

Secreted semaphorin 5A suppressed pancreatic tumour burden but increased metastasis and endothelial cell proliferation

A Sadanandam^{*,1,2,3}, SS Sidhu⁴, S Wullschleger³, S Singh¹, ML Varney¹, C-S Yang¹, AE Ashour⁵, SK Batra⁶ and RK Singh^{*,1,6}

¹Department of Pathology and Microbiology, University of Nebraska Medical Center, 985900 Nebraska Medical Center, Omaha, NE 68198-5900, USA; ²Swiss Institute of Bioinformatics (SIB), Quartier Sorge, Batiment Genopode, Lausanne 1015, Switzerland; ³Swiss Institute for Experimental Cancer Research (ISREC), Swiss Federal Institute of Lausanne (EPFL), Station 19, Batiment SV, Lausanne 1015, Switzerland; ⁴Institut National de la Santé et de la Recherche Médicale U753, Laboratoire Immunologie des Tumeurs Humaines, Interaction Effecteurs Cytotoxiques-Système Tumoral, Institut Fédératif de Recherche-54, Villejuif, France; ⁵Department of Pharmacology, College of Pharmacy, King Saud University, Riyadh, Saudi Arabia; ⁶Department of Biochemistry and Molecular Biology, University of Nebraska Medical Center, Omaha, NE 68198, USA

BACKGROUND: Our earlier reports demonstrated that membrane-bound semaphorin 5A (SEMA5A) is expressed in aggressive pancreatic cancer cells and tumours, and promotes tumour growth and metastasis. In this study, we examine whether (1) pancreatic cancer cells secrete SEMA5A and (2) that secreted SEMA5A modulates certain phenotypes associated with tumour progression, angiogenesis and metastasis through various other molecular factors and signalling proteins.

METHODS AND RESULTS: In this study, we show that human pancreatic cancer cell lines secrete the extracellular domain (ECD) of SEMA5A (SEMA5A-ECD) and overexpression of mouse Sema5A-ECD in Panc1 cells (not expressing SEMA5A; Panc1-Sema5A-ECD; control cells - Panc1-control) significantly increases their invasion *in vitro* via enhanced ERK phosphorylation. Interestingly, orthotopic injection of Panc1-Sema5A-ECD cells into athymic nude mice results in a lower primary tumour burden, but enhances the micrometastases to the liver as compared with Panc1-control cells. Furthermore, there is a significant increase in proliferation of endothelial cells treated with conditioned media (CM) from Panc1-Sema5A-ECD cells and a significant increase in microvessel density in Panc1-Sema5A-ECD orthotopic tumours compared with those from Panc1-control cells, suggesting that the increase in liver micrometastases is probably due to increased tumour angiogenesis. In addition, our data demonstrate that this increase in endothelial cell proliferation by Sema5A-ECD is mediated through the angiogenic molecules – interleukin-8 and vascular endothelial growth factor.

CONCLUSION: Taken together, these results suggest that a bioactive, secreted form of Sema5A-ECD has an intriguing and potentially important role in its ability to enhance pancreatic tumour invasiveness, angiogenesis and micrometastases.

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Pancreatic adenocarcinoma is one of the leading causes of cancer-related deaths in the United States. The prognosis of patients suffering from pancreatic adenocarcinoma is extremely poor. More than 80% of patients suffer from metastatic disease at the time of diagnosis and almost all the patients die either because of the local disease or metastases. The overall survival of pancreatic cancer patients is <6 months from the time of diagnosis and current therapeutic strategies are inadequate (Stathis and Moore, 2010). This low survival of pancreatic cancer patients is due to lack of early detection and/or metastatic markers. This urges for a better understanding of the molecular mechanism(s) behind pancreatic adenocarcinoma tumour progression and metastasis to identify novel metastatic markers for the disease.

Semaphorins are a family (containing eight subfamilies) of axonal growth molecules that contain secreted, glycosylphosphatidylinositol-anchored and transmembrane signalling proteins. They

share a conserved ~500 amino acids semaphorin domain in their N-terminal regions and have subfamily-specific domains in their carboxy-terminal regions (Chen *et al*, 1998; Yazdani and Terman, 2006). They were originally identified as bi-functional axonal growth molecules involved in the development of the nervous system (Chen *et al*, 1998). Semaphorins exert their cellular functions by binding to two different families of proteins called plexins and neuropilins (Kolodkin *et al*, 1993; Chen *et al*, 1998; Yazdani and Terman, 2006; Roth *et al*, 2009). In addition to their function during normal physiological processes such as the development of the nervous, cardiac, skeletal and vascular systems as well as the immune response, semaphorins also have major roles during pathogenesis of tumour formation, angiogenesis and metastasis (Chen *et al*, 1998; Christensen *et al*, 1998; Brambilla *et al*, 2000; Tamagnone and Comoglio, 2000; Kusy *et al*, 2005; Roth *et al*, 2009; Sadanandam *et al*, 2010b). Among the different semaphorins, the role of membrane-bound semaphorins in cancer has recently drawn more attention.

Previously, we identified semaphorin 5A (SEMA5A), a transmembrane bound semaphorin, as a putative metastatic marker for pancreatic cancer (Sadanandam *et al*, 2007). Semaphorin 5A is

*Correspondence: Dr A Sadanandam; E-mail: Anguraj.sadanandam@epfl.ch or Dr RK Singh; E-mail: rsingh@unmc.edu

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characterised by five thrombospondin-1-specific repeats in its extracellular domain (ECD) in addition to the semaphorin and plexin domains (Adams *et al*, 1996). Semaphorin 5A is known to induce inhibitory responses during optic nerve development (Oster *et al*, 2003) and also has major roles in innate immunity (Sugimoto *et al*, 2006) and idiopathic autism (Melin *et al*, 2006). For the first time, we observed the expression of SEMA5A in human pancreatic tumours, but not in normal pancreatic tissue. In addition, we showed that SEMA5A is expressed by pancreatic cancer cell lines established from metastases, but not those from primary tumours. We also demonstrated the membrane localisation of SEMA5A particularly to regions of cell–cell adhesion in aggressive pancreatic cancer cell lines. Overexpression of membrane-bound mouse Sema5A in pancreatic cancer cells has been demonstrated to increase migration and aggregation *in vitro* and tumorigenesis and metastasis *in vivo* (Sadanandam *et al*, 2010b). These results suggest that membrane-bound SEMA5A functions as a paracrine molecule involved in pancreatic tumour growth and metastasis. However, different reports on various cancer types either endogenously expressing SEMA5A or treated with the ECD of mouse Sema5A have contrasting tumour phenotypes (Li and Lee, 2010; Lu *et al*, 2010; Pan *et al*, 2010).

Recently, others and we have demonstrated the interaction of Sema5A with Plexin B3 in different mouse tissues and various cell lines (ectopically expressing Plexin B3; Artigiani *et al*, 2004; Sadanandam *et al*, 2008). Moreover, we have demonstrated that Sema5A is a proangiogenic molecule that regulates various steps of angiogenesis including endothelial cell proliferation, survival and migration *in vitro* and sprouting of blood vessels *in vivo* through its functional receptor, Plexin B3 (Sadanandam *et al*, 2010a). However, SEMA4D, a membrane-bound protein and a close relative with structural similarity to SEMA5A (Sadanandam *et al*, 2007), has been shown to be secreted (Wang *et al*, 2001) and can regulate tumour angiogenesis and metastasis (Basile *et al*, 2006). In this report, we demonstrate that pancreatic cancer cells secrete ECD of SEMA5A (SEMA5A-ECD) and that secreted Sema5A enhances invasiveness and metastasis of tumour cells via ERK phosphorylation and proliferation of endothelial cells through upregulation of angiogenic factors.

MATERIALS AND METHODS

Cell culture

Human pancreatic cancer cell lines (Panc1 and T3M4) and immortalised human dermal microvascular endothelial cells (HMEC-1; obtained from the Centre for Disease Control and Prevention, Atlanta, GA, USA; Ades *et al*, 1992) were maintained in culture as an adherent monolayer with RPMI-1640/DMEM with 5% Foetal Calf Serum (Mediatech, Herndon, VA, USA) supplemented with 1X nonessential amino acids, 2 mM L-glutamine and 1X vitamin solution.

Transfection of pancreatic cancer cells and purification of recombinant protein

Panc1 cells were stably transfected with either 1 μg of a cDNA construct containing the mouse Sema5A-ECD conjugated to human IgG Fc γ cloned in to the pcDNA3.1 vector (generous gift from Dr David Stretavan, University of California San Francisco at San Francisco, CA, USA) or pcDNA3.1 control vector (Stratagene, La Jolla, CA, USA) using LipofectAMINE (Invitrogen, Carlsbad, CA, USA) according to the protocol provided by the manufacturer. Panc1 cells transfected with Sema5A-ECD (Panc1-Sema5A-ECD) or control vector (Panc1-control) were selected and maintained with 400 $\mu\text{g ml}^{-1}$ G418 sulphate (Mediatech). Fc-tagged Sema5A-ECD was isolated by protein-A affinity chromatography from the supernatant of Panc1-Sema5A-ECD expressing cells and the protein purity

was tested using silver staining (Fischer Chemical, Fairlawn, NJ, USA) as described previously (Sadanandam *et al*, 2010a).

Tumorigenic and metastatic assays

The tumorigenic and metastatic assays were performed using male athymic BALB/c nude mice (NCI-nu, 6–8 weeks old, purchased from the National Cancer Institute). In accordance with approval from the American Association of Laboratory Animal Care and current regulations and standards of the US Department of Agriculture, Department of Health and Human Services and National Institute of Health, the mice used for this study were maintained under specific pathogen-free conditions in facilities. All procedures performed in mice were in accordance with the institutional guidelines and were approved by the University of Nebraska Medical Center Institution Animal Care and Use Committee.

For tumour growth and metastasis, the preparation of cells and the orthotopic injections in to the pancreas of the mice were performed as described in our previous study (Sadanandam *et al*, 2010b).

Immunohistochemistry (IHC)

Fixed paraffin-embedded tissue samples on slides were deparaffinised in a series of xylene and alcohol treatments (two times xylenes, one time each in 100%, 95%, 80% and 70% alcohol) for 5 min each. The slides were washed with PBS for 5 min for three times and quenched for endogenous peroxidase activity by incubating with 3% H₂O₂ in PBS for 5 min. Slides were again washed thoroughly with PBS and blocked with blocking buffer (10% horse/donkey serum) for 30 min. Without further washing, the samples were incubated with primary antibody (anti-PCNA (5 $\mu\text{g ml}^{-1}$; Santa Cruz Biotechnology, Santa Cruz, CA, USA), anti-SEMA5A (5 $\mu\text{g ml}^{-1}$; Sadanandam *et al*, 2010b), anti-interleukin (IL)-8 (2 $\mu\text{g ml}^{-1}$), anti-vascular endothelial growth factor (VEGF; 2 $\mu\text{g ml}^{-1}$; R&D System, Minneapolis, MN, USA) or anti-CD31 (1:100, Abcam, Cambridge, MA, USA) in blocking buffer for overnight at 4°C and then with biotinylated secondary antibody for 30 min to 1 h after washing with PBS. After washing the slides in PBS, the samples were incubated with ABC reagent (Vector Laboratories, Burlingame, CA, USA) for 15–30 min. Finally, DAB substrate (Vector Laboratories) was added until a brown colour developed and counter stained with hematoxylin. The quantitation of the number of cells stained for a particular antibody was determined by counting at least five different random fields in the same section at $\times 200$ resolution using a Nikon microscope (Nikon Inc., Melville, NY, USA). The average number of positive cells was calculated.

mRNA analysis

Total cellular RNA isolation and reverse transcription based polymerase chain reaction (RT-PCR) specific to SEMA5A and β -actin was performed as described (Sadanandam *et al*, 2010b).

In vitro proliferation assay

For endothelial cell proliferation, cells (1×10^3) were seeded in individual wells of 96-well flat-bottomed plates in triplicate and treated with serum-free supernatant from Panc1-Sema5A-ECD or Panc1-control cells. The proliferative activity of the cells after 72 h of incubation was measured using the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay using a microtiter plate reader (Bio-Tek Instruments, Inc., Winooski, VT, USA) at 570 nm as described (Sadanandam *et al*, 2010a).

Invasion and migration assay

For invasion assay, cells (1×10^5) were plated onto the upper chamber of transwell chambers and coated with matrigel (6.5 mm;

Corning Costar Corp., Cambridge, MA, USA) on the bottom of the upper chamber. MTT was added to the wells during the last 2 h of incubation. Using a cotton swab, the residual cells from the top of the transwell chambers were removed. The inserts were transferred to a carrier plate and dimethyl sulfoxide was added to solubilise the crystals. The migratory activity was measured by reading at 570 nm using a microplate reader.

For migration assay using MEK1 inhibitor (PD98059, Millipore, Billerica, MA, USA), cells (1×10^5) along with $10 \mu\text{M}$ MEK1 inhibitor or media alone were plated onto the upper chamber of transwell chambers and media on the bottom chamber. The cells were incubated for 4 h at 37°C and cells that did not pass through the membrane pores were removed using a cotton swab. Migrated cells were stained using the Hema 3 kit (Fisher Scientific Company L.L.C., Kalamazoo, MI, USA) as per the manufacturer's instructions. Cells were counted in 10 random fields ($\times 200$) and expressed as the average number of cells per field of view. The data are represented as the average of three independent experiments.

Immunoprecipitation and western blot analysis

Cells (1×10^6) were plated on 10-cm cell culture dishes and serum-starved for 24 h. Serum-free supernatants or conditioned media (CM) and lysates from T3M4 cells were collected, concentrated using Amicon Ultra-30 Centrifugal Filter Units (Millipore) and were directly subjected to western blot as described (Sadanandam *et al*, 2010b).

For immunoprecipitation, CM from Panc1-Sema5A-ECD or control cells were collected and were immunoprecipitated using Protein A/G (Invitrogen) and anti-SEMA5A antibody and incubated at 4°C overnight. Later, the immunoprecipitates were washed three times using RIPA buffer followed by SDS-PAGE and western blot using anti-SEMA5A antibody as described (Sadanandam *et al*, 2008).

For ERK phosphorylation, pancreatic cancer cells were seeded in 10-cm cell culture dishes. After the cells reached 80–90% confluence, cells were treated with Sema5A-ECD recombinant protein (10 ng ml^{-1}) for 5 or 30 min or media alone. Cells were washed three times with cold PBS and lysed in membrane lysis buffer and the lysates ($20 \mu\text{g}$ per lane) were separated on SDS-PAGE and subjected to western blot as described (Sadanandam *et al*, 2010b). Densitometric quantitation of intensity of the bands was performed using the ImageQuant (GE Healthcare, Piscataway, NJ, USA).

Enzyme-linked immunosorbent assay (ELISA)

IL-8 or VEGF protein levels in culture supernatants were determined using enzyme-linked immunosorbent assay (ELISA) matched-pair antibodies for IL-8 (Endogen Inc., Woburn, MA, USA) or VEGF (R&D Systems) according to the manufacturer's instructions with modification as described previously (Varney *et al*, 2003). Briefly, $50 \mu\text{l}$ of culture supernatant from treated or untreated cells was assayed in duplicate using the sandwich ELISA technique. A standard curve of the absorbance vs the concentration of protein (IL-8 or VEGF) in the standard wells was generated. We determined the concentration of protein in the unknown samples by comparing the absorbance of the sample to the standard curve.

Statistical analysis

The significance of the data was determined by the Student's *t*-test (two-tailed) for all the studies. $P < 0.05$ was deemed significant. All statistical analyses were done using the SPSS software (SPSS, Inc., Chicago, IL, USA).

RESULTS

Secretion of SEMA5A in pancreatic cancer cells

To examine whether pancreatic cancer cells secrete SEMA5A, we performed western blots for SEMA5A using serum-free supernatant (CM, see Materials and Methods for supernatant preparation) and protein lysates collected from T3M4, a pancreatic cancer cell line expressing high levels of SEMA5A (Sadanandam *et al*, 2010b). The results shown in Figure 1A demonstrate a band at $\sim 110 \text{ kD}$ in the T3M4 CM whereas the protein lysate shows bands at $\sim 135 \text{ kD}$ (membrane-bound SEMA5A) and $\sim 110 \text{ kD}$ (equivalent to the molecular weight of its extracellular domain). These results suggest that SEMA5A is secreted in pancreatic adenocarcinoma cell lines.

Overexpression of Sema5A-ECD in pancreatic cancer cells increases their invasive potential

As the $\sim 110\text{-kD}$ band in T3M4 supernatant (Figure 1A) is predicted to be equivalent to SEMA5A-ECD, we studied the role of SEMA5A-ECD in pancreatic cancer. We stably transfected Panc1 cells, which do not express SEMA5A, with a cDNA construct containing mouse Sema5A-ECD (Panc1-Sema5A-ECD) fused with human IgG Fc γ region at the C-terminus or with empty vector (Panc1-control). Panc1-Sema5A-ECD cells expressed Sema5A mRNA whereas Panc1-control cells did not show detectable levels of Sema5A mRNA (Figure 1Ba). In addition, we performed immunoprecipitation followed by western blot both using anti-SEMA5A antibody and supernatant from Panc1-Sema5A-ECD and Panc1-control. We observed a band between ~ 180 and $\sim 220 \text{ kD}$ that represents the Sema5A-ECD plus with human IgG Fc γ region (Figure 1Bb) similar to that shown by Oster *et al* (2003).

Next, we examined whether expression of Sema5A-ECD in pancreatic cancer cells modulates their invasive potential. Significantly more Panc1-Sema5A-ECD cells invaded through Matrigel as compared with control cells ($P < 0.05$) in the absence of serum (Figure 1C). This result demonstrates that Sema5A-ECD modulates the invasive potential of pancreatic cancer cells.

Expression and phosphorylation of ERK in Sema5A treated Panc1 cells

Downregulation of *D-Sema5c* has been shown to decrease phosphorylation of ERK in a *Drosophila* model for metastasis (Woodhouse *et al*, 2003). To identify a signalling network mediating migration of pancreatic cancer cells by secreted SEMA5A, we examined ERK phosphorylation after treatment with recombinant Sema5A-ECD. Increased levels of p42 ERK phosphorylation were observed in Panc1 cells treated with 10 ng ml^{-1} of Sema5A-ECD for 5 and 30 min compared with cells treated with media alone (Figures 1D and E) as well as a slight increase in the level of p44 ERK phosphorylation (Figure 1D and Supplementary Figure 1A). To verify if ERK signalling is important for SEMA5A-ECD induced migration of pancreatic cancer cells, we performed transwell chamber migration assays by treating Panc1-Sema5A-ECD and Panc1-control cells with $10 \mu\text{M}$ concentration of MEK1 inhibitor (PD98059) or media alone. A significant decrease in migration of Panc1-Sema5A-ECD cells with MEK1 inhibitor was observed compared with the treatment with media alone whereas there was not much change in the Panc1-control cells (Figure 1F). This data supports the hypothesis that Sema5A-dependent cellular migration is mediated through the increased ERK phosphorylation.

Sema5A-ECD overexpression enhances spontaneous metastasis

To examine the role of Sema5A-ECD in tumour cells in an orthotopic pancreatic microenvironment, we injected

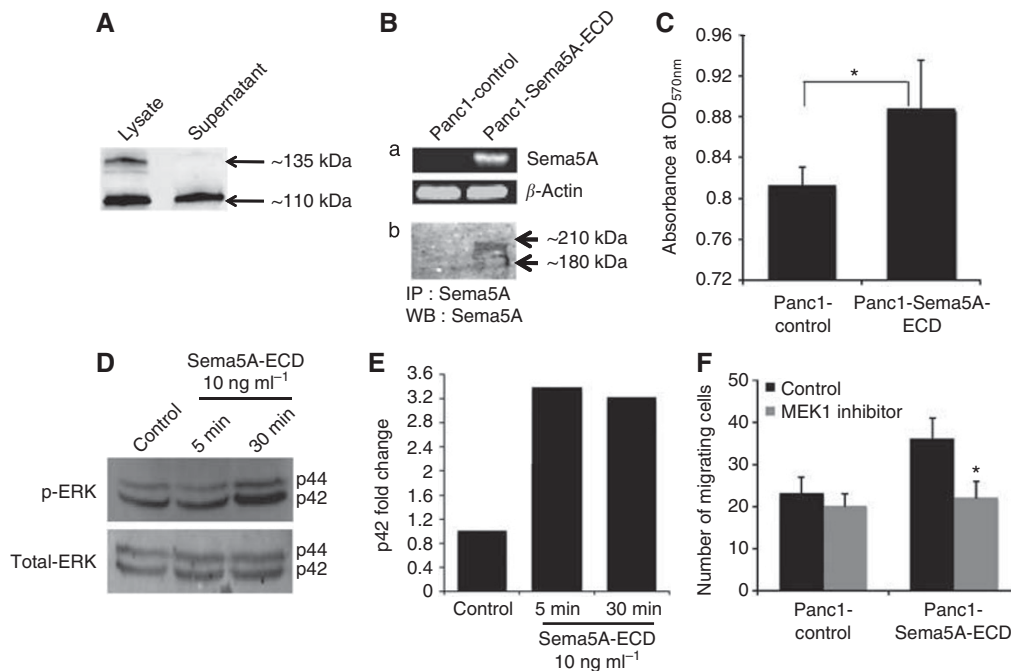


Figure 1 Secretion of SEMA5A and its invasive role in pancreatic cancer cells. **(A)** A representative western blot analysis showing expression of membrane-bound SEMA5A in protein lysates and secretion of SEMA5A in supernatant of human pancreatic cancer cell line – T3M4. **(B)** A standard (a) RT-PCR (using specific primers) and (b) immunoprecipitation and western blot analysis (using anti-SEMA5A antibody) of Panc1 cells ectopically expressing mouse Sema5A-ECD (Panc1-Sema5A-ECD) and control vector (Panc1-control) showing expression of Sema5A. β -Actin was used as an internal control for RT-PCR. **(C)** Cellular invasion in Panc1-Sema5A-ECD or Panc1-control after 1 h was measured using Matrigel-coated transwell chambers and MTT assay. Invasion was measured as mean optical density (OD) at 570 nm (OD_{570nm}) \pm standard deviation (s.d., bars) of triplicate culture and *a statistical significance of $P < 0.05$. **(D)** Western blot analysis of phosphorylated and total ERK in protein lysates of Panc1 cells treated with 10 ng ml^{-1} of recombinant Sema5A-ECD protein at different time points (5 or 30 min) or media alone using appropriate antibodies. **(E)** Densitometric quantitation of phosphorylated p44 ERK that was normalised to total p44 ERK from western blots from **(D)** using the ImageQuant software. **(F)** Cellular migration in Panc1-Sema5A and Panc1-control after 4 h was measured after treating the cells either with $10 \mu\text{M}$ MEK1 inhibitor (PD98059) or media alone (control) using uncoated transwell chambers and the number of migrated cells was counted at $\times 10$ magnification using a Nikon microscope. The values are number of migrated cells \pm s.d. (bars) of triplicate culture and *a statistical significance of $P < 0.05$ using Student *t*-test.

Panc1-Sema5A-ECD or Panc1-control (5×10^5) cells into the pancreas of athymic nude mice. Mice were killed after 10 weeks. The presence of primary tumour in the pancreas and metastatic cells in different organs was confirmed by H&E staining (data not shown). Surprisingly, no difference in tumour incidence was observed between mice injected with Panc1-Sema5A-ECD or Panc1-control cells (Figure 2A). However, in mice injected with Panc1-Sema5A-ECD cells, tumour burden was lower and yet there was an increased incidence of micrometastases to the liver (Figure 2A). Though there were no obvious macrometastases in either Panc1-Sema5A-ECD mice or control mice, there was a significant increase in number of micrometastatic cells to the liver (Figure 2B). To examine the proliferation of orthotopic tumours, we examined PCNA staining in Panc1-Sema5A-ECD and control tumours by IHC analysis using an anti-PCNA antibody and then counting the number of positive cells. No significant difference in PCNA staining was observed between Panc1-Sema5A-ECD and control tumours (Supplementary Figure 1B).

Next, we performed IHC for CD31 marker for microvessels in orthotopic tumours. Here, we identified a significant increase in microvessel density in orthotopic tumours expressing Sema5A-ECD as compared with control tumours (Figures 2C and D). These data suggest that Sema5A-ECD stimulates aggressiveness in pancreatic tumour cells leading to increased micrometastases via increased tumour angiogenesis, but with lower primary tumour burden.

Sema5A-ECD increases endothelial cell proliferation and upregulates expression of pro-angiogenic factors

To investigate whether Sema5A-ECD mediates increased metastasis by increasing endothelial cell proliferation (one of the steps involved in tumour angiogenesis), we performed MTT proliferation assays after treating HMEC-1 endothelial cells with conditioned serum-free supernatants from Panc1-Sema5A-ECD or Panc1-control cells (Figure 3A). Significantly increased proliferation was observed in HMEC-1 cells treated with serum-free conditioned supernatant from Panc1-Sema5A-ECD as compared with that from Panc1-control cells (Figure 3A). This result demonstrates that Sema5A-ECD increases proliferation of endothelial cells.

Next, we examined the expression of well-known pro-angiogenic regulators IL-8 and VEGF in the serum-free supernatants of Panc1-Sema5A-ECD or control cells *in vitro* by ELISA. Interleukin-8 and VEGF were significantly increased in the supernatant of Panc1-Sema5A-ECD cells as compared with control cells (Figures 3B and C). We further validated the regulation of IL-8 and VEGF expression by Sema5A-ECD by treating Panc1-Sema5A-ECD cells with neutralising antibody to Sema5A for 72 h and then measuring the expression levels of IL-8 and VEGF in the supernatant using ELISA. A significant decrease ($P < 0.05$) in IL-8 and VEGF protein expression was seen in the supernatant of Panc1-Sema5A-ECD treated with 10 ng of neutralising antibody to Sema5A (Figures 3D and E). Upregulation of IL-8 expression was also seen in

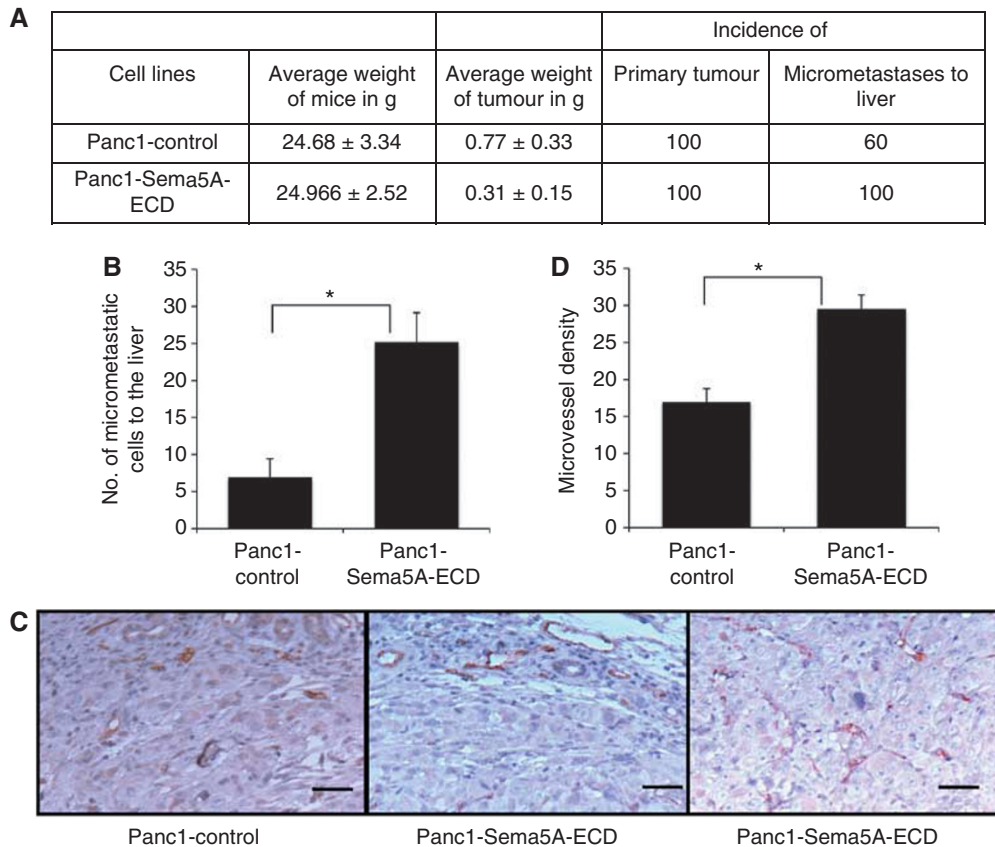


Figure 2 Orthotopic assay, micrometastasis and microvessel density of Panc-1 Sema5A-ECD. Athymic nude mice ($n = 5$) were injected orthotopically with Panc1-control or Panc1-Sema5A-ECD and were euthanised after 10 weeks. **(A)** Average weight of mice and tumour (tumour burden) as well as incidence of primary tumour and metastasis to the liver is shown. **(B)** There are a significantly enhanced number of micrometastatic cells per field of $\times 200$ resolution of the Nikon microscope in the liver in Panc1-Sema5A-ECD tumour-bearing mice as compared to control mice. **(C)** Immunohistochemistry using CD31 staining showing microvessels in Panc1-Sema5A-ECD and control orthotopic tumours. There is an increased angiogenesis in Panc1-Sema5A-ECD compared to the Panc1-control orthotopic tumours. **(D)** Densitometric quantitation of microvessel density from C, per field at $\times 200$ resolution of the Nikon microscope. The values are number of microvessels \pm s.d. (bars) of five different areas per field or different slides and *a statistical significance of $P < 0.05$.

Panc1-Sema5A-ECD orthotopic tumours as compared with control tumours by IHC ($P < 0.05$; Figures 3F and G). However, we did not observe a significant change in VEGF staining between orthotopic tumours from Panc1-Sema5A-ECD and Panc1-control (Supplementary Figure 1C). These results demonstrate that Sema5A-ECD increases the protein expression of IL-8 and to a certain extent VEGF in pancreatic cancer cells and tumours.

DISCUSSION

Previously, we found that pancreatic tumours and aggressive pancreatic cancer cell lines express membrane-bound SEMA5A, however, it is not expressed in normal pancreatic tissues or less aggressive pancreatic cancer cell lines (Sadanandam *et al*, 2007, 2010b). In this study, we sought to determine if pancreatic cancer cells secrete SEMA5A and whether secreted SEMA5A has a role in pancreatic tumour progression, metastasis and angiogenesis by regulating other molecules.

Many secreted forms of membrane-associated proteins such as CD44 (Stickeler *et al*, 2000) and transforming growth factor- β type III receptor (Bandyopadhyay *et al*, 2002) were reported to be involved in tumour metastasis. One transmembrane protein of the semaphorin family, SEMA4D, which displays structural similarity to SEMA5A (Sadanandam *et al*, 2007), has also been reported to be released as a functional secreted protein and has been shown to be

involved in tumorigenesis and metastasis (Wang *et al*, 2001; Basile *et al*, 2004, 2006). In this report, we detected SEMA5A protein in the serum-free supernatant of a human pancreatic cancer cell line (T3M4) and the ~ 110 -kD band was predicted to be similar to that of the Sema5A-ECD. To our knowledge, this is the first report showing that SEMA5A is secreted.

In our previous study, we demonstrated that membrane-bound Sema5A increases pancreatic cancer invasion *in vitro* and metastasis *in vivo* (Sadanandam *et al*, 2010b). In this report, we studied the role of secreted SEMA5A in pancreatic tumour progression and metastasis by overexpressing mouse Sema5A-ECD in Panc1 cells not expressing SEMA5A. Sema5A-ECD expression in pancreatic cancer cells demonstrated increased invasion *in vitro*. This increase in invasion of pancreatic cancer cells induced by Sema5A-ECD especially in serum-free conditions suggests that this molecule is a motogenic factor secreted by tumour cells.

Previously, we have shown that orthotopic injections of pancreatic cancer cells overexpressing membrane-bound Sema5A in athymic nude mice are tumorigenic and metastatic (Sadanandam *et al*, 2010b). Here, we demonstrate that orthotopic injection of Sema5A-ECD-expressing cells resulted in reduced tumour burden, but enhanced micrometastases to the liver as compared with control transfected cells. It has already been reported that small primary tumours could lead to metastasis in different cancer types (Menz *et al*, 1985; Sho *et al*, 1998; Chiang

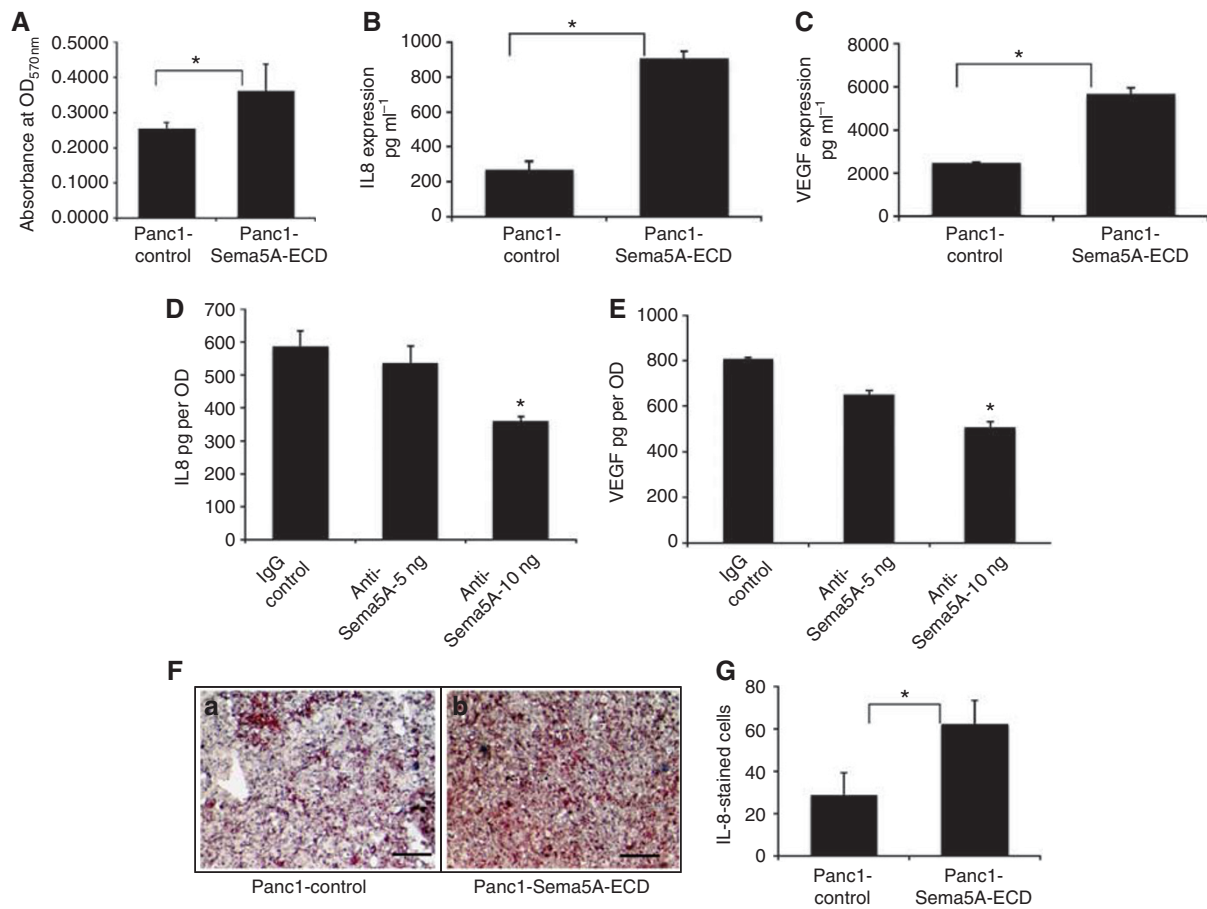


Figure 3 Sema5A-ECD stimulates endothelial cell proliferation and increases the expression of angiogenic factors. **(A)** MTT proliferation assay after 72 h using 1000 HMEC-1 cells per well that were treated with conditioned medium (no serum) from Panc1-Sema5A-ECD or Panc1-control. There is a significant ($P < 0.05$) increase in proliferation of HMEC-1 in the presence of supernatant from Panc1-Sema5A-ECD cells compared with control cells with no serum. Proliferation was measured as mean optical density (OD) at 570 nm ($OD_{570\text{nm}}$) \pm s.d. (bars) of triplicate culture and *statistical significance of $P < 0.05$. Enzyme-linked immunosorbent assay for **(B)** IL-8 and **(C)** VEGF protein in the supernatant of Panc1-Sema5A-ECD and Panc1-control cells using appropriate antibodies. There is increased expression of IL-8 and VEGF in the supernatants of Panc1-Sema5A-ECD compared with control. Data are mean pg mg^{-1} \pm s.e.m. (bars) of triplicate samples and *statistical significance of $P < 0.05$. Neutralisation of Sema5A-ECD in supernatants of Panc1-Sema5A-ECD using 10 ng ml^{-1} of anti-Sema5A or rabbit-IgG antibody followed by ELISA for **(D)** IL-8 or **(E)** VEGF. There is a significant decrease in IL-8 and VEGF using 10 ng ml^{-1} of anti-Sema5A antibody compared with rabbit-IgG. Data are mean pg/OD \pm s.d. (bars) of triplicate samples and *statistical significance of $P < 0.05$. **(F)** Immunohistochemistry showing IL-8 staining in (a) Panc1-control and (b) Panc1-Sema5A-ECD orthotopic tumours using anti-IL-8 antibody. The magnification is $\times 200$ resolution using a Nikon microscope. **(G)** The number of cells positive for IL-8 from **F**. The values are mean number of cells \pm s.d. (bars) of three slides. *Statistical significant difference of $P < 0.05$.

et al, 2008). The mechanism by which secreted Sema5A induces increased metastasis with reduced tumour burden is intriguing. Based on our previous findings that Panc1 cells express Plexin B3, a functional receptor for SEMA5A (Sadanandam *et al*, 2008), we suggest that the secretion of SEMA5A in pancreatic cancer induces an autocrine mechanism via the ERK phosphorylation to metastasise to distant organs but with smaller primary tumour, unravelling this cell signalling mechanism might be the basis of our future studies.

We have previously shown that Sema5A can promote angiogenesis, in general (Sadanandam *et al*, 2010a). In this study, we show that Panc1-Sema5A-ECD is biologically active and also has the ability to promote tumour angiogenesis as shown by its ability to significantly increase *in vitro* proliferation of HMEC-1 endothelial cells and significantly increase microvessel density in orthotopic Panc1-Sema5A-ECD tumours compared with control tumours. In addition, supernatants from Panc1-Sema5A-ECD cells show increased expression of pro-angiogenic factors, IL-8 and VEGF.

We speculate that secreted Sema5A derived from the membrane-bound protein could be the reason for the increased IL-8 and VEGF expression. The direct effect of Sema5A-ECD on endothelial cells to secrete IL-8 and VEGF was shown using neutralising antibody against Sema5A in endothelial cells. Overall, the increase in micrometastasis in Sema5A-ECD-expressing pancreatic cancer cells is due to increased tumour angiogenesis via IL-8 and VEGF.

In conclusion, in this study, for the first time we show that malignant pancreatic cancer cells secrete SEMA5A and that this secreted form demonstrates the ability to drive micrometastatic spread via the enhancement of angiogenesis and migration of pancreatic cancer cells via increased ERK signalling and expression of angiogenic factors, but with reduced primary tumour burden. These results suggest a potentially important and hitherto unsuspected role of the secreted form of SEMA5A in pancreatic cancer metastasis. Further study is warranted to examine the presence of secreted SEMA5A in pancreatic cancer patients and to understand the autocrine mechanism of its action.

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