

Anti-Tumor Activity of a Novel HS-Mimetic-Vascular Endothelial Growth Factor Binding Small Molecule

Basappa^{1,5,6}, Kazuyuki Sugahara¹, Kuntebommanahalli N. Thimmaiah², Hemant K. Bid³, Peter J. Houghton³, Kanchugarakoppal S. Rangappa⁴*

1 Faculty of Advanced Life Sciences, Hokkaido University, Sapporo, Japan, 2 Chemical Biology and Therapeutics, St. Jude Children's Research Hospital, Memphis, Tennessee, United States of America, 3 Center for Childhood Cancer, Nationwide Children's Hospital, Columbus, Ohio, United States of America, 4 Department of Studies in Chemistry, University of Mysore, Mysore, India, 5 Singapore-MIT Alliance for Research and Technology, Singapore, Singapore, 6 Department of Chemistry, Bangalore University, Bangalore, India

Abstract

The angiogenic process is controlled by variety of factors of which the vascular endothelial growth factor (VEGF) pathway plays a major role. A series of heparan sulfate mimetic small molecules targeting VEGF/VEGFR pathway has been synthesized. Among them, compound 8 (2-butyl-5-chloro-3-(4-nitro-benzyl)-3H-imidazole-4-carbaldehyde) was identified as a significant binding molecule for the heparin-binding domain of VEGF, determined by high-throughput-surface plasmon resonance assay. The data predicted strong binding of compound 8 with VEGF which may prevent the binding of VEGF to its receptor. We compared the structure of compound 8 with heparan sulfate (HS), which have in common the functional ionic groups such as sulfate, nitro and carbaldehyde that can be located in similar positions of the disaccharide structure of HS. Molecular docking studies predicted that compound 8 binds at the heparin binding domain of VEGF through strong hydrogen bonding with Lys-30 and Gln-20 amino acid residues, and consistent with the prediction, compound 8 inhibited binding of VEGF to immobilized heparin. In vitro studies showed that compound 8 inhibits the VEGF-induced proliferation migration and tube formation of mouse vascular endothelial cells, and finally the invasion of a murine osteosarcoma cell line (LM8G7) which secrets high levels of VEGF. In vivo, these effects produce significant decrease of tumor burden in an experimental model of liver metastasis. Collectively, these data indicate that compound 8 may prevent tumor growth through a direct effect on tumor cell proliferation and by inhibition of endothelial cell migration and angiogenesis mediated by VEGF. In conclusion, compound 8 may normalize the tumor vasculature and microenvironment in tumors probably by inhibiting the binding of VEGF to its receptor.

Citation: Basappa, Sugahara K, Thimmaiah KN, Bid HK, Houghton PJ, et al. (2012) Anti-Tumor Activity of a Novel HS-Mimetic-Vascular Endothelial Growth Factor Binding Small Molecule. PLoS ONE 7(8): e39444. doi:10.1371/journal.pone.0039444

Editor: Ramani Ramchandran, Medical College of Wisconsin, United States of America

Received January 5, 2012; Accepted May 24, 2012; Published August 15, 2012

Copyright: © 2012 Basappa et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Funding: This work was supported in part by the INSA (Indian National Science Academy)-JSPS (Japan Society for the Promotion of Science) program between India (to K.S.R.) and Japan (to K.S.), Future Drug Discovery and Medical Care Innovation Program (K.S.) from the Ministry of Education, Culture, Sports, Science, and Technology of Japan (MEXT), and a Grant-in-aid (19-07194) from JSPS, VGST from government of Karnataka (B.) and USPHS award CA23099 (P.J.H.). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

1

Competing Interests: The authors have declared that no competing interests exist.

* E-mail: rangappaks@gmail.com

Introduction

There is a need for anticancer agents with novel mechanisms of action [1]. Recently identified molecular targets for new anticancer agents include, inducers of cell differentiation, cell cycle arrest, and apoptosis, as well as inhibitors of mitogenic signaling pathways elicited by growth factors and cytokines [2,3,4]. A largely unexplored opportunity is the use of carbohydrate mimetics as drugs [5]. The development of new synthetic methods for sugar mimetics has led to a wide variety of novel structures, such as oxazines, triazole-imines, isoxazolines, and imidazole derivatives [6,7,8] that may exert anti-tumor activity. The cell surface and extracellular matrix heparan sulfate (HS) is known to play a major role in tumor metastasis, and acts as a storage site for various proteins [9]. Heparan sulfate glycosaminoglycans (HSGAGs) have been found to play regulatory roles in many biological functions; both normal physiological processes (e.g., embryogenesis) as well as pathological processes (e.g., tumorigenesis and metastasis).

The approach of modulating tumor progression with drugs that inhibit growth of tumors/endothelial cells or the activities of angiostatic factors represents a new concept in adjuvant chemotherapy when used alone or in combination with classic antitumor drugs. This approach is being explored either with the use of monoclonal antibodies directed to the factors [10] or with growth factor-complexing molecules of different origin such as suramin [11], polysaccharide [12], and anionic compounds [13]. The sulphonated derivative of distamycin A, FCE 26644, has been found to inhibit angiogenesis and showed anti-tumor activity by complexing with the growth factors [14,15]. Most recently, a VEGF-A receptor neuropilin 1 binding ligand (EG00229) was designed and synthesized that reduced the viability of A549 lung carcinoma cells [16]. The proliferation and growth of solid tumors in vivo is dependent on various growth factors and cytokines, either through a direct stimulus to cell division and/or through activation of neovascularization, an essential event in tumor progression and metastasis [17]. The increased production of vascular endothelial growth factor (VEGF), fibroblast growth factor-2 (FGF-2),

heparin-binding epidermal growth factor-like growth factor (HB-EGF), and tumor necrosis factor-α (TNF-α), as well as the overexpression of their receptors has been reported in a variety of human tumors [17]. It is well known that both VEGF and FGF-2 are capable of stimulating angiogenesis in vitro and in vivo [18]. During angiogenesis, the proliferation, migration and remodelling of fully differentiated endothelial cells to form microvessels are regulated by growth factors such as VEGF or FGF secreted by tumor cells [19]. Therefore, the inhibition of VEGF or FGF expression or blocking their mitogenic response on endothelial cells might have therapeutic value in the treatment of a variety of cancers [20].

The enormous structural diversity of HSGAGs makes it possible for them to interact specifically with a wide variety of proteins and other ligands. Synthesis of such HS mimetic compounds with antitumor and antiangiogenic properties are currently an active area of research. In order to dissect a variety of the pathological roles of HSGAGs and specifically to identify and develop novel anti-tumor agents, we explored virtual libraries of HSGAGmimetic compounds using database search techniques. The literature revealed the anti-tumor and anti-heparanase activities of a non-sugar-based HS mimetic compound 2-[3-nitro-4-(phenylthio)benzoyl]benzoic acid (KI-105) [21]. We herein, synthesized these types of compounds, which bear HS mimetic structure in order to evaluate their efficacy in preventing the invasion and proliferation of tumor cells. Here, we prepared the hybrid of HS-mimetic core structure of KI-105 by replacing the benzene group by imidazole, since the chemistry of imidazole occupies an extremely important niche within the family of 5membered heterocyclic compounds. 2-butyl-5-chloro-3-(4-nitrobenzyl)-3H-imidazole-4-carbaldehyde (**compound 8**) is considered to mimic the HS non-structurally. We report that the **compound 8**, directly binds to the heparin binding domain of VEGF, as detected by high throughput surface plasmon resonance (SPR) analysis, as well as by in silico binding analysis, and showed promising antitumor activity in experimental model of liver matastasis.

Results and Discussion

Several studies have shown that HS mimetics act as antitumor agents [21]. However, the pleiotropic effects and interactions of such mimetics with heparin-binding proteins might elicit off-target effects associated with toxicity. Low molecular weight HS mimetics which perform multiple biological functions *in vivo* and *in vitro* with high specificity are rare. To this end, we sought to synthesize a series of novel non-sugar-based compounds which can mimic the HS non-structurally.

Synthesis

Synthesis of small molecules (1–9) is shown in **Figure 1**, which depicts the synthesis of 1,3-oxazine derivatives, 1,2,4-triazole derivatives, and imidazole derivatives. 1,3-oxazine derivatives 1–4, were prepared by the cyclization of 1-[2-amino-1-(4-methoxyphenyl)-ethyl]-cyclohexanol monoacetate with aromatic or aliphatic aldehydes in the presence of potassium carbonate [22,23]. 4-Amino-4,5-disubstituted-[1,2,4] triazole-3-thiols 5–7, were synthesized by condensation reaction of 4-amino-5-methy/ethyl/phenyl-4H-[1,2,4]triazole-3-thiols with 1,6-difluorobenzaldehyde in presence of catalytic amount of concentrated sulphuric acid in ethanolic media (**Figure 1**). Microwave-assisted synthesis of N-substituted 2-butyl-5-chloro-3H-imidazole-4-carbaldehyde derivatives **8** and **9**, were synthesized as reported [24,25].

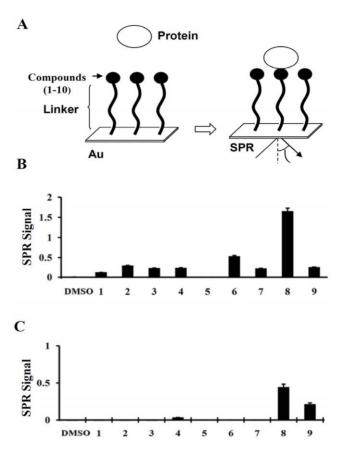


Figure 1. Reagents and condition: a) R₁-CHO, K₂CO₃, Methanol, RT; b) 2,6-difluorobenzaldehyde, Conc. H₂SO₄, ethanol, reflux; c) R₁-Cl, K₂CO₃, DMF, RT. doi:10.1371/journal.pone.0039444.q001

High throughput surface plasmon resonance (SPR) screening of small molecules binding to growth factors

We immobilized the HS-mimetic small molecule libraries (around 60) in order to determine their binding ability with heparin-binding growth factors such as VEGF, FGF-2, TNF-α, midkine, pleotrophin, or HB-EGF by conducting a novel SPR assay (Basappa et al, Cancer Letter, 2010; Supplementary Fig. 1-3). An overview of the SPR analysis is shown in **Figure 2A**. Compounds were immobilized on the photoaffinity-linker-coated gold substrates (PGSs) as reported previously [26,27]. In our in vitro experimental conditions, strong SPR signals for the direct binding of the selected compounds (1-9) with VEGF (Figure 2B) or FGF-2 (Figure 2C) were found. Compound 2 bound to VEGF moderately when compared to other oxazine molecules tested during the assay. Among the triazole compounds like 5, 6, and 7, compound 6 bound to VEGF significantly. More importantly, the imidazole derivative, compound 8, bound to VEGF very strongly, when compared to other screened molecules by SPR assay. It also interacted with FGF-2 significantly, although with weaker affinity compared to its VEGF binding. Some of the synthesized compounds weakly bound to other heparin-binding growth factors such as HB-EGF or TNF- α (data not shown). In addition, none of the compounds showed any binding to pleiotrophin and midkine (data not shown). The assay was found to be specific in terms of binding; hence these results indicate the binding specificity of **compound 8** towards VEGF and led us to

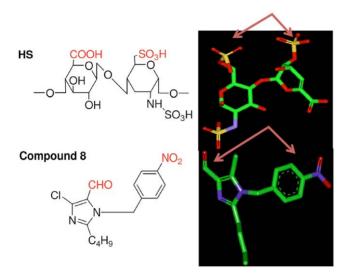


Figure 2. Interaction between the small molecules and growth factors. An overview of the SPR analysis showing the interaction between the sugar mimetics and growth factors/cytokines (A). Small molecules (1–9) were immobilized on the photoaffinity-linker-coated gold substrates. Interactions were detected between the small molecules (1–9) and the growth factors in solution by SPR imaging as described under "Experimental Procedures". The maximum SPR signal strength observed between the small molecules (1–9) and VEGF (B), or FGF-2 (C) are presented. doi:10.1371/journal.pone.0039444.g002

speculate that this molecule may be used to modulate the cellular processes that are mediated by VEGF.

Molecular interaction of compound 8 with the heparin binding site of $VEGF_{165}$

We initially prepared the energy minimized structures of the synthesized molecules. The results of the calculations were compared those for HexUA-GlcNAc(6S). Finally, we selected compound 8 as the core structure (Figure 3A). The energyminimized structures of HexUA(2S)-GlcNAc(6S) and compound 8 revealed that anionic groups, such as sulfate, carboxylic acid, carbaldehyde and nitro functional groups, are able to locate in similar positions and directions (Figure 3B). Because of structural similarity (by functional groups) to HS, **compound 8** could bind to VEGF₁₆₅ in the heparin binding site. **Compound 8** interacts with VEGF strongly, and FGF-2 weakly as determined by SPR assay, hence we performed the molecular docking processes using CDOCKER, which docks ligands to the heparin binding site of VEGF (27). The molecular docking studies predicted that compound 8 binds to the heparin binding domain of VEGF (Figure 4A) with high binding affinity while some other compounds do not show affinity towards the heparin binding site of VEGF (data not shown). Compound 8 showed the CDOCKER score (-CDOCKER ENERGY) of 24.7, a high value, which indicates more favourable binding. This score includes internal ligand strain energy and receptor-ligand interaction energy, and is used to sort the different conformations of each input ligand. Visual analysis of the docked **compound 8** shows the imidazole and benzene nucleus resides in the pocket formed by Val-19, Phe-18, His-15, Lys-16, and Pro-9. In addition, the **compound 8** bound through hydrogen bonding with Lys-30 and Gln-20 residues at the heparin binding pocket of VEGF (Figure 4B). On the other hand, we found the hydrophobic interaction of the butyl group of compound 8 with the heparin binding domain of FGF-2 with a CDOCKER score of 4.8, which is lower when compared to VEGF interaction CDOCKER score (Fig. S1). This revealed that the binding of compound 8 with VEGF is stronger when compared to FGF-2. A similar observation has been reported previously for the compound KI-105 that bound to the HS binding site of heparanase [21]. We know that HS binds to heparin-binding growth factors and cytokines such as VEGF and FGF-2. We analysed the interaction between VEGF₁₆₅ and bound heparin using the BIAcore system. This study revealed that **compound 8** inhibited the binding of VEGF₁₆₅ to the immobilised heparin (Figure 4C). Although the concentration of compound 8 needed to inhibit the binding of VEGF to the heparin was high, these data show that the molecular interaction takes place. In addition, the structurally related compound 7 that does not bind VEGF, failed to inhibit the binding of VEGF to heparin, suggesting selectivity. The results indicate that compound 8 initially competes or is bound initially to the heparin binding domain of VEGF and thereby decreases the binding of VEGF to the immobilized heparin.

Effect of small molecules on the proliferation of tumor cells

Small-molecules that intervene in biological systems have been sought as tools to perturb signaling pathways and of course as therapeutic agents for disease. Heparin and HS structural mimetics have been reported as antitumor agents that significantly inhibit tumor growth, metastasis, and angiogenesis. To validate the binding ability of these compounds with VEGF and to further elucidate the biological functions of these molecules, we initially examined the effects of compounds 1-9 on the proliferation of murine osteosarcoma cell lines, LM8 and LM8G7, which are known to the secrete proangiogenic factor VEGF [28]. Compounds 1-4, 6, 7, and 9 did not show promising inhibitory activity towards the proliferation of osteosarcoma cancer cells (**Table 1**). Compounds **5** and **6** inhibited moderately the proliferation of LM8 and LM8G7 cells. In contrast to other compounds, compound 8, which bears an imidazole moiety, showed more potent inhibition towards the proliferation of VEGFsecreting LM8 and LM8G7 cells with an IC50 values of 7 µM and 4.5 µM, respectively. In comparison cisplatin was less potent with IC₅₀ concentrations of 30 and 15 μM, respectively (Table 1). The inhibitory activity of suramin was comparable with the inhibitory activity of **compound 8** on the proliferation of tumor cells.

Real-time monitoring of the effect of compound 8 on the proliferation of LM8G7 cells

Elimination or arrest of tumor cell proliferation in the target organ are the eventual objectives of anticancer therapy. We monitored the effect of **compound 8** (2 to 50 μ M) on the proliferation of VEGF secreting LM8G7 cells using real-time cell electronic sensing system TM (RT-CES) to confirm the results obtained through TetraColor One assay. **Compound 8** inhibited the proliferation of LM8G7 cells in a concentration-dependent manner with an IC₅₀ value 5 μ M, confirming its anti-proliferative effect on LM8G7 cells (**Figure 5A**).

Effect of small molecules on the proliferation of endothelial cells

The proliferation and migration of endothelial cells to form microvessels is critical for angiogenesis, and inhibition of angiogenic factor-mediated proliferation of endothelial cells has been shown to be an effective antiangiogenic therapy [29]. Hence, we tested the effects of these compounds on the VEGF-induced

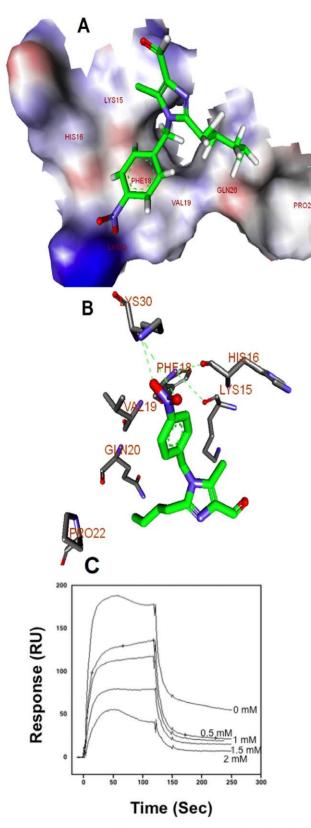


Figure 3. Discovery of New HS mimetic small molecule. *Left.* Chemical structure of a HS disaccharide unit (1), HexUA(2S)-GlcNAc(6S), and **compound 8** are also shown. *Right.* Energy-minimized structures of HS and **compound 8**. The arrows show the anionic functional groups. The carbon, oxygen, nitrogen, and sulfur atoms are represented in green, red, blue, and yellow, respectively. The hydrogen atoms are not shown. doi:10.1371/journal.pone.0039444.g003

proliferation of endothelial (UVQ2) cells. Except for compounds ${\bf 6}$, the remaining compounds had no significant effects on the proliferation of UVQ2 cells, whereas compound ${\bf 6}$ which had weak binding with VEGF moderately inhibited its proliferation.

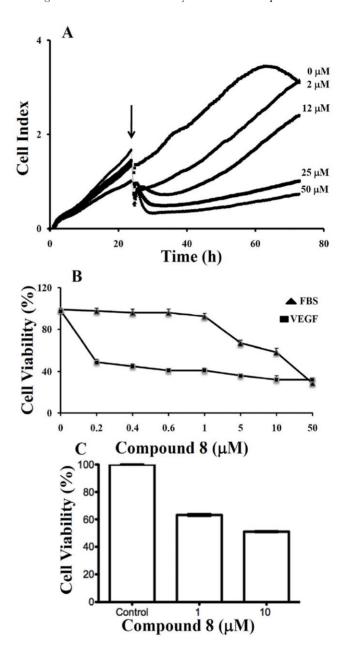


Figure 4. Molecular basis for the interaction between compound 8 and heparin binding domain of VEGF₁₆₅. A. Binding mode of **compound 8** with heparin binding site of VEGF₁₆₅. The amino acid residues of heparin binding site of VEGF₁₆₅ are shown in stick models. B. Interactions of the **compound 8** within the heparin binding pocket of VEGF₁₆₅ with putative hydrogen bonds shown as green dotted lines. The **compound 8** is shown in green color. The bonding between the **compound 8** and the heparin binding domain of VEGF₁₆₅ are shown in yellow color. All the hydrogen atoms are not shown. The picture is rendered in Discovery Studio, version 2.5 (right panel). C. Binding of VEGF₁₆₅ to immobilised-heparin. Various concentrations of **compound 8** and VEGF₁₆₅ (250 nM) mixture in running buffer was injected onto the surface of the heparin-immobilised sensor chip. Sensograms obtained were overlaid using a BIA evaluation software (version 3.1).

doi:10.1371/journal.pone.0039444.g004

Table 1. Inhibition of the proliferation of tumor and endothelial cells by compounds 1–9 (IC_{50} in μ M).

·				
Compound	R1/R2/R3	LM8	LM8G7	UVÇ2
1	Н	NS ^a	NS	NS
2	4-Fluoro-phenyl	64±0.5	86 ± 0.1	NS
3	4-Chloro-phenyl	NS	>100	NS
4	pyridin-3-yl	95±6.7	NS	>100
5	Methyl	36±6.4	NS	45±2.1
6	Ethyl	64 ± 6.7	$23\!\pm\!0.5$	14±4.6
7	Phenyl	>100	NS	NS
8	4-Nitro-phenyl	6.8 ± 0.3	5 ± 2.8	$42\!\pm\!0.1$
9	6-methyl-benzo[1,3] dioxol-5-yl methyl)-	46±6.7	NS	NS
Suramin		12 ± 2.8	14±24	$34\!\pm\!1.4$
Cisplatin		30±24	15±0.6	12±0.4

^aNS, not significant.

doi:10.137 1/journal.pone.0039444.t001

Compound 8, which strongly binds to VEGF, markedly inhibited the proliferation of UVQ2 cells with an IC $_{50}$ value of 42 μ M (Table 1). These results show that **compound 8** suppressed the proliferation of endothelial cells, but the concentration of **compound 8** required to suppress cell proliferation was high compared with that required to suppress proliferation of LM8G7 osteosarcoma cells.

In contrast to UVQ2 cells treated with **compound 8** under serum-containing conditions (10%FBS), **compound 8** showed a potent inhibitory effect on the VEGF-stimulated proliferation with an IC $_{50}$ value 0.3 μ M (**Figure 5B**). These results indicate that **compound 8** inhibited endothelial cell proliferation through inhibition of VEGF receptor function.

We next examined the effect of **compound 8** to suppress VEGF-stimulated proliferation of HUVECs. As shown in **Figure 5C**, proliferation was inhibited approximately 40% at 1 μ M, and by 50% at 10 μ M. Taken together, these data suggest that **compound 8** can block the VEGF-mediated angiogenic response in endothelial cells.

Compound 8 suppresses VEGF-induced migration and tube formation of endothelial cells

Endothelial cell migration is critical events for angiogenesis. To examine whether **compound 8** inhibited migration of UVQ2 endothelial cells the effect of compound 8 against the VEGFinduced migration across the porous membrane of BD BioCoatTM chambers was determined. VEGF significantly enhanced migratory activity and cotreatment with compound 8 resulted in significant inhibition of VEGF-induced migration. Cells were treated with VEGF, drug vehicle (0.5% DMSO, Control) or compound 8 at 0.5 and 1.0 µM. Representative photomicrographs are shown in Figure 6A-C. Quantification of the reduction in migration showed that **compound 8** significantly inhibited VEGF-induced migration of endothelial cells by 52, and 83%, at 0.5 and 1.0 μ M, respectively (**Figure 6D**). Further, compound 8 at $1 \mu M$ was non-toxic as is evident from the TetraColor One assay (Table 1); hence, the inhibitory effect could not be attributed to cytotoxic activity.

Stimulation of endothelial cells on ECMatrixTM by angiogenic factors such as VEGF promotes differentiation to form capillary-like tubes. We performed the tube formation assay to assess any

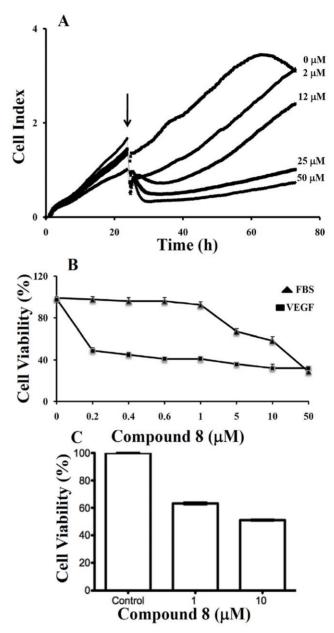


Figure 5. A. Real-time monitoring of the effects of compound 8 on the proliferation of LM8G7 cells. Cells were seeded in ACEA's $96 \times \text{e-plate}^{\text{TM}}$ at a density of 5×10^3 cells per well, and continuously monitored using the RT-CES system up to 24 h, at which point Compound 8 (2 to 50 µM) was added. The cell index is plotted against time. The arrow indicates the time of the addition of compound 8. Data represent the mean values \pm S.D. for three identical wells from three independent experiments. B, compound 8 poorly inhibited proliferation of UVQ2 under serum-replete conditions (FBS), but potently inhibited VEGF-dependent proliferation of UVQ2 cells as analysed by TetraColor One assay. Results were normalised to DMSO controls. Data represent mean values \pm SD for three independent experiments. * P<0.05 versus P<0.01 versus control. C, compound 8 inhibits VEGFstimulated proliferation of human vascular endothelial cells. HUVEC cells were seeded on 6-well plates at a density of approximately 1×10^5 cells/ well in M200 medium supplement with LSGS (Low serum growth Supplement). The next day cells were stimulated with 10 ng/mL of VEGF in the presence or absence of 1 μM and 10 μM compound 8. After 48 hrs, Alamar Blue was added directly into culture media at a final concentration of 10% and the plate was returned to the incubator. Optical density (OD) of the plate was measured at 540 and 630 nm. As a negative control, AB was added to medium without cells. doi:10.1371/journal.pone.0039444.g005

specific effect of **compound 8** on this process. The results showed that VEGF significantly increased tube formation (**Figure 6E**) and cotreatment of endothelial cells with **compound 8** at 0.5 to 1 μ M resulted in strong inhibition of VEGF-induced tube formation in a concentration-dependent manner (**Figure 6F and G**). At 1 μ M **compound 8** resulted in complete inhibition of the tube formation by the endothelial cells. The combined results show that **compound 8** suppressed the proliferation of endothelial cells, but the concentration of **compound 8** required to suppress the proliferation was high compared with that required to suppress the tube formation and cell migration. However, the compound **8** failed to inhibit the FGF-2-induced tube formation by the endothelial cells at the tested concentrations (**Figure S2**). Taken together, we observed that **compound 8** may block the VEGF-mediated angiogenic response in endothelial cells.

Effect of compound 8 on the invasion of LM8G7 cells

The effect of **compound 8** on the invasion of highly metastatic LM8G7 cells across MatrigelTM-coated porous membranes was studied. Control (DMSO-treated) LM8G7 cells were highly invasive in the assay (**Figure 7A**). **Compound 8** at 0.5 and 1 μ M (**Figure 7B and C**) effectively inhibited the invasion of LM8G7 cells by 46, and 75% (**Figure 7D**), respectively, when compared to control. Compounds **1–7** and **9** failed to inhibit the invasion of LM8G7 cells at 1 μ M concentration effectively (data not shown).

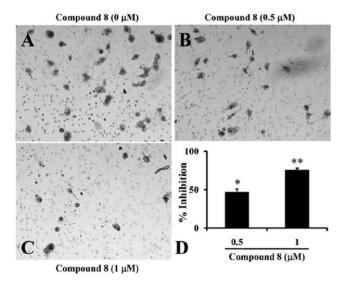


Figure 6. Compound 8 suppresses VEGF-induced migration and angiogenesis. UVQ2 cells (2.5×10⁴) were seeded along with VEGF (2 ng/ml) in Boyden-chambers and incubated for 24 h with DMEM (A) or a medium containing 0.5 μM of $\boldsymbol{compound}$ 8 (B) or $1~\mu M$ compound 8 (C); the photographs (100×) shows the cell on the lower surface of the filter (migrated) stained with the Diff-Quick solution. (**D**) Inhibition rate of the **compound 8** on UV \bigcirc 2 migration was presented as described under "Experimental Procedures". Data represent the mean values \pm S.D. for three identical wells from three independent experiments. Bars represent the mean \pm SD of three independent experiments. * P<0.05 versus control. ** P<0.01 versus control. The UVQ2 cells were seeded in 6-well plates pre-coated with MatrigelTM (ECM 625, Chemicon) was added to a 24-well plate in a final volume of 100 μ L and allowed to solidify at 37°C for 30 min. UVQ2 cells (4×10⁵ cells) were seeded to the ECMatrixTM-coated wells in the presence or absence of compound 8 (0.5 (F) or 1 µM (G) along with 2 ng/ml VEGF. After 18 h of culture, the reorganization of the subconfluent monolayer of UV $_2$ cells in 3-dimensional Matrigel was monitored and photographed under an Olympus FX380 microscope attached to a 3CCD camera. doi:10.1371/journal.pone.0039444.g006

Compound 8 prevents experimental liver metastasis

Because **compound 8** potently inhibited endothelial cell migration and tube formation, as well as LM8G7 cell proliferation, we examined the in vivo anti-tumor effect of compound 8. Following intravenous injection of LM8G7 cells mice developed metastatic nodules in the liver within 30 days (Figure 8A). In contrast, mice treated intravenously with **compound 8** (0.5 or 1.0 mg/kg, days 3, 5, and 10) or heparin (5 mg/kg, used on the same schedule of administraion as a positive control) were completely free of metastatic nodules in the liver. Figure 8B and C, respectively. Compound 8 at 0.25 mg/kg treatment, failed to inhibit the tumor formation by the LM8G7 cells (Figure 8D). The animals tolerated the dosages of compound 8 (0.5 to 1.0 mg/kg), with no signs of toxicity or weight loss during the experiments (data not shown). Unlike heparin, compound 8 did not show anti-coagulant activity in vitro (data not shown). In addition, heparin at 2.5 mg/kg dose failed to inhibit the metastasis of LM8G7 cells (Figure 8D). We also evaluated the effect of **compound 8** (0.25 mg/kg) in combination with heparin (2.5 mg/kg). As shown in **Figure 8D**, the combination treatment resulted in complete abrogation of metastatic tumor nodules.

Summary

We demonstrated here that the compound 8 strongly interacted with VEGF as evidenced by a novel SPR assay. Molecular docking calculations revealed that hydrogen bonding interactions may play crucial role in demonstrating biological activity. Concerning the mechanism of action, we observed that **compound 8** may displace VEGF from its specific binding sites on HUVECs. It is noteworthy that HS mimetic **compound 8** might therefore behave as an antagonist or partial agonist. Since heparin can interfere with VEGF action either by binding to a specific domain on VEGF or by interacting with its cellular binding sites, KDR or NRP1, Consequently, compound 8 might affect both the ligand and its cellular binding sites. We show here that **compound 8** interacts directly with VEGF. This is in agreement with our previous findings that compound \mathbf{v} bound to FGF2, and thus altering the conformation of ligand receptor complexes. Compound 8 showed in vitro antiproliferative activity against LM8G7 osteosarcoma cells (endogenously expressing VEGF), and inhibited their invasiveness. Compound 8 inhibited the VEGF induced endothelial tube formation by UVQ2 cells and suppressed VEGF-stimulated proliferation and tube formation of HUVEC cells. *In vivo*, **compound 8** potently inhibited formation of liver metastases following intravenous injection of LM8G7 osteosarcoma cell. Collectively, these data indicate that com**pound 8** may prevent tumor growth through a direct effect on tumor cells by blocking VEGF paracrine or autocrine stimulated proliferation and by inhibition of endothelial cell migration and angiogenesis mediated by VEGF. In conclusion, the effects of compound 8 are probably through inhibiting the binding of VEGF to its receptor. Furthermore, to the best of our knowledge, this is the first report on a non-structural HS mimetic, which interacts directly with VEGF. Our data suggest that compound 8 has potential use to build a therapeutic drug for minimizing tumor growth and angiogenesis in VEGF related models.

Materials and Methods

Cisplatin, recombinant human (rh)-FGF-2, and rh-VEGF $_{165}$ were purchased from Wako Pure Chemicals Co. (Osaka, Japan). $100\times$ non-essential amino acids, b-mercaptoethanol, $100\times$ sodium pyruvate and L-glutamine were from GIBCO (Auckland, New Zealand). The cell proliferation assay kit Tetracolor One was

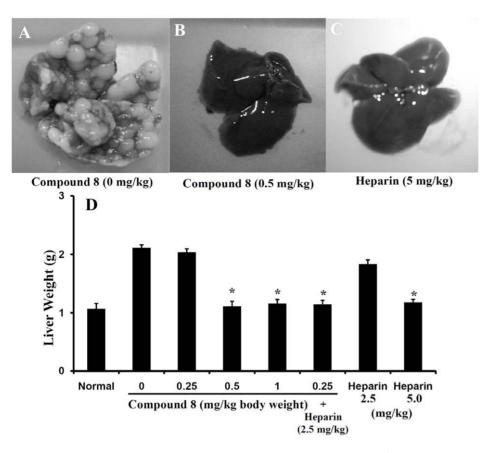


Figure 7. Effect of compound 8 on the invasion of LM8G7 cells. Cells (2.5×10^4) were seeded in MatrigelTM-coated Boyden chambers and were incubated for 24 h with DMEM (A) or a medium containing 0.5 μ M of compound 8 (B) or 1 μ M compound 8 (C); the photographs $(100 \times)$ shows the cell on the lower surface of the filter (invaded) stained with the Diff-Quick solution. (D) Inhibition rate of the compound 8 on LM8G7 invasion was presented as described under "Experimental Procedures". Bars represent the mean \pm SD of three independent experiments. * P<0.05 versus control. ** P<0.01 versus control. doi:10.1371/journal.pone.0039444.g007

obtained from Seikagaku Corp. (Tokyo, Japan) and the Diff-Quick solution was from International Reagent Corp. (Kobe, Japan). All other chemicals and reagents used were of the highest commercial grade available.

The melting points were determined on a SELACO-650 hot stage apparatus and are uncorrected. IR (KBr) spectra were recorded on a Jasco FT/IR-4100 Fourier transform infrared spectrometer, $^1\mathrm{H}$ NMR were recorded on a Shimadzu AMX 400 spectrometer using CDCl3 as solvent and TMS as an internal standard (chemical shift in ppm). Elemental analyses were obtained on a vario-EL instrument and were within $\pm 0.4\%$ of calculated values. Thin layer chromatography (TLC) was conducted on 0.25 mm silica gel plates (60F254, Merck). Visualization was made with ultraviolet light. All extracted solvents were dried over anhydrous Na2SO4 and evaporated with a BUCHI rotary evaporator. Silica gel (200–400 mesh) from Aldrich, Inc., was used for column chromatography. All other chemicals were obtained from Aldrich, Inc.

Preparation of the compounds 1-4, 8, and 9

Compounds 1, 8, and 9 were prepared by the methods described in our previous papers [25,30].

The compounds 5 to 7 were prepared by the following procedure

Equimolar amount of 4-amino-5-substituted-4H-[1,2,4]triazole-3-thiols and 2,6-difluoro-benzaldehyde were heated in ethanol

(10 ml) until a clear solution was obtained. Then a few drops of conc. $\rm H_2SO_4$ were added and the solution was refluxed for 3–4 h on a water bath. After completion of the reaction, the solvent was evaporated, water was added, and cooled to 10–15°C. The product was filtered at the same temperature, and the residue washed with chilled n-hexane, dried and recrystallised with ethyl acetate to obtain the pure compounds.

4-[(2,6-Difluoro-benzylidene)-amino]-5-methyl-4H-[1,2,4]triazole-3-thiol 5

The product was obtained from 4-amino-3-methyl-4H-[1,2,4]triazole-5-thiol (1 g, 7.6 mmol) and 2,6-difluoro-benzaldehyde (1.09 g, 8.3 mmol) as white crystalline solid (0.94 g, 94%). mp: 225–228°C. IR $\nu_{\rm max}$ (KBr): 1625 (s), 1285 (m), 892 cm⁻¹. ¹H NMR (CDCl₃ 400 MHz) δ : 13.8 (s, 1H, S-H), 10.6 (s, 1H, CH = N-), 7.65–7.74 (m, 2H, Ar-H), 7.27–7.33(t, 1H, Ar-H), 2.32 (s, 3H, CH₃). Anal. (C₁₁H₉F₂N₃S) C, H, N.

4-[(2,6-Difluoro-benzylidene)-amino]-5-ethyl-4H-[1,2,4]triazole-3-thiol 6

The compound was obtained from 4-amino-3-ethyl-4H-[1,2,4]triazole-5-thiol (1 g, 6.9 mmol) and 2,6-difluoro-benzalde-hyde (0.98 g, 6.76 mmol) as white powder (0.75 g, 75%) mp: 185–187°C. IR $\nu_{\rm max}$ (KBr): 1615 (s), 1245 (m), 875 cm⁻¹. ¹H NMR (CDCl₃ 400 MHz) δ : 13.8 (s, 1H, S-H), 10.5 (s, 1H, CH=N-),

$$R_2$$
 R_2
 R_2
 R_3
 R_4
 R_5
 R_4
 R_5
 R_5
 R_4
 R_5
 R_5
 R_6
 R_7
 R_8
 R_8
 R_8
 R_9

Figure 8. Effect of the compound 8 against the metastatic potential of LM8G7 cells. C3H/HeN mice were intravenously injected with LM8G7 cells via tail. Some mice received an intravenous injection of compound 8 (0.25, 0.5, or 1.0 mg/kg) on day 3, 5 and 10. After 4 weeks, the mice were sacrificed, the number of liver nodules was counted macroscopically, and the liver weight was measured in the control and compound 8-treated animals. Representative livers from mice injected with LM8G7 cells treated with DMEM (A), compound 8 (0.5 mg/kg) (B), and heparin (5 mg/kg) (C) are shown. The average liver weight of the mock and the **compound 8** treated mice (D). Heparin from porcine intestinal mucosa was used as a positive control. The possible synergistic or additive effects of compound 8 and heparin were investigated by subsequent administration of compound 8 (0.25 mg/kg) and heparin (2.5 mg/kg) as described above. Data represent mean values \pm SD for three independent experiments and each experiment was conducted with 6 mice per group. * P<0.05 versus control. Mann-Whitney U test. doi:10.1371/journal.pone.0039444.g008

7.65–7.74 (m, 2H, Ar-H), 7.23–7.34 (t, 1H, Ar-H), 1.23 (t, 3H, -CH₃), 2.68–2.76 (q, 2H, -CH₂-). Anal. ($C_{12}H_{11}F_2N_3S$) C, H, N.

4-[(2,6-difluoro-benzylidene)-amino]-5-phenyl-4H-[1,2,4]triazole-3-thiol 7

The product was obtained from 4-amino-3-phenyl-4H-[1,2,4]triazole-5-thiol (1 g, 5.2 mmol) and 2,6-difluoro-benzaldehyde (0.74 g, 3.8 mmol) as crystalline solid. Weight: 0.875 g; yield 87%; mp: 195–200°C; IR $\nu_{\rm max}$ (KBr): 1612 (s), 1236 (m), 842; $^1{\rm H}$ NMR (CDCl $_3$ 400 MHz) δ : 14.35 (s, 1H, S-H), 10.2 (s, 1H, CH = N-), 7.89–7.96 (dd, 2H, Ar-H), 7.66–7.76 (m, 1H, Ar-H), 7.46–7.56 (dd, 2H, Ar-H), 7.28–7.34 (t, 1H, Ar-H); Anal. (C $_{16}H_{11}F_2N_3S$) C, H, N.

SPR assay

The gold-coated glass chip (TOYOBO, Osaka, Japan) by immersing in ethanol solution, containing $0.1~\mathrm{mM}$ of the photo-affinity linker and $0.9~\mathrm{mM}$ dummy linker for $12~\mathrm{h}$ as reported

previously [26]. The chip was washed successively with ethanol, water, and ethanol. Then the chip is air dried to obtain the photo reactive chip. The compounds 1–9 were spotted on the chip (10 mM), washed, and dried in *vacuo*, and then irradiated at 365 nm under a UV transmission filter (Sigma-Koki, Japan). The compounds were immobilized onto the photo-reactive linker that bound to the gold-surface of the glass chip was set into an SPR imaging instrument (TOYOBO), and each growth factor (50 $\mu g/$ mL) was injected onto the array surface at 0.1 ml/min and incubated for 10 min. The SPR image and signal data were collected with an SPR analysis program (TOYOBO). The SPR difference image was constructed by using the Scion Image program (Scion, MD).

Molecular docking study

As a model for docking simulations, we chose the solution structure of the 55-residue heparin-binding domain of VEGF₁₆₅, which has been solved using data from two-dimensional homonuclear and three-dimensional heteronuclear NMR spectroscopy (PDB entry: 2vgh) [31]. We also considered the FGF-2 from the crystal structure of a ternary FGF-FGFR-heparin complex that reveals a dual role for heparin in FGFR binding and dimerization [32]. Molecular docking was carried out using discovery studio program (Accelrys, California, USA). To dock compound 8 with VEGF₁₆₅, a CDOCKER protocol was used, which is an implementation of the algorithm CDOCKER [33] that allows us to run a refinement docking of any number of ligands with a single protein receptor, which is a grid-based molecular docking method that employs CHARMm. The receptor is held rigid while the ligands are allowed to flex during the refinement. MOLECULAR docking was done by specifying the ligand placement in the active site using a binding site sphere.

Interaction Analysis

The interaction of VEGF₁₆₅ with heparin in the presence of **compound 8** was examined using a BIAcore J system (BIAcore AB, Uppsala, Sweden). The heparin-immobilized sensor chip was prepared as reported earlier [34]. The interaction of **compound 8** with the heparin binding domain of VEGF₁₆₅ was analysed by injecting the various concentrations of **compound 8** and VEGF₁₆₅ (250 nM) onto the surface of the sensor chip in running buffer, pH 7.4 (HBS-EP; BIAcore AB), containing 10 mM HEPES, 0.15 M NaCl, 3 mM EDTA, and 0.005% (w/v) Tween 20. The flow rate was kept at a moderate speed (30 ml/min) according to the manufacturer's recommendations. Each mixture of the VEGF₁₆₅ and **compound 8** was allowed to interact with the heparin-immobilized sensor chip for 2 min allowing association and dissociation.

Cell lines

LM8G7 (a gift from M. Miyasaka, Department of Pharmaceutical Chemistry, Osaka University) [23], a highly metastatic murine osteosarcoma cell line with the potential to invade the liver, was cloned from LM8G5 cells [35] as described [28] and cultured in Dulbecco's Modified Eagle's Medium (DMEM) supplemented with 10% (v/v) fetal bovine serum (FBS) (Thermo Trace, Melbourne, Australia), streptomycin (100 µg/mL), penicillin (100 units/mL), $100\times$ non-essential amino acids, b-mercaptoethanol (50 µM), $100\times$ sodium pyruvate, and L-glutamine (2 mM) at 37°C in a humidified 5% CO₂ atmosphere. The cells were harvested after incubation with 0.1% trypsin/1 mM EDTA in PBS for 5 min at 37°C followed by gentle flushing with a pipette, and subcultured three times a week. Mouse normal vascular

endothelial cells (UVQ2) were maintained in DMEM supplemented with 10% (v/v) FBS.

Proliferation assay

Two tumor cell lines such as LM8, LM8G7, and UVQ2 endothelial were used to evaluate the anti-proliferative activity of the compounds. UVQ2 $(1\times10^4~\text{cells})$ or LM8 or LM8G7 cells $(5\times10^3~\text{cells})$ were seeded in 96-well plate and incubated overnight at 37°C. The tumor cells or endothelial cells stimulated with VEGF (2 ng/ml) were treated with various concentrations of compounds **1–9** for an additional 48 h. 5 μ L of TetraColor One reagent was added and incubated for an additional 2–4 h and the absorbance at 450 nm (Bio-Rad) was measured. Further, the effect of **compound 8** on VEGF-mediated proliferation of UVQ2 endothelial was measured. The results were expressed as a percentage of viable cells relative to cells treated with DMSO. The viability of the cells was expressed in percentage terms and IC₅₀ value was calculated.

Human umbilical vein endothelial cells (HUVEC) were obtained from the American Type Culture Collection (ATCC). All experiments were done using endothelial cells between passages 3 and 8. HUVECs were maintained in endothelial cell growth medium M200 (Invitrogen) high glucose supplemented with 15% fetal bovine serum, endothelial cell growth supplements (LSGS Medium, Cascade Biologics), and glutamine at 37°C with 5% CO₂. All cells were maintained as sub confluent cultures and split 1:3, 24 h before use. HUVECs were seeded on 6-well plates at a density of approximately 1×10⁵ cells/well in M200 medium supplement with LSGS (Low Serum Growth Supplement). The next day cells were stimulated with 10 ng/mL of VEGF in the absence or presence of 1 and 10 µM of compound 8. After 48 hrs, AB (Alamar Blue) was added directly into culture media at a final concentration of 10% and the plate was returned to the incubator. Optical density (OD) of the plate was measured at 540 and 630 nm with a standard spectrophotometer at 3-4 h after adding AB. As a negative control, AB was added to medium without cells.

Real-time proliferation assay

The cell proliferation assay was done using the Real-Time Cell Electronic Sensing (RT-CES) system (ACEA Biosciences, San Diego, CA). LM8G7 $(5\times10^3~\text{cells/well})$ cells were seeded in ACEA's $96\times~\text{e-plate}^{TM}$ in a final volume of $150~\mu l$ [36]. Approximately 24 h after seeding, when in log growth phase, the cells were incubated with $150~\mu l$ of DMEM containing various concentrations of **compound 8** (final concentration 2 to $50~\mu M$) or DMEM containing DMSO as control. The effects of **compound 8** on the proliferation of LM8G7 cells were monitored dynamically every 10 min. A cell index (quantitative measurement of cell proliferation) was plotted against time. The IC50 values were calculated from concentration-response curves by a non-linear regression analysis using the GraphPad Prism (GraphPad Prism Software Inc., San Diego).

In vitro cell migration and invasion assays

The ability of LM8G7 and UVQ2 cells to migrate and invade was assessed using the BD BioCoatTM chamber with or without Matrigel (BD Biosciences) *in vitro*. The single cell suspensions of LM8G7 cells or UVQ2 cells were prepared by detaching and resuspending in serum-free DMEM. Before the cells were added, the chambers were rehydrated for 2 h in an incubator at 37°C. In the upper chamber (8 μ m PET pores), LM8G7 (6×10⁴ cells/ml) or UVQ2 (1×10⁵ cells/ml) cells were added along with **compound 8** (0.5 or 1.0 μ M) in 500 μ l DMEM and the lower

chambers were filled with DMEM containing 10% fetal bovine serum. The MatrigelTM invasion chambers were incubated for a further 22 h at 37°C. After incubation for 24 h, the cells that had migrated or invaded through the membrane alone or the Matrigel-coated membrane remained bound to the underside of the membranes. In some instances, VEGF (2 ng/ml) was added exogenously to understand the effect of **compound 8** on VEGF-induced migration of UVQ2 cells. These cells were stained with Diff-Quik staining kit and counted in five random microscopic fields/filter. The percent inhibition of the invasion or migration of LM8G7 cells or UVQ2 cells by **compound 8** were calculated.

In vitro angiogenesis assay

MatrigelTM (ECM 625, Chemicon) was added to a 24-well plate in a final volume of 100 μ L and allowed to solidify at 37°C for 30 min. UVQ2 cells (4×10⁵ cells) were seeded to the ECMatrix TM-coated wells in the presence or absence of **compound 8** (0.5 or 1 μ M) along with 2 ng/ml VEGF of FGF-2. After 18 h of culture, the reorganization of the sub-confluent monolayer of UVQ2 cells in 3-dimensional ECMatrix TM was monitored and photographed under an Olympus FX380 microscope attached to a 3CCD camera.

Liver metastasis assay

Nine-week-old female C3H/HeN mice were obtained from Japan SLC (Hamamatsu, Japan) and kept in standard housing. All the experiments were performed according to a protocol approved by the local animal care and use committee of Hokkaido University. C3H/HeN mice were intravenously injected with 1×10⁶ LM8G7 cells in 200 μl of DMEM *via* the tail on day 0. Some mice received an intravenous injection of **compound 8** (0.25, 0.5, or 1.0 mg/kg) suspended in 200 μL of DMEM, on day 3, 5 and 10 after the tumor cell injection. After 4 weeks, the mice were sacrificed and the number of liver nodules was counted macroscopically in the control and **compound 8**-treated animals. Heparin (5 mg/kg) was used as a positive control. The synergetic effects of **compound 8** and heparin were investigated by subsequent administration of **compound 8** (0.25 or 0.5 mg/kg) and heparin (2.5 or 5.0 mg/kg) as described above.

Ethics statement

All animal experiments were conducted in accordance with protocols specifically approved for this study by the animal care and use committee (IACUC) at Hokkaido University, designed to minimize the numbers of mice used and to minimize any pain or distress.

Statistical analysis

The statistical analysis was done using software Origin 8 (OriginLab). The Mann-Whitney U test was used to determine P-values.

Supporting Information

Figure S1 Interactions of the compound 8 within the heparin binding pocket of FGF-2 (PDB ID: 1FQ9). The compound 8 carbons are shown in grey color. The heparin binding domain of FGF-2 amino acids carbons are shown in pink color.

(TIF)

Figure S2 Effect of Compound 8 on FGF-2-induced tube formation. The UVQ2 cells were seeded in 6-well plates precoated with MatrigelTM (ECM 625, Chemicon) and allowed to

solidify the presence or absence of compound 8 (0.5 or 1 μ M) at 37°C for 30 min. along with 2 ng/ml FGF-2. After 18 h of culture, the reorganization of the sub-confluent monolayer of UVQ2 cells in 3-dimensional ECMatrix TM was monitored and photographed. The number of intact tubes were counted in five randomly chosen regions and expressed as the percentage of the control, and the results are expressed as mean \pm S.D. Inhibition rates of compound 8 on the tube formation of UVQ2 cells was presented. (TIF)

Acknowledgments

We thank Prof. M. Miyasaka for providing the LM8 and LM8G7 cells. We also thank Prof. H. Osada, Dr. Y. Kondoh and Dr. S. Akiko for providing

References

- Goss PE, Baker MA, Carver JP, Dennis JW (1995) Inhibitors of carbohydrate processing: A new class of anticancer agents. Clin Cancer Res 1: 935–944.
- Marks PA (2007) Discovery and development of SAHA as an anticancer agent. Oncogene 26: 1351–1356.
- Tiligada E, Miligkos V, Delitheos A (2002) Cross-talk between cellular stress, cell cycle and anticancer agents: mechanistic aspects. Curr Med Chem Anticancer Agents 2: 553–566.
- 4. Dawson MI, Xia Z, Jiang T, Ye M, Fontana JA, et al. (2008) Adamantyl-substituted retinoid-derived molecules that interact with the orphan nuclear receptor small heterodimer partner: effects of replacing the 1-adamantyl or hydroxyl group on inhibition of cancer cell growth, induction of cancer cell apoptosis, and inhibition of SRC homology 2 domain-containing protein tyrosine phosphatase-2 activity. J Med Chem 51: 5650–5662.
- Compain P, Martin OR (2001) Carbohydrate mimetics-based glycosyltransferase inhibitors. Bioorg Med Chem 9: 3077–3092.
- Slee DH, Romano SJ, Yu J, Nguyen TN, John JK, et al. (2001) Development of potent non-carbohydrate imidazole-based small molecule selectin inhibitors with antiinflammatory activity. J Med Chem 44: 2094–2107.
- Perion R, Ferrieres V, Garcia-Moreno MI, Mellet CO, Duval R, et al. (2005) 1,2,3-Triazoles and related glycoconjugates as new glycoside inhibitors. Tetrahedron 61: 9118–9128.
- Gallos JK, Koumbis AE (2003) 1,3-Dipolar cycloadditions in the synthesis of carbohydrate mimics. Part 1: Nitrile oxides and nitronates. Current Org Chem 7: 397–426.
- Vlodavsky I, Bar-Shavit R, Ishai-Michaeli R, Bashkin P, Fuks Z (1991) Extracellular sequestration and release of fibroblast growth factor: a regulatory mechanism? Trends Biochem Sci 16: 268–271.
- Hori A, Sasada R, Matsutani E, Naito K, Sakura Y, et al. (1991) Suppression of solid tumor growth by immunoneutralizing monoclonal antibody against human basic fibroblast growth factor. Cancer Res 51: 6180–6184.
- Stein CA (1993) Suramin: a novel antineoplastic agent with multiple potential mechanisms of action. Cancer Res 53: 2239–2248.
- Wellstein A, Zugmaier G, Califano JA III, Kern F, Paik S, et al. (1991) Tumor growth dependent on Kaposi's sarcoma-derived fibroblast growth factor inhibited by pentosan polysulfate. J Natl Cancer Inst 83: 716–720.
- Benezra M, Vlodavsky I, Yayon A, Bar-Shavit R, Regan J, et al. (1992) Reversal of basic fibroblast growth factor-mediated autocrine cell transformation by aromatic anionic compounds. Cancer Res 52: 5656–5662.
- Ciomei M, Pastori W, Mariani M, Sola F, Grandi M, et al. (1994) New sulfonated distamycin A derivatives with bFGF complexing activity. Biochem Pharmacol 47: 295–302.
- Sola F, Farao M, Pesenti E, Marsiglio A, Mongelli N, et al. (1995) Antitumor activity of FCE 26644 a new growth-factor complexing molecule. Cancer Chemother Pharmacol 36: 217–222.
- Jarvis A, Allerston CK, Jia H, Herzog B, Garza-Garcia A, et al. (2010) Small molecule inhibitors of the neuropilin-1 vascular endothelial growth factor A (VEGF-A) interaction. J Med Chem 53: 2215–2226.
- 17. Folkman J, Klagsbrun M (1987) Angiogenic factors. Science 235: 442-447.
- Liotta LA, Steeg PS, Stetler-Stevenson WG (1991) Cancer metastasis and angiogenesis: an imbalance of positive and negative regulation. Cell 64: 327– 336.
- Cross M, Dexter TM (1991) Growth factors in development, transformation, and tumorigenesis. Cell 64: 271–280.

the SPR imaging data. We wish to thank C. V. Kavitha, S. L. Gaonkar, and S. Nanjundaswamy, University of Mysore, for their help in preparing the small molecules. We would like to acknowledge Dr. Purushothaman Anurag, University of Alabama at Birmingham, USA and Dr. Sengottuvelan Murugan, Hokkaido University, Japan, for their fruitful discussion during the preparation of this manuscript.

Author Contributions

Conceived and designed the experiments: B KS KNT HKB PJH KSR. Performed the experiments: B KS KNT HKB KSR. Analyzed the data: B KS KNT HKB PJH KSR. Contributed reagents/materials/analysis tools: B KS HKB KSR. Wrote the paper: B KS KNT HKB PJH KSR.

- Roskoski R Jr. (2007) VEGF signaling in tumor progression. Crit Rev Oncol Hematol 62: 179–213.
- Ishida K, Wierzba MK, Teruya T, Simizu S, Osada H (2004) Novel heparan sulfate mimetic compounds as antitumor agents. Chem Biol 11: 367–377.
- Kavitha CV, Lakshmi S, Basappa (2006) Synthesis and molecular structure analysis of venlafaxine intermediate and its analog. J Chem Cryst 35: 957–963.
- Basappa, Murugan S, Kavitha CV, Purushothaman A, Nevin KG, et al. (2010)
 A small oxazine compound as an anti-tumor agent: a novel pyranoside mimetic that binds to VEGF, HB-EGF, and TNF-α. Cancer letters 297: 231–243.
- Gaonkar SL, Lokanatha Rai KM, Shetty SN (2008) Microwave-assisted synthesis and evaluation of anti-inflammatory activity of new series N-substituted 2-butyl-5-chloro-3H-imidazole-4-carbaldehyde derivatives. Med Chem Res 18: 221–230.
- Priya BS, Basappa, Swamy SN, Rangappa KS (2005) Synthesis and characterization of novel 6-fluoro-4-piperidinyl-1,2-benzisoxazole amides and 6-fluoro-chroman-2-carboxamides: antimicrobial studies. Bioorg Med Chem 13: 2623–2628.
- Saito A, Kawai K, Takayama H, Sudo T, Osada H (2008) Improvement of photoaffinity SPR imaging platform and determination of the binding site of p62/SQSTM1 to p38 MAP kinase. Chem Asian J 3: 1607–1612.
- Basappa, Murugan S, Sugahara KN, Lee CM, ten Dam GB, et al. (2009) Involvement of chondroitin sulfate E in the liver tumor focal formation of murine osteosarcoma cells. Glycobiology 19: 735–742.
- Asai T, Ueda T, Itoh K, Yoshioka K, Aoki Y, et al. (1998) Establishment and characterization of a murine osteosarcoma cell line (LM8) with high metastatic potential to the lung. Int J Cancer 76: 418–422.
- Folkman J, Klagsbrun M (1987) Vascular physiology. A family of angiogenic peptides. Nature 329: 671–672.
- Basappa, Kavitha CV, Rangappa KS (2004) Simple and an efficient method for the synthesis of 1-[2-dimethylamino-1-(4-methoxy-phenyl)-ethyl]-cyclohexanol hydrochloride: (+/-) venlafaxine racemic mixtures. Bioorg Med Chem Lett 14: 3279–3281.
- Fairbrother WJ, Champe MA, Christinger HW, Keyt BA, Starovasnik MA (1998) Solution structure of the heparin-binding domain of vascular endothelial growth factor. Structure 6: 637–648.
- Schlessinger J, Plotnikov AN, Ibrahimi OA, Eliseenkova AV, Yeh BK, et al, (2000) Crystal structure of a ternary FGF-FGFR-heparin complex reveals a dual role for heparin in FGFR binding and dimerization. Mol Cell 6, 743–50.
- Wu G, Robertson DH, Brooks CL III, Vieth M (2003) Detailed analysis of gridbased molecular docking: A case study of CDOCKER-A CHARMm-based MD docking algorithm. J Comput Chem 24: 1549–1562.
- Bao X, Mikami T, Yamada S, Faissner A, Muramatsu T, et al. (2005) Heparinbinding growth factor, pleiotrophin, mediates neuritogenic activity of embryonic pig brain-derived chondroitin sulfate/dermatan sulfate hybrid chains. The Journal of biological chemistry 280: 9180–9191.
- Lee CM, Tanaka T, Murai T, Kondo M, Kimura J, et al. (2002) Novel chondroitin sulfate-binding cationic liposomes loaded with cisplatin efficiently suppress the local growth and liver metastasis of tumor cells in vivo. Cancer Res 69: 4289–4288.
- Xing JZ, Zhu L, Jackson JA, Gabos S, Sun XJ, et al. (2005) Dynamic monitoring of cytotoxicity on microelectronic sensors. Chem Res Toxicol 18: 154–161.