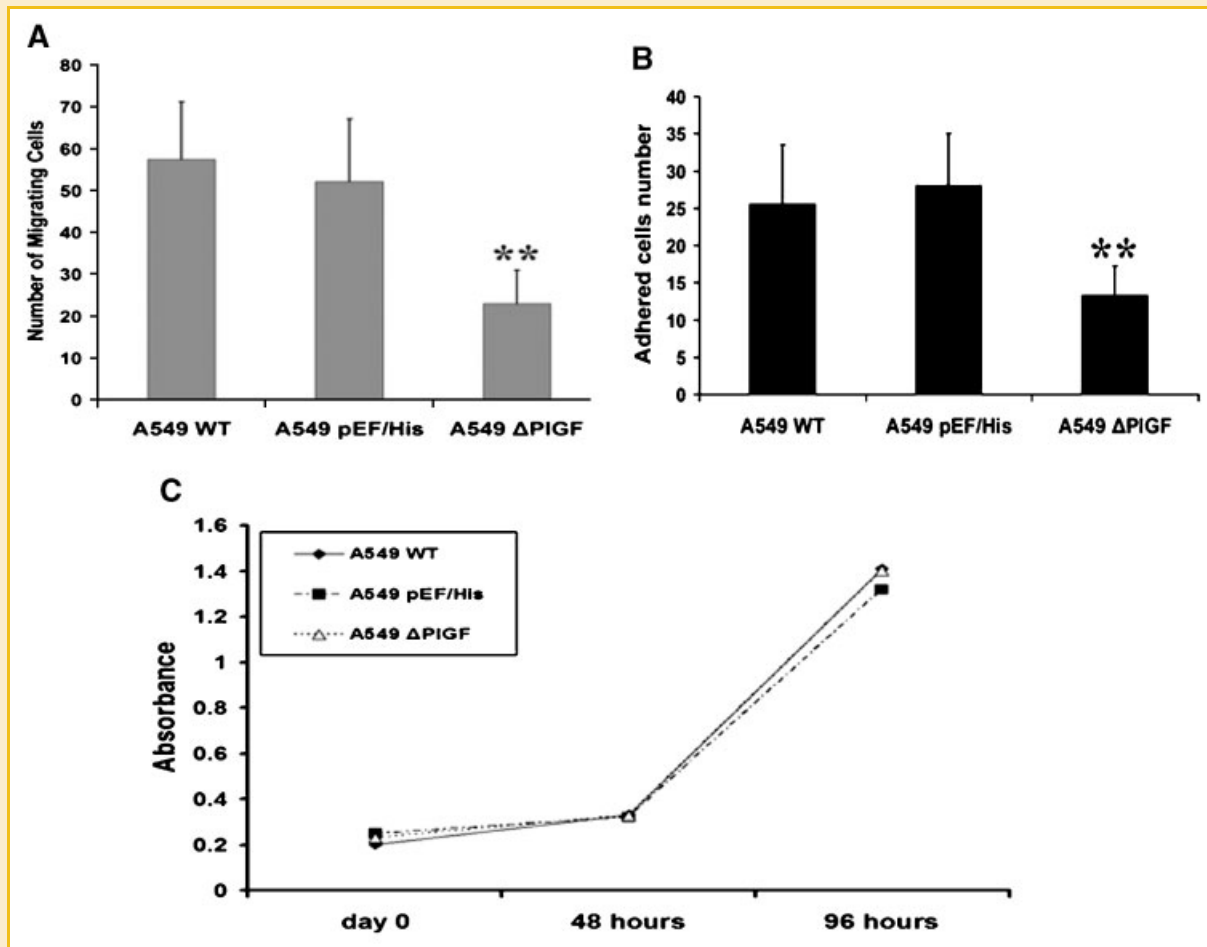


Fig. 3. The effect of ribozyme transgene targeting PIGF on the biological functions of A549 cells. A: motility of 549^{ΔPIGF} cells was reduced remarkably compared with A549 wild-type cells. Assays were performed in triplicate in three preparations each. Asterisk indicates $P < 0.001$ versus A549^{WT} and A549^{pEF/His} cells. B: A decrease of cell–matrix adhesion was seen in 549^{ΔPIGF} cells. Cell matrix adhesion assay was done in quadruplicate per cell line per experiments, $n = 3$. Asterisk indicates $P < 0.001$ versus A549^{WT} and A549^{pEF/His} cells. C: There was no significant change in the rate of cell growth of 549^{ΔPIGF} cells compared with A549^{WT} and A549^{pEF/His} cells. Growth assay was done in six wells per cell line. Error bars represent SD.

DISCUSSION

The present study provides the first line evidence that reduction of PIGF directly affects the motility of lung cancer cell, thus indicating that PIGF has a biological effect on lung cancer cells and that this effect is likely to be via a ROCK dependent pathway. The study has provided further evidence to indicate a therapeutic implication of PIGF in lung cancer.

Levels of PIGF have a demonstrable correlation with tumour progression and poor prognosis in human solid tumours [Wei et al., 2005; Zhang et al., 2005], a connection thought to be linked to PIGF induced angiogenesis [Dull et al., 2001; Castellon et al., 2002; Khurana et al., 2005]. The function of PIGF in tumour cells is poorly understood, although a direct effect on cancer cells has been suggested in recent years. The current study assessed the changes of cellular behaviour with the loss of PIGF expression in A549 cell, a non small cell line that endogenously expresses PIGF. The decrease in endogenous PIGF in A549 cell by way of ribozyme transgenes

resulted in reduction of cellular motility and adhesion. This indicates that endogenous PIGF plays an important role in controlling the functions that are critical to the aggressiveness of the lung cancer cell. In the last few years, there has been limited number of reports which demonstrate that PIGF acts on non-endothelial cells.

Hollborn et al. [2006] demonstrated that high concentration of exogenous PIGF (at 100 ng/ml) decreased the cell proliferation of retinal pigment epithelial cells. When pre-mixed with soluble HSPGs (Heparin Sulphate Proteoglycans), PIGF-1 can induce concentration-dependent proliferation stimulation of extravillous trophoblasts [Athanasziades and Lala, 1998]. To date, the effect on the growth of cancer cells and indeed tumours in vivo, remains controversial. Xu et al. [2006] showed that over-expression of PIGF can decrease the amount of homodimeric VEGF and thus contribute to the inhibition of tumour growth in vivo. In contrast, Marcellini et al. [2006] reported that melanoma cells over-expressing PIGF grew at a higher rate in transgenic mice compared with the wide type

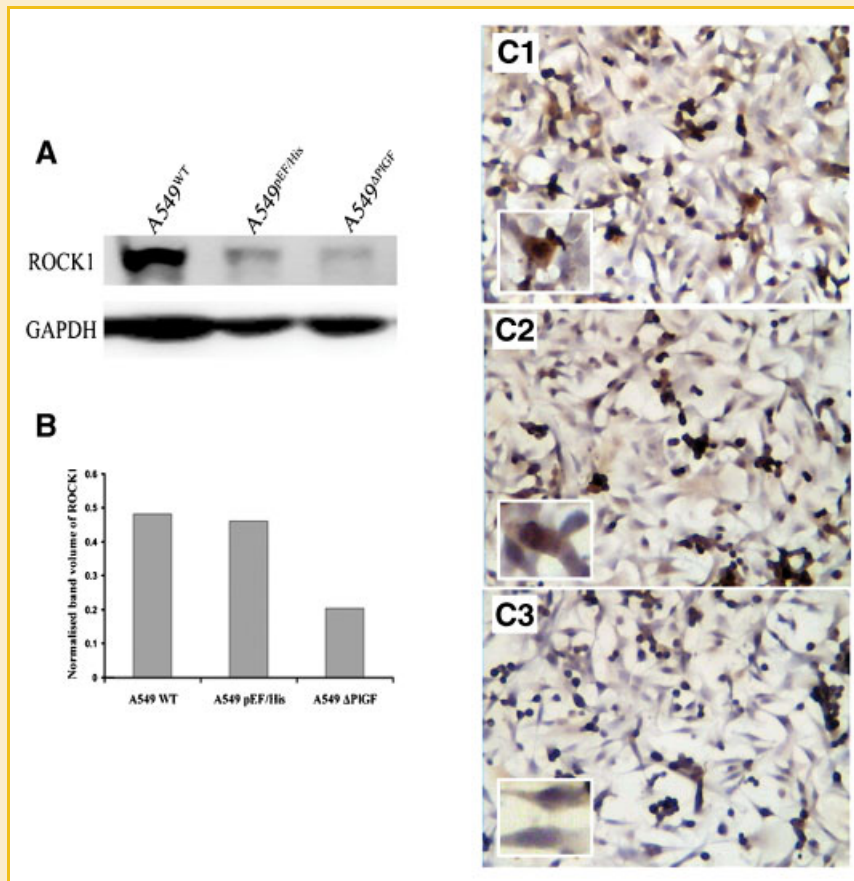


Fig. 4. Knock-down of PIGF in A549 cells resulted in reduction of ROCK1 expression. A: The expression of ROCK1 protein was decreased after loss of PIGF in A549^{ΔPIGF} cells, as shown by Western blot analysis. B: Quantified changes of ROCK1 protein bands. The bands of PIGF were quantified and the volume of bands was normalised against the corresponding GAPDH bands. C: Immunocytochemical staining also demonstrated that ROCK 1 protein level was reduced by knock-down of PIGF. C1: Immunostaining of ROCK1 in A549^{WT}. C2: ROCK1 staining in A549^{pEF/His} control cells. C3: ROCK1 staining in 549^{ΔPIGF} cells. C1/C2/C3: magnifications were 100×; Insets 400×. [Color figure can be viewed in the online issue, which is available at www.interscience.wiley.com.]

melanoma cells. The present study has clearly added weight to the reports that PIGF does indeed have a direct effect on non-small lung cancer cells in vitro. This, together with previous studies has clearly shown that PIGF acts via both autocrine and paracrine routes in cancer: directly on cancer cells themselves or on the tumour related endothelial cells and, in doing so, PIGF acts as a powerful protein factor in driving the spread of lung cancer.

Rho-associated kinase (ROCK) is a serine/threonine kinase that mediates some of the downstream signalling of RhoA [Riento and Ridley, 2003]. It has two isoforms, namely ROCK1 and ROCK2. The serine/threonine kinase activities of ROCKs have an important role in cell migration, cell death and survival [Shi and Wei, 2007]. ROCKs are known to regulate the metastatic process in cancer such as pancreatic cancer and small cell lung cancer [Kaneko et al., 2002; Li et al., 2006]. Our study clearly demonstrated that there was a decreased ROCK1 expression in protein level with the loss of PIGF expression in A549 cell. ROCK inhibitor could reduce the motility of A549^{WT} and A549^{pEF/His} cells. The effect of the ROCK inhibitor was significantly weaker with the A549^{ΔPIGF} cells which have lost their endogenous PIGF as the result of ribozyme transgene. This has

provided strong evidence that ROCK1 at least partly contributes to the increase of motility in lung cancer with highly expressed PIGF. It is further extrapolated from this result that ROCK1 has a potential therapeutic value in lung cancer.

In summary, the present study shows that ribozyme designed to target human PIGF is an effective approach to inhibit the action of PIGF in lung cancer cells. It can specifically reduce the expression of PIGF at both mRNA and protein level in the cancer cells. The lung cancer cells carrying the ribozyme transgene exhibit low level of migration and adhesion ability, providing the evidence that PIGF is an important molecule in the control of dissemination of lung cancer cells by directly promoting the aggressiveness of cancer cells. These results indicate that PIGF is an important biological regulator of lung cancer cells and may be a valuable target in the therapeutic development in lung cancer.

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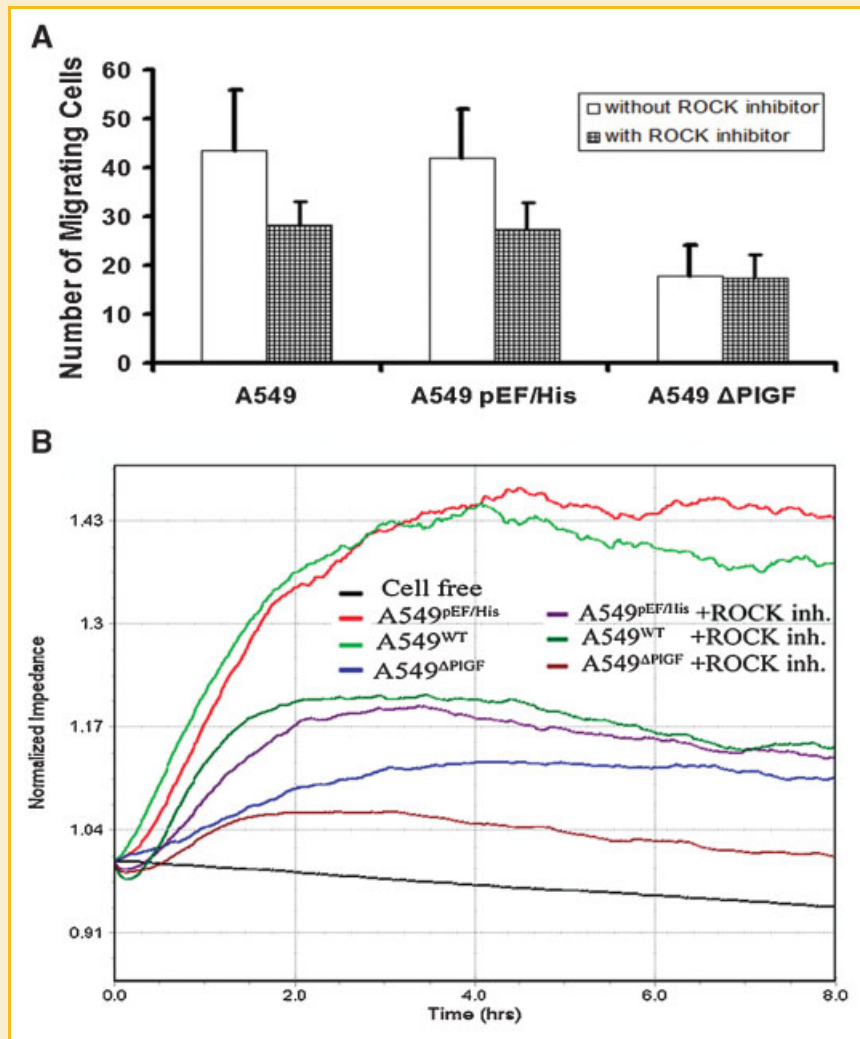


Fig. 5. The effects of ROCK inhibitor and the ribozyme transgene targeting on cell motility. A: Cytodex carrier assay. ROCK inhibitor (10 μ M) significantly reduced motility of A549^{WT} and A549^{pEF/His} cells, but only had a marginal effect on A549^{ΔPIGF} cells. B: ECIS-based wounding motility assay. A549^{WT} and A549^{pEF/His} cells had significantly reduced migration in response to the ROCK inhibitor. The A549^{ΔPIGF} cells failed to show a significant change. [Color figure can be viewed in the online issue, which is available at www.interscience.wiley.com.]

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