

Papers

Plasminogen activator system, vascular endothelial growth factor, and colorectal cancer progression

E A Baker, F G Bergin, D J Leaper

Abstract

Aims—The plasminogen activator system (PAS) consists of the plasminogen activators (urokinase (uPA) and tissue-type (tPA) plasminogen activators), the uPA receptor (uPAR), and the plasminogen activator inhibitors (PAI-1 and PAI-2). Plasminogen activators activate plasminogen to plasmin, which can break down extracellular matrix (ECM) components. Vascular endothelial growth factor (VEGF) is a mitogen for endothelial cells and is involved in angiogenesis. VEGF has been shown to upregulate uPA and this may facilitate tumour angiogenesis further.

Methods—PAS components and VEGF were determined by enzyme linked immunosorbent assay (ELISA) in paired colorectal tumour and normal tissue (n = 50) and correlated with pathological staging.

Results—uPA, uPAR, PAI-1, and VEGF values were significantly higher in tumour tissue (for example, tumour uPA: median, 2.3 (range, 0.1–6.7) ng/mg protein *v* normal uPA: median, 0.2 (range, 0–2.6) ng/mg protein). tPA was significantly higher in normal mucosa and there was no difference in PAI-2. uPA, uPAR, PAI-1, and VEGF values significantly correlated with each other and with Dukes’s staging (uPA in adenomas: median, 0.42 (range, 0.1–1.2) ng/mg protein; uPA in Dukes’s B tumours: median, 2.1 (range, 0.4–4.3) ng/mg protein; and uPA in Dukes’s D tumours: median, 4.0 (range, 3.7–4.2) ng/mg protein) and lymphatic invasion. In addition PAI-1 also correlated with tumour size and differentiation.

Conclusion—The involvement of the PAS and VEGF in colorectal cancer appears to be complex. uPA, uPAR, PAI-1, and VEGF

were upregulated in tumour tissue and this correlated with Dukes’s staging and lymphatic invasion.

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Angiogenesis is important for tumour growth and metastasis.^{1,2} The formation of tumour microvessels is stimulated by angiogenic factors such as vascular endothelial growth factor (VEGF) and basic fibroblast growth factor (bFGF).² VEGF is a 45 kDa glycoprotein that is mitogenic for endothelial cells. It binds to specific receptors on endothelial cells, where it induces endothelial proliferation and capillary tube formation, enhancing tumour neovascularisation or angiogenesis. Previous studies have shown that overexpression of VEGF correlated with the progression of human malignancies, including colorectal cancer.^{3,4} The properties of VEGF have been reviewed elsewhere.^{5,6}

Tumour invasion and metastasis is a complex process involving many sequential, interrelated events including angiogenesis, intravasation, and extravasation. Many of these stages, including angiogenesis, involve the controlled degradation of extracellular matrix (ECM) components by proteinases. The extracellular proteinases involved in ECM degradation are the matrix metalloproteinases (MMPs) and the serine proteinases.

The plasminogen activator system (PAS) consists of two serine proteinases, plasminogen activators, urokinase-type (uPA) and tissue-type (tPA), the plasminogen activator inhibitors 1 and 2 (PAI-1 and PAI-2), and the receptor for urokinase type plasminogen activator (uPAR). The plasminogen activators, as the name suggests, convert inactive plasminogen to the active serine proteinase, plasmin. Plasmin is

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Table 1 Conditions for performing the plasminogen activator system and vascular endothelial growth factor (VEGF) ELISAs

ELISA	Dilution	Capture antibody	Incubation	Detection antibody	Conjugate	Standard curve
uPA	1/10	Mouse antihuman	Overnight 4°C	Biotinylated antihuman uPA	Streptavidin–HRP	0.1–1.0 ng/ml
tPA	None	Goat anti-tPA	1 hour RT shaking	–	HRP anti-tPA	0.5–10.0
uPAR	1/5	Mouse antihuman	Overnight 4°C	Biotinylated antihuman UpaR	Streptavidin–HRP	0.1–3.0 ng/ml
PAI-1	1/10	Mouse antihuman	Overnight 4°C	Biotinylated antihuman PAI-1	Streptavidin–HRP	0.1–10.0 ng/ml
PAI-2	None	Polyclonal antihuman	2 hours RT	Biotinylated antihuman PAI-2	Streptavidin–HRP	0.5–10 ng/ml
VEGF	None	Mouse antihuman	2 hours RT	–	HRP anti-VEGF	31.2–2000 pg/ml

ELISA, enzyme linked immunosorbent assay; HRP, horseradish peroxidase; PAI, plasminogen activator inhibitor; RT, room temperature; tPA, tissue type plasminogen activator; uPA, urokinase plasminogen activator.

Table 2 Differences in plasminogen activator, PAI and uPAR (ng/mg protein), and VEGF (pg/mg protein) values in colorectal tumour and normal tissue samples

	Tumour	Normal
uPA	2.3* (0.1–6.7)	0.2 (0–2.6)
tPA	3.3 (0.1–23.8)	6.8* (1.1–37.5)
uPAR	2.2* (0.2–14.0)	0.6 (0.2–4.1)
PAI-1	9.8* (0.2–62.4)	2.0 (0.1–20.7)
PAI-2	1.0 (0–7.3)	0.7 (0–4.7)
VEGF	189* (30–4156.0)	27 (3–108)

PAI, plasminogen activator inhibitor; PAR, plasminogen activator receptor; tPA, tissue type plasminogen activator; uPA, urokinase plasminogen activator; VEGF, vascular endothelial growth factor.

Values are expressed as median with the range in parentheses.

* $p < 0.0001$, Mann Whitney.

involved in ECM degradation directly because it is a broad substrate proteinase that can degrade most proteins within the ECM (for example, fibronectin, laminin, and proteoglycans).⁷ Plasmin also acts on the ECM indirectly by activating the latent forms of several MMPs (for example, MMP-1).

The function and regulation of PAS components has been reviewed elsewhere.^{7–9} In brief, the two plasminogen activators differ in biological function and distribution; uPA is important in tissue degradation in both physiological and pathological processes,^{7, 10} whereas tPA is important in fibrinolysis.⁸ The plasminogen activators also differ in inhibitory functions. PAI-1 inhibits both uPA and tPA; however, PAI-2 inhibits only uPA. In vivo it is the balance between plasminogen activators and their inhibitors that determines matrix

degradation by plasmin. The function of the uPA receptor is to focus the proteolytic potential of the tumour cells, thereby enhancing invasion.¹¹

The PAS has been studied extensively in various human cancers. Combinations of these components, in particular uPA and PAI-1, have been found to be of prognostic relevance in several human cancers, including colorectal^{12, 13} and breast cancer.^{14, 15}

Most previous studies in colorectal cancer have determined the expression of individual components,^{13, 16, 17} or related the expression of PAS components to specific histological grades.^{18, 19}

In addition to VEGF expression correlating with tumour progression, several studies have also found VEGF to increase both uPA and uPAR on endothelial cells in vitro,^{20, 21} and to correlate with uPA²² and uPAR²³ expression in vivo.

Our study aims to determine the concentrations of VEGF and the five components of the PAS in paired colorectal tumour and normal tissue samples and correlate these with the clinical pathological grade of the tumour.

Methods

MATERIALS

Enzyme linked immunoassay (ELISA) kits for uPA, tPA, uPAR, PAI-1, and PAI-2 were purchased from American Diagnostica (Greenwich, USA) and for VEGF from R&D Systems (Abingdon, UK). Phosphate buffered

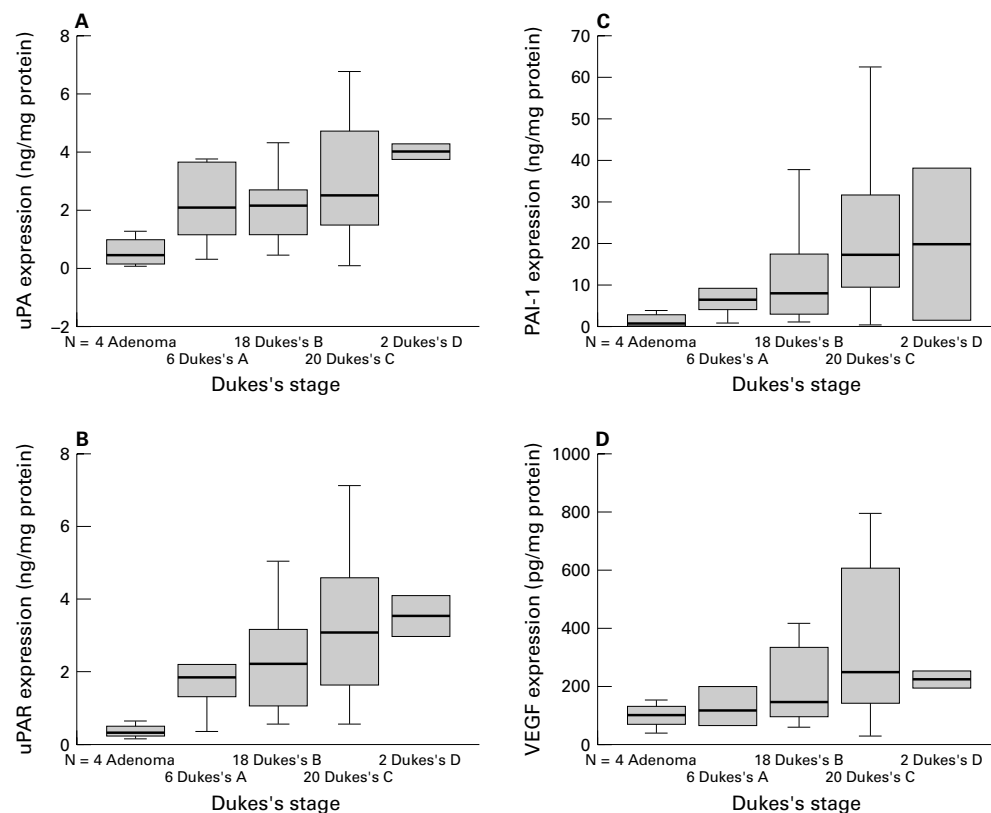


Figure 1 Box plots showing the positive correlation between (A) urokinase plasminogen activator (uPA), (B) the uPA receptor (uPAR), (C) plasminogen activator inhibitor 1 (PAI-1), and (D) vascular endothelial growth factor (VEGF) values with the Duke's stage of the tumour (median and interquartile range) in colorectal cancer tissue (* $p < 0.05$, Spearman's correlation). The line within the box plot corresponds to the median value.

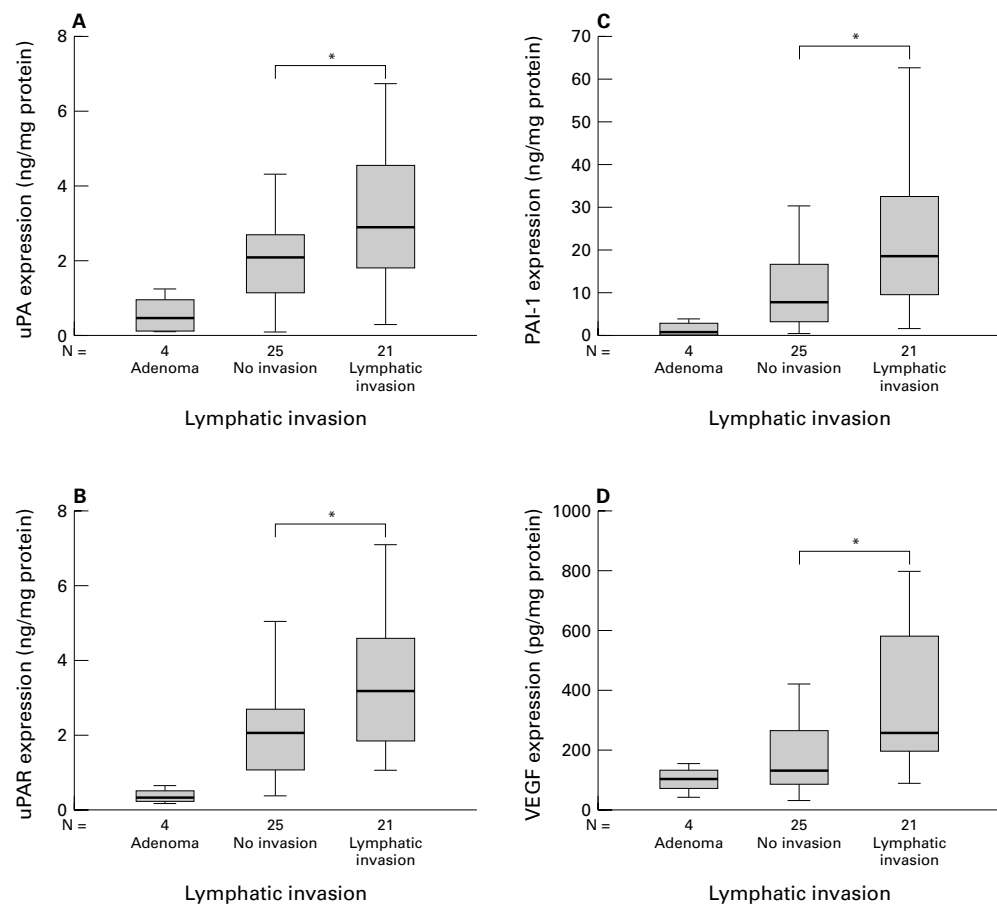


Figure 2 Box plots showing the positive correlations between (A) urokinase plasminogen activator (uPA), (B) the uPA receptor (uPAR), (C) plasminogen activator inhibitor 1 (PAI-1), and (D) vascular endothelial growth factor (VEGF) values and lymphatic invasion (median and interquartile range). Expression was significantly greater in tumours that had undergone lymphatic invasion ($*p < 0.05$, Spearman's correlation). The line within the box plot corresponds to the median value.

saline (PBS), total protein assay, and sulphuric acid were from Sigma Aldrich (Poole, Dorset, UK).

PATIENTS AND TISSUE SAMPLES

Colorectal specimens were collected from the operating theatre and taken to the department of pathology, North Tees General Hospital. Paired tissue samples (tumour margin and normal colon) were obtained by a consultant histopathologist within 30 minutes of surgical resection. Fifty paired colorectal tissue samples were obtained from consecutive patients between April 1998 and June 1999.

Each tissue sample was finely diced with a scalpel, homogenised (Polytron PT 2,100; Kinematica; Philip Harris Scientific, Lichfield, UK) in PBS for five minutes, and centrifuged for 15 minutes at $750 \times g$. The homogenate was removed and stored in aliquots at -80°C until analysis. The total protein concentration of each tissue sample was determined by the method of Ohnishi and Barr.

All the colorectal tumours were classified according to their pathological staging: Dukes's stage, differentiation, tumour depth, and whether the tumour had undergone lymphatic and/or vascular invasion.

ENZYME LINKED IMMUNOSORBENT ASSAYS

The concentrations of uPA, tPA, uPAR, PAI-1, PAI-2, and VEGF were measured by enzyme immunoassay kits. Table 1 summarises the required dilutions, antibodies, conditions, and detection ranges for each ELISA. The concentration of each protein was measured at 450 nm on a microplate reader (Dynex Technologies, Billingham, UK). Values for the PAS components (ng/ml) and VEGF (pg/ml) were determined for each sample from a standard curve using Revelation Software (Dynex Technologies, Billingham, UK). Final tissue values were expressed as ng/mg protein or pg/mg protein for PAS components and VEGF, respectively.

STATISTICAL ANALYSIS

For comparisons between concentrations in colorectal tumour and normal tissue samples the Mann Whitney U test for non-parametric data, with 95% confidence limits, was performed. Spearman's correlation coefficient was used to determine whether a correlation existed between these components and clinical pathological staging (Dukes's stage, tumour depth, differentiation, and vascular and lymphatic invasion). The data were considered to be significant at $p < 0.05$.

Results

In our study, paired colorectal tissues (tumour and normal mucosa) from 50 patients were analysed for expression of VEGF and components of the PAS. Histologically, four tumours were tubular adenomas, six were Dukes's A, 18 Dukes's B, 20 Dukes's C, and two Dukes's D tumours. Eleven patients had well differentiated, 28 moderately, and six poorly differentiated tumours. Twenty one of 46 tumours had undergone lymphatic invasion (lymphatic vessel invasion at the site of the primary tumour), and 13 of 46 had undergone vascular invasion at the time of resection.

EXPRESSION OF PAS COMPONENTS

uPA

uPA values were significantly higher in tumour tissue (median, 2.3 ng/mg protein; range, 0.1–6.7) than in the corresponding normal tissue (median, 0.2 ng/mg protein; range, 0–2.6; $p < 0.0001$, Mann Whitney; table 2).

tPA

tPA values were significantly higher in normal colorectal tissue (median, 6.8 ng/mg protein; range, 1.1–37.5) than in the tumour tissues (median, 3.3 ng/mg protein; range, 0.1–23.8; $p < 0.0001$, Mann Whitney U; table 2).

uPAR

Values for the uPA receptor were significantly higher in colorectal tumour tissue (median, 2.2 ng/mg protein; range, 0.2–14) than in normal colon tissue (median, 0.6 ng/mg protein; range, 0.2–4.1; $p < 0.0001$, Mann Whitney U; table 2).

PAI-1

PAI-1 values were significantly higher in tumour tissue (median, 9.8 ng/mg protein; range, 0.2–62.4) than in normal colorectal tissue (median, 2.0 ng/mg protein; range, 0.1–20.7; $p < 0.0001$, Mann Whitney; table 2).

PAI-2

Although PAI-2 values were greater in tumour tissue (median, 1.0 ng/mg protein; range, 0–7.3) than in normal mucosa (median, 0.7 ng/mg protein; range, 0–4.7) the difference was not significant (table 2).

VEGF

VEGF values were significantly higher in tumour tissue (median, 189 pg/mg protein; range, 307–4156) than normal colorectal tissue (median, 27.3 pg/mg protein; range, 3–108; $p < 0.0001$, Mann Whitney U; table 2).

CORRELATIONS BETWEEN PAS COMPONENTS AND VEGF

uPA, uPAR, PAI-1, and VEGF values within colorectal tumour tissue correlated significantly with each other (Spearman's correlation, $p < 0.05$). In addition, VEGF expression correlated with PAI-2 expression.

PAS, VEGF, AND PATHOLOGICAL STAGE

The expression of VEGF and the components of the PAS were correlated with the clinical

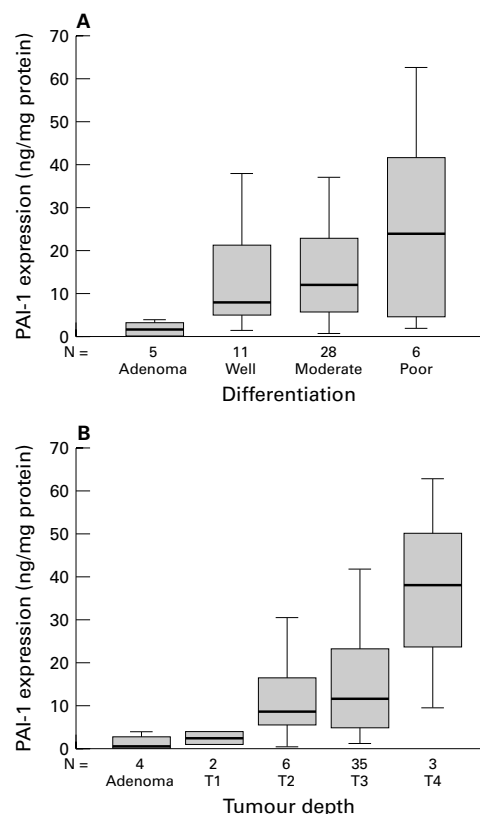


Figure 3 Box plots showing the positive correlation between plasminogen activator inhibitor 1 (PAI-1) values and (A) tumour differentiation and (B) tumour size (median and interquartile range). * $p < 0.05$ Spearman's correlation. The line within the box plot corresponds to the median value.

pathological stage of the tumour; that is, Dukes's stage, tumour differentiation, tumour depth (T₁–T₄), and finally whether the tumour had undergone lymphatic and/or vascular invasion.

Three components of the PAS (uPA, uPAR, and PAI-1), as well as VEGF, significantly correlated with tumour stage. These four factors correlated with the Dukes's stage of the tumour (fig 1) and lymphatic invasion (fig 2). In addition, the expression of PAI-1 also correlated with tumour differentiation and tumour depth (fig 3).

Discussion

The involvement of the PAS in colorectal cancer progression appears to be complex. In agreement with previous studies, values for uPA,^{19 25–29} its receptor uPAR,¹⁹ and the inhibitor, PAI-1,^{19 29} were higher in colorectal tumours than in the corresponding normal mucosa. However, tPA values were higher in normal colorectal tissue.^{19 29 30} This differential expression of uPA and tPA in tumour and normal tissue confirms their different roles in plasminogen activation in vivo: uPA is involved in tissue degradation and tPA in fibrinolysis.

The balance between the expression of the activators and inhibitors is important in vivo in determining whether matrix degradation and activation of other proteinases is likely to occur. In our present study, both uPA and the inhibitor PAI-1 were greater in colorectal

tumour tissue than in normal tissue. However, previous studies have shown that this increased PAI-1 expression does not significantly inhibit uPA activity in primary tumours; therefore, the balance will favour proteolysis.⁹

Within colorectal tumours there is a differential production and expression of PAS components: uPA and the PAIs are produced by stromal cells, whereas uPAR expression has been demonstrated on tumour cells. The increased expression of the receptor for uPA seen in colorectal tumours is thought to localise the proteolytic potential of the tumour cells and therefore their invasion.¹¹

Our study has also demonstrated a significant correlation between uPA, uPAR, and PAI-1 and the pathological staging of the tumour; in particular, Dukes's staging and lymphatic invasion. Previous studies have shown conflicting results on the correlations between PAS components and tumour pathology. Previously, Suzuki and colleagues demonstrated that uPAR expression increased from adenomas through to invasive carcinomas.¹⁷ Another study found that uPA values at the tumour-host interface in Dukes's C tumours were twice those of Dukes's A tumours; uPAR was also significantly greater in Dukes's C tumours, but there was no significant difference in PAI-1 values.²⁷ However, other studies found no correlation between uPA expression and tumour stage or the site of the tumour.^{16, 28}

As well as their involvement in tumour progression, PAS components have also been implicated in cancer prognosis in colorectal¹³ and other cancers.^{14, 15} In colorectal cancer, uPA,^{18, 19} uPAR,¹³ and PAI-1¹² expression in plasma has been associated with shorter survival. Furthermore, PAI-2 values in carcinomas and tPA concentrations in normal mucosa have been shown to have prognostic value on overall survival.¹⁹

In agreement with our results, it has been reported that VEGF expression (protein and mRNA) is significantly greater in colorectal tumour tissue than in normal tissue.^{31, 32} VEGF expression was also found to be significantly greater in tumours with lymphatic invasion than in those without.²³ It has been suggested that the involvement of VEGF in lymphatic invasion is similar to vascular angiogenesis, in that lymphatic endothelial cells may respond to VEGF by increasing angiogenesis within the lymphatics, thereby inducing metastasis via the lymphatics as well as via the blood vessels.³³ Our study also found that VEGF values within tumour tissue significantly correlated with the Dukes's stage of the tumour. In addition, one previous study has shown a correlation between VEGF expression and tumour size.³¹

Also in agreement with our results, the few studies that have compared the expression of VEGF with PAS components found a positive correlation between VEGF and both uPA²² and uPAR expression.²³

Previously, studies have determined the expression of individual PAS components in colorectal cancer progression and or survival;

however, there have been few studies determining the association of all PAS components with VEGF. Our results have demonstrated an increased coexpression of uPA, uPAR, PAI-1, and VEGF in colorectal tumour tissue compared with normal mucosa. A positive correlation was found between these PAS components (uPA, uPAR, PAI-1) and VEGF values, as well as with the pathological staging of colorectal cancer. uPA and VEGF are known to be involved in tumour angiogenesis, one of the initial stages of the metastatic cascade. However, further research is needed to determine their involvement in colorectal cancer invasion and metastasis. In addition, future follow up of the patients involved is required to determine the prognostic relevance of these factors.

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