

# Expression of pigment epithelial derived factor is reduced in non-small cell lung cancer and is linked to clinical outcome

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**Abstract.** Angiogenesis is under the exquisite control of a network of angiogenic factors and anti-angiogenic factors. PEDF (pigment epithelial derived factor) is one of the known anti-angiogenesis factors and is naturally occurring in the body. There has been studies to show that the factor plays an important role in negating the angiogenic process in pathological conditions in the eye. However, little is known about its expression in solid tumors. The current study examined PEDF expression at protein and message levels and investigated its critical link with cancer progression and prognosis in patients with non-small cell lung cancer (NSCLC). We used immunohistochemistry to examine the protein expression of PEDF and to evaluate the micro-vessel density (MVD) in a cohort of 91 NSCLC patients. In addition, real-time quantitative PCR was used to measure levels of the PEDF transcript. PEDF was positively stained in cytoplasm of cancer cells, but at a lower level, compared with normal cells in the lung tissues. Low levels of PEDF were seen in 57.1% patients. The levels of PEDF appeared to be associated with MVD, in that patients with reduced PEDF had a significantly high MVD count (28.50), compared with patients with high levels of PEDF who had a 16.98 MVD count ( $P<0.0005$ ). In univariate but not multivariate analysis PEDF was an independent prognostic factor. In real-time PCR analysis, PEDF mRNA level of cancer tissue was significantly lower than normal tissue ( $0.55\pm0.36$  vs  $0.72\pm0.26$ ,  $P=0.024$ , paired t-test). PEDF mRNA level in cancer tissue was negatively associated with TNM stage and the tumor size ( $P<0.05$ , independent t-test). Finally, low levels of PEDF in lung tumor tissues was associated with a significantly shorter survival ( $P=0.038$ ) using Kaplan-Meier and Cox analyses. In this first study, PEDF was reduced at both protein and mRNA level in NSCLC tumors compared with normal lung tissues. This reduction is associated with an increase in micro-

vessel density in tumors and significantly associated with TNM stage, tumor size and the overall survival. PEDF is an important factor in NSCLC development and may be a of prognostic value for NSCLC patients.

## Introduction

PEDF (pigment epithelial derived factor) is a protein with MW 50 kDa, encoded by the *EPC-I* gene (1). It is a glycoprotein composed of 418-aa, of which four (glutamate-101, isoleucine-103, leucine-112, and serine-115) in the 44-aa region, has been found critical for its anti-vasopermeability activity (2), i.e. ability to counteract VEGF-induced vascular permeability. It is expressed in normal tissue throughout the body (3,4). It was first discovered in 1989 as a neurotrophic serpin (serine protease inhibitor) that was secreted by retinal pigment epithelial cells (5,6). As a secreted glycoprotein, PEDF is expressed in adult liver, testis, ovaries, placenta, brain and pancreas (7). PEDF has been previously shown to have neurotrophic activity. In sequence and structure, PEDF resembles members of the serine protease inhibitor family, but lacks protease inhibitor activity. PEDF therefore has been seen as a member of the serine protease inhibitor (serpin) family (8).

PEDF has been shown to be a potential and novel anti-angiogenic factor in a limited number of studies. It can inhibit endothelial cell migration and proliferation and reduce choroidal and retinal neovascularization (5,9-11). Thus far, PEDF has been shown to have strong anti-angiogenesis effect on at least a comparable basis with factors, such as angiostatin, thrombospondin-1 and endostatin (5,12). PEDF has been shown to regulate blood vessel growth in the eye by creating a permissive environment for angiogenesis when oxygen supply is limited, and creating an inhibitory environment when oxygen concentrations are normal or high. In human solid tumors, there has been very little study on the expression and clinical significance of PEDF. Using immunohistochemistry to investigate the expression of PEDF in ductal pancreatic adenocarcinoma patients, a correlation between PEDF and MVD, and clinical pathological factors and prognosis was reported (13). Although there has been no studies on the expression of PEDF in lung cancer, we have recently reported that PEDF is actively involved in the regulation of lung cancer induced angiogenesis *in vitro* (14).

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In the present study, we used immunohistochemical staining and a real-time quantitative PCR assay to measure PEDF protein and message, respectively, in lung cancer, and attempted to establish if a link existed between PEDF and non-small cell lung cancer and patients clinical outcome.

## Materials and methods

**Patients and samples.** A total of 91 patients with non-small cell lung cancer, who did not receive any adjuvant therapy prior to surgical operation, at Beijing Cancer Hospital from July 2000 to August 2003, were included in this study. Histological types of these lung cancers included squamous carcinoma (42 cases), adenocarcinoma (33 cases), adeno-squamous carcinoma (6 cases), large cell carcinoma (2 cases), carcinoid (2 cases), alveolar carcinoma (6 cases). No other previous or concomitant primary cancer was present. Clinicopathological characteristics were defined according to TNM criteria of UICC (15). Clinical and pathological information including age, sex, histological types of tumors, tumor cell grade, TNM stage, vessel embolism, lymph node metastasis, were reviewed and stored in a database. Patients were followed up from the day of their operation to August 2004. The follow-up intervals were calculated as survival intervals after surgery. Ethical approval and informed consents were obtained from the local ethics committee and patients respectively.

Fresh tissue samples were obtained from 21 primary non-small cell lung cancers, resected surgically from patients at Clinical Oncology School of Peking University from 2002 to 2003. The tissues were immediately collected after surgery, frozen in liquid nitrogen and stored at  $-80^{\circ}\text{C}$  for RNA extraction. Surrounding normal lung tissues ( $>5$  cm away from the tumor margin) were also collected. These cohorts include 13 men and 8 women, with a mean age of  $56.2 \pm 6.4$  years. The histological type of lung cancer was classified based on the World Health Organization method (16). Tumor staging was performed according to the TNM staging criteria of the UICC (15). The histological type of this collection included 10 squamous cell carcinomas, 8 adenocarcinomas, and 3 undifferentiated carcinomas. Tumor staging was I in 3 cases, II in 7 cases, IIIA or IIIB in 10 cases, and IV in 1 case.

**Materials.** A goat polyclonal antibody of PEDF and a mouse monoclonal antibody of  $\beta$ -actin were purchased from Santa Cruz Biotechnology Inc. (Santa Cruz, CA, USA). Mouse monoclonal antibody of CD31 was purchased from Beijing Zhongshan Biotechnology Co. Ltd. (Beijing, China). The biotin conjugated anti-goat IgG, anti-mouse IgG antibodies were purchased from Sigma (Poole, Dorset, UK). The Target Retrieval Solution was purchased from Dako Corp. RNA extraction and reverse transcription kits and PCR mix were purchased from Bio-Rad Corp. Primers were synthesized by BioAsia Corp. (Shanghai, China).

**PEDF staining and micro-vessel counting.** The paraffin-embedded tissue sections of 91 patients were cut at  $4\ \mu\text{m}$  and mounted on polylysine-coated glass slides for immunohistochemistry. Deparaffinized sections were first heated at  $60^{\circ}\text{C}$  for 1 h. Antigen retrieval was performed by heating the

samples without boiling in Target Retrieval Solution, pH 6.7 (200 ml) in a microwave oven for 10 min. After endogenous peroxidase was blocked with 3% hydrogen peroxide, each section was incubated with non-immunised horse serum (Sigma) for 15 min, in order to block the non-specific antigen site. The immunohistochemical staining procedure was performed according to manufacturer's protocol. The primary anti-PEDF and anti-CD31 antibodies were used at a dilution 1:100 from the stock. Following incubation at  $4^{\circ}\text{C}$  overnight, the sections were extensively washed and then incubated with the respective link antibody. Following extensive washing, bound antibodies were linked to avidin-biotin-peroxidase according to the instruction of Dako Corp., followed by developing colour using a DAB (diaminobenzidine tetrahydrochloride) solution for 5 min. The slides were counterstained with Mayers hematoxylin blue in 0.3% ammonia. For negative controls, sections were stained in the same manner, except that the primary antibody was absent from the solution.

**Evaluation of PEDF staining and the micro-vessel density counting.** PEDF staining in lung cancer cells was independently assessed by two observers using a modification of the system of grading the relative intensity of immunoreactivity for the respective antibodies (17,18). PEDF immunohistochemical staining of a tissue sample was graded as either low level (patchy and weak or negative immunoreactivity) or high level (uniformly intense immunoreactivity).

Micro-vessel density (MVD) was evaluated according to a method recently described (18,19). After screening the areas with intense neovascularized spots at low power field ( $\times 100$ ), microvessels in the area with the highest number of discrete microvessels were counted in a  $\times 400$  field. Three separate areas with intense neovascularisation were assessed for each 'hot' spot, and the mean was calculated as MVD of each tumor evaluated. The MVD levels were graded as low with MVD number  $<26$ , and as high with MVD number  $>26$ .

**Generation of cDNA from fresh frozen NSCLC tissue and normal tissue and RT-PCR.** RNA was extracted from tumor and the matched normal surrounding tissues in RNA extraction buffer using the standard guanidine isothiocyanate method by following the manufacturer's protocol and as we previously reported (20). Reverse transcription was performed from  $1\ \mu\text{g}$  of total RNA using oligo dt primer according to the manufacturer's instructions.

Conventional PCR primers were designed using primer 3 ([http://frodo.wi.mit.edu/cgi-bin/primer3/primer3\\_www.cgi](http://frodo.wi.mit.edu/cgi-bin/primer3/primer3_www.cgi)), to allow amplification of regions that have no overlap with other known genes and span at least one intron. Sequences for the PEDF primers are: 5'-CGATGAGATCAGCATTC TCC-3' and 5'-ATTCTGGGTCACTTTCAGGG-3' (product size 256 bp); and for the GAPDH primers: 5'-AGGTCGGAG TCAACGGATTTG-3' and 5'-GTGATGGCATGGACTG TGGT-3' (product size 532 bp). Conventional PCR was with the following conditions:  $95^{\circ}\text{C}$  5 min,  $94^{\circ}\text{C}$  for 1 min,  $55^{\circ}\text{C}$  for 45 sec,  $72^{\circ}\text{C}$  for 30 sec and a final extension phase at  $72^{\circ}\text{C}$  for 7 min for 40 cycles. The PCR products were separated on a 2% agarose gel and stained with  $5\ \mu\text{l}$  ethidium bromide prior to examination under UV light and photographs taken.

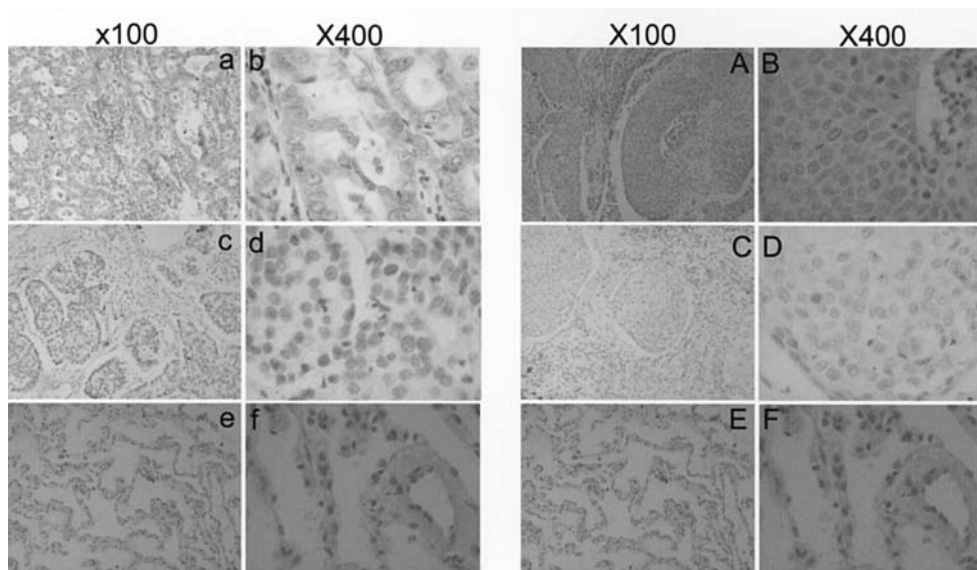


Figure 1. Immunohistochemical staining for PEDF in adenocarcinoma (a-d) and squamous carcinoma (E-D) and in normal lung tissues (e and f). (a/A and b/B) Showed a diffuse cytoplasmic staining of PEDF in adenocarcinoma of lung. (c/C and d/D) Showed a negative staining in adenocarcinoma of lung. (e/E and f/F) Showed a positive staining status of PEDF in alveoli. Original magnification: x100 for the left panel and x400 for the right panel.

**Preparation of real-time PCR standard and real-time quantitative RT-PCR analysis for PEDF transcript.** The procedure was similar to a method recently described (21). The PCR product from the above reaction was gel-excised and purified using a gel purification kit (Tianwei Corp., Beijing). It was subsequently quantified on a gel with lambda molecular weight standards and in a spectrophotometer. The number of copies of target template was later calculated. The DNA sample was serially diluted to yield a concentration range between  $10^2$  and  $10^8$  copies, which was subsequently used as an internal standard for quantitation. This was finally prepared in elution buffer, aliquated and stored at  $-80^{\circ}\text{C}$  until use.

The iCycler iQ™ system incorporates a gradient thermocycler and a 96-channel optical unit. SYBR Green, which is a fluorescence dye only for dsDNA, was used to detect the PCR product of PEDF and GAPDH (22,23). Quantitative PCR was carried out in a 96-well plate with 10 pmol forward and reverse primers, and the working solution of SYBR Green, using a customer PCR master mix, with the following conditions:  $95^{\circ}\text{C}$  for 5 min, followed by 40 cycles at  $95^{\circ}\text{C}$  for 1 min,  $55^{\circ}\text{C}$  for 45 sec,  $72^{\circ}\text{C}$  for 30 sec. Detection of fluorescence signal was at extension stage. The copy number of each transcript was calculated as the relative copy number normalized by GAPDH copy number. Every assay included test cDNA samples, 10-fold serial dilutions of the standard qualification, and controls (no template).

The optimum temperature and the specificity of the analysis based on SYBR Green method were pre-determined using the melting temperature curve. The specificity was further confirmed by agarose gel electrophoresis.

**Statistic analysis.** Chi-square analysis was used to test the association of PEDF expression level with standard pathological variations. Clinical pathological parameters and PEDF expression status were correlated with survival time in both univariate and multivariate analyses. Variables included in

univariate analysis were MVD, PEDF, gender, TNM stage, grade of differentiation, vessel cancer embolus, lymph nodal status and use of postoperative adjuvant therapy. Variables included in multivariate analysis were MVD, PEDF expression status, gender, TNM stage, grade of differentiation, vessel cancer embolus, lymphatic nodal status and use of post-operative adjuvant therapy. The log-rank test was used to test equality across categorical factors in univariate analysis, and the level of significance was set at  $P \leq 0.05$  based on two-sided test. The multivariate analysis was performed using Cox proportional hazards model, and the level of significance was set at  $P \leq 0.05$  based on two-sided test.

Paired-sample analysis was used (Student's t-test) to determine the difference of PEDF mRNA expression level observed between matched cancer tissue and the normal tissue. Independent-samples analysis was used (Student's t-test) to determine the differences observed between PEDF expression level in NSCLC tissue and the clinicopathological characteristics. Statistical tests were performed using the software SPSS 10.0 (SPSS Inc., Chicago, IL).

## Results

**Immunohistochemical analysis and the localization of PEDF in lung cancer.** The staining pattern for PEDF was mainly in the cytoplasm. In both adenocarcinoma (Fig. 1a and b) and squamous cell carcinoma of lung (Fig. 1A and B) the staining was diffuse and weaker, when compared with normal lung tissues (Fig. 1e and f, Fig. 1E and F).

**PEDF and the clinical correlation.** The immunohistochemical staining results of PEDF in NSCLC are shown in Table I. Low level expression of PEDF in NSCLC was found in 48 (52.7%) cases. Amongst the available clinical parameters, tumor grade was the only one showing a significant correlation with PEDF ( $P=0.016$ ), while others failed to reach a significant correlation ( $P>0.05$ ).

Table I. Tumor expression of PEDF vs clinicopathologic features in the complete series (n=91).

Variable	Cases (%)	PEDF expression		P-value ( $\chi^2$ )
		Low expression	Over-expression	
Sex				
Men	63 (69.2)	35	28	0.498
Women	28 (30.8)	13	15	
Age				
≤60	43 (47.3)	21	22	0.532
>60	48 (52.7)	27	21	
Histological type				
Squamous carcinoma	42 (46.2)	24	18	0.054
Adenocarcinoma	33 (36.3)	15	18	
Adenosquamous carcinoma	6 (6.6)	6	0	
Large cell carcinoma	2 (2.2)	0	2	
Carcinoid	2 (2.2)	0	2	
Alveolar carcinoma	6 (6.6)	3	3	
Grade of differentiation				
Poor	11 (12.1)	5	6	0.016
Moderate	48 (51.7)	32	16	
Well	32 (35.2)	11	21	
Tumor size				
T1	9 (9.9)	4	5	0.822
T2	60 (65.9)	32	28	
T3	19 (20.9)	11	8	
T4	3 (3.3)	1	2	
Nodal status				
N (-)	49 (53.8)	22	27	0.141
N (+)	42 (46.2)	26	16	
Vessel cancer embolus				
V (-)	67 (73.6)	34	33	0.635
V (+)	24 (26.4)	14	10	
TNM stage				
I	40 (43.9)	16	24	0.191
II	20 (22.0)	11	9	
IIIa	28 (30.8)	19	9	
IIIb	1 (1.1)	1	0	
IV	2 (2.2)	1	1	

Table II. Micro-vessel density counting vs clinicopathologic features in the complete series (n=91).

Variable	Cases (%)	MVD		P-value ( $\chi^2$ )
		Low	High	
Sex				
Men	63 (69.2)	37	26	0.654
Women	28 (30.8)	15	13	
Age				
≤60	43 (47.3)	24	19	0.835
>60	48 (52.7)	28	20	
Histological type				
Squamous carcinoma	42 (46.2)	24	18	0.312
Adenocarcinoma	33 (36.3)	20	13	
Adenosquamous carcinoma	6 (6.6)	2	4	
Large cell carcinoma	2 (2.2)	2	0	
Carcinoid	2 (2.2)	0	2	
Alveolar carcinoma	6 (6.6)	4	2	
Grade of differentiation				
Poor	11 (12.1)	7	4	0.345
Moderate	48 (52.7)	24	24	
Well	32 (35.2)	21	11	
Tumor size				
T1	9 (9.9)	5	4	0.988
T2	60 (65.9)	34	26	
T3	19 (20.9)	11	8	
T4	3 (3.3)	2	1	
Nodal status				
N (-)	49 (53.8)	29	20	0.678
N (+)	42 (46.2)	23	19	
Vessel cancer embolus				
V (-)	67 (73.6)	39	28	0.812
V (+)	24 (26.4)	13	11	
TNM stage				
I	40 (44.0)	27	13	0.389
II	20 (22.0)	10	10	
IIIa	28 (30.8)	14	14	
IIIb	1 (1.1)	0	1	
IV	2 (2.2)	1	1	

*Micro-vessel density in lung cancer and its correlation with PEDF.* Micro-vessel endothelial cells were positively stained for CD31. The correlation between MVD number and the clinicopathologic features in NSCLC are shown in Table II.

Low level of MVD in NSCLC was found in 52 (57.1%) cases. Amongst the available clinical parameters, no clinical factors showed significant correlation with CD31 as shown in Table II ( $P>0.05$ ).



Table III. The correlation between PEDF and microvascular density.

PEDF expression	Cases	Mean $\pm$ SD	Standard error	P-value
Low	48	28.50 $\pm$ 8.22	1.19	<0.0005
High	43	16.98 $\pm$ 6.15	0.94	

Table III shows the correlation between PEDF and MVD. The mean MVD counts were 28.5 $\pm$ 8.2 (mean $\pm$ 8SD) for the 43 patients who had low-level PEDF staining. MVD counts for the 43 high-level PEDF was 16.98 $\pm$ 6.15. The difference between these two groups was highly significant (P<0.0005).

*Clinical outcome and prognostic value of variables.* The median time of follow-up for the cohort (n=91) was 24.6 months (range 2-49 months). The mean (SD) survival time was 23.1 $\pm$ 14.4 months. Cumulative survival curves were calculated using the Kaplan-Meier method. Univariate analysis of the impact of histological types on PEDF expression status is presented in Table IV. Longer survival time was found to be significantly correlated with the following factors: low TNM stage (P=0.0005), node negative (mean survival time, 41.04 months vs 31.25 months for node positive, P=0.0051), low MVD (39.96 months vs 31.85 months with high MVD, P=0.0434), high-level PEDF (40.56 months vs 32.41 months vs low PEDF, P=0.0378) (Fig. 2, top A).

A multivariate prognostic analysis based on the Cox proportional hazard model showed that amongst all the available indicators, TNM stage was the only independent prognostic factors (P=0.004, respectively).

Survival analysis using Kaplan-Meier survival curve based on PEDF is shown in Fig. 2 (top), in that cumulative survival time for patients with high-level PEDF (Fig. 2, top A; n=43) was significantly longer than those with low-level PEDF (n=43) (Fig. 2, top B; P=0.0378). Significantly shorter survival time was also seen in patients with node positive tumors (n=42), P=0.0051, vs node negative tumors (n=49) (Fig. 2, middle). Furthermore, survival difference between patients with low MVD (n=52) and high MVD (n=43) was significant (P=0.0434) (Fig. 2, bottom).

*PEDF mRNA expression level in NSCLC and normal lung tissue assessed by real-time RT-PCR.* We have generated an internal standard curve for the quantitative analysis of PEDF and GAPDH. The regression coefficients for the regression lines plotting the threshold cycle ( $C_T$ ) against the starting amount of the standard sample were 0.997, 0.997, for GAPDH and PEDF respectively. Both the reaction curve and the melting curve confirmed that the amplification products for these molecules were specific. This standard was subsequently used to deduce the transcript numbers of PEDF which is shown in the current report as copy number of PEDF mRNA/copy number of GAPDH mRNA.

The relative copy number for PEDF in 21 samples of lung cancer tissue ranged from 0.11 to 1.11, with a mean  $\pm$  SD of

Table IV. Potential prognostic factors using univariate analysis.

Characteristics	Pts.	Mean survival time (months)	P-value from log-rank test
Sex			
Male	63	36.42 (31.77-41.08)	0.8860
Female	28	37.08 (30.73-43.44)	
TNM stage			
I	40	41.61 (37.35-45.87)	0.0005
II	20	38.44 (31.18-45.70)	
IIIa	28	30.31 (22.76-37.86)	
IIIb	1	6.00 (6.00-6.00)	
IV	2	13.50 (3.11-23.89)	
Grade of differentiation			
Well	32	37.43 (31.17-43.69)	0.4984
Moderate	48	34.76 (29.86-39.67)	
Poor	11	39.73 (28.34-51.12)	
Nodal status			
N (-)	49	41.04 (36.64-45.45)	0.0051
N (+)	42	31.25 (25.64-36.86)	
Vessel cancer embolus			
V (-)	67	37.50 (33.41-41.58)	
V (+)	24	33.89 (25.75-42.03)	
Postoperative adjuvant therapy			
No	26	37.47 (29.70-45.24)	0.7225
Chemotherapy or radiation	65	36.03 (31.57-40.49)	
MVD			
High	39	31.85 (25.68-38.03)	0.0434
Low	52	39.96 (35.48-44.44)	
PEDF staining			
Low expression	48	32.41 (26.72-38.09)	0.0378
Over-expression	43	40.56 (35.92-45.21)	

0.55 $\pm$ 0.36, while the corresponding values in the matched normal lung tissue ranged from 0.29 to 1.31, with a mean  $\pm$  SD of 0.72 $\pm$ 0.26. PEDF mRNA expression level in normal tissue was significant higher than in the matched cancer tissue (95% CI: 2.530E-02-0.3176, P=0.024, paired t-test), in that 16/21 (76%) of tumor samples had lower levels of PEDF transcript than their matched healthy lung tissues and 5/21 (26%) had higher levels.

*Relationship between PEDF mRNA expression and clinicopathologic variables.* Table V shows the relationship between PEDF mRNA expression and the clinicopathological characteristics. PEDF mRNA expression level was

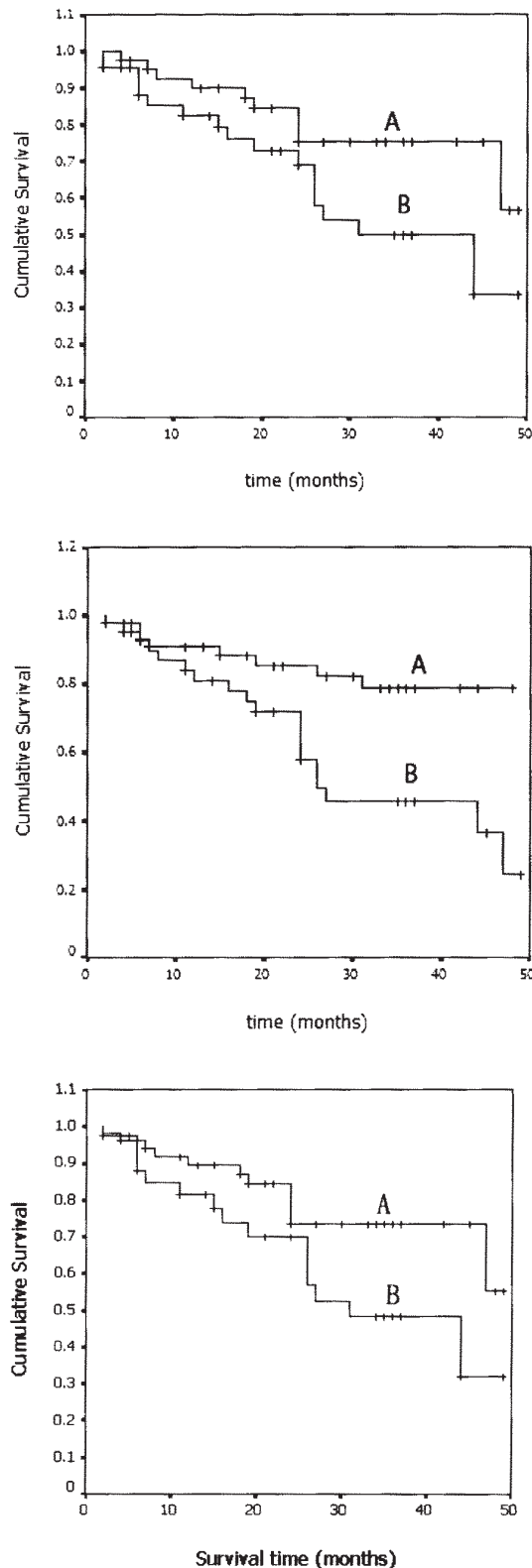


Figure 2. Survival analysis. Top, overall survival of the NSCLC patients based on PEDF levels (n=91). A, PEDF over-expressing tumors (n=43). B, PEDF low expressing tumors (n=48). The survival curves are significantly separated ( $P=0.0378$ , log-rank test) and the patients with PEDF over-expression have longer survival time than those with low expression. Middle, overall survival based on nodal status of NSCLC patients (n=91). The difference between nodal negative patients (A, n=49) and nodal positive patients (B, n=42) is significant ( $P=0.0051$ , log-rank test). Bottom, overall survival based on MVD status of NSCLC patients (n=91). The difference between low level of MVD (A, n=52) and high level MVD patients (B, n=43) is significant ( $P=0.0434$ , log-rank test).

Table V. Relationship between PEDF mRNA expression relative level (PEDF/GAPDH) and clinicopathologic characteristics.

Variables	No.	PEDF mRNA expression <sup>a</sup> (mean $\pm$ SD)	P-value
Sex			
Male	13	0.47 $\pm$ 0.35	0.198
Female	8	0.68 $\pm$ 0.35	
Histological type			
Squamous	10	0.60 $\pm$ 0.35	0.601
Non-squamous	11	0.51 $\pm$ 0.382	
Stage			
I-II	10	0.75 $\pm$ 0.39	0.010
III-IV	11	0.37 $\pm$ 0.22	
Tumor status			
T1	6	0.87 $\pm$ 0.30	0.007
T2-4	15	0.42 $\pm$ 0.31	
Lymph node status			
N0	8	0.68 $\pm$ 0.40	0.216
N1-3	13	0.47 $\pm$ 0.32	
Tissue			
Tumor	21	0.55 $\pm$ 0.36	0.024 <sup>b</sup>
Adjacent normal tissue	21	0.72 $\pm$ 0.26	

<sup>a</sup>PEDF mRNA expression derived from real-time quantitative RT-PCR: PEDF/GAPDH. <sup>b</sup>P-value derived from paired t-test; others derived from independent t-test.

significantly higher in stage I-II patients than in stage III-IV patients (0.75 $\pm$ 0.39 vs 0.37 $\pm$ 0.22;  $P=0.010$ , independent t-test). The levels were also significantly higher in small tumors (T1) than in large tumors (T2-T4) (0.87 $\pm$ 0.30 vs 0.42 $\pm$ 0.31;  $P=0.007$ , independent t-test). The relationship between PEDF mRNA expression and sex, histologic type, and lymph node status was not statistically significant ( $P>0.05$ ).

## Discussion

PEDF is one of the most potent naturally occurring angiogenesis inhibitor, which is a key factor associated with avascularity of the cornea, vitreous, and the outer retinal layer of the eye (5). PEDF is also an important negative regulator of angiogenic activity of aqueous humour. The current study is the first to report that PEDF was reduced at both protein and mRNA level in non-small cell lung cancer and bore a clinical significance to the patients.

First, the current study has demonstrated that PEDF presents in lung cancer cells, and that it primarily located in

the cytoplasmic region of the cells, which is in accordance with literature reports. Stromal cells and endothelial cells displayed little staining. Second, PEDF is significantly linked to MVD. Low levels of PEDF is significantly linked to high MVD. The formation of neovasculature is under tight regulation in most healthy tissue and is most probably controlled by the balance between angiogenic and anti-angiogenic factors. The disruption of such a balance might play an essential role in the dysregulated development of neovasculature which form the one essential part of cancer development. PEDF was first purified from the conditioned media of human retinal pigment epithelial cells as a neurotrophic factor. The most convincing indication that PEDF may regulate angiogenesis is probably coming from studies on retinopathy, in which PEDF and VEGF dually regulate the development of neovascularization (5). When oxygen is sufficient, the VEGF level decreases but the PEDF level is upregulated to prevent new blood vessels growth (24). The levels of PEDF in diabetic patients can predict the development and progression of retinopathy (25). The regulation of PEDF gene expression and the anti-angiogenic mechanisms of PEDF remain unclear, although Dawson *et al* demonstrated that hypoxia may reduce the level of PEDF proteins (5). Thus, it appears that this balance is the evolutionary result of adaptation to environment change in normal biology situation. The role of PEDF in solid tumors has not been previously described. The current study suggests that low level of PEDF in human lung tumor cells is inversely associated with MVD and that low level of PEDF is linked to longer survival. This observation has put PEDF on a 'good' list of protein factors in the control of angiogenesis, in lung cancer.

In the current study, we have provided additional evidence that the message level of PEDF is also reduced in NSCLC tumors. We have first successfully developed a quantitative analysis method for PEDF. Using matched tumor and normal tissue samples, it was found that PEDF mRNA expression level in matched normal tissue was significant higher than the cancer tissue (Table V,  $P < 0.05$ ). Furthermore, levels of PEDF message were significantly higher in stage III and stage IV tumors than lower stage tumors. This information provides further supporting data for protein based analysis, in that PEDF protein levels were lower in aggressive tumors, and suggest that the aberration of PEDF expression may be at both protein and transcription level. The second key point here is the provision of direct comparison between matched normal and tumor tissues. The study has clearly shown a reduced level in tumor tissues than in the matched normal tissues. However, the regulation of PEDF gene expression and the anti-angiogenic mechanisms of PEDF are largely unknown. Dawson *et al* demonstrated that hypoxia could reduce the level of PEDF proteins (5), and that this occurs at the translational or post-translational level.

PEDF has a very interesting dual function feature. In the eyes, it acts as a potent anti-angiogenic factor. It is also an extremely potent inhibitor that is active at low nanomolar concentrations against a wide variety of angiogenic factors, including VEGF (26). It acts directly on endothelial cells and can halt the forming of new vessels by inducing the apoptotic death of endothelial cells that have been stimulated during the formation of new vessels (10).

Targeting angiogenesis has also been postulated to have potential in cancer therapies, including lung cancer. Indeed, encouraging reports have demonstrated that the anti-VEGF antibody, Avastin, can prolong the survival of patients with advanced colon cancer (27) and non-small cell lung cancers (28). These early successes have suggested that targeting angiogenesis has some important role in NSCLC therapies. Given the potential anti-angiogenesis effect and the profound reduction of the factor in lung cancer, particularly in aggressive lung cancer, it is suggested that PEDF may have a good therapeutic value, if it can be successfully engineered and delivered.

In summary, we have shown the PEDF protein and mRNA expression was significantly lower in cancer tissue than in normal tissue. We also found that PEDF expression level was negatively correlated with NSCLC stage and tumor size, and patient survival. Therefore, PEDF to has an anti-angiogenesis role in NSCLC.

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