Heparan Sulfate Proteoglycans Mediate the Angiogenic Activity of the Vascular Endothelial Growth Factor Receptor-2 Agonist Gremlin

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**Objective**—Heparan sulfate proteoglycans (HSPGs) modulate the interaction of proangiogenic heparin-binding vascular endothelial growth factors (VEGFs) with signaling VEGF receptor-2 (VEGFR2) and neuropilin coreceptors in endothelial cells (ECs). The bone morphogenic protein antagonist gremlin is a proangiogenic ligand of VEGFR2, distinct from canonical VEGFs. Here we investigated the role of HSPGs in VEGFR2 interaction, signaling, and proangiogenic capacity of gremlin in ECs.

**Methods and Results**—Surface plasmon resonance demonstrated that gremlin binds heparin and heparan sulfate, but not other glycosaminoglycans, via $N$, $2$-O, and $6$-O-sulfated groups of the polysaccharide. Accordingly, gremlin binds HSPGs of the EC surface and extracellular matrix. Gremlin/HSPG interaction is prevented by free heparin and heparan sulfate digestion or undersulfation following EC treatment with heparinase II or sodium chlorate. However, at variance with canonical heparin-binding VEGFs, gremlin does not interact with neuropilin-1 coreceptor. On the other hand, HSPGs mediate VEGFR2 engagement and autophosphorylation, extracellular signaling-regulated kinase1/2 and p38 mitogen-activated protein kinase activation, and consequent proangiogenic responses of ECs to gremlin. On this basis, we evaluated the gremlin-antagonist activity of a panel of chemically sulfated derivatives of the *Escherichia coli* K5 polysaccharide. The results demonstrate that the highly $N,O$-sulfated derivative K5-N,OS(H) binds gremlin with high potency, thus inhibiting VEGFR2 interaction and angiogenic activity in vitro and in vivo.

**Conclusion**—HSPGs act as functional gremlin coreceptors in ECs, affecting its productive interaction with VEGFR2 and angiogenic activity. This has allowed the identification of the biotechnological K5-N,OS(H) as a novel angiostatic gremlin antagonist. (Arterioscler Thromb Vasc Biol. 2011;31:e116-e127.)

**Key Words:** angiogenesis ■ endothelium ■ gremlin ■ heparan sulfate proteoglycans

Angiogenesis plays a key role in various physiological and pathological conditions, including embryonic development, wound repair, inflammation, and tumor growth. Numerous inducers of angiogenesis have been identified, including the members of the vascular endothelial growth factor (VEGF) family. Different isoforms of mammalian VEGFs interact with tyrosine kinase VEGF receptors (VEGFRs) expressed on the surface of endothelial cells (ECs) and with heparan sulfate (HS) proteoglycan (HSPG) and neuropilin (NRP) coreceptors, thus activating a proangiogenic response.

HSPGs consist of a core protein and of glycosaminoglycan (GAG) chains represented by unbranched heparin-like polysaccharides. They are found in free forms, in the extracellular matrix (ECM), or associated with the plasma membrane where they regulate the function of a wide range of ligands. In particular, endothelial HSPGs modulate angiogenesis by affecting bioavailability and interaction of heparin-binding VEGFs with signaling VEGFRs and NRP coreceptors. Heparin/HS interaction with angiogenic growth factors depends on the degree/distribution of sulfate groups and length of the GAG chain, distinct oligosaccharide sequences mediating its binding activity. Moreover, the study of the biochemical bases of this interaction has been exploited to design angiostatic heparin-like compounds, chemically modified harpins, and biotechnological heparins.

Gremlin belongs to the CAN (Cerberus and Dan) family of cystine-knot–secreted proteins. Gremlin binds various bone morphogenic proteins (BMPs), thus preventing their interaction with cell surface receptors. The capacity of gremlin to bind BMPs and to inhibit their interaction with the cognate transforming growth factor-$\beta$ (TGF-$\beta$) family receptors is thought to play a role in embryonic development of various organ systems, such as bone, kidney, and lung. Also, gremlin has been implicated in the pathogenesis of human diseases, including pulmonary hypertension and idiopathic pulmonary fibrosis and diabetic nephropathy. Moreover,
Gremlin is produced by human tumors\textsuperscript{25-26} and is expressed by proangiogenic ECs in vitro and tumor endothelium in vivo.\textsuperscript{27} Gremlin stimulates EC intracellular signaling and migration in vitro, leading to a potent angiogenic response in vivo.\textsuperscript{27,28} This is due to its capacity to bind and activate VEGFR2, the main transducer of VEGF-mediated angiogenic signals, in a BMP-independent manner.\textsuperscript{29} Thus, gremlin acts as a VEGFR2 agonist distinct from VEGF family members that may play paracrine/autocrine functions in tumor neovascularization.

Scattered experimental evidence indicates that gremlin may act as a heparin-binding angiogenic factor: gremlin binds to heparin-Sepharose beads;\textsuperscript{27} it associates to the surface of producing cells;\textsuperscript{10} and a heparin-binding domain has been identified in sclerostin, a BMP antagonist structurally related with gremlin.\textsuperscript{31,32} Here, we investigated the capacity of gremlin to interact with heparin and cell-associated HSs and the role of endothelial HSs in mediating gremlin-dependent VEGFR2 activation and angiogenic activity. The results demonstrate that gremlin binds heparin/HS. However, at variance with canonical heparin-binding VEGFR2 ligands, gremlin interacts with HSs but not with NRP-1 in ECs. Nevertheless, cell surface HSs play a nonredundant role in mediating the ability of gremlin to bind and activate VEGFR2 in endothelium, thus triggering an NRP-1-independent proangiogenic response. Accordingly, we have identified a chemically modified, heparin-like derivative of the capsular \textit{Escherichia coli} K5 polysaccharide\textsuperscript{16} as a potent angiostatic gremlin antagonist.

### Methods

#### Chemicals

Recombinant gremlin, anti-gremlin antibody, and human recombinant VEGF-A165 were from R&D Systems (Minneapolis, MN); anti-phospho (P)-VEGFR2 antibody was from Cell Signaling Technology (Boston, MA); anti-tubulin antibody, heparinase II, and phorbol myristate acetate (PMA) were from Sigma-Aldrich (St Louis, MO); anti-P-ERK\textsubscript{1/2}, anti-VEGFR2, anti-ERK\textsubscript{2}, anti-p38, and anti-phospho (P)-VEGFR2 antibody was from Cell Signaling Technology (Billerica, MA); anti-focal adhesion kinase (clone 2A7) was from Upstate, Millipore; sodium chlorate (50 and 30 mmol/L, respectively). Alternatively, cells were incubated for 2 hours at 37°C with heparinase II (15 μg/mL) in phosphate-buffered saline (PBS), Sigma-Aldrich) before experimentation.

#### Human Pancreatic Carcinoma

Human pancreatic carcinoma (PANC-1) cells were grown in Dulbecco’s modified Eagle medium containing 10% FCS, vitamins, and essential and nonessential amino acids. Gremlin-overexpressing PANC-1 cells (PANC-1-gremlin cells) were generated with infection by the pRRLsin.cPT.PKG lentiviral vector (kindly provided by Dr. Naldini, San Raffaele Institute, Milan, Italy) harboring the rat gremlin cDNA,\textsuperscript{40} Green fluorescent protein (GFP)–overexpressing PANC-1 cells (PANC-1-GFP cells) were generated with the same procedures and used as a negative control. PANC-1-GFP and PANC-1-gremlin cells were grown for 48 hours in presence of sodium chloride (50 and 30 mmol/L, respectively).\textsuperscript{39} Surface plasmon resonance (SPR) measurements were performed on a BIAcore X instrument (GE Healthcare). For the study of gremlin/heparin interaction, heparin was immobilized onto a CM3 sensor chip (GE Healthcare) as described.\textsuperscript{39} A streptavidin-coated sensor chip was used as reference and for blank subtraction. Increasing concentrations of gremlin in 10 mmol/L HEPES buffer (HBS-EP), pH 7.4, containing 150 mmol/L NaCl, 3 mmol/L EDTA, and 0.005% surfactant P20 (HBS-EP) (GE Healthcare) were injected over the heparin-coated or streptavidin surfaces for 4 minutes and then washed until dissociation was observed. After each run, the sensor chip was regenerated by injection of HBS-EP containing 2.0 mol/L NaCl.

In all experiments, association and dissociation rates ($k_a$ and $k_d$, respectively) were obtained by fitting the raw sensorgrams with the 1:1 Langmuir binding model using the Biaevaluation software (BIAcore). Equilibrium affinity constants (expressed as dissociation constant, $k_d$) were either derived from the kinetic parameters ($k_d=k_d(k_m)$) or determined from Scatchard plot analysis of the equilibrium binding data.

#### Cell Cultures

Human umbilical vein endothelial (HUVE) cells were used at early passages (passages I–IV) and grown on plastic surface coated with porcine gelatin (Sigma-Aldrich) in M199 medium (Invitrogen, Carlsbad, CA) supplemented with 20% fetal calf serum (FCS) (Invitrogen), EC growth factor (100 μg/mL) (Sigma-Aldrich), and porcine heparin (100 μg/mL, Sigma-Aldrich). Fetal bovine aortic endothelial GM7373 cells\textsuperscript{35} were grown in Dulbecco’s modified Eagle medium (Invitrogen) containing 10% FCS, vitamins, and essential and nonessential amino acids. These cells were transfected according to standard protocols with a pcDNA3.1 expression vector harboring the mouse VEGFR2 cDNA (provided by G. Breier, Max Planck Institute, Bad Nauheim, Germany) to generate stable GM7373-VEGFR2 transfectants. Wild-type Chinese hamster ovary cells (CHO)-K1 cells and GAG-deficient A745 CHO cells\textsuperscript{38} were kindly provided by J.D. Esko (University of California, La Jolla, CA) and grown in Ham’s F-12 medium supplemented with 10% FCS.

To inhibit the sulfation of cell-associated HS chains, GM7373 and HUVE cells were grown for 48 hours in presence of sodium chloride (50 and 30 mmol/L, respectively).\textsuperscript{39}
with gremlin and analyzed for HSPG-bound gremlin as described above.

**VEGFR2 Cross-Linking Assay**

Confluent GM7373-VEGFR2 cells were incubated for 2 hours at 4°C with gremlin dissolved in PBS containing the bis(sulfosuccinimidyl) suberate cross-linking reagent (BS3, Pierce). Then, cells were lysed in 50 mMol/L Tris-HCL buffer (pH 7.4) containing 150 mMol/L NaCl, 1% Triton X-100, 1.0 mMol/L Na3VO4, and protease inhibitors. Lysates were immunoprecipitated and separated by SDS-PAGE followed by Western blotting with anti-gremlin or anti-VEGFR2 antibodies.

**ELISA**

Ninety-six-well plates were coated for 16 hours at room temperature with the extracellular domain of soluble VEGFR2E1-7 (soluble VEGFR2, Calbiochem) or rat NRP-1 receptor (R&D Systems) in PBS followed by a 3-hour blocking step with 1% bovine serum albumin. Next, gremlin or VEGF-A165 (in PBS containing 0.1% bovine serum albumin, 5 mMol/L EDTA, 0.004% Tween 20 [PBET]) were sequentially incubated for 1 hour at 37°C and for 1 hour at room temperature. For competitive ELISA, gremlin was added in presence of K5 derivatives. Then, wells were further incubated for 1 hour at 37°C and for an additional 1 hour at room temperature with an anti-gremlin antibody or anti-VEGF antibody (R&D Systems), both at 100 ng/mL in PBET. Finally, wells were incubated for 1 hour at room temperature with a secondary anti-goat or anti-mouse horseradish peroxidase–conjugated antibody (Santa Cruz Biotechnology).

**VEGFR2/NRP-1 Coimmunoprecipitation Analysis**

Confluent HUVE cells were made quiescent by a 16-hour starvation in M199 medium containing 5% FCS. Cells were then stimulated with gremlin (50 ng/mL) or VEGF-A165 (30 ng/mL) and lysed in 50 mMol/L Tris-HCI buffer, pH 7.4, containing 150 mMol/L NaCl, 1% Triton X-100, 10% glycerol, and a protease/phosphatase inhibitor mix (Sigma-Aldrich). After centrifugation (15 minutes, 10,000g), supernatants were preclarified by 1 hour of incubation with protein A–Sepharose. Samples (1 mg of proteins) were incubated overnight at 4°C with rabbit polyclonal anti-VEGFR2 antibody (Santa Cruz Biotechnology), and immune complexes were recovered on protein A–Sepharose. Samples (1 mg of proteins) were washed 4 times with lysis buffer, twice with the same buffer without Triton X-100 and once with TBS. Proteins were solubilized under reducing conditions, separated by SDS-PAGE (8%), and analyzed by Western blotting with anti-NRP-1 and anti-P-VEGFR2 (Tyr1175) antibodies. Uniform immunoprecipitation was confirmed using an anti-VEGFR2 antibody.

**Immunofluorescence Analysis**

Cells were seeded on glass coverslips in M199 medium containing 2% FCS. Cells were treated with gremlin (100 ng/mL) or PMA (100 ng/mL) in M199 containing 2% FCS and incubated overnight at 4°C with rabbit polyclonal anti-VEGFR2 antibody (Santa Cruz Biotechnology), and immune complexes were recovered on protein A–Sepharose. Samples (1 mg of proteins) were washed 4 times with lysis buffer, twice with the same buffer without Triton X-100 and once with TBS. Proteins were solubilized under reducing conditions, separated by SDS-PAGE (8%), and analyzed by Western blotting with anti-NRP-1 and anti-P-VEGFR2 (Tyr1175) antibodies. Uniform immunoprecipitation was confirmed using an anti-VEGFR2 antibody.

**Western Blot Analysis**

Confluent HUVE cells were seeded at 150 cells/cm2 in 24-well plates for 2 hours and then stimulated with gremlin (100 ng/mL). Constant temperature (37°C) and pCO2 (5%) were maintained throughout the experimental period by means of an heatable stage and climate chamber. Cells were observed under an inverted photomicroscope (Zeiss Axiovert 200M), and phase-contrast snap photographs (1 frame every 8 minutes) were digitally recorded for 360 minutes. Cell paths (25–30 cells per experimental point) were generated from centroid positions and migration parameters were analyzed with the Chemotaxis and Migration tool of ImageJ software (http://rsweb.nih.gov/ij).

**Cell Motility Assay**

Cell motility was assessed by time-lapse videomicroscopy. Cells were seeded at 150 cells/cm2 in 24-well plates for 2 hours and then stimulated with gremlin (100 ng/mL) or PMA (100 ng/mL) in the absence or in the presence of K5 derivatives. Formation of radially growing cell sprouts was observed during the next 24 hours and photographed at a ×200 magnification using an Axiovert 200M microscope equipped with a ×20 objective (LD A Plan 20×/0.30PH1, Zeiss).

**Chick Embryo Chorioallantoic Membrane Assay**

Chick embryos were incubated overnight at 4°C with rabbit polyclonal anti-VEGFR2 antibody (Santa Cruz Biotechnology), and immune complexes were recovered on protein A–Sepharose. Samples (1 mg of proteins) were washed 4 times with lysis buffer, twice with the same buffer without Triton X-100 and once with TBS. Proteins were solubilized under reducing conditions, separated by SDS-PAGE (8%), and analyzed by Western blotting with anti-NRP-1 and anti-P-VEGFR2 (Tyr1175) antibodies. Uniform immunoprecipitation was confirmed using an anti-VEGFR2 antibody.

**Human Artery Ring assay**

One-millimeter human umbilical artery rings were embedded in fibrin gel28 in the presence of the conditioned medium from PANC-1-GFP or PANC-1-gremlin cells in the absence or in the presence of 100 ng/mL K5 derivatives. After 6 days, EC sprouts, morphologically distinguishable from scattering fibroblasts/smooth muscle cells, were counted. Then, rings were photographed at a ×100 magnification using an Axiovert 200M microscope equipped with a ×20 objective (LD A Plan 20×/0.30PH1, Zeiss).

**Statistical Analysis**

Data were expressed as mean ± SD. The analysis was performed using ANOVA models with probability values correction for multiple comparisons.

**Results**

**Characterization of Gremlin/Heparin Interaction**

Real-time SPR analysis was exploited to characterize gremlin/heparin interaction. Increasing concentrations of gremlin were injected over a streptavidin-activated BIAcore sensor
chip coated with biotinylated heparin (Figure 1A). Analysis of kinetic parameters ($k_{on}$ and $k_{off}$), $k_{on} = 1.86 \times 10^5$ [mol/L]^{-1} per second^{-1} and $3.66 \times 10^{-3}$ per second^{-1}, respectively), and Scatchard plot analysis of steady-state SPR (data not shown) demonstrates that gremlin/heparin interaction occurs with high affinity ($K_d = 20$ nmol/L).

The capacity of different GAGs to inhibit the binding of gremlin to immobilized heparin was then evaluated in a SPR competition assay (Figure 1B). Free heparin and HS prevented the binding of gremlin to immobilized heparin with ID$_{50}$ values of 0.12 and 0.7 g/mL, respectively (data not shown), whereas dermatan sulfate, chondroitin-4-sulfate, chondroitin-6-sulfate, and hyaluronic acid were ineffective. Thus, GAG/gremlin interaction depends, at least in part, on differences in GAG structure and degree of sulfation. On this basis, selectively desulfated heparins were evaluated in the SPR competition assay (Figure 1C). No competition was exerted by totally O-desulfated heparin, whereas 2-O-desulfated, 6-O-desulfated, and N-desulfated/N-acetylated heparins show a significant reduction in the capacity to inhibit the binding of gremlin to the heparin-coated sensor chip ($ID_{50} = 0.6, 0.6, and 0.9 \mu g/mL$, respectively) compared with unmodified heparin ($ID_{50} = 0.12 \mu g/mL$). Taken together, the data indicate that at least some 2-O-, 6-O-, and N-sulfate groups must be organized on the same heparin chain to allow an optimal interaction with gremlin.

Also, the influence of the length of the heparin chain on its binding activity was investigated by using tinzaparin, a therapeutic low-MW heparin whose MW ranges between 5.5 and 7.5 kDa, and 3 different very-low-MW heparins (average MWs of 4.0, 3.0, and 2.1 kDa, respectively). On a molar basis, tinzaparin binds gremlin with an affinity that is similar to unmodified heparin, whereas the capacity of very-low-MW heparins to bind gremlin is progressively reduced as a function of size, with 2.1-kDa heparin showing the weakest activity (Figure 1D).

**Gremlin Binds EC-Associated HSPGs**

We then evaluated the capacity of gremlin to interact with cell-associated HSPGs. In a first set of experiments, HSPG-bearing CHO-K1 cells and GAG-deficient A745 CHO-K1 cells were incubated with gremlin (100 ng/mL) for 2 hours at 4°C. At the end of incubation, nonpermeabilized cells were decorated with anti-gremlin antibody (in green) and 4',6-diamidino-2-phenylindole (in blue). z-Stack sections and orthogonal z reconstitution were analyzed by confocal microscopy.

![Figure 2. Confocal analysis of gremlin binding to cell-surface heparan sulfate proteoglycans (HSPGs). CHO-K1 cells (A) and glycosaminoglycan-deficient A745 CHO-K1 cells (B) were incubated with gremlin (100 ng/mL) for 2 hours at 4°C. At the end of incubation, nonpermeabilized cells were decorated with anti-gremlin antibody (in green) and 4',6-diamidino-2-phenylindole (in blue). z-Stack sections and orthogonal z reconstitution were analyzed by confocal microscopy.](image-url)
to bind the surface of CHO-K1 cells but not of GAG-deficient A745 CHO-K1 cells. The binding of gremlin to CHO-K1 cell surface was inhibited by free heparin or HS, whereas hyaluronic acid was ineffective. Also, in keeping with a postulated HSPG interaction, a 1.5 mol/L NaCl wash detached gremlin bound to the CHO-K1 cell surface. Similar results were obtained when cell-surface bound gremlin was analyzed by Western blotting of the cell monolayers (Figure 3B).

We then evaluated the capacity of gremlin to bind EC-associated HSPGs. In keeping with the data obtained with CHO-K1 cells, immunofluorescence analysis revealed that gremlin binds the surface of bovine aortic endothelial GM7373 cells and that the binding is competed by heparin or abrogated by a 1.5 mol/L NaCl wash (data not shown). Accordingly, digestion of cell-associated HSPGs by heparinase II or inhibition of the sulfation of the HS chains by pretreatment of ECs with sodium chlorate both prevented the binding of gremlin to the EC surface (Figure 3C). Similar results were obtained with HUVE cells (see below).

Finally, we analyzed the capacity of gremlin to bind HSPGs associated with the cell-free subendothelial ECM deposited by GM7373 cells (Figure 3C). Also in this case, binding of gremlin to subendothelial ECM was prevented by digestion with heparinase II, pretreatment of ECs with chlorate before ECM isolation, or coincubation with heparin. Again, a 1.5 mol/L NaCl wash was able to remove gremlin from subendothelial ECM-associated HSPGs.

**HSPGs Are Required for VEGFR2 Binding and Activation by Gremlin**

Previous observations had implicated HSPGs in the interaction of heparin-binding VEGF isoforms with tyrosine kinase-
VEGFR2 and NRP-1 coreceptor in ECs. However, at variance with the prototypic heparin-binding VEGFR2 ligand VEGF-A165 and similar to VEGF-A121 that lacks the heparin/NRP-1 binding domain, gremlin does not induce VEGFR2/NRP-1 complex formation when administered to HUVE cells (Figure 4A), thus indicating that HSPG interaction is not per se sufficient to drive NRP-1 engagement. Also, gremlin does not directly bind recombinant NRP-1 produced in mammalian cells in ELISA (Figure 4B) or NRP-1 expressed on HUVE cells in cross-linking experiments (data not shown). Thus, gremlin appears to differ from canonical heparin-binding VEGFs in the ability to co-opt VEGFR2 coreceptors.

On this basis, we investigated whether HSPG coreceptors play any role in mediating a productive interaction of gremlin with VEGFR2. In a first set of experiments, we exploited VEGFR2-overexpressing endothelial GM7373 cells (GM7373-VEGFR2 cells) that express HSPGs and high levels of VEGFR2 on their cell surface (Figure 5). These cells were incubated at 4°C with gremlin in the presence of the bis(sulfosuccinamide)suberate cross-linking reagent BS3. After incubation, cell lysates were immunoprecipitated with anti-gremlin antibodies, and immunocomplexes were analyzed by Western blotting with anti-VEGFR2 antibodies. When cross-linked to the GM7373-VEGFR2 cell surface, gremlin forms a 250-kDa VEGFR2 complex whose formation is abrogated when the cross-linking reaction was performed in chlorate-treated cells (Figure 6A). Accordingly, chlorate pretreatment abolished the capacity of gremlin to induce VEGFR2 autophosphorylation in chlorate-treated HUVE cells, as shown by immunostaining of intact cells and Western blotting of the cell extracts using an anti-P-VEGFR2 antibody (Figure 6B and 6C). Also, the capacity of gremlin to induce ERK1/2 and p38 mitogen-activated protein kinase phosphorylation was hampered in chlorate-pretreated HUVE cells (Figure 6C). Similar results were obtained when HUVE cells were incubated with heparinase II before gremlin stimulation (data not shown). Also, addition of increasing concentrations of unmodified heparin to the cell culture medium restored the capacity of gremlin to induce VEGFR2 autophosphorylation in chlorate-pretreated HUVE cells (Figure 6D). It must be pointed out that even though chlorate pretreatment prevented the binding of gremlin to the HUVE cell surface (Figure 7A), it did not affect cell viability and the capacity of the protein kinase C activator PMA to induce ERK1/2 phosphorylation in these cells (Figure 7B and 7C), confirming the specificity of the effect. Taken together, these observations demonstrate that HSPGs are required for a productive interaction of gremlin with VEGFR2.

HSPGs Mediate the Angiogenic Activity of Gremlin

Stimulation of EC motility is part of the VEGFR2-dependent angiogenic program activated by gremlin. To assess the
role of HSPGs in mediating the angiogenic activity of gremlin, chlorate-pretreated HUVE cells were tested in a Boyden chamber assay for their capacity to migrate through a gelatin-coated filter in response to gremlin. Despite a similar capacity to adhere to immobilized gelatin (Figure 7D) and to migrate in response to PMA (Figure 8A), chlorate-pretreated HUVE cells displayed a dramatic reduction in their chemotactic response to gremlin compared with control cells (Figure 8A). Again, HSPG digestion following heparinase II incubation mimicked the effect of chlorate, thus abolishing the chemotactic activity of gremlin in HUVE cells (data not shown). In keeping with these observations, chlorate pretreatment abolished the motogenic response of HUVE cells to gremlin as assessed by time-lapse microscopy, without affecting EC motility induced by PMA (Figure 8B). Moreover, as observed for gremlin-induced VEGFR2 autophosphorylation (see Figure 6D), addition of heparin to chlorate-treated cells restored their motogenic response to gremlin stimulation (Figure 8C).

To confirm the role of HSPGs in mediating EC activation by gremlin, spheroids of chlorate or heparinase II–treated HUVE cells were embedded in a 3-dimensional fibrin gel and stimulated with gremlin.29 As shown in Figure 8D and 8E, HSPG digestion or undersulfation caused a significant reduction in the number and length of EC sprouts induced by gremlin.

Heparin-Like K5 Derivatives as Gremlin Antagonists

The capsular E. coli K5 polysaccharide has the same structure as the heparin precursor N-acetyl heparosan.16 K5 derivatives can be generated by chemical sulfation in the N- or O-position, leading to heparin-like molecules with defined sulfation patterns48 and able to hamper the angiogenic activity of the heparin-binding fibroblast growth factor-2.33 To assess their ability to act as gremlin antagonists, a panel of selectively sulfated K5 derivatives (including the N-sulfated derivative K5-NS; the O-sulfated derivatives with low and high degrees of sulfation, K5-OS(L) and K5-OS(H); and the N,O-sulfated derivatives with low and high degrees of sulfation, K5-N,OS(L) and K5-N,OS(H))33 were evaluated for their capacity to prevent gremlin/heparin interaction in a SPR competition assay. As shown in Figure 9A, K5-N,OS(H) and K5-N,OS(L) competed for the binding of gremlin to immobilized heparin with high potency (ID50 of 0.05 and 0.2

Figure 6. Endothelial heparan sulfate proteoglycans (HSPGs) mediate vascular endothelial growth factor receptor-2 (VEGFR2)/gremlin interaction and signaling. A, Control and chlorate-treated GM7373-VEGFR2 cells were incubated for 2 hours at 4°C with gremlin (100 ng/mL). After a further incubation with the cross-linking reagent BS3, cell lysates (1.0 mg) were immunoprecipitated with anti-gremlin antibody and probed with the same antibody or with an anti-VEGFR2 antibody in a Western blot. B, Serum-starved control and chlorate-treated human umbilical vein endothelial (HUVE) cells were stimulated for 10 minutes at 37°C with gremlin (100 ng/mL), decorated with anti-VEGFR2 or anti-P-VEGFR2 antibodies (in green) followed by nuclear counterstaining with 4',6-diamidino-2-phenylindole (in blue), and observed under a confocal microscope. C, Serum-starved control and chlorate-treated HUVE cells were stimulated for 10 minutes at 37°C with gremlin (100 ng/mL) and increasing concentrations of heparin. Then, cell lysates (50 μg) were analyzed by Western blotting with anti-P-VEGFR2, anti-P-ERK1/2, or anti-p38 antibodies. D, Serum-starved control and chlorate-treated HUVE cells were incubated for 10 minutes at 37°C with gremlin (100 ng/mL) and increasing concentrations of heparin. Then, cell lysates (50 μg) were analyzed by Western blotting with anti-P-VEGFR2 antibody. Uniform loading of the gels was confirmed by incubation of the membranes with antibodies directed against the unphosphorylated form of the antigen. The data are representative of 2 or 3 independent experiments that gave similar results.
whereas K5-OS(H) and K5-OS(L) were less effective (IC50 of 0.5 and 1.0 μg/mL, respectively). K5-NS and unmodified K5 did not exert a significant competition.

Then, the most active derivative K5-N,OS(H) was tested for the capacity to affect the angiogenic activity exerted by gremlin in vitro and in vivo. As shown in Figure 9B and 9C, K5-N,OS(H) inhibits the binding of gremlin to the extracellular domain of VEGFR2 in a competitive ELISA and its ability to stimulate the sprouting of HUVE cell aggregates in fibrin gel. Also, K5-N,OS(H) exerts a potent inhibitory activity on blood vessel formation triggered by gremlin delivered on the top of the chick embryo chorioallantoic membrane via an alginate pellet implant (Figure 9D and 9E). Accordingly, K5-N,OS(H) inhibits the angiogenic activity exerted by gremlin in a tumor-driven ex vivo angiogenesis assay in which fibrin-embedded human umbilical cord artery rings68 are grown in the presence of heparin (1 μg/mL), washed with PBS or with PBS plus 1.5 mol/L NaCl (salt wash).

Figure 7. Characterization of chlorate-treated human umbilical vein endothelial (HUVE) cells. A, HUVE cell monolayers were left untreated or treated with chlorate. Then, cells were incubated for 2 hours at 4°C with gremlin (100 ng/mL) in the absence or in the presence of heparin (1 μg/mL), washed with PBS or with PBS plus 1.5 mol/L NaCl (salt wash). Then, cell lysates were analyzed by Western blotting with an anti-gremlin antibody. Uniform loading of the samples was confirmed by incubation of the membrane with an anti-tubulin antibody. B, Control and chlorate-treated HUVE cells were assessed for cell viability by the tetrazolium salt (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) colorimetric assay according to standard procedures. C, Control and chlorate-treated HUVE cells were stimulated for 10 minutes with phorbol myristate acetate (PMA) (10 ng/mL). Then, 50 μg- aliquots of cell lysates were analyzed by Western blotting with an anti-P-extracellular signaling–regulated kinase (P-ERK1/2) antibody. Uniform loading of the gel was confirmed by incubation of the membrane with an anti-ERK2 antibody. D, Control and chlorate-treated HUVE cells were allowed to adhere onto substrate-immobilized gelatin for 2 hours at room temperature, and attached cells were photographed under an inverted microscope.
Heparan sulfate proteoglycans (HSPGs) mediate the angiogenic activity of gremlin in endothelial cells (ECs). A and B, Control (black bars) and chlorate-treated (open bars) human umbilical vein endothelial (HUVE) cells were assessed for their capacity to migrate in response to gremlin or phorbol myristate acetate (PMA) in a modified Boyden chamber assay (A) or by time-lapse videomicroscopy (B). C, Control (black bars) and chlorate-treated (open bars) HUVE cells were incubated with gremlin in the absence or in the presence of heparin (0.5 μg/mL) and assessed for their motility by time-lapse microscopy. D, HUVE cell spheroids obtained from control, chlorate-treated, or heparinase II–treated cells were embedded in fibrin gel and incubated with gremlin. Formation of radially growing cell sprouts was evaluated after 24 hours of incubation. E, Representative microphotographs of control and chlorate-treated HUVE cell spheroids stimulated with gremlin. In A to D, each data are the mean±SD of 2 or 3 independent experiments. Significantly different from gremlin alone under the same experimental conditions: *P<0.004, **P<0.001. FCS, fetal calf serum.

Discussion

This is the first characterization of the interaction of gremlin with heparin/HSPGs and of its role in mediating the angiogenic response triggered by this noncanonical VEGFR2 ligand in ECs. Compared with other sulfated GAGs, heparin (SO$_3^-$/COO$^-$=2.14)$^{34}$ binds gremlin with high affinity ($K_D=20$ nmol/L). In HS (SO$_3^-$/COO$^-$=1.0),$^{34}$ sulfate groups are arranged in high-charge-density clusters intercalated with low-charge-density regions,$^{49}$ thus resulting in a lower gremlin-binding activity. On the other hand, HS is a much more effective gremlin interactor than dermatan sulfate, chondroitin-4-sulfate, or chondroitin-6-sulfate, even though they all share a similar charge, no significant binding being observed with the nonsulfated hyaluronic acid and K5 polysaccharide. Also, similar to other heparin-binding angiogenic proteins,$^{10,14}$ the affinity for gremlin is affected by the size of the polysaccharide chain. Indeed, whereas low-MW heparin tinzaparin binds gremlin with an affinity similar to unfractionated heparin, very-low-MW heparins showed a reduced interaction, and a negligible binding was observed for 2.1-kDa heparin. Interestingly, when compared with different low-MW heparin preparations, tinzaparin has demonstrated significant antiangiogenic and anticancer properties.$^{44}$ Several sets of experimental evidence demonstrate that physicochemical and biological properties of different low-MW heparins are not alike.$^{44}$ Even though additional studies are required to fully characterize the interaction of gremlin with different heparin preparations, our data clearly indicate that the degree of sulfation, charge distribution, disaccharide composition, and size are all important in determining the capacity of sulfated GAGs to bind gremlin.

Previous observations had shown that fibroblast growth factor-2 binds avidly to a pentasaccharide region in which N-sulfate groups and a single 2-O-sulfate group are essential for interaction. In contrast, N-sulfation and 6-O-sulfation are crucial for VEGF-A165 binding, whereas 2-O-sulfate groups are less essential.$^{50}$ Our data demonstrate that selective 2-O-, 6-O-, or N-desulfation significantly reduces gremlin/heparin interaction, indicating that they are all required for high affinity binding. These findings support the hypothesis that heparin-binding cytokines may bind sulfated GAGs in a distinct manner. Even though specific factor binding sequences may be hidden in heparin because of its high degree
of sulfation, the high heterogeneity in HS structure allows a more refined tailoring of selective binding regions that may influence the biological activity and bioavailability of heparin/HS-binding growth factors.

In keeping with its heparin/HS binding capacity, gremlin binds the surface of CHO-K1 cells but not of GAG-deficient A745 CHO-K1 cells. Accordingly, gremlin binds HSPGs exposed on the surface of ECs and deposited in the subendothelial ECM, the binding being prevented by HS chain digestion with heparinase II or by inhibition of HS chain sulfation following EC treatment with chlorate. Similar to the canonical heparin-binding VEGFR2 ligand VEGF-A165, HSPG interaction appears to be essential for the binding of gremlin to VEGFR2 and receptor activation in ECs. Indeed, chlorate treatment or heparinase II digestion both hamper gremlin/VEGFR2 interaction in HUVE cells. Accordingly, VEGFR2 autophosphorylation triggered by gremlin is abolished following HS digestion or undersulfuration, yet is restored by the addition of exogenous heparin. Similarly, inhibition of HSPG interaction suppresses the ability of gremlin to induce the activation of intracellular signaling pathways in ECs, including ERK1/2 and p38 mitogen-activated protein kinase phosphorylation. Also, it inhibits EC chemotactic migration, sprouting, and motility, all activities that are mediated by a productive interaction of gremlin with VEGFR2.20

NRP-1 is a transmembrane glycoprotein involved in the interaction with a plethora of binding partners, including class-3 semaphorin receptors, heparin-binding growth factors, and HSPGs.51 Also, NRP-1 core protein is modified with either a single HS or chondroitin sulfate chain depending on the expressing cell type, thus increasing the complexity of NRP-1 extracellular interactions.52 Several sets of experimental evidence indicate that the binding of mammalian heparin-binding VEGF isoforms to HS and NRP-1 is interdependent.4 Gremlin does not contain the amino acid sequence CDKPRR present in the VEGF-A165 carboxyl terminus and involved in the interaction with b1b2 ectodomains of the NRP-1 core protein.51 Even though we could not rule out the possibility that gremlin may bind the HS chain of NRP-1 in target cells, our data indicate that, at variance with VEGF-A165, gremlin does not interact with recombinant NRP-1 and does not trigger the formation of a VEGFR2/NRP-1 complex in ECs. Also, the structural bases for heparin/HSPG interaction may differ between VEGF isoforms and gremlin. Indeed, VEGFs bind HS via a highly basic carboxyl-terminal region distinct from the cystine-knot domain of the protein and encoded by alternatively spliced exons.4 At variance, gremlin may interact with heparin/HS via a linear basic patch along the third loop of its cystine-knot domain, as suggested by homology modeling prediction studies based on the cystine-knot BMP antagonist sclerostin.31 32 Thus, gremlin differs from canonical heparin-binding VEGFs in the ability to co-opt VEGFR2 coreceptors and appears to exert a VEGFR2-dependent proangiogenic activation of ECs with mechanisms that are, at least in part, different from VEGFs. For instance, HSPGs contribute to VEGF-triggered angiogenesis by a mechanism dependent on the modulation of α5β1 integrin expression on the EC surface.54 At variance with VEGF-A, gremlin does not upregulate the expression of α5 integrin in ECs (data not shown), further underscoring the different mechanism of action of the 2 angiogenic factors.

Chemical or enzymatic modifications of the E. coli capsular K5 polysaccharide allow the production of “biotechnological” semisynthetic heparin/HS-like compounds with selected chemical features, with potential implications in the therapy of various human pathologies, including thromboembolic diseases, viral infection, and neoplasia.16 55 Here we evaluated the gremlin-binding capacity of a series of specific N-, O-, or N,O-sulfated nonanticoagulant K5 derivatives.33 Among the compounds tested, highly sulfated K5-N,OS(H) binds gremlin with higher affinity compared with the corresponding derivative K5-OS(H) devoid of sulfate group in the N position but bearing an identical total negative charge (SO3/COO of 3.8 for both compounds34). K5-OS(H) consists of the virtually homogeneous repeat of GlcA2,3SO3−-GlcNAc3,6SO3− disaccharide units, whereas most of K5-N,OS(H) sequences are represented by GlcA2,3SO3−-GlcNSO−,6SO3− disaccharide units.33 Our data indicate that the requirement for N-sulfation present in K5-N,OS(H) cannot be fully overcome by the high density charge present in the highly O-sulfated K5-OS(H). Similarly, low sulfated K5-N,OS(L) binds gremlin more efficiently than
K5-OS(L), even though both compounds have a \( \text{SO}_3^-/\text{COO}^- \) value of 1.7.\(^{33}\) On the other hand, K5-NS devoid of sulfate groups in \( O \) positions (\( \text{SO}_3^-/\text{COO}^- = 1.0 \)) does not have a significant interaction with gremlin. Thus, in agreement with the results obtained with selectively desulfated heparins, \( N \) and \( O \)-sulfate groups are required for gremlin interaction with sulfated K5 derivatives.

Gremlin exerts both BMP-dependent and BMP-independent functions in different physiopathological conditions by inhibiting BMP-mediated TGF-\( \beta \) receptor activation or by a direct activation of VEGFR2 signaling, respectively. In this respect, it is interesting to note that a complex, not yet fully elucidated cross-talk may exist among TGF-\( \beta \)/BMP, VEGFR2, and canonical and noncanonical Wnt (Wnt) signaling pathways during angiogenesis,\(^{56}\) making unpredictable the impact of the modulation of a single agonist/antagonist on the neovascularization process in different pathological conditions. Here, we demonstrate that K5-N,OS(H) is a potent antagonist of gremlin, inhibiting VEGF2 interaction and angiogenic activity in vitro and in vivo. Also, K5-N,OS(H) may prevent gremlin-driven tumor neovascularization, as indicated by its capacity to suppress the angiogenic potential of gremlin-overexpressing human tumor PANC-1 cells in an ex vivo human artery ring assay. Previous observations had shown the ability of K5-N,OS(H) to inhibit the angiogenic activity of the heparin-binding molecules fibroblast growth factor-2 and HIV-1 Tat protein\(^{16}\) and to reduce tumor invasion by hampering heparanase activity\(^{16}\) and tumor metastasis.\(^{37}\) Thus, K5-N,OS(H) may represent the basis for the design of novel nonanticoagulant biotechnological heparins endowed with a multtarget-antiproliferative action by interacting with different angiogenic growth factors and limiting tumor cell invasion and metastasis. Also, preclinical and clinical studies identify gremlin as a potential therapeutic target in human fibrotic diseases characterized by alterations of TGF-\( \beta \)/BMP signaling and vascular defects in different organs as liver,\(^{58}\) kidney,\(^{24}\) and lung.\(^{23}\) Given the role of gremlin in physiological development and different pathological conditions, additional preclinical in vivo studies will be required to define the therapeutic capability and possible side effects of gremlin antagonists, including K5-N,OS(H) and its derivatives.

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Disclosures

None.

References

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