Semaphorin-3C signals through Neuropilin-1 and PlexinD1 receptors to inhibit pathological angiogenesis

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Abstract

Retinopathy of prematurity causes visual impairment due to destructive neoangiogenesis after degeneration of the retinal microvasculature. This study was aimed at analyzing whether local delivery of Semaphorin-3C (Sema3C) suppresses pathological retinal angiogenesis. Sema3C exerted potent inhibiting effects in cellular models of angiogenesis. In an endothelial cell xenotransplantation assay, Sema3C acted primarily on immature microvessels by inducing endothelial cell apoptosis. Intravitreal administration of recombinant Sema3C disrupted endothelial tip cell formation and cell–cell contacts, which led to decreased vascular bed expansion and vessel branching in the growing retinal vasculature of newborn mice, while not affecting mature vessels in the adult retina. Sema3C administration strongly inhibited the formation of pathological pre-retinal vascular tufts during oxygen-induced retinopathy. Mechanistically, Sema3C signaled through the receptors Neuropilin-1 and PlexinD1, which were strongly expressed on vascular tufts, induced VECadherin internalization, and abrogated vascular endothelial growth factor (VEGF)-induced activation of the kinases AKT, FAK, and p38MAPK. This disrupted endothelial cell junctions, focal adhesions, and cytoskeleton assembly resulted in decreased cell migration and survival. Thus, this study identified Sema3C as a potent and selective inhibitor of pathological retinal angiogenesis.

Keywords angiogenesis; semaphorin; retinopathy of prematurity; Sema3C

Subject Categories Cardiovascular System; Vascular Biology & Angiogenesis

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Introduction

Angiogenesis is a prerequisite for embryonic development. Reactivation of this embryo–fetal program occurs in the adult, for example, during wound healing, tissue growth, or the menstrual cycle. However, excessive or uncontrolled vessel sprouting is seen in a number of diseases, most notably malignant tumors and proliferative eye diseases (Chung & Ferrara, 2011). Several drugs, which interfere with vascular endothelial growth factor (VEGF) signaling, have been approved for the treatment of advanced carcinomas and the wet form of macular degeneration (Barakat & Kaiser, 2009; Welti et al., 2013). While these drugs only slow down tumor growth for a while, they have revolutionised treatment of the wet form of age-related macular degeneration. In this disease, the local application of anti-VEGF compounds prevents further outgrowth and leakage of new, immature blood vessels.

Similar therapeutics would also be highly desirable to treat preterm babies suffering from retinopathy of prematurity (ROP). The human retinal vasculature develops between the 16\textsuperscript{th} week of gestation and birth. Therefore, preterm born babies who receive intensive neonatal care therapy with increased oxygen supply are at high risk of developing ROP, which is the major ocular disorder of the neonate and the dominant cause of visual impairment in childhood. In phase 1 of ROP, relative hyperoxia and decreased VEGF levels cause vessel regression in the retina. Phase 2 of ROP involves relative hypoxia and exaggerated VEGF, Epo, and Angiopoietin-2 secretion from neurons and astrocytes leading to excessive neovascularization, which is often misguided toward the vitreous body and lens. This can cause hemorrhage and retinal detachment leading to vision loss (Sapieha et al., 2010). The use of anti-VEGF agents is an emerging treatment for acute ROP, but studies analyzing long-term effects are still rare (Mutlu & Sarici, 2013).

Vascular endothelial growth factor acts through VEGF receptors and Neuropilin receptors on endothelial cells. It induces the formation of tip cells, which are motile and guide new vessel sprouts. In
addition to VEGF, the further growth of new vessel sprouts is also regulated by attractive and repulsive cues. This is similar to neurogenesis, where the outgrowth of axons is coordinated by neuronal guidance molecules of the Netrin, Slit, Ephrin, and Semaphorin families (Larrivee et al, 2009; Adams & Eichmann, 2010).

Class-3 Semaphorins (Sema3A to Sema3G) are secreted proteins that were originally identified as factors controlling axonal branching and pathfinding in the nervous system. They signal mainly through a receptor complex in which a Neuruplin (Nrp-1 or Nrp-2) associates with a Plexin receptor. The Plexin receptors modulate activity of GTPases, which cause changes in the cytoskeletal assembly and cell adhesive forces (Kruger et al, 2005; Negishi et al, 2005).

Among Sema3 proteins, Sema3C is still a poorly understood molecule. It is expressed during development in neurons, neural crest cells, myocardium, and kidney (Feiner et al, 2001; Niquille et al, 2009; Reidy & Tufro, 2011). Sema3C-deficient mice show a defective cardiac outflow tract (Feiner et al, 2001), and decreased ureteric bud branching (Reidy & Tufro, 2011). The cardiac phenotype is partially phenocopied by loss of PlexinD1 expression in endothelial cells (Gütler et al, 2004), suggesting an essential role of paracrine Sema3C signaling during cardiogenesis.

The function of class-3 Semaphorin proteins is regulated by furin or furin-like endoproteinasises. These enzymes can recognize two or three consensus RXK/RR motifs in the Plexin-Sema-Ig (PSI) domain and C-terminal basic domain (Adams et al, 1997). Metalloproteinases also cleave Sema3C at the C-terminus, and it was assumed that this cleavage of the last 13 amino acids releases Sema3C from the extracellular matrix (Esselens et al, 2010). The consequence of cleavage within the PSI domain that significantly shortens the protein (Sema3Cp60 isoform) is not fully understood, but the repulsive activity of Sema3C during the outgrowth of sympathetic axons is increased when this processing is blocked (Adams et al, 1997).

Sema3C is not expressed by human endothelial cells (Serini et al, 2003), and there are no data indicating a role for physiological angiogenesis. However, Sema3C expression is high in gastric and biliary epithelial and cancer cells (Adams et al, 2003; Miyato et al, 2012). These findings together with the known anti-angiogenic functions of Sema3A and Sema3F (Gu & Giraud, 2013) indicate a potential role of Sema3C on blood vessels under pathological conditions. This study was aimed at elucidating how Sema3C regulates angiogenesis and exploring its potential for anti-angiogenic therapy.

Results

Sema3C inhibits endothelial tube formation and sprouting angiogenesis

We first evaluated Sema3C expression in different human primary endothelial cells and mural cells. We found that endothelial cells freshly isolated from the vein of umbilical cord (HUVEC) and human brain microvascular endothelial cells express very little Sema3C mRNA. On the contrary, mural cells in culture express higher levels of Sema3C mRNA and protein (Supplementary Fig S1A and B). Based on these finding, we used brain-derived pericytes as producer cells for Sema3C-conditioned medium. We generated adenoviral expression vectors for full-length Sema3C and the two isoforms: the truncated Sema3CA13 that lacked the last 13 amino acids in the C-terminal basic domain and corresponds to metalloproteinase-dependent cleavage (Esselens et al, 2010), and Sema3Cp60 that resembles furin cleavage within the PSI domain (Fig 1A). These proteins were expressed in pericytes (Supplementary Fig S1C). Forced expression of Sema3C and Sema3CA13 cDNA led to secretion of proteins with the same molecular weight (81 kDa) (Fig 1B). Sema3C was further cleaved into the p60 isoform and this could be prevented by the furin convertase inhibitor Decanoyl-RVKR-CMK but not by the metalloproteinase inhibitor Batimastat (Supplementary Fig S1D).

To assess the effect of Sema3C proteins in angiogenesis, HUVECs were treated with freshly prepared conditioned medium. VEGF induced the formation of a tube-like HUVEC network on Matrigel. The treatment with Sema3C- and Sema3CA13-conditioned medium strongly inhibited endothelial tube formation. However, Sema3Cp60 had no effect (Fig 1C).

Next, we validated these findings in a three-dimensional angiogenesis assay, which better resembles the process of endothelial sprouting. VEGF, VEGF-C, and FGF2 stimulated the outgrowth of capillary-like structures, while Sema3C- or Sema3CA13-conditioned medium significantly impaired endothelial sprouting and tube elongation (Fig 1D and Supplementary Fig S1E). HUVEC spheroids treated with Sema3Cp60 showed unaltered sprouting (Fig 1D). This indicated that cleavage within the PSI domain abrogated Sema3C functions.

In order to rule out the possibility that other factors within the conditioned media may also be responsible for the observed effects, we employed a purified, recombinant human Sema3C-Fc fusion protein (Supplementary Fig S1F). This protein carried point mutations that prevent cleavage within the PSI domain and has a deletion of C-terminal basic amino acid residues. Application of recombinant Sema3C-Fc, but not IgG-Fc, to HUVEC spheroids similarly inhibited sprouting angiogenesis (Supplementary Fig S1F). Thus, the data indicated that Sema3C acted as a potent anti-angiogenic protein in vitro.

Sema3C inhibits vessel formation of transplanted endothelial cells in mice

We next examined whether Sema3C could also inhibit de novo vessel formation in vivo. We applied an established model using HUVEC spheroids transplanted within a growth factor-rich matrix into nude mice (Alajati et al, 2008; Wustehube et al, 2010). We modified the established protocol to co-graft HUVEC and human brain-derived pericytes (Fig 2A). The pericytes were stably transduced with lentiviral expression vectors for Sema3C or GFP as control. The plugs were dissected after 4 weeks, and HUVEC-derived vessels were stained with an antibody that specifically recognizes human CD34. The data showed that pericyte-secreted Sema3C strongly inhibited vessel formation (Fig 2B and C). However, the remaining vessels were well covered by αSMA-positive mural cells (Fig 2D and E) and perfused to a similar extent as those in the control group (Fig 2F). This experiment demonstrated that Sema3C was able to inhibit de novo vessel formation in vivo.
Figure 1. Sema3C inhibits endothelial tube formation and sprouting angiogenesis.
A Sema3C domain structure with two consensus motifs for furin and furin-like proteases. Sema3CA13 and Sema3Cp60 expression vectors were generated by replacing the amino acids in the consensus motifs with stop codons (asterisk).
B Conditioned medium was collected from pericytes 48 h after adenoviral transduction. Western blotting verified the secretion of Sema3CA13 (81 kDa) and Sema3Cp60 (60 kDa) isoforms.
C HUVECs were suspended in conditioned medium with VEGF (25 ng/ml) and plated on Matrigel. Branch points of the capillary-like network were counted after 18 h. n = 4 independent experiments; scale bar, 200 μm. Control vs. Sema3C, *P = 0.0375 and control vs. Sema3CA13, *P = 0.0162.
D HUVEC spheroids in collagen were treated with conditioned medium with or without VEGF (25 ng/ml). After 24 h, the cumulative sprout length per spheroid was quantified. Data are representative of three independent experiments with ten spheroids per group. Scale bar, 100 μm. Control vs. Sema3C (− VEGF), *P = 0.0276; control vs. Sema3CA13 (− VEGF), *P = 0.0443; control vs. Sema3C (+ VEGF), *P = 0.028 and control vs. Sema3CA13 (+ VEGF), *P = 0.032.
Data information: (C, D) Mean ± s.e.m; n.s., non-significant using paired Student’s t-test.

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Figure 2.

A

HUVEC

Sema3C- or GFP-expressing pericytes

Mixed spheroids

Spheroids in a fibrinogen/thrombin/Matrigel were s.c. implanted into SCID mice

28 days

Analysis

B

Control

Sema3C

C

Vessel area [%]

Control  Sema3C

2.5  *

1.5

1.0

0.5

0.0

D

Control

Sema3C

E

Vessel coverage [%]

Control  Sema3C

100  n.s.

80

60

40

20

0

F

Vessel perfusion [%]

Control  Sema3C

50  n.s.

30

20

10

0

Figure 2.
The relative VEGF-164 mRNA levels in whole retinal explants were favor stalk cell selection. Consistently, the retinae showed an aberrantly decreased number of tip cell-enriched genes not DLL4, was strongly diminished upon Sema3C treatment. Sema3C also led to slight induction of the Notch target genes EFNB2, HEY1, and DLL4 (Supplementary Fig S2C). Increased Notch activity is well known to inhibit tip cell differentiation (Adams & Eichmann, 2010); therefore, Sema3C appeared to favor stalk cell selection. Consistently, the retinae showed an aberrant vascular network with enlarged intercapillary spaces and fewer vessel branches after Sema3C treatment (Fig 3Aii, iv and E). Importantly, the astrocytes that form a template under the vascular bed and that secrete VEGF were not affected by Sema3C (Fig 3F). The relative VEGF-164 mRNA levels in whole retinal explants were not significantly altered between control and Sema3C treatment (1.099 ± 0.5 vs. 1.20 ± 0.37; n = 4). As such, Sema3C specifically inhibited the formation of new vascular sprouts during retinal angiogenesis.

In a second experiment, we investigated whether such a short-term treatment would also affect mature blood vessels in the eye of adult mice. We injected recombinant Sema3C protein into the vitreous humor of 6-week-old mice and analyzed the retinae 2 days later. This manipulation did not change the morphology of the fully developed vascular network at all (Fig 3G), indicating that Sema3C acted predominantly on immature blood vessels.

This was further supported by the finding that Sema3C was unable to disrupt a HUVEC monolayer cultured on Xellulin (Supplementary Fig S2D), a hydrogel matrix that allows long-term culture of endothelial cells in a quiescent state.

Sema3C inhibits adherens junction integrity leading to endothelial cell death

The observed anti-angiogenic effects of soluble Sema3C in the growing vasculature prompted us to investigate its cellular mechanisms, in particular VEGF-induced endothelial cell proliferation and survival. HUVECs were treated with Sema3C-conditioned medium or control-conditioned medium from GFP-expressing pericytes. Sema3C did not interfere with endothelial cell proliferation as measured by BrdU incorporation into newly synthesized DNA (Fig 4A). Consistently, the activation of extracellular signal-regulated kinase (ERK), which promotes endothelial cell proliferation, was not changed by Sema3C treatment (Supplementary Fig S3A). However, Sema3C impaired endothelial cell survival as evidenced by increased levels of active caspase-3 and caspase-7 in HUVEC (Fig 4B). The elevated endothelial cell apoptosis rate was accompanied by a marked decrease in phosphorylated AKT (Fig 4C and D).

Figure 2. Sema3C inhibits vessel formation in a xenotransplantation model.

- A Scheme of the experimental procedure. Pericytes were lentivirally transduced with Sema3C or GFP expression vectors. HUVEC/pericyte spheroids (1:1 ratio) were injected s.c. into SCID mice.
- B Microvessels within the plugs stained against human-specific CD34. Cell nuclei were stained with DAPI. Scale bar, 100 μm.
- C The area of CD34-positive microvessels was quantified and normalized to the total area of plug. Results represent the means ± s.e.m of 5 plugs (control) and 7 plugs (Sema3C), *P = 0.018.
- D Vessel coverage was assessed by sMA staining. Scale bar, 100 μm.
- E Vessel coverage was quantified as the ratio of endothelial cell-associated mural cells to total amount of vessels (n = 5 [control] and n = 7 [Sema3C]).
- F Vessel perfusion was evaluated by i.v. injection of FITC-lectin (0.15 mg) 30 min before mice were sacrificed. The data show the percentage of perfused vessels in the plug and normalized to total vessel area (n = 5 [control] and n = 7 [Sema3C]).

Data information: (C, E, F) Mean ± s.e.m; n.s., non-significant using unpaired Student’s t-test.

Figure 3. Intravitreal injection of Sema3C inhibits retinal vessel formation.

- A C57BL/6N mice pups at P3 received 1 μg of mouse Sema3C-Fc (right eye) or control IgG-Fc (left eye). The retinal vasculature was examined at P5 by whole-mount immunostaining and CD31 immunofluorescent staining. Representative images show the total retinal vasculature and sprouting tips at the vascular front normalized to control IgG-Fc treatment, ***P = 0.0009 (control) and ***P = 0.0002 (Sema3C)).
- B (C) Vessel density was quantified by CD31-immunostained vessels and normalized to the total vessel area. **P = 0.0001. (Mean ± s.e.m; n = 23 retinae in each group from four independent experiments; unpaired Student’s t-test.)
- D, E (D) Number of sprouting tips per retina, *P < 0.0001. (E) Branch points in the capillary plexus represented as the ratio to the control IgG-treated retinae, ***P < 0.0001. (Mean ± s.e.m; n = 18 retinae in each group from three independent experiments; unpaired Student’s t-test.)
- F Pups received same treatment as described in (A). The astrocyte layer was examined at P5 by GFAP staining. Scale bar, 100 μm (upper panel) and 20 μm (lower panel). Representative images of six retinae per group.
- G Six-week-old C57BL/6N male mice were intravitreally injected with Sema3C-Fc (right eye) or control IgG-Fc (left eye), and retinae were collected 2 days later. The retinal vessel network was examined with CD31 immunostaining. Scale bar, 100 μm (upper panel) and 20 μm (lower panel). Representative images of four retinae per group.
Figure 3.

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Figure 4.

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AKT acts downstream of VEGF signaling and protects endothelial cells from apoptosis (Gerber et al., 1998). Therefore, the data showed that Sema3C mainly suppresses endothelial cell survival but not cell proliferation.

We also observed pronounced changes of endothelial cell morphology upon the addition of Sema3C-conditioned medium. This included the breakdown of adherens junctions with less colocalization of VE-cadherin and β-catenin. This was not observed by treatment with the p60 isoform (Supplementary Fig S3B). Also cell fractionation showed less VE-Cadherin at the cell membrane (Supplementary Fig S3C). To better visualize this, HUVECs were adenovirally transduced with GFP-tagged VE-cadherin and observed by time-lapse microscopy (Fig 4E and Supplementary Movies S1 and S2). VE-cadherin is an endothelial cell-specific adherens junction protein that associates with the VEGF receptor-2 (VEGFR2) at cell junctions. This interaction is essential for VEGF-induced cell survival signaling (Carmeliet et al., 1999). Treatment with control-conditioned medium did not alter the localization of VE-cadherin, which was predominantly found at cell-cell contacts. In contrast, Sema3C induced dissociation of VE-cadherin and VEGFR2 from the cell junctions and internalization into the cytosol. Subsequently, this resulted in the formation of intercellular gaps within the endothelial monolayer (Fig 4E and Supplementary Fig S3D).

VE-cadherin is integral to maintain vessel integrity. Intravitreal administration of recombinant Sema3C at postnatal day 5 led to a breakdown of VE-Cadherin at cell junctions as observed 6 h later (Fig 4F). This was accompanied by increased endothelial cell apoptosis (Fig 4G and H). Increased endothelial cell apoptosis after Sema3C delivery to endothelial cells was also observed in the HUVEC/pericyte xenotransplantation experiment in nude mice (Fig 2A). Notably, Sema3C overexpression in pericytes had no influence on pericyte cell death (Supplementary Fig S4A). However, increased rates of cleaved caspase-3 were detected in CD34-positive endothelial cells when co-grafted with Sema3C-secreting pericytes (Supplementary Fig S4B and C).

Sema3C impairs VEGF-induced endothelial cell migration through inhibiting focal adhesion kinase and p38MAPK signaling

VE-cadherin associates with endothelial cell migration through adherens junctions, which are linked to the actin cytoskeleton. To assess the effect of Sema3C on actin fibers, HUVECs were transduced with the fluorescent F-actin binding peptide LifeAct-RFP (Riedl et al., 2008) and imaged by time-lapse microscopy. Sema3C induced not only the internalization of VE-cadherin, but also the depolymerization of the F-actin bundles. This finally led to endothelial cell detachment from the extracellular matrix (Fig 5A and Supplementary Movies S3 and S4), which causes anoikis.

The findings prompted us to investigate endothelial cell migration since this process relies on dynamic changes of actin filaments and adhesive structures with the extracellular matrix. Sema3C treatment delayed the closure of a defined gap within a HUVEC monolayer, indicating decreased endothelial cell migration. This could not be rescued by the addition of 25 ng/ml VEGF (Fig 5B and C). Additionally, Sema3C treatment impaired cell adhesion and haptotaxis on different matrix proteins and poly-L-lysine (Supplementary Fig S5A and B).

VEGF signaling activates p38MAPK and focal adhesion kinase (FAK), which are both essential for endothelial cell migration (Rousseau et al., 1997; Avraham et al., 2003). VEGF-mediated phosphorylation of p38MAPK at residues threonine-180 and tyrosine-182 promotes the assembly of actin stress fibers. Sema3C significantly inhibited the activation of p38MAPK in HUVEC (Fig 5D). Sema3C also inhibited phosphorylation of FAK at tyrosine-397 (Fig 5E). This protein modification is required for its full activation, which leads to the recruitment of focal adhesion proteins like vinculin and paxillin. Vinculin associates with FAK in elongated dot-like structures and couples focal adhesions with actin fibers. This structure could be observed in HUVEC treated with control medium. In contrast, Sema3C almost abolished the formation of such focal adhesion complexes (Fig 5F). The recruitment of paxillin, another focal adhesion protein, was also disturbed by Sema3C treatment (Supplementary Fig S5C). Collectively, the data indicate that Sema3C impaired the formation of focal adhesions that are necessary for cell adhesion and migration.

Nrp-1 and Plexin-D1 act as receptors for Sema3C in endothelial cells

Lastly, we pursued experiments aimed at identifying the receptor(s) through which Sema3C exerts its functions on endothelial cells. Previous studies showed that Sema3C induces growth cone collapse...
Figure 5.

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in sympathetic neurons through both Nrp-1 and Nrp-2 receptors (Chen et al., 1997) and that PlexinD1 promotes the binding of Sema3C to Nrp-1 in COS cells (Gitter et al., 2004). PlexinA2 appeared as a potential receptor for Sema3C during heart development (Brown et al., 2001). To identify the functional receptors for Sema3C on endothelial cells, HUVECs were transfected with siRNA against PlexinD1 or lentivirally transduced with shRNA targeting Nrp-1, Nrp-2, or PlexinA2. qPCR analysis and Western blot analysis confirmed a sufficient downregulation of gene expression (Fig 6A and Supplementary Fig S6A and B). The cells were then treated with Sema3C-conditioned medium or recombinant protein for 30 min and changes of cell morphology were analyzed. Sema3C-conditioned medium or recombinant protein for 30 min and changes of cell morphology were analyzed. Sema3C-conditioned medium or recombinant protein for 30 min and changes of cell morphology were analyzed. Sema3C-conditioned medium or recombinant protein for 30 min and changes of cell morphology were analyzed.

Figure 5. Sema3C suppresses endothelial cell adhesion and migration by inhibiting FAK and p38MAPK signaling pathways.

A HUVECs were transduced with adenoviral vector expressing RFP-conjugated F-actin binding peptide (LifeAct). Time-lapse imaging of F-actin in HUVEC was obtained at 20× magnification (Supplementary Movies 53 and 54). Representative images are shown at 0 and 2 h. Arrowheads indicate detached cells after Sema3C treatment. Scale bar, 100 μm.

B Endothelial cell migration was determined using a scratch wound assay with a co-culture model. A wound was generated in a confluent HUVEC monolayer, and cells were cultured in basal medium with or without VEGF (25 ng/ml) in a 24-well plate. At the same time, the transwells containing Sema3C or GFP-expressing pericytes. Representative pictures show remaining gaps within the HUVEC monolayer. Scale bar, 200 μm.

C Migration distance was measured after 24 h and normalized to control treatment (mean ± s.e.m.; n = 3 independent experiments; control vs. Sema3C (−VEGF), *P = 0.021 and control vs. Sema3C (+VEGF), *P = 0.039 using unpaired Student's t-test).

D, E (D) Serum-starved HUVECs were treated with Sema3C or control-conditioned medium with or without VEGF (50 ng/ml) for 10 min. Cell lysates were subjected to SDS–PAGE and probed with anti-phosphoY1092/1093 YAP2 p38MAPK antibody. Quantification of p38MAPK activation was normalized to total p38MAPK and presented as the ratio to the cells with control medium without VEGF treatment. Control vs. Sema3C (−VEGF), *P = 0.046 and control vs. Sema3C (+VEGF), *P = 0.043 (E). HUVECs were treated as described before and the activation of FAK detected by anti-phosphoY939 YAP2 FAK antibody and normalized to the total amount of FAK. Control vs. Sema3C (−VEGF), *P = 0.033 and control vs. Sema3C (+VEGF), *P = 0.047. (Mean ± s.e.m.; n = 3 independent experiments; unpaired Student's t-test).

F Serum-starved HUVECs were treated with conditioned medium for 15 min. Focal adhesion proteins were examined with anti-phosphoY397 YAP2 FAK and anti-vinculin antibodies. F-actin was stained with Alexa Fluor 546-conjugated phalloidin. Merged images show the coupling of stress fibers with focal adhesion complexes in the cells treated with control medium. Inset, unphospho-Y397 YAP2 FAK and vinculin formed focal adhesions that were stained in an elongated dot-like structure (i). Lower panel (ii), Sema3C treatment induced F-actin disassembly and membrane ruffling. Phospho-Y397 YAP2 FAK and vinculin were randomly distributed at the peripheral membrane. Scale bars, 10 μm.

Source data are available online for this figure.

Figure 6. Sema3C, but not Sema3Cp60, acts through Neuropilin-1 and PlexinD1 to suppress endothelial cell adhesion.

A HUVECs were transduced with siRNA against PlexinD1 or control NS (non-silence) siRNA qPCR analyses and Western blotting were performed after 48 h (n = 3; paired Student's t-test, NS siRNA vs. PlexinD1 siRNA, ***P = 0.0004). Cell lysates were subjected to SDS–PAGE and probed with anti-PlexinD1.

B HUVECs were transduced with siRNA against PlexinD1 and plated on gelatin-coated coverslips. Confluent monolayers were treated with Sema3C-, Sema3Cp60- or control-conditioned medium for 30 min. Cells were fixed and stained with Alexa Fluor 488-conjugated phalloidin to visualize F-actin. Arrows indicate Sema3C-induced cell membrane ruffling. Scale bar, 20 μm. Cell repellence was quantified as the number of cells with membrane ruffling normalized to the number of nuclei (n = 5, mean ± s.e.m., paired Student's t-test, **P = 0.001).

C Silencing of Nrp-1 by lentiviral particles expressing shRNA. After culturing 72 h in selection medium, RNA was harvested for qPCR analysis. NS shRNA vs. Nrp-1 shRNA, #P = 0.037 and #P = 0.02, n = 3; mean ± s.e.m. Cell lysates were subjected to SDS–PAGE and probed with anti-Nrp-1.

D Depletion of Nrp-1 abolished the response to Sema3C. Sema3Cp60 had no effect on HUVEC adhesion. Arrows indicate membrane ruffling. Scale bar, 20 μm. Cell repellence was quantified as the number of cells with membrane ruffling normalized to the number of nuclei (n = 5, mean ± s.e.m., paired Student's t-test, ***P = 0.0003).

E HUVECs were treated with Sema3C-conditioned medium for 10 min and lysed, and Nrp-1 was immunoprecipitated. Bound Sema3C was detected by Western blotting.

F Schematic model of Sema3C signaling through Nrp-1 and PlexinD1 receptors in endothelial cells.

Source data are available online for this figure.

Sema3C isoforms for their ability to directly interact with Nrp-1. To this end, HUVECs were first treated with Sema3C-, Sema3CA13-, and Sema3Cp60-conditioned media and Nrp-1 was immunoprecipitated. The bound Sema3C proteins were detected by immunoblotting. This revealed that Sema3C, but not the Sema3CA13 and Sema3Cp60 isoforms, interacted with Nrp-1 (Fig 6E and Supplementary Fig S6F). This was in line with the result that the short Sema3Cp60 isoform had no anti-angiogenic activity (Fig 1C and D), suggesting that cleavage within the PSI domain abrogated binding of Sema3C to the Nrp-1 receptor.

Taken together, these results demonstrate that Sema3C signals via Nrp-1 and PlexinD1 receptors on endothelial cells to impair the integrity of adherens junctions, the assembly of actin cytoskeleton, and focal adhesions. The inhibition of these critical cellular functions strongly impairs VEGF-promoted endothelial cell survival, cell adhesion, and cell migration during angiogenesis (Fig 6F).

Sema3C suppresses pathological angiogenesis in an oxygen-induced retinopathy model

The human disease retinopathy of prematurity (ROP) is characterized by the rapid outgrowth of pathological blood vessels from the retina to the vitreous body. These pre-retinal tufts can lead to hemorrhage or retinal detachment, which often results in vision...
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Figure 6.

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Figure 7. Sema3C inhibits pathological angiogenesis in an oxygen-induced retinopathy model in mice.

A Scheme of the experiment. C57BL/6N mice pups at P7 with nursing mothers were subjected to hyperoxic conditions leading to vessel obliteration in the central zone of the retina. At P12, mice pups were returned to room air, which induces vessel regrowth but also pathological neovascularization, characterized by pre-retinal neovascular tufts. At P15, mice pups were injected intravitreally with 1 μg control IgG-Fc (left eye) and Sema3C-Fc (right eye). Retinae were analyzed 2 days later after intravitreal injection.

B RNA scope analyses showed Sema3C and PlexinD1 expression in pre-retinal tufts (OIR model, P19). The retinae were subsequently stained against collagen IV. Sema3C mRNA was detected in peri-endothelial cells. Scale bar: 20 μm.

C Nrp-1 expression in pre-retinal tufts (OIR model, P17). Abundant expression of Nrp1 in endothelial cells. Pericytes at vascular tufts were immunostained with anti-NG2. Scale bar, 20 μm.

D H&E staining shows neovascular tufts (arrows). Representative of five retinae per group in two independent experiments. Scale bar, 50 μm.

E Whole-mount CD31 immunostaining. The white line defines the whole retinal area, and red line outlines the vaso-obliterative zone. Inset image shows large numbers of peri-retinal tufts (i, arrow heads) in control IgG-Fc-treated retinae compared to Sema3C-treated retinae (ii). Scale bar, 500 μm (upper panel) and 100 μm (lower panel).

F The avascular area is presented as percentage of the whole retinal area (n = 10 retinae in each group; n.s., non-significant using paired Student’s t-test).

G The formation of pre-retinal tufts was quantified in the superficial layer of the retinal vasculature and normalized to whole retinal area (n = 6 retinae in each group; *P = 0.033 using paired Student’s t-test).
Discussion

Sprouting angiogenesis requires dynamic changes in endothelial cell shape, orientation, and adhesion. This is similar to the outgrowth of axons and it is not surprising that axonal guidance molecules coordinate both processes. This study showed for the first time that the secreted class-3 Semaphorin protein Sema3C inhibits sprouting angiogenesis by causing a substantial rearrangement of the actin cytoskeleton, breakdown of adherens junctions, and impairment of focal adhesion formation. Our data are in line with the functions of other semaphorins which act through Rho GTPases, FAK, and Src kinase family members in regulating cell spreading during neurite outgrowth, tumor progression, and vascular barrier integrity (Barberis et al., 2005; Franco & Tamagnone, 2008; Shimizu et al., 2008; Le Guellec et al., 2012). Here, we identified Sema3C functions that impeded VEGF-mediated endothelial cell survival and cell migration and finally caused cell detachment and apoptosis. This was seen in cellular models as well as in vivo. However, the few vessels that were formed by transplanted endothelial cells in the presence of Sema3C were covered by mural cells as in control animals. This is similar to the function of Sema3A, which acts anti-angiogenic but also increases pericyte coverage in a pancreatic tumor model (Maione et al., 2012). Notably, Sema3C did not influence cell survival of pericytes. It was reported that PlexinD1 is only expressed in the growing vasculature (within tumors) but not in non-malignant mature blood vessels (Roodink et al., 2009). This is in line with our expression data of Nrp-1 and PlexinD1 in the retina; hence, we suggest that Sema3C is only capable of driving such vessels into apoptosis that are undergoing angiogenesis and remodeling. This was further supported by the finding that quiescent HUVEC on Xellulin hydrogels were protected from Sema3C-induced cell repellation.

Exaggerated secretion of VEGF, Epo, Angiopoietin-2, and metabolic factors like succinate is responsible for the excessive growth of retinal blood vessels toward vitreous and lens during retinopathy of prematurity (ROP) (Sapieha et al., 2010). This disease is difficult to treat since on the one hand revascularization of the vaso-obliterative zone is desirable, whereas sprouting angiogenesis toward the vitreous body may lead to retinal detachment, hemorrhage, and blindness. The local injection of a single dose of recombinant Sema3C protein into the vitreous body in a mouse model of oxygen-induced retinopathy did not change the size of the vaso-obliterative zone within the retina. However, Sema3C effectively prevented the formation of pathological pre-retinal tufts that are responsible for major retinal damage. This inhibition was stronger than that observed after the local administration of Nrp-1 blocking antibodies (Oh et al., 2002). Importantly, Sema3C did not cause any changes in vascular morphology of mature retinal vessels, a crucial safety factor for any therapeutic agent targeting pathological angiogenesis. This may be due to the fact that our data and also previous study showed that the expression of both Sema3C receptors Nrp-1 and PlexinD1 is highly expressed in the growing vasculature (within tumors) but not in non-malignant mature blood vessels (Roodink et al., 2009). This is in line with our expression data of Nrp-1 and PlexinD1 in the retina; hence, we suggest that Sema3C is only capable of driving such vessels into apoptosis that are undergoing angiogenesis and remodeling (Fukushima et al., 2011). Therefore, our study suggests that Sema3C/Nrp-1/PlexinD1 signaling represents an attractive target for anti-angiogenic therapy.

Semaphorin signal transduction requires the association of Plexin and Neuropilin receptors. The presence of different receptor combinations mediates differential functions of the ligands. As such, Sema3C repels cortical neurons through Plexin/Nrp-1 but attracts neurons expressing Plexin/Nrp-2 receptor complexes (Ruediger et al., 2013). Though it was reported that Sema3C could act via both Neuropilins in sympathetic neurons (Chen et al., 1998), our data revealed that Sema3C selectively acted through PlexinD1 and Nrp-1 on endothelial cells. This is compatible with mouse knockout studies. Sema3C−/− mice die perinatally with cardiac defects (Feiner et al., 2001), which is similar to the phenotype observed in endothelial-specific Nrp-1-deficient or PlexinD1-deficient mice (Gu et al., 2003; Gilter et al., 2004). However, PlexinD1−/− or Nrp-1−/− mouse embryos show additional vascular patterning defects and several
other abnormalities (Kawasaki et al., 1999; Gitler et al., 2004). This may be due to the fact that both receptors also mediate the signals of other ligands including VEGF (Soker et al., 1998).

This study also demonstrated that proteolytic processing controls binding of Sema3C to its receptor. We found that the short Sema3Cp60 isoform had no anti-angiogenic activity, which is similar to the effect of cleaved Sema3B (Varshavsky et al., 2008). Mechanistically, we found that the Sema3Cp60 isoform did not bind to the Nrp-1 receptor indicating that the other domains of Sema3C are of utmost functional importance. The Sema3CA13 isoform, which may result from metalloprotease cleavage, showed, however, strong anti-angiogenic activity. This is similar to a study that describes strong activity of the Sema3CΔ13 isoform in a model of cancer cell migration (Esselens et al., 2010). Furin cleavage of semaphorins can lead to the exposure arginine/proline residues at the C-terminus, which are critical for ligand–receptor engagement (c-end rule) (Parker et al., 2013). The impaired binding of Sema3CΔ13, which lacks this C-terminal motif, to Nrp-1 in a co-immunoprecipitation experiment supports the importance of the C-end rule. However, it was also shown that the Ig domain of Sema3C could bind both Nrp-1 and Nrp-2 (Chen et al., 1998). Therefore, it may well be that Sema3CA13 executes its activity through binding to Nrp-1 with lower affinity, which could not be demonstrated in this experimental setup.

In summary, this study demonstrates for the first time that Sema3C can act as a potent anti-angiogenic agent that suppresses sprouting angiogenesis through dual inhibition of endothelial junction integrity and focal adhesion with the extracellular matrix. Moreover, administration of Sema3C effectively inhibits the formation of pathological vessel growth, such as pre-retinal neovascular tufts in the ROP model. Therefore, our study shows for the first time that Sema3CA13 is a promising agent to selectively interfere with pathological angiogenesis in the retina.

Materials and Methods

Plasmids, shRNA, and RNAi

Sema3C cDNA (NCBI DQ890817) vector from Genomic and Proteomics Core Facility of German Cancer Research Center was shuttled into pAD/CMV/V5 and pLenti6.2/V5-DEST (Invitrogen) by Gateway cloning for virus production. Sema3CA13 and Sema3Cp60 isoforms were generated using the QuickChange Lightening Site-Directed Mutagenesis Kit (Agilent Technologies) to introduce stop codons. PCR primers for the Sema3CA13 isoform: (5′-CAAGTAAAGGCCTCTC ATCAATAGTTAGTAAAAGTGAACAGGAGAAATCGT-3′), (5′-A ACTGATTCCTCGCTTCTTTTACTTNTAATTATGAGGCCTCTT TAACCTTTG-3′); and for Sema3Cp60: (5′-GAAACGAGGAGACCGA AATAATTGGTGGACATGAAACACAC-3′) and (5′-GTGGGTTCCT ATGTCCTACATATTACCTTGCCTCCTCGTTCT-3′). The translated protein sequences are shown in Fig 1A. Control siRNA (AM4636) and siRNA against human plexinD1 (AM16708, clone ID: 108678) were from LifeTechnologies. Transfection of HuVEC was performed with Oligofectamine according to the transfection protocol (Invitrogen). siRNA against Nrp-1 (clone ID: 16466, 333723) and Nrp-2 (clone ID: 34687, 176015) were from Thermo Scientific. shRNA against PlexinA2 (clone ID: 389145, 389146) was from Dharmacon.

RNA isolation and qPCR analysis

RNA was purified with the RNeasy Kit (Promega) and transcribed into cDNA (High Capacity cDNA Reverse Transcription Kit; Life Technologies). Real-time PCR was performed using the STEPone-Plus real-Time PCR system (Applied Biosystems). GAPDH and OAZ1 served as housekeeping genes for normalization. Primer sequences are listed in Table 1.

Cell culture

HBMEC (Cell Systems) were maintained in BMEC growth media (PELO Biotech). HUVECs were grown and maintained until passage 5 in Endopan3 Growth Medium containing supplements (Pan-Biotech). Human brain pericytes (ScienCell Research Lab) were cultured in pericyte medium containing supplements (SciCell Research Lab) and used between passages 3 and 7. Sema3C-conditioned medium was prepared 24 h after adenoviral infection of pericytes. The medium was replaced with Endopan3 basal medium.

Table 1. Primer sequences for real-time PCR analysis.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Forward Primer</th>
<th>Reverse Primer</th>
</tr>
</thead>
<tbody>
<tr>
<td>APLN</td>
<td>forward 5′-3′</td>
<td>reverse 5′-3′</td>
</tr>
<tr>
<td>PDCFB</td>
<td>forward 5′-3′</td>
<td>reverse 5′-3′</td>
</tr>
<tr>
<td>ESM1</td>
<td>forward 5′-3′</td>
<td>reverse 5′-3′</td>
</tr>
<tr>
<td>ANGPT2</td>
<td>forward 5′-3′</td>
<td>reverse 5′-3′</td>
</tr>
<tr>
<td>DLL4</td>
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</tr>
<tr>
<td>HEY1</td>
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<tr>
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</tr>
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<td>GAPDH</td>
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</tr>
<tr>
<td>OAZ1</td>
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</tr>
<tr>
<td>NRP1</td>
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<tr>
<td>PLXND1</td>
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<td>reverse 5′-3′</td>
</tr>
<tr>
<td>PLXNA2</td>
<td>forward 5′-3′</td>
<td>reverse 5′-3′</td>
</tr>
</tbody>
</table>
(0.5% FCS) for additional 48 h. Conditioned medium was centrifuged at 201 g for 5 min and directly applied to HUVEC.

**Antibodies and reagents**

Recombinant human VEGF165 (Ala27-Arg191, no. 293-VE), human Antibodies and reagents

HUVECs (6 l

**Immunofluorescent staining**

HUVECs (6 x 10^6 cells) were cultured on 0.2% gelatin-coated coverslips for 48 h. Cells were starved in basal medium containing 0.5% FCS for 16 h before treatment with conditioned medium. Cells were fixed in 4% PFA, permeabilized with 0.3% Triton X-100 in PBS, and incubated in 3% BSA in PBS for 1 h. pY397FAK, vinculin, or pY118

**Cell proliferation, migration, and apoptosis**

HUVECs were labeled by BrdU labeling solution (Roche) and treated with conditioned medium for 24 h. The level of incorporated BrdU was measured at OD_{450} with an ELISA reader (TECAN). To measure cell migration, HUVECs (20,000 cells) were seeded into the lower compartment of Matrigel. The assay was terminated after 18 h by adding 4% PFA. The vessel area was quantified as the CD34-positive area divided to the total plug area. Vessel coverage was quantified as the number of co-localized cleaved caspase-3 and CD34 stained cells divided to the vessel area. The next day, cells were observed using a wide-field microscope (Zeiss Cell Observer) equipped with an incubation chamber (5% CO_2 at 37°C). The serial images were analyzed using ImageJ (Fiji) software.

**Tube formation and spheroid-based sprouting angiogenesis**

Growth factor-reduced Matrigel (BD Biosciences) was added to 48-well plates to polymerize at 37°C for 30 min. HUVECs (2.5 x 10^4 cells) were suspended in 260 μl conditioned medium. Cells were mixed with VEGF (25 ng/ml), and 250 μl solution was plated on top of Matrigel. The assay was terminated after 18 h by adding 4% PFA.

**Endothelial cell xenotransplantation**

Human brain pericytes were stably transduced with lentivirus expressing Sema3C or GFP at MOI = 10. After selection with 10 μg/ml blasticidin, an equal number of pericytes and HUVECs were mixed to generate spheroids. These were mixed with 300 l conditioned medium with or without 50 ng/ml VEGF164 (R&D Systems). Cells were lysed with 0.5% NP-40 containing protease and phosphatase inhibitors (Roche) on ice. Denatured protein lysates were then applied for SDS–PAGE. For immunoprecipitation of endogenous Nrp-1, starved HUVECs were treated with conditioned medium for 10 min, washed with ice-cold PBS, and lysed with 0.5% NP-40 lysis buffer containing protease and phosphatase inhibitors on ice for 1 h. After centrifugation, the supernatant was pre-incubated with Protein-G PLUS-agarose (Santa Cruz) for 30 min. The supernatant was then incubated with 2 μg Nrp-1 antibody and Protein-G PLUS-agarose at 4°C overnight. Precipitated agarose beads were washed with ice-cold PBS and denatured in Laemmli sample buffer at 95°C for 5 min. Samples were then subjected to Western blotting.

**Time-lapse imaging**

HUVECs were transduced with viral vectors expressing GFP-VE-cadherin or RFP-LifeAct overnight, and 3 x 10^5 cells were seeded into ibidi wells (ibidi). The next day, cells were observed using a wide-field microscope (Zeiss Cell Observer) equipped with an incubation chamber (5% CO_2 at 37°C). The serial images were analyzed using ImageJ (Fiji) software.
was counted manually and normalized to the number of CD34-positive cells.

**Whole-mount in situ hybridization (ISH) and RNAscope**

For PlexinD1 ISH, the procedure was described previously (Fukushima et al., 2011). Briefly, enucleated retinas were fixed in 4% PFA at RT for 15 minutes and post-fixed in 4% PFA and 0.2% glutaraldehyde. Samples were hybridized with digoxigenin-labeled cRNA probes at 65°C overnight. The templates for cRNA probe: mouse PlexinD1, nt 5,184–6,055 of GenBank AK129175. The retinas were washed and incubated with alkaline phosphatase-conjugated sheep anti-digoxigenin antibody (Roche) at 4°C overnight. Bound antibodies were visualized using nitro blue tetrazolium and 5-bromo-4-chloro-3-indolylphosphate (NBT/BCIP, Roche) as substrate. Images were taken with an Axiosoplan 2 microscope (Zeiss) equipped with associated software (AxioVision, version 3.1; Zeiss). RNAscope ISH (Advanced Cell Diagnostics) was modified from the protocol described previously (Gross-Thebing et al., 2014). Retinas were fixed in 4% PFA for 2 h at RT. Samples were hybridized with RNA target probes (PlexinD1, nt 1,555–2,414 of NM_026376.3, Channel 1; Sema3C, nt 906–1,841 of NM_013657.5, Channel 3) overnight at 40°C. The retinae were further immunostained with anti-collagen IV antibody. Images were taken with a confocal microscope (Zeiss LSM 700).

**Retinal angiogenesis and oxygen-induced retinopathy**

Intravitreal injection of 1 μl recombinant mouse Sema3C fused to IgG-Fc (1 μg/μl) into the right eye and 1 μl of control human IgG-Fc into the left eye was performed in C57BL/6N mice pups at postnatal day P3 or in 6-week-old male mice. Two days later, the retinae were dissected and fixed in ice-cold methanol overnight. Samples were permeabilized in permeabilizing buffer (0.5% Triton X-100 and 10% donkey serum in PBS) for 1 h and blocked in blocking buffer (0.2% Tween-20, 10% donkey serum) for 1 h. Samples were then stained with anti-CD31 antibodies (1:100) in PBS overnight at 4°C. After washing with PBS (0.2% Tween-20), the retinae were incubated with Alexa Fluor 488 donkey anti-rat IgG (H + L) antibody for 1 h. After washing steps, samples were mounted using fluorescence mounting medium (DAKO). For evaluating cell apoptosis, retinae were collected 8 h after injection of recombinant proteins and fixed in ice-cold methanol. Samples were incubated with anti-cleaved caspase-3 (1:50) and anti-CD31 (1:100). For investigating endothelial cell junctions, retinae were fixed in methanol 6 h after injection. Z-stack images were acquired using a confocal microscope (Zeiss LSM 700).

Oxygen-induced retinopathy was induced by placing C57BL/6N mice pups with the nursing mother at day P7 in an Oxycycler (BioSpherix) with 75% oxygen for 5 days. Mice returned to room air at P12 and received intravitreal injections of 1 μg recombinant Sema3C (right eye) or IgG-Fc (left eye) at P15. The retinae were collected and analyzed 2 days later. All immunofluorescence images were taken using a confocal microscope (Zeiss LSM 700) and analyzed with Fiji software. For H&E staining, eyes were fixed in 4% PFA overnight, dehydrated, and embedded in paraffin. Staining was performed using 6-μm-thick slides. Images were taken with a wide-field microscope (Zeiss Cell Observer) at 20× magnification.

All animal experiments were performed according to the guidelines of the local institution (German Cancer Research Center Heidelberg) and the local government (Regierungspräsidium Karlsruhe, Germany).

**Supplementary information** for this article is available online: http://embomolmed.embopress.org

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**Author contributions**

WJY performed most of the experiments. JH conducted intravitreal retinal injections. AU performed in situ hybridization and some immunostaining. WJY, JH, FT, AU, H.G.A., and AF interpreted data and wrote the manuscript. AF conceived the project.

**Conflict of interest**

The authors declare that they have no conflict of interest.
References


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