CD44-related Chondroitin Sulfate Proteoglycan, a Cell Surface Receptor Implicated with Tumor Cell Invasion, Mediates Endothelial Cell Migration on Fibrinogen and Invasion into a Fibrin Matrix

Craig A. Henke,* Urvashi Roongta,* Daniel J. Mickelson,‡ Jennifer R. Knutson,‡ and James B. McCarthy‡
Departments of *Medicine and ‡Laboratory Medicine and Pathology, University of Minnesota, Minneapolis, Minnesota 55455

Abstract

Microvascular endothelial cell invasion into the fibrin provisional matrix is an integral component of angiogenesis during wound repair. Cell surface receptors which interact with extracellular matrix proteins participate in cell migration and invasion. Malignant cells use CD44-related chondroitin sulfate proteoglycan (CSPG) as a matrix receptor to mediate migration and invasion. In this study, we examine whether cell surface CSPG can mediate similar events in nonmalignant wound microvascular endothelial cells or whether use of CSPG for migration and invasion is a property largely restricted to malignant cells. After inhibiting CSPG synthesis with p-nitrophenyl β-d xylopyranoside (β-d xyloside), wound microvascular endothelial cells were capable of attaching and spreading on the surface of a fibrin gel; however, their ability to invade the fibrin matrix was virtually eliminated. To begin to examine the mechanism by which endothelial cells use CSPG to invade fibrin matrices, cell adhesion and migration on fibrinogen was examined. Endothelial cell adhesion and migration on fibrinogen were inhibited by both β-d xyloside and after cleavage of chondroitin sulfate from the core protein by chondroitinase ABC. We have determined that wound microvascular endothelial cells express the majority of their proteoglycan as CSPG and that the CSPG core protein is immunologically related to CD44. PCR studies show that these cells express both the “standard” (CD44H) isoform and an isoform containing the variably spliced exon V3. In addition, anti-CD44 antibody blocks endothelial cell migration on fibrinogen. Affinity chromatography studies reveal that partially purified microvascular endothelial cell CSPG binds fibrinogen. These findings suggest that CD44-related CSPG, a molecule implicated in the invasive behavior of tumor cells, is capable of binding fibrinogen/fibrin, thereby mediating endothelial cell migration and invasion into the fibrin provisional matrix during wound repair. (J. Clin. Invest. 1996, 97:2541–2552.) Key words: wound healing • angiogenesis • cell movement • cell surface receptor • CD44-related chondroitin sulfate proteoglycan

Introduction

After tissue injury, plasma fibrinogen extravasates from blood vessels into the extravascular space where it becomes clotted to cross-linked fibrin forming a provisional matrix in the wound (1). In response to migration promoting signals present within the provisional matrix (2, 3), microvascular endothelial cells invade into the fibrin matrix and form an extensive capillary network, providing an integral component of granulation tissue during wound healing (4, 5). The initiation of endothelial cell invasion involves a series of interdependent events, including detachment from the physiological matrix, migration, and attachment to the provisional matrix comprised of fibrin and other plasma-derived components.

Endothelial cells adhere and migrate on fibrin and its precursor fibrinogen via cell surface receptors that interact with distinct domains on extracellular matrix (ECM) proteins. One well-characterized mechanism of cell adhesion to ECM proteins involves integrins. Endothelial cells express αβ integrin which binds fibrinogen via an RGD dependent mechanism (6–9). Despite the recognized importance of integrins in mediating cell adhesion, there are additional cell surface molecule(s) that play an important role in mediating endothelial cell migration and invasion. For example, there is increasing evidence that endothelial cell migration can be associated with the expression of distinct cell surface matrix receptors that mediate migration (10). Additionally, wounded large vessel endothelial cells in culture rapidly change proteoglycan synthesis from heparan sulfate to chondroitin sulfate as migration is initiated, linking chondroitin sulfate expression with the process of cell motility (11).

Studies evaluating tumor cell motility link cell surface chondroitin sulfate proteoglycan (CSPG) with migration and invasion. Malignant cells synthesize increased levels of CSPG which can be found on microspikes on the cell surface (12). Microspikes are a specific cell surface microdomain believed to facilitate the initial contact of the cell with the ECM. Malignant cell surface CSPG, which contains a core protein recognized by CD44 antibodies, plays an important role in mediating melanoma cell motility and invasion into type I collagen gels (13). Although the mechanism by which cell surface CD44-related CSPG mediates tumor cell migration and invasion remains to be elucidated, CD44 has been implicated in mediating cell–cell and cell–ECM interactions and is associated with the cell cytoskeleton (14, 15) and plays a role in cell migration (16). CD44 is a transmembrane glycoprotein with extracellular, membrane, and cytoplasmic domains (17). The

Abbreviations used in this paper: α-d xyloside; p-nitrophenyl α-d xylopyranoside; β-d xylose, p-nitrophenyl β-d xylopyranoside; CSPG, chondroitin sulfate proteoglycan; ECM, extracellular matrix; GAG, glycosaminoglycan.
extracellular domain of CD44 is capable of binding a variety of ECM molecules (18) and contains attachment sites for chondroitin sulfate (19–21). Studies using tumor cells link CD44 with cell motility (22). Importantly, the expression of specific splice variants of CD44 (23, 24) and the overexpression of CD44 cDNA clones in tumor cells (25) have been associated with tumor growth and metastasis.

The present study intends to further examine the role of cell surface CSPG in mediating microvascular endothelial cell adhesion, migration, and invasion into the fibrin provisional matrix. To determine whether CSPG influences endothelial cell adhesion, migration, and invasion on fibrinogen/fibrin, we examined the effect of β-xylosidase (β-xyloside), an agent that competitively inhibits the coupling of chondroitin sulfate to the core protein (26), and chondroitinase ABC, an agent that cleaves the chondroitin sulfate from the core protein, using standard adhesion, migration, and invasion assays. Our results show that β-xyloside inhibits the invasion of microvascular endothelial cells into three-dimensional fibrin gels. Furthermore, haptotactic migration and adhesion of endothelial cells on fibrinogen was inhibited by both β-xyloside and chondroitinase ABC. We have determined that wound microvascular endothelial cells express the majority of their proteoglycan as CSPG and that the CSPG core protein is immunologically related to CD44. PCR studies reveal that wound microvascular endothelial cells express both the “standard” (CD44H) isoform and an isoform containing the variably spliced exon V3. Thus, CD44-related CSPG, a molecule implicated in the invasive behavior of tumor cells, may play a role in mediating endothelial cell migration and invasion into the fibrin provisional matrix during wound healing.

**Methods**

**Cell culture and endothelial cell characterization.** Wound microvascular endothelial cells were isolated from sponges implanted into the backs of rabbits, as previously described (2). The cells were identified as endothelial cells by positive staining for vWF, angiotensin-convert ing enzyme, and Factor VIII–related antigen, and by selective uptake of acetylated LDL. The microvascular endothelial cells were cultured in M199 (Gibco Laboratories, Grand Island, NY) supplemented with 20% heat-inactivated FCS (Sigma Chemical Co., St. Louis, MO), and subcultivated weekly at a split ratio of 1:2. Wound microvascular endothelial cells were used for subsequent experiments at 50–75% of their saturation density unless otherwise specified, and before the 10th subcultivation.

**Cell adhesion and migration assays.** Cell adhesion to protein coated substrata was performed as previously described (2, 27) using 0.005–5 μM fibrinogen. Endothelial cell cultures were radiolabeled (18 h) with [3H]thymidine (2 μCi/ml), released with enzyme solution (0.5% collagenase, 0.2% DNase I in PBS, pH 7.4), washed, and resuspended to a final concentration of 1 × 10^6 cells/ml in DME, 20 mM Hepes, and 5 mM/ml BSA. Microvascular endothelial cells were released with enzyme solution (0.5% collagenase, 0.2% DNase I in PBS, pH 7.4), washed, and resuspended to a final concentration of 1 × 10^6 cells/ml in DME, 20 mM Hepes, and 5 mM/ml BSA. Cells were added to the upper wells at 4.5 × 10^4 cells/well and migration to the underside of the precoated filter was measured after 4 h at 37°C.

**Invasion gels.** Gels composed of fibrin were prepared under sterile conditions using a modification of the protocol described by Dvorak et al. (4). Human thrombin (0.2 U/ml; Sigma Chemical Co.) was added to HPLC-DEAE purified rabbit fibrinogen (3 mg/ml; Sigma Chemical Co.) dissolved in DME containing 20 mM Hepes, pH 7.4, and 0.5 ml of this solution was added to chambers (Falcon 24-well plates; Becton Dickinson Labware, Lincoln Park, N.J.) before polymerization. The solution was gently swirled to allow uniform distribution and allowed to solidify (30 min, 37°C). Cultures of wound microvascular endothelial cells were seeded with enzyme solution (0.5% collagenase, 0.2% DNase I in PBS, pH 7.4), washed, and resuspended in DME containing 2.5% heat-inactivated calf serum. A cell suspension containing 5 × 10^4 cells was added to the top of the fibrin gels and the plates were gently swirled to obtain an even distribution of the cells on the gel surface. The gels were incubated (37°C, 5% CO_2) in a humidified incubator with media replenished three times per wk.

Cell invasion was measured by quantifying the number of tubular networks formed by invading microvascular endothelial cells using an inverted phase contrast microscope (IMT-2; Olympus Corp., Lake Success, NY). By focusing into the gel, invading cells forming tubular networks could be clearly identified by their position below the original starting plane containing the monolayer of cells on the top surface of the gel. Quantification of the number of tubular networks formed by invading microvascular endothelial cells was performed on days 4, 7, 10, and 14. Data are presented as the total number of tubular networks formed by invading cells per gel. Results are reported as the means ± SEM of quadruplicate gels.

**β-xyloside and chondroitinase ABC pretreatment.** For adhesion, migration, and invasion assays, cells were treated with β-xyloside (p-nitrophenyl-β-xylopyranoside; Sigma Chemical Co.) for 48 h to inhibit CSPG synthesis by replacing the medium of cell cultures with DME containing 2.5% heat-inactivated calf serum, with or without β-xyloside. α-xyloside (p-nitrophenyl-α-xylopyranoside; Koch-Light Ltd., Suffolk, England) does not inhibit CSPG synthesis and was used as a control. α- or β-xyloside was also included in the medium during the assays. For invasion assays, cells were seeded onto the gels as described above in the presence of varying concentrations of α- or β-xyloside. Additionally, fibrin gels used in the α- and β-xyloside experiments were composed of fibrinogen dissolved in DME, 20 mM Hepes, containing varying concentrations of either α- or β-xyloside.

For chondroitinase ABC treatment, cells were pretreated with varying concentrations of chondroitinase ABC (15 min, 37°C; Sigma Chemical Co.) before adhesion and migration assays as previously described (13). Chondroitinase ABC was also included in the cell medium during the adhesion and migration assays.

**Inhibition assay: anti-CD44 mAb.** An inhibition bioassay using a mouse anti-human CD44 mAb (BU52; The Binding Site, Inc., San Diego, CA) was performed to assess the role of CD44 in mediating microvascular endothelial migration on fibrinogen. Briefly, for the migration assay, microvascular endothelial cells were preincubated (30 min, 37°C) with the anti-CD44 mAb (0.02–20 μg/ml) and added to the upper wells of the modified Boyden chambers. Migration to the underside of the precoated filter was measured after 4 h at 37°C. Control assays were performed substituting normal mouse IgG antibody for the anti-CD44 mAb.

**Extraction and purification of microvascular endothelial cell proteoglycans.** Microvascular endothelial cell proteoglycans were preferentially labeled with [35S]sulfate (18 h, 37°C) by replacing the medium in cultures with low sulfate DME (0.1 mM Na_2SO_4, Gibco Laborato-
ries) containing 50 μCi/ml Na\(^{35}\)SO\(_4\) (Carrier free, sp act 43 Ci/mg; ICN Biomedicals, Irvine, CA). Proteoglycans were extracted by adding 30 ml cellular extraction buffer (0.15 M NaCl, 10 mM Tris, 5 mM MgCl\(_2\), 2 mM EDTA, 0.25 mM DTT, 1 mM PMSF, 1% Triton X-100, pH 7.2) to each culture flask (75 cm\(^2\); Corning Glass Works, Corning, NY) and incubated (20 min, 37°C). The cellular extracts were centrifuged (1,500 rpm, 5 min) to remove insoluble material. The extracts were dialyzed (mol wt cut off 3,500) against successive changes of acetic buffer (0.5 M C\(_6\)H\(_5\)O\(_2\)Na, pH 6.8, 0.1 Na\(_2\)SO\(_4\), 10 mM EDTA, 0.1 mM PMSF, 10 mM 6-amino-hexanoic acid) until the amount of radioactivity in the dialysate buffer was at background levels (13).

Anion exchange chromatography of detergent-extracted CSPG. Detergent-extracted \(^{35}\)S-proteoglycans were dialyzed into DEAE buffer (0.15 M Tris, 6.0 M urea, 0.1 M NaCl, 0.01 M EDTA, pH 7.0) and purified by HPLC (Model 110 A; Beckman Instruments) with a buffer (0.15 M Tris, 6.0 M urea, 0.1 M NaCl, 0.01 M EDTA, 0.01 M Na\(_2\)SO\(_4\), 10 mM EDTA, 0.1 mM PMSF, 10 mM 6-amino-hexanoic acid) until the amount of radioactivity in the dialysate buffer was at background levels (13).

Characterization of glycosaminoglycan chains. \(^{35}\)S-glycosaminoglycans (GAG) were released from proteoglycan core protein by alkaline borohydride reduction, neutralized, and desalted on Sephadex G-50 (Sigma Chemical Co.) columns as previously described (13). Under these conditions, greater than 95% of added GAGs were recovered from the V\(_c\) fractions. The heparan sulfate and chondroitin sulfate content of the recovered GAGs were determined by sequential nitrous acid deaminative cleavage (22

Immunoprecipitation and SDS-PAGE. Protein after chondroitinase ABC digestion. Gel sample buffer was

PCR reactions were carried out in a total vol of 100 μl containing 20 μl of cdNA, 8 μl of each 10 mM dNTP’s, 1 μl of 20 pmol/μl of each primer, 8 μl 10× PCR buffer, and 2.5 U of Taq polymerase (Fisher Scientific Co., Pittsburgh, PA). Primers were added when the reaction mixture was at 95°C. The cdNA was amplified in 35 cycles. The PCR amplification program was as follows: denaturation (0.5 min, 95°C), annealing (1.5 min, 56°C), extension (1 min, 73°C), and enzyme inactivated (1 min, 73°C). The oligonucleotides used as PCR primers were the CD4E3-3P and the CD4E16-RP primers as described by Bennett et al. (29). The CD4A variant exon primers were the p2v through p10 primers as previously described by van Weering et al. (30).

Fibrinogen affinity chromatography. For affinity chromatography, HPLC-DEAE purified fibrinogen was covalently coupled to Reacti-Gel according to manufacturers instructions (Pierce Chemical Co., Rockford, IL). Columns (30-mI bed vol) were equilibrated with 50 mM Tris, pH 6.8, 0.5% CHAPS, 0.05 M NaCl, 0.01 M 6-amino-hexanoic acid, 0.1 mM PMSF, 1.0 mM NEM, and 0.02% azide. Detergent-extracted HPLC-DEAE partially purified \(^{35}\)S-CSPG was applied to the fibrinogen affinity column, washed, and eluted with a linear salt gradient (0.05–1.0 M NaCl). To assure that the binding of CSPG was specific to fibrinogen. \(^{35}\)S-CSPG was applied to a duplicate column prepared without fibrinogen. Recovery of radioactivity from this column was > 95%.

Statistical analysis. All data were expressed as mean±SEM performed in triplicate unless otherwise indicated.

Results

Cell surface CSPG mediates microvascular endothelial cell invasion into a fibrin matrix. To examine the role of cell surface CSPG in mediating microvascular endothelial cell invasion, we used three-dimensional fibrin gels, using a modification of the technique described by Dvorak et al. (5). Microvascular endothelial cells layered on to the top surface of the fibrin formed a monolayer. After 4–7 d, cells invaded below the plane of the monolayer and formed extensive tubular networks within the fibrin matrix (Fig. 1 A). Microscopic sections of the tubular networks within the fibrin gel revealed that the cells formed vascular-like structures containing lumens (Fig. 1 B).

To evaluate the potential role of cell surface CSPG in mediating cell invasion into fibrin matrices, microvascular endothelial cell invasion was observed in the presence of 1.0 mM β-d xyloside. Microvascular endothelial cells cultured in the presence of β-d xyloside attached to the substratum and spread, forming monolayers on the surface of the fibrin gel. Analysis of their ability to invade the gel revealed that this function was
inhibited by ~95% at day 14 of gel culture (Fig. 2A). Wound microvascular endothelial cell invasion into fibrin was inhibited by β-D xyloside in a concentration dependent fashion (Fig. 2B). No inhibition of endothelial cell invasion was observed in the presence of α-D xyloside, an inactive analogue of β-D xyloside which does not affect CSPG synthesis (Fig. 2A). Additionally, endothelial cell invasion into fibrin gels made from HPLC-DEAE purified fibrinogen and into hyaluronidase treated gels were also inhibited by β-D xyloside. These studies were performed to ensure that disruption of endothelial cell interaction with fibrin and not other ECM proteins contaminating the fibrin gels was the mechanism by which β-D xyloside inhibited invasion (data not shown).

As a control to determine whether β-D xyloside treatment was toxic to microvascular endothelial cells, the cells were pre-incubated with 1.0 mM β-D xyloside for 48 h, enzyme released, washed, and layered onto the fibrin gels in the absence of additional β-D xyloside. The number of tubular networks formed by invading endothelial cells that had been exposed to β-D xyloside followed by withdrawal of the agent was actually increased compared to cells that had not been exposed to this drug (150±28 vs. 80±10). Additionally, β-D xyloside treated cells used for invasion assays were determined to be greater than 95% viable by trypan blue exclusion.

Cell surface CSPG mediates microvascular endothelial cell adhesion and migration to fibrinogen. Microvascular endothelial cells pretreated with β-D xyloside exhibited a 43% reduction in adhesion to fibrinogen as compared to untreated cells (Fig. 3A). α-D xyloside, which does not affect CSPG synthesis, had no affect on endothelial cell adhesion. Endothelial cell ad-
Proteoglycan in Endothelial Cell Motility

Adhesion was inhibited by β-D xyloside in a concentration-dependent fashion. To further investigate the role of endothelial cell surface CSPG in mediating cell adhesion, we measured microvascular endothelial cell adhesion after pretreatment with chondroitinase ABC, which cleaves the chondroitin sulfate sidechains from the core protein. Endothelial cell adhesion to fibrinogen after this treatment was inhibited by 33% (Fig. 3B).

To examine the role of cell surface CSPG in endothelial cell migration, the effects of β-D xyloside and chondroitinase ABC on microvascular endothelial cell motility were evaluated using the modified Boyden chamber. Endothelial cell motility on filters precoated with fibrinogen was inhibited by 65% in the presence of 1.0 mM β-D xyloside (Fig. 4A) and not affected by the inactive analogue, α-D xyloside. Cell migration after treatment with chondroitinase ABC (0.1 U/ml) was decreased by 70% on fibrinogen (Fig. 4B). Both β-D xyloside and chondroitinase ABC inhibited endothelial cell migration in a concentration dependent-fashion. These data suggest an important role for cell surface CSPG in mediating microvascular endothelial cell motility on fibrinogen.

Wound microvascular endothelial cell expression of CSPG. Based on the results of the functional assays implicating cell surface CSPG in endothelial cell adhesion and migration on fibrinogen and invasion into a fibrin matrix, wound microvascular endothelial cell proteoglycans were isolated and partially characterized. Proteoglycans were extracted from both wound microvascular endothelial cells in log phase growth (50–75% confluent) and from cells confluent for 3 d to determine if the expression of proteoglycans were altered when cells became confluent.

Detergent extracts of 35SO4-labeled cell cultures were chromatographed twice by HPLC-DEAE column chromatography. One predominant peak was obtained from both confluent and log phase wound microvascular endothelial cells which...
eluted from the HPLC-DEAE column at 0.38 M NaCl (Fig. 5 A). Fractions from this peak were pooled and rechromatographed on HPLC-DEAE. $^{35}$S-CSPG eluted at 0.38 M NaCl and contained 80–90% chondroitin sulfate as determined by chondroitinase ABC sensitivity and nitrous acid resistance and 10–20% heparan sulfate as determined by nitrous acid sensitivity and chondroitinase ABC resistance (Fig. 5 B–D). The remaining minor peak eluting at 0.2 M NaCl contained $^{35}$S-glycoproteins and was not characterized further. These data indicate that CSPG is the predominant proteoglycan isolated from wound microvascular endothelial cells.

**Microvascular endothelial cell CSPG core protein analysis.** The relationship of microvascular endothelial cell CSPG to the previously described mouse melanoma cell CSPG core protein (CD44) was analyzed by Western blot analysis of cell lysates and by immunoprecipitation of $^{125}$I-surface proteins digested with chondroitinase ABC using Hermes 1 and Hermes 3 anti-CD44 mAbs. The cell lysates and immunoprecipitated $^{125}$I-labeled protein(s) were analyzed by 6–15% SDS-PAGE under nonreducing conditions. Western analysis of undigested cell lysates revealed the presence of two broad smears ranging from $\sim$ 95 to 110 kD and $\sim$ 150 to $> 200$ kD (Fig. 6 A, lane A; brackets). After digestion of the cell lysates with chondroitinase ABC there was a downward shift in immunoreactivity with two bands present at 85 and 110 kD (Fig. 6 A, lane B; arrowheads). This data is consistent with the interpretation that two CD44 core proteins are present which are modified by chondroitin sulfate. No immunoreactivity was present when the blots were incubated with normal mouse serum in place of the anti-CD44 antibody (data not shown).

In addition, two predominant proteins with molecular masses of 85 and 110 kD were immunoprecipitated with the Hermes 1 and 3 anti-CD44 mAbs indicating an immunological relationship to CD44 (Fig. 6 B, lane B, data shown for Hermes 3 antibody only) and confirming the Western analysis. A faint third band at $\sim$ 200 kD was also immunoprecipitated. However, this protein was not immunoprecipitated by the anti-proteoglycan core protein antibodies (see below) and therefore probably does not represent a CSPG core protein. Multiple faint bands are visible in the undigested lane of CSPG proteins immunoprecipitated by anti-CD44 antibody (Fig. 6 B, lane A) with the two most predominant at $\sim$ 97 and 150 kD. These two bands roughly correspond to the smears present on Western analysis (Fig. 6 A, lane A) and are therefore consistent with CD44 modified by chondroitin sulfate. However, the 97 kD band is present in all lanes in the immunoprecipitation experiment, and therefore we are not certain whether this band represents CD44 without chondroitin sulfate, incompletely chondroitinase digested CD44, or nonspecific binding. Several other very faint bands at $< 85$ were immunoprecipitated with both the anti-CD44 antibody and anti-proteoglycan core antibodies and were present in all lanes and likely represent nonspecific binding. As a negative control, rabbit anti–mouse IgG antibody coupled to protein A agarose beads failed to bind the proteoglycan core proteins (data not shown). Of note, $\sim$ 25% of radiolabeled CSPG was immunoprecipitated by the anti-CD44 antibodies indicating that at a minimum, 25% of total CSPG is CD44-related CSPG.

To confirm that the two proteins immunoprecipitated by anti-CD44 antibodies were proteoglycan core proteins, $^{125}$I-surface-labeled proteins were detergent extracted and immunoprecipitated using two mouse mAbs that recognize the unsaturated bonds of 4-O and 6-O sulfated chondroitin sulfate residues (stubbs) that remain attached to the proteoglycan core protein after chondroitinase ABC digestion. The resulting autoradiogram showed that the two CD44-related CSPG proteins were recognized by the monoclonal proteoglycan core antibodies after chondroitinase ABC digestion with exactly the same molecular masses of 85 and 110 kD (Fig. 6 B, lane C). As mentioned above, a faint band at 97 kD is present in the lane showing undigested CSPG immunoprecipitated with the mAbs that recognize digested core protein (Fig. 6 B, lane D). We are unsure whether this band represents some reactivity of the antibodies with core protein or nonspecific binding. These data demonstrate that two predominate CD44 core proteins are present in wound microvascular endothelial cells.

**Analysis of CD44 mRNA transcripts in wound microvascular endothelial cells.** RT-PCR was performed to determine which mRNA transcripts encoding for different CD44 isoforms are present in wound microvascular endothelial cells. Consistent with the Western analysis and immunoprecipita-
tion studies showing two predominant CD44 isoforms, RT-PCR using PCR primers located in the common exons E3 and E16 generated two PCR products of ~450 and 550 bp (Fig. 7). The approximate 450-bp product is the predicted size (469 bp) for the “standard” isoform of CD44 (CD44H), which corresponds to the 85-kD protein.

RT-PCR was also used to examine expression of CD44 transcripts containing variant exons. The variant exon primers pv2 through pv10 were used as forward primers and the E16 PCR primer was used as the reverse primer. Using the pv3-2 primer which is downstream of the alternate acceptor site in exon 3, a PCR product of ~200 bp was generated. No other transcripts were present using other variant exon primers. The approximate 200-bp product is the predicted size for variant exon 3 (V3 alternate acceptor site) plus E16. This corresponds to the 550-bp product present using the E3 and E16 primers and represents a PCR product which begins in constant exon 3, ends in constant exon 16, and contains the variant exon 3 (alternate acceptor site). Importantly, variant exon 3 contains the Ser-Gly-Ser-Gly sequence which is a potential glycosaminoglycan attachment site (29). Negative controls consisting of PCR of primers only (no DNA) and PCR of RNA (no reverse transcriptase reaction) yielded no PCR products. This indicates that the higher mol wt CD44 isoform present in the immunoprecipitation experiments (110-kD protein) is encoded by a mRNA transcript containing variant exon 3.

Anti-CD44 antibody blocks endothelial cell migration on fibrinogen. Anti-CD44 mAb was used to assess the role of CD44 in mediating endothelial cell motility on fibrinogen. Wound microvascular endothelial cells preincubated with anti-CD44 mAb exhibited a dose-dependent reduction in migration on fibrinogen (Fig. 8). At the two highest concentrations (10 and 20 μg/ml) of mAb tested, microvascular endothelial cell migration was completely inhibited. As controls, normal mouse IgG
antibody and an anti-\(\alpha_5\beta_1\) integrin mAb (mediates binding to collagen) were substituted for the anti-CD44 antibody. No reduction in microvascular endothelial cell migration on fibrinogen occurred when the microvascular endothelial cells were preincubated with control antibodies.

HPLC-DEAE partially purified microvascular endothelial cell CSPG binds fibrinogen. To directly assess whether microvascular endothelial cell CSPG binds fibrinogen, detergent-extracted HPLC-DEAE partially purified \(^{35}\)S-CSPG was applied to a fibrinogen affinity column. Rabbit fibrinogen was purified by HPLC-DEAE chromatography to remove small quantities of contaminating fibronectin. HPLC-DEAE partially purified CSPG bound to the fibrinogen affinity column and eluted in a relatively broad band from the column by 0.45 M

Figure 6. Identification of two CSPG core proteins with molecular masses of 85- and 110-kD by Western analysis and immunoprecipitation with CD44 mAbs. To determine the molecular mass of the CSPG core protein(s), and to analyze the core proteins for antigenic similarity to CD44, Western analysis of microvascular endothelial cell lysates was performed (A). To confirm Western analysis data, \(^{125}\)I surface-labeled proteins were detergent-extracted and immunoprecipitated using anti-CD44 mAbs (B, lanes A and B) and mAbs that recognize chondroitin sulfate residues attached to the core protein after chondroitinase treatment (B, lanes C and D). The cell lysates and immunoprecipitated proteins were analyzed by 6–15% SDS-PAGE under nonreducing conditions. (A, lane A) Western analysis of undigested cell lysates. (A, lane B) Western analysis of chondroitinase ABC digested cell lysates. B, lane A contains undigested CSPG Hermes 3 (mouse anti-human CD44 mAb) immunoprecipitated proteins. B, lane B contains proteins immunoprecipitated by Hermes 3 after chondroitinase ABC digestion. B, lane C contains proteoglycan core proteins immunoprecipitated by the monoclonal antibodies that recognize the unsaturated bonds of chondroitin sulfate-associated uronic acid residues that remain on the core protein after chondroitinase ABC digestion. B, lane D contains undigested CSPG proteins immunoprecipitated with the anti-proteoglycan core protein antibodies.

Figure 7. Expression of CD44 RNA transcripts in wound microvascular endothelial cells. (A) Line drawing representation of the extracellular domains of CD44 based on the exon map of Screaton et al. (31). E1-E5 and E15-E16 represent constant CD44 exons. E6-E14 represent potential alternatively spliced exons. E3 represents the constant exon forward primer and E16 represents the constant exon reverse primer used to generate the PCR products in lane 2. The arrow above E7 represents the alternate acceptor site in variant exon V3. PV2-PV10 represent forward primers for each of the variable exons that were used to generate RT-PCR bands. (B) RT-PCR was performed on wound microvascular endothelial cells. PCR products were fractionated on a 2% agarose gel and stained with ethidium bromide. Lanes 1 and 10 contain the 1-kb DNA ladder. Lane 2 contains the PCR products for the constant to constant exon primers (E3–E16). Lanes 3–12 contain the PCR products for each of the variant exon primers. Lanes 14 and 15 are negative controls.
NaCl (Fig. 9). CSPG eluting from the column was verified as authentic by nitrous acid resistance and chondroitinase ABC sensitivity. In addition, 35S-CSPG was applied to a sham column prepared without fibrinogen. Recovery of radioactivity from this column was 95% indicating that the binding of CSPG to fibrinogen was specific.

![Figure 8. Wound microvascular endothelial cell migration on fibrinogen is inhibited by anti-CD44 mAb. Microvascular endothelial cells were preincubated for 30 min with varying concentrations of a mouse anti-human CD44 mAb. Endothelial cell migration to the undersides of filters coated with varying concentrations of fibrinogen (0.1–5 μM) were assessed using modified Boyden chambers in the continued presence of CD44 antibody (●). Normal mouse IgG antibody (△) and anti-α2β1 integrin mAb (□) were used as controls. Shown is wound microvascular endothelial cell migration to 0.5 μM fibrinogen quantified as the number of cells migrating per ×100 field.](image)

![Figure 9. Wound microvascular endothelial cell CSPG binds to a fibrinogen affinity column. Detergent extracted HPLC-DEAE partially purified 35S-CSPG was applied to a fibrinogen affinity column equilibrated with 50 mM Tris, pH 6.8, 0.5% CHAPS, 0.05 M NaCl, 0.01 M 6-aminohexanoic acid, 0.1 mM PMSF, 1.0 mM NEM, 0.02% azide. Rabbit fibrinogen was purified by HPLC-DEAE chromatography to remove small amounts of contaminating fibronectin. The column was eluted with a linear salt gradient. Radioactivity was monitored for each 1-ml fraction (○), and the salt gradient was monitored by conductivity measurements (●).](image)

Discussion

Microvascular endothelial cell invasion into the fibrin provisional matrix is an integral component of angiogenesis during wound and tissue repair after injury. Cell surface matrix receptors, which interact with ECM proteins like fibrin, mediate the adhesion and migration of the cell into the ECM. The central finding in this paper is that a CD44-related CSPG cell surface matrix receptor is capable of mediating wound microvascular endothelial cell migration on fibrinogen and invasion into a fibrin matrix. By inhibiting the synthesis of CSPG with β-D xyloside, a reduction in microvascular endothelial cell adhesion and migration on fibrinogen and invasion into a three-dimensional fibrin gel was observed. Similarly, microvascular endothelial cell adhesion and migration on fibrinogen was significantly decreased by pretreatment with chondroitinase ABC which enzymatically cleaves chondroitin sulfate from the core protein. We have also determined that wound microvascular endothelial cells express CSPG as their predominant proteoglycan and that the CSPG contains two core proteins with molecular masses of 85 and 110 kD. Each of these core proteins can be immunoprecipitated with two different CD44 mAbs indicating antigenic similarity with CD44. PCR studies indicate that wound microvascular endothelial cells express both the standard (CD44H) isoform and an isoform containing variably spliced exon V3. In addition, anti-CD44 antibody blocks endothelial cell migration on fibrinogen. Furthermore, partially purified wound microvascular endothelial cell CSPG is capable of binding purified fibrinogen. These data suggest that wound microvascular endothelial cell CD44-related CSPG is capable of binding fibrinogen/fibrin thereby mediating endothelial cell adhesion, migration, and invasion into the fibrin matrix.

CD44-related CSPG mediates cell motility. CD44 is a transmembrane glycoprotein that may mediate tumor cell migration. The CD44 gene contains 19 exons, 12 of which may be alternatively spliced leading to the existence of multiple isoforms of CD44 (31). The diversity of CD44 is further amplified by the differential use of GAG attachment sites on its extracellular domain. Various isoforms of CD44 have been associated with enhanced migration and invasion of transformed cells (22–25). For example, we have previously shown that mouse melanoma cell invasion into type I collagen gels is mediated by CSPG containing an 110-kD CD44 core protein (13).

Several lines of evidence link cell surface CSPG with the motility of endothelial cells during wound/tissue repair. First, wounded large vessel endothelial cell cultures rapidly alter their proteoglycan synthesis from heparan to chondroitin sulfate, which coincidently occurs with the onset of migration (11). Second, immunohistochemical analysis of granulation tissue shows intense staining for CSPG. Third, proteoglycans extracted from healing wounds contain more CSPG than normal dermis and had longer glycaminoglycan chondroitin sulfate sidechains. Finally, alterations of proteoglycan synthesis associated with granulation tissue formation correlates with cell proliferation and migration (32). These data suggest that alterations in CSPG synthesis in granulation tissue may play an important role in regulating such cellular functions as migration and invasion during wound/tissue repair. This is in agreement with our finding that wound microvascular endothelial cells express the majority of their cell surface proteoglycan as...
CSPG and that cell surface CSPG is important in mediating migration and invasion into the fibrin matrix.

Acquisition of an endothelial cell migratory phenotype may be important for angiogenesis to occur. Studies indicate that endothelial cells increase their proteolytic activity (33–36), alter their fibronectin synthesis (37), and express specific migration-associated cell surface glycoproteins (11) during wound repair. These modifications are thought to facilitate cell movement. In support of the hypothesis that endothelial cells express a specific phenotype during migration are a series of in vivo and in vitro experiments. First, in vivo work indicates that fibrin gels (but not collagen or agarose) implanted into the subcutaneous tissue of guinea pigs are capable of inducing angiogenesis (4). Presumably those endothelial cells invading the fibrin matrix from the connective tissue space express the migratory phenotype. In contrast, prior in vitro investigations as well as our own unpublished observations have shown that large vessel endothelial cells, in the absence of exogenous stimuli, are incapable of invading the fibrin gel (38). However, as shown in the current study, wound microvascular endothelial cells, which express CSPG as their predominant cell surface proteoglycan, are fully capable of invading into the fibrin matrix and forming vascular-like networks without exogenous stimulation. Our finding that microvascular endothelial cell CSPG is capable of binding fibrinogen coupled with Dvorak’s work indicating that fibrin is capable of eliciting an angiogenic response suggests that CD44-related CSPG/fibrin interaction may play an important role in directing endothelial cell migration into the fibrin provisional matrix during wound repair. Our migration assays, which demonstrate that anti-CD44 antibody blocks migration on fibrinogen, support the concept of a specific CD44-related CSPG-fibrinogen interaction. However, since CD44-related CSPG represents a portion of total microvascular endothelial cell CSPG we cannot be certain that CD44-related CSPG binds fibrinogen. This requires further investigation. These data suggest that wound microvascular endothelial cells assume a migratory phenotype.

It is possible that the expression of CD44-related CSPG on the cell surface may be an important component of the migration phenotype. Our data indicate that microvascular endothelial cells express predominantly CSPG and that the CSPG contains two CD44 core proteins, the standard or 85-kD isoform and an ~ 110-kD isoform. Western analysis shows two broad smears of immunoreactivity at ~ 95–110 kD and 150 to > 200 kD in undigested cell lysates which shift downward after chondroitinase digestion resulting in the 85- and 110-kD bands. This data is consistent with the interpretation that CD44 is modified by chondroitin sulfate. Immunoprecipitation data indicate that the chondroitinase digested 85- and 110-kD CD44 proteins are proteoglycan core proteins. Interestingly, the 97-kD band, which is present as a prominent broad smear in the chondroitinase undigested lane in the Western blot, is present in all lanes (both chondroitinase digested and undigested) in the immunoprecipitation experiment. Because there is a downward shift in immunoreactivity after chondroitinase digestion in the Western analysis, this suggests that the 97-kD band is CD44 modified by chondroitin sulfate. However, because the 97-kD band is present in all lanes in the immunoprecipitation experiment, this band could also represent CD44 without chondroitin sulfate, CD44 partially modified by chondroitin sulfate, or non-specific binding. Further work is necessary to clarify this issue. In addition, our PCR studies demonstrate that microvascular endothelial cells express both the standard (CD44H) isoform encoding the 85-kD isoform of CD44 and an isoform containing the variably spliced exon V3 encoding the ~ 110-kD CD44 isoform. Importantly, exon V3 contains the Ser-Gly-Ser-Gly sequence which serves as a potential glycosaminoglycan attachment site (29). Additional glycosaminoglycan attachment sites are also present in the constant regions of CD44. Therefore, since wound microvascular endothelial cells express CSPG on their surface it seems plausible that these glycosaminoglycan attachment sites are occupied by chondroitin sulfate. In contrast however, a prior study has determined that large vessel endothelial cells express only the standard (CD44H) isoform of CD44 encoding the 85-kD protein (29). Additionally, large vessel endothelial cells express predominantly heparan sulfate on their surface and only switch their proteoglycan synthesis from heparan to chondroitin sulfate as migration is initiated after wounding (11). Collectively these data support the hypothesis that the wound microvascular endothelial cell migratory phenotype is characterized by the presence of the 85- and 110-kD CD44 isoforms which are modified by the addition of chondroitin sulfate.

Insights into the mechanism of CD44-related CSPG-mediated cell motility. The precise mechanism by which CD44-related CSPG mediates cell motility is not completely understood. However, as demonstrated in this study, anti-CD44 mAbs block endothelial cell migration, suggesting that CD44 core protein is involved in cell migration. The ability of CD44 to bind multiple extracellular proteins and therefore function as a matrix adhesion receptor is likely important in facilitating cell movement. CD44 is a matrix receptor for hyaluronate (39), fibronectin (40), type I collagen (13, 41), and the adhesion and migration assays in our current study provide evidence that CD44-related CSPG also interacts with fibrinogen. The importance of CD44 as an ECM adhesion receptor is illustrated by a recent study in which mouse fibroblasts transfected with a primate CD44 gene acquired a new adhesive phenotype (42). Furthermore, CD44 has been shown to associate with the cytoskeleton (14, 15, 43, 44) further linking it with cell adhesion and migration.

Chondroitin sulfate on the CD44 core protein may also play a role in facilitating cell migration. We have previously shown that chondroitin sulfate binds ECM proteins such as collagen (13). Other work has shown that CSPG is present on microspores on the cell surface which mediate cell attachment to the ECM (12) Our adhesion and migration assays in which chondroitin sulfate is cleaved from the core protein by chondroitinase resulting in reduced adhesion and migration suggest that chondroitin sulfate is involved in cell attachment and motility. Further support for this hypothesis comes from the finding that endothelial cell adhesion and migration to fibrinogen were inhibited to a similar degree by both chondroitinase and β-D xyloside in a concentration dependent fashion, suggesting that the degree of removal of chondroitin sulfate from the core protein corresponded with the observed decreases in adhesion and migration.

Interestingly, we have found that when CSPG function is blocked by either β-D xyloside or chondroitinase ABC treatment in the short term adhesion assay, the degree of inhibition of endothelial cell adhesion to fibrinogen was the same, 50%. The fact that only 50% inhibition of adhesion was seen despite high concentrations of β-D xyloside or enzyme suggests that other receptors such as integrins are involved in microvascular
endothelial cell adhesion to fibrinogen. Furthermore, in the invasion assay, the cells were capable of adhering to the surface of the fibrin gel even in the presence of β-d xyloside but were unable to invade the gel. Of note, the apparent discrepancy between the short term adhesion assay where endothelial cell adhesion to fibrinogen can be partially inhibited vs the long term invasion assay where endothelial cells are capable of attaching to fibrin may be explained by the use of alternate adhesion receptors. Therefore, in addition to CSPG, other receptors such as αβ integrin, which mediates large vessel endothelial cell adhesion to fibrinogen, may mediate endothelial cell adhesion to the fibrin gel matrix. In support of this concept, we have previously shown that the coordinate interaction of CSPG and αβ integrin facilitates cell adhesion to fibronectin (45). These data suggest that the coordinate interaction of matrix receptors may be important in mediating such cellular behavior as adhesion and migration.

The action of xylosides on protein synthesis is complex. β-d xyloside is a specific inhibitor of xylose-linked proteoglycan assembly and inhibits the synthesis of xylose-linked proteoglycans such as CSPG (26). α-d xyloside, the control enantiomer of β-d xyloside used in this study, does not inhibit xylose linked proteoglycan assembly. However, both xylosides have been shown to inhibit total protein synthesis and specifically decrease fibronectin synthesis in large vessel endothelial cells (46). This leaves open the possibility that the effect of β-d xyloside on endothelial cell invasion may not only be due to interruption of CSPG function, but also due to modulation of fibronectin synthesis. However, in support of the role of CD44-related CSPG in mediating endothelial cell migration and invasion are the inhibition assays using chondroitinase ABC and anti-CD44 antibodies. The migration of endothelial cells on fibrinogen were significantly decreased using these agents indicating that the CD44-related CSPG molecule itself is important in mediating locomotion. In further support of the role of CSPG in mediating invasion is the finding that α-d xyloside does not significantly block cell invasion. α-d xyloside inhibits both total protein and fibronectin synthesis but does not affect CSPG synthesis. This implies that the inhibition of CSPG synthesis by β-d xyloside is largely responsible for the inhibition of invasion. Furthermore, to ensure that disruption of endothelial cell interaction with fibrin and not other matrix components contaminating the fibrin matrix was the mechanism by which β-d xyloside inhibited invasion, HPLC-DEAE purified fibrinogen was used to make the fibrin gels. Additionally, invasion assays were performed using hyaluronidase treated gels. Despite these treatments, endothelial cell invasion was inhibited indicating that disruption of CSPG interaction with fibrin and not other ECM proteins was the likely mechanism of inhibition of invasion by β-d xyloside. Further investigations aimed at examining whether CD44 core protein and/or chondroitin sulfate bind fibronectin and studies which define whether specific domains on the fibrinogen molecule mediate receptor interaction are needed to fully understand the role of this cell surface matrix receptor in mediating microvascular endothelial cell migration and invasion.

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