

HHS Public Access

Author manuscript

Polym Chem. Author manuscript; available in PMC 2021 July 21.

Published in final edited form as:

Polym Chem. 2020 July 21; 11: 4454–4463. doi:10.1039/d0py00495b.

Hydrogen sulfide-releasing micelles for promoting angiogenesis

Jerry J. Y. Chena, **A. J. van der Vlies**b, **U. Hasegawa**b,*

aOsaka University, Department of Applied Chemistry, 2-1 Yamadaoka, Suita 565-0871, Osaka, Japan.

^bKansas State University, Tim Taylor Department of Chemical Engineering, 1005 Durland Hall, 66506, Manhattan Kansas, USA.

Abstract

Hydrogen sulfide $(H₂S)$, an important gaseous signalling molecule in the human body, has been shown to be involved in many physiological processes such as angiogenesis. Since the biological activities of H_2S are known to be significantly affected by the dose and exposure duration, the development of H_2S delivery systems that enable control of H_2S release is critical for exploring its therapeutic potential. Here, we prepared polymeric micelles with different $H₂S$ release profiles, which were prepared from amphiphilic block copolymers consisting of a hydrophilic poly(Nacryloyl morpholine) segment and a hydrophobic segment containing H_2S -releasing anethole dithiolethione (ADT) groups. The thermodynamic stability of the micelles was modulated by altering the ADT content of the polymers. The micelles with higher thermodynamic stability showed significantly slower H_2S release. Furthermore, the sustained H_2S release from the micelles enhanced migration and tube formation in human umbilical vein cells (HUVECs) and induced vascularlization in the *in ovo* chick chorioallantoic membrane (CAM) assay.

Graphical Abstract

H2S-releasing polymeric micelles promote angiogenesis

^{*} hasegawa@ksu.edu.

Conflicts of interest

There are no conflicts to declare.

[†]Electronic Supplementary Information (ESI) available: 1H NMR spectra, MTT assay data, AFM cross section curves and size distribution, cellular uptake fluorescent micelles, rates of cell migration and quantitative analysis capillary-like tube formation in HUVECs. See DOI: 10.1039/x0xx00000x

Introduction

In the human body, hydrogen sulfide (H_2S) plays important roles as a endogenous signalling molecule in the cardiovascular, immune and nervous systems.¹ Of particular interest are its roles in angiogenesis, $²$ the process of new blood vessel formation from pre-existing ones,</sup> which is a vital process during growth and development.³ The significance of endogenous H2S in angiogenesis was shown by impaired microvessel formation in aortic rings and wound healing in cystathionine-γ-lyase (CSE)-deficient mice.⁴ Therefore, H₂S has been linked to pathophysiological conditions such as ischemia injury and diabetes underscoring its therapeutic potential. Indeed, exogenously delivered H2S, in the form of the CSE substrate propargylcysteine, has been shown to induce angiogenesis in a mouse model of hindlimb ischemia and a rat model of myocardial ischemia.⁵ Furthermore, H_2S was also shown to promote angiogenesis after hindlimb ischemia in rats⁶ and improve wound-healing in type 2 diabetic mice.⁷ Further studies have shown that H_2S stimulates angiogenesis by activating the VEGF receptor and ATP-sensitive potassium (K_{ATP}) channels in endothelial cells.5,8,9

Due to difficulties in handling and dosing of H₂S gas, most studies employ sodium salts of H_2S , i.e., Na₂S or NaHS, that generate H_2S upon dissolution in water.^{4,10} However, because of their instantaneous H_2S release, which is unlikely to be relevant to the physiological H_2S generation, these compounds are not suited for studying the biological functions of H2S as well as exploring its pharmaceutical applications.⁶ Instead, approaches using substrates for H₂S-producing enzymes⁵ or small organic molecules that serve as prodrugs of H₂S (H₂S donors) are more viable.¹¹⁻¹³ Especially the latter allows for design of diverse classes of organosulfur compounds with tailored H2S release kinetics in response to endogenous factors such as hydrolysis, reaction with reducing agents like cysteine or enzymatic digestion. However, their biological effects must be evaluated carefully as these low molecular weight organic compounds and their decomposition products can cause side effects that are not due to H_2S released from these molecules.^{14, 15}

The use of polymeric materials has emerged as an alternative approach that enables exogenous administration of H2S in a controlled manner with minimal side effects. In our previously studies, we showed that chemically incorporating the $H₂S$ donor anethole

dithiolethione (ADT, Scheme 1A) into polymers, a poly(ethylene glycol) (PEG-ADT, Scheme 1B) conjugate¹⁴ and polymeric micelles of an amphiphilic block copolymer consisting of a hydrophilic poly(ethylene glycol) (PEG) segment and a hydrophobic ADTcontaining segment (PEG-PADT, Scheme 1C), led to altered intracellular trafficking thereby reducing toxic side effects of the small ADT donor.¹⁶ These polymeric H₂S donors showed a slower and continuous H_2S release compared to ADT. Furthermore, the advantage of the polymeric micelle approach has been confirmed in an in vitro ischemia model where micelles were better in protecting cardiomyocytes from apoptosis.17 The benefits of polymeric H_2S donors have also been recognized by others.^{18,19} Recently, reports have appeared tuning H_2S release by controlling mobility in the micelle core²⁰ and by shapecontrolled micellar structures.²¹

Here, we present polymeric micelles with different H_2S release profiles, which were prepared from amphiphilic block copolymers consisting of a hydrophilic poly (N-acryloyl morpholine) segment and a hydrophobic segment containing H₂S-releasing anethole dithiolethione (ADT) groups. By changing the ADT content in the polymers, the thermodynamic stability and H_2S release profiles from the micelles were modulated. In addition, the proangiogenic effects of the polymeric micelles with different H_2S release profiles was evaluated in the in vitro human umbilical vein endothelial cells (HUVECs) migration and tube formation assays as well as the in ovo chick chorioallantoic membrane (CAM) assay.

Experimental

Instrumentation

¹H NMR spectra were measured on a Bruker DPX400 NMR instrument at room temperature. For each sample a total of 32 scans was collected, and the D1 was set to 10 s. Chemical shifts are referenced to the residual undeuterated solvent signal at 2.50 ppm $(DMSO-d₆)$ or 7.26 ppm $(CDCl₃)$.

Size Exclusion Chromatography (SEC) was done on a Tosoh 8020 instrument equipped with a photo diode array (PDA) detector operating at a wavelength of 310 nm and a differential refractometer operating at 50°C. Polymer solutions (15 μL) were injected onto a KD-803 (Shodex) column kept at 50°C and eluted with 0.1 M LiCl solution in DMF. The number average molecular weight (M_n) , weight average molecular weight (M_w) , and dispersity (M_w/M_n) were calculated from the elution profiles of a PEG standard.

UV/Vis spectra were measured either with a Nanodrop 2000 or with a Tecan infinite M200 instrument in transparent polystyrene 96 well plates.

Fluorescence spectroscopy.—Fluorescence intensities were measured on a Tecan infinite M200 instrument in black polystyrene 96 well plate.

Dynamic light scattering (DLS) data were acquired on an Otsuka ELSZ-2 instrument in 4.5 mL disposable poly(methyl methacrylate) cuvettes. The diameter of the micelles in nm and the polydispersity index were calculated by the cumulant method.

Atomic force microscopy (AFM) images were acquired on a Seiko SPI300 instrument in the dynamic mode using a Seiko SI-DF20 Si probe. The micelle solutions were placed onto a fresh mica surface and water was removed by air blower before measurements.

Cell concentration was measured with the countess automated cell counter (Invitrogen) with disposable cell counting chamber slides using the trypan blue method.

An Olympus MVX10 macro zoom microscope was used to obtain bright field images of the Matrigels on the CAMs and the fluorescent images of HUVECs in the *in vitro* migration and tube formation assays.

An Olympus FV1000-D confocal laser scanning fluorescent microscope (CLSFM) was used to obtain fluorescent images of intracellular H2S release from the micelles in HUVECs.

Materials

Chemistry.—*N*,*N*-dicyclohexylcarbodiimide (DCC), diethyl ether (Et₂O), *N*,*N*dimethylformamide (DMF) (super dehydrated and peptide grade), sodium hydrogen sulfate (NaHSO₄), lithium chloride (LiCl), phosphorus pentoxide (P₄O₁₀), acetic acid (CH₃COOH), n-hexane, ninhydrin, triphenylphosphine (PPh₃), trifluoroacetic acid (TFA), azobisisobutyronitrile (AIBN), diisopropyl azodicarboxylate (DIAD), dry tetrahydrofran (THF) and anetholtrithione (ADT) were purchased from Wako Pure Chemical Industries (Osaka, Japan). 2-(tert-butoxycarbonylamino)-1-ethanol and benzyldimethyltetradecylammonium chloride (BTDACl) were purchased from Tokyo Chemical Industry (Tokyo, Japan). Triethylamine (Et₃N), ethyl acetate (EtOAc), Nmethylpyrrolidone (NMP), potassium hydroxide (KOH), 3A molecular sieves and calcium hydride (CaH2) were purchased from Nacalai Tesque (Kyoto, Japan), 1-ethylpiperidine hypophosphite (EPHP), N -hydroxysuccinimide (NHS), alumina (Al₂O₃), polyethylene oxide (PEG) GPC standard, triisopropyl silane (TIS) and phenylmethanesulfonyl fluoride (PMSF) were purchased from Sigma-Aldrich (Missouri, USA). Deuterated chloroform $(CDCl₃)$ and dimethylsulfoxide (DMSO- d_{θ}) were purchased from Cambridge Isotope Laboratories (Massachusetts, USA). Dialysis tubing with a molecular weight cut off (MWCO) 2 kDa was purchased from Spectrum Labs. Sephadex LH-20 was purchased from GE Healthcare (Little Chalfont, UK). WSP-1 was purchased from Cayman Chemical (Michigan, USA). All chemicals were used as received unless indicated otherwise. Et₃N was refluxed for 2 h over ninhydrin, filtered and distilled from KOH pellets under argon. 1,4-Dioxane was distilled from CaH2 and stored over molecular sieves in the presence of Xpell pellets to prevent peroxide formation. 3A molecular sieves were heated in a vacuum oven at 200°C for 6 h to remove water. AIBN was recrystallized from MeOH and kept at −20°C until use. CH₂Cl₂ was dried over molecular sieves overnight before use. Acryloyl morpholine was passed through a plug of Al_2O_3 to remove inhibitor. ADT-OH²² and PEG-Gly-ester-ADT¹⁴ were prepared following literature procedures.

Biology.—1-(4,5-Dimethyl-2-thiazolyl)-3,5-diphenylformazan (MTT) was purchased from Tokyo Chemical Industry (Tokyo, Japan). Slide-A-Lyzer G2 dialysis cassettes (MWCO 2 kD), acetoxymethyl calcein (calcein AM), Dulbecco's phosphate-buffered saline (D-PBS), Hank's balanced salt solution (HBSS) were purchased from Life Technologies Corporation

(Tokyo, Japan). Normal human umbilical vein endothelial cells (HUVECs), endothelial cell growth medium EGM-2 MV SingleQuot kit supplements and growth factors and reagent pack subculture reagents containing trypsin/EDTA, trypsin neutralizing solution (TNS), and HEPES buffered saline (HEPES-BSS) were purchased from Lonza (New Jersey, USA). Water used for preparation of sterile micelle solutions and cell experiments was purchased from Otsuka Pharmaceutical Factory. μ-Slide angiogenesis for the tube formation assay and the culture-inserts 2 well for self-insertion were purchased from Ibidi (Martinsried, Germany). The μ-Angiogenesis activation assay kit was purchased from Millipore (Massachusetts, USA). Fibrinogen and thrombin from human plasma were purchased from Sigma-Aldrich (Missouri, USA). Recombinant human vascular endothelial growth factor- A_{121} (VEGF₁₂₁), 16% paraformaldehyde solution (methanol free), and 25% glutaraldehyde solution were purchased from Wako (Osaka, Japan). Triple-well glass base dish, 24 and 96 well plates, culture flasks, Falcon tubes, dishes were purchased from Iwaki (Tokyo, Japan). CellTracker Green (5-chloromethylfluorescein diacetate, CMFDA) was purchased from ThermoFisher Scientific (Massachusetts, USA).

Synthesis of the PAM-PADT block copolymers—The PAM-PADT block copolymers were prepared as outlined in Scheme 2. Block copolymers **6a** and **6b** were prepared by RAFT polymerization following our previous report with modifications.²³

Synthesis of polymer 3.—Monomer **1** (185.3 mg, 1.0 mmol), 2-

(dodecylthiocarbonothioylthio)-2-methylpropionic acid **2** (14.7 mg, 0.04 mmol) and AIBN (1.32 mg, 0.008 mmol) were dissolved in 1,4-dioxane resulting in a final monomer concentration of 2 M. The clear solution was degassed by five argon freeze-pump-thaw cycles and placed in an oil bath at 70°C. After 24 h the reaction was stopped by cooling in liquid nitrogen and opening to air. The clear yellow solution was added to 200 mL hexane to precipitate the polymer. The polymer was recovered by filtration, washed with hexane and dried under vacuum to yield 169.1 mg (84%) of the homopolymer as a yellow solid. The ${}^{1}H$ NMR spectrum in CDCl₃ is shown in Figure S1 supporting information. The degree of polymerization was calculated to be 21.

Synthesis of polymers 4a and 4b.—As an example, the synthesis of **4b** is described. Acryloyl morpholine (568 mg, 4 mmol), polymer 3 (171.2 mg, 0.04 mmol) and AIBN (0.66 mg, 0.004 mmol) were dissolved in 1,4-dioxane resulting in a final monomer concentration of 2 M. The clear solution was degassed by five argon freeze-pump-thaw cycles and placed in an oil bath at 70°C. After 24 h the reaction was stopped by cooling in liquid nitrogen and opening to air. The clear yellow solution was added to 200 mL cold $Et₂O$ to precipitate the polymer. The polymer was recovered by filtration, washed with $Et₂O$ and dried under vacuum to yield 614 mg (82%) of the block copolymer as a yellow solid. The ¹H NMR spectrum in CDCl₃ is shown in Figure S2 supporting information. The degree of polymerization was calculated to be 100.

Synthesis of polymers 5a and 5b.—As an example, the synthesis of polymer **5b** is described. Polymer **4b** (384.2 mg, 0.02 mmol), 1-ethylpiperidine hypophosphite (234.1 mg, 1.3 mmol) and AIBN (2.2 mg, 0.013 mmol) were dissolved in 1.4 mL 1,4-dioxane. The

clear colourless solution was degassed by five argon freeze-pump-thaw cycles and placed in an oil bath at 95°C. After 24 h the reaction was stopped by cooling in liquid nitrogen and opening to air. The clear solution was diluted with water causing the precipitation of a small amount of solid that was removed by filtering through a 0.8 μm syringe filter. The clear solution was dialyzed (2 kDa dialysis tubing) against 4 L milliQ water for 3 d with regularly

replacing the water to remove 1-ethylpiperidine hypophosphite-related impurities. The polymer was recovered by lyophilization to yield end group-removed polymer **5b** (334.0 mg, 88%) as a white solid.

Synthesis of polymers 6a and 6b.—As an example, the synthesis of **6b** is described. Polymer **5b** (276 mg) was dissolved in 3.0 mL of TFA/H₂O (9:1) and stirred at room temperature. After 24 h the clear solution was evaporated by passing a stream of nitrogen and the residue dried under reduced pressure. The residue was dissolved in water and lyophilized to yield polymer **6b** (219 mg, 84%) as a white solid.

Synthesis of 7.—This compound was prepared in two steps with improved overall yield compared to that reported¹⁴ employing a Mitsunobu reaction (Scheme S1 supporting information). Under argon flow 560 mg (2.1 mmol) of PPh₃ in a dried Schlenk tube was dissolved in 20 mL THF. While maintaining an argon flow the solution was put in a NaCl/ice bath and cooled to −20°C. After 5 min at −20 °C 440 μL (2.2 mmol) of DIAD was added under argon. After 15 min 400 mg (1.8 mmol) of ADT-OH was added to the formed yellow suspension followed by 290 mg (1.8 mmol) of 2-(tert-butoxycarbonylamino)-1ethanol dissolved in 0.5 mL dry THF. The mixture was kept for another hour at −20°C and stirred overnight at room temperature. The mixture was concentrated under reduced pressure and purified by A_2O_3 column chromatography eluting with EtOAc/hexane 1:1. Fractions containing product were concentrated and the solid triturated with 50 mL of hexane. After decanting the hexane and drying this yielded 489 mg (1.3 mmol, 74%) of an orange solid. The 1 H NMR spectrum in CDCl₃ is shown in Figure S3 supporting information.

Deprotection was carried out following our previous report¹⁴ with the difference that a solution of Boc-protected amine was added to a solution of TIS in TFA: 118 mg (0.3 mmol) was dissolved in 6 mL CH₂Cl₂ and added dropwise to a mixture of 1 mL TIS in 3 mL CH_2Cl_2 that had been cooled at 0°C for 10 min. After everything was added the resulting mixture was stirred for 1 h at 0°C before warming to room temperature. After 1 h TLC $(SIO₂, EtOAc/hexane 1:1)$ showed the absence of the starting material. The reaction mixture was then placed in a water bath at 30° C and the TFA, TIS and CH₂Cl₂ were evaporated by gently passing a flow of air. To the oily residue was added 2-3 mL $CH₂Cl₂$ and again evaporated three time total. To the orange solid was added 25 mL Et₂O and the suspension was stirred for 10 min and the fine solid isolated by filtration on a glass filter. The solid was washed on the filter with Et₂O (2 x 25 mL) and CH₂Cl₂ (8 x 2 mL) and finally dissolved in 10 mL MeOH. After removing MeOH on the rotavapor and drying under vacuum this yielded 90.1 mg, $(0.23 \text{ mmol}, 75\%)$ of the amine as an orange solid. The ¹H NMR is shown in Figure S4 supporting information.

Synthesis of the PAM-PADT polymers 8a-c.—As an example, the synthesis of **8c** is described: 70.8 mg (4.2 μmol, 0.088 mmol COOH groups), 15.0 mg (0.13 mmol) of NHS

were dried overnight under vacuum at 60°C before dissolving in 2.1 mL of DMF. To the solution was added 21.5 mg (0.10 mmol) of DCC and 1.0 mg (0.008 mmol) DMAP in 400 μL of DMF. The reaction mixture was stirred for 24 h at room temperature during which time the solution became turbid due to the formation of dicyclohexl urea. To the mixture was then added 17.9 mg (0.047 mmol, 0.5 eq) of **7** followed by 13 μ L (0.093 mmol) of Et₃N. After 24 h acetic acid was added (1 mL) and the mixture filtered over a plug of glass wool. The residue was washed with DMF (4 x 0.5 mL) and the orange filtrate diluted with water and dialyzed against 500 mL of water (MWCO 2 kDa) for 3 days with regularly replacing the water. After lyophilizing this yielded an orange solid. The crude polymer was dissolved in DMF and purified by size exclusion chromatograph on Sephadex LH20 eluting with DMF. Fractions containing polymer were combined and diluted with 50 mL $Et₂O$ to precipitate the polymer. The ${}^{1}H$ NMR is shown in Figure S5 supporting information.

Micelle Formation—The micelle solutions were prepared in a laminar cell culture hood by adding solutions of the polymers in 200 μL NMP (25 mg/mL) drop wise to 1800 μL milliQ water under vigorous stirring at RT. After stirring for 30 min, the solution was transferred to a slide-a-lyzer G2 (MWCO 2 kDa) and dialyzed against 5 L of milliQ water for 1 day with replacing the water two times. Micelle solutions were then sterile filtered under laminar flow. The micelle concentration, after diluting 1:1 with DMF, was calculated from the absorbance at 436 nm and a standard curve of ADT in the same solvent mixture.

Critical micelle concentration (CMC) measurement—Micelle solutions at different concentrations (2 μL) were placed onto a parafilm-coated surface and the contact angle was measured using a Kyowa Interface Science Drop Master DM 300 instrument. From a plot of the contact angle versus polymer concentration the critical micelle concentration was calculated. Measurements were done in triplicate.

Cell culture—HUVECs were cultured in endothelial cell growth medium EGM-2 supplemented with 2% FBS, 0.04% hydrocortisone, 0.4% hFGF, 0.1% VEGF, 0.1% R3- IGF-1, 0.1% ascorbic acid, 0.1% hEGF, 0.1% GA-1000, and 0.1% heparin in 5% $CO₂$ inside an incubator at 37°C. Cells were subcultured when reaching 70-80% confluency. Cells were washed with HEPES-BBS and 2 mL of trypsin/EDTA was added. Periodically, the trypsinization process was checked under the microscope and after 2-3 min the trypsin was neutralized by adding 4 mL of TNS. The cell suspension was centrifuged at 1200 rpm for 5 min and after removing the supernatant resuspended in 3 mL of medium. This cell suspension was then diluted to the concentration needed for the experiment.

Preparation of HUVEC cell lysate for the H2S release experiments—HUVECs were detached and the cell suspension was centrifuged at 500 rpm for 5 min. After washing the cell pellet with cold PBS, the cells were resuspended in PBS supplemented with 1 mM PMSF (Final concentration: 1.4×10^6 cells/mL). The cell suspension was sonicated for 1 min with a Taitec VP-050N ultrasonic homogenizer and centrifuged. The supernatant was collected and stored at −80°C until use.

H2S release from the micelles in HUVEC lysate with the WSP-1 H2S fluorescent detection probe—Cell lysate was diluted with PBS at a volume ratio of 1:4

and mixed with WSP-1 and BTDACl (Final concentrations: 10 μM for WSP-1 and 2.5 μM for BTDACl). This solution was mixed with the micelle solutions in a 96 well plate (Final concentration of ADT groups: 100 μM). The fluorescence intensity was measured as function of time on the plate reader (λ_{ex} = 465 nm, λ_{em} = 515 nm).

H2S release of the micelles in HUVECs—HUVECs were seeded in a 96 well plate at 7.5 x 10^3 cells/well and cultured for 2 d. After removing the medium, cells were incubated with 25 μM WSP-1/PBS at room temperature for 30 min. After the WSP-1 solution was removed, cells were cultured in medium containing the micelles **8a**, **8b** and **8c** (ADT concentration: 50 μM). At the indicated time points, the fluorescence intensity was measured $(\lambda_{\text{ex}}=465 \text{ nm}, \lambda_{\text{em}}=515 \text{ nm})$ using the bottom reading mode.

Intracellular H2S detection by CLSFM—HUVECs were seeded into a triple-well glass base dish at 1×10^4 cells/well and cultured for 24 h in a CO₂ incubator. The medium was replaced with 100 μL fresh medium containing ADT or the micelles (Final ADT concentration: 100 μM). After culturing for 6 h, the cells were washed with FBS-free medium and incubated in 100 μL FBS-free medium containing 10 μM WSP-1 and 2.5 μM BTDACl for 30 min at 37°C. The cells were washed with PBS and observed by CLSFM.

MTT assay—HUVECs were seeded in a 96 well plate at 1×10^4 cells/well and cultured for 24 h. After replacing the medium with 100 μL of fresh medium containing ADT or the micelles at different concentrations, cells were cultured for 24 h. The medium was replaced with 100 μ L of fresh medium containing 0.5 mg/mL MTT. After culturing for 4 h, 100 μ L/ well of 0.1 g/mL sodium dodecyl sulfate in 0.01 M HCl (aq) was added to lyse the cells and solubilize the formazan crystals. The absorbance at 570 nm was measured and the cell viability is expressed as the percentage of the absorbance for cells that had not received any treatment.

Migration Assay—HUVECs at 70-85% confluency were stained with Cell Tracker solution (1 μg/mL) in HBSS for 15 min at 37°C. Cells were detached by trypsinization, resuspended in EBM-2 medium containing 2% FBS and seeded at 3.5×10^4 cells (70 µL medium) in an ibidi Culture-Insert 2 well, which is placed in a 24-well plate. After culturing for 24 h, the culture-inserts were removed to create a confluent cell monolayer with a cellfree gap of 500 μm. To each well was added 300 μL medium containing $VEGF₁₂₁$, ADT or the micelles (Final concentrations: 100 ng/mL for $VEGF₁₂₁$, 100 μM for ADT or ADT groups of the micelles). Cells were cultured in a $CO₂$ incubator and observed under a fluorescence microscope at the indicated time points. The MetaMorph software was used to analyze the images and to calculate the rate of cell migration.²⁴

Tube formation assay—Frozen solutions of fibrinogen and thrombin were thawed inside an icebox while cooling the μ-Slide Angiogenesis plates inside the fridge. To the wells on the cold μ-Slide Angiogenesis plate was then added 5.5 μL of fibrinogen solution followed by 5.5 μL thrombin solution. The resulting fibrinogen-thrombin solution mixture was gently mixed by pipetting. The μ-Slide Angiogenesis plate was then placed inside the cell incubator at 37˚C for 1 h to form the fibrin gel. To each well was added 45 μL of a HUVEC cell suspension (2.3 \times 10⁶ cells/mL) and 5 µL of either water (NT), VEGF₁₂₁, ADT or the

micelle solutions (Final concentrations: 100 ng/ mL for VEGF₁₂₁, 330 μM for ADT or ADT groups of the micelles). After culturing for 24 h, the cells were stained with calcein AM and observed under a macro zoom fluorescence microscope.

Preparation of the Matrigels with VEGF and the H2S donors for the CAM assay —PBS solutions of the micelles **8a-c**, PEG-Gly-ester-ADT and VEGF₁₂₁ (5 μL) were mixed with cold growth factor reduced Matrigel (30 μL) and kept at 37°C for 30 min to form the

Matrigels (Final concentrations: 11 μg/mL for $VEGF₁₂₁$, 400 μM for ADT or ADT groups of the micelles).

Chick Chorioallantoic Membrane (CAM) Assay.—On embryonic day 1, the eggs were cleaned with tissue paper wetted with 70% ethanol (aq) and put inside the incubator at 37.6°C. On day 3 the eggs were candled with a LED light source to locate the air sac. The egg was then punctured at this side with an egg hole puncher and through the hole about 2-3 mL albumen was removed using a 21 G x $11/2$ plastic syringe.²⁵ The hole was closed with Opsite tape (Smith and Nephew). Then a square window of about 2 cm x 2 cm was cut in the centre of the egg and the shell and inner shell membrane carefully removed with a pincer. The cut window was covered with Opsite and the area above the window cut away with scissors and covered with Tegaderm tape (3M). The eggs were then placed back inside the incubator. At day 9, eggs were candled and those containing a live embryo (60-77%) were used for the experiment. The Tegaderm covering the window on the egg was cut away and the Matrigels placed onto the CAM using sterilized tweezers. After resealing the window with Tegaderm tape the egg was put back in the incubator. On day 11, the CAM was fixed by adding 2 mL of fixation solution containing methanol-free paraformaldehyde (16 w/v%) solution and glutaraldehyde (25 w/v%) solution, in 20 mM PBS pH7.4 at room temperature for 1 d. The CAM was taken out using a pincer and scalpel, washed with PBS and placed inside a 6 well plate. The images were collected using a macro zoom microscope and analysed by blind test using the reported method.²⁶

Results and discussion

Synthesis of ADT-containing diblock copolymers (PAM-PADT)

The amphiphilic PAM-PADT block copolymers consisting of a hydrophilic poly (N-acryloyl morpholine) (PAM) segment and a hydrophobic ADT-containing (PADT) segment were synthesized as outlined in Scheme 2. We first prepared block copolymers consisting of a carboxylic acid-containing block and PAM block using the reversible addition fragmentation transfer (RAFT) polymerization technique. RAFT polymerization of tert-butyl glycine acrylamide **1** using thiocarbonate **2** as the chain transfer agent (CTA) gave homopolymer **3**. This polymer had a dispersity of 1.06 as determined by gel permeation chromatography (GPC). The homopolymer **3** was then used as a macro CTA to polymerize N-acryloyl morpholine to yield diblock copolymers **4a** and **4b**. The dispersity of both polymers was 1.07 and 1.12 respectively, showing the successful synthesis of the diblock copolymers with narrow size distributions.

Then, the CTA end-group of polymers **4a** and **4b** was removed by radical-induced reduction using 1-ethylpiperidine hypophosphite as the hydrogen source.²⁷ Successful thiocarbonate

group removal was confirmed by the absence of the absorbance at 310 nm as measured by the PDA detector connected to the GPC instrument. After removing the *tert*-butyl esters groups with trifluoroacetic acid, the carboxylic acid-containing polymers **6a** and **6b** were first activated by preparing the corresponding N-hydroxysuccinimide esters and used to conjugate with the amine linker-modified ADT **7** to yield the PAM-PADT polymers (**8a**-**c**).

We obtained the polymers with different ADT contents by changing the molar feed ratio of ADT (**7**) to -COOH groups of polymers **6a** and **6b** (21 per polymer). The number of ADT groups per polymer (x) of **8a**, **8b** and **8c** were 21, 18 and 12, respectively, as determined from the 1 H NMR spectra (Table 1).

Micelle preparation and characterization

The micelles were prepared by dispersing the PAM-PADT diblock copolymers (**8a**-**c**) in water. Due to the hydrophobic nature of the ADT groups, the block copolymers selfassembled in water to form micelles with an average hydrodynamic diameter (D_h) of 34-37 nm as measured by dynamic light scattering (DLS) as shown in Table 1. Furthermore, the spherical structures of the micelles were confirmed by atomic force microscopy (AFM) as shown in Figure 1. Average size distribution of the micelles as well as cross-section curves can be found in Table S1 and Figure S6 (supporting information).

To see how the ADT content of the polymers affects the thermodynamic stability of the micelles, the critical micelle concentration (CMC), the polymer concentration above which micelles form, were determined by the surface tension method.16 As shown in Table 1, the CMC values follow the order: $8a < 8b < 8c$. This result shows that the micelles prepared from the polymers with a higher ADT content (i.e., **8a**) has a higher thermodynamic stability, which is likely to be due to the increased hydrophobicity of the PADT segment.

H2S release from the micelles in presence of HUVEC's cell lysate

The effect of the thermodynamic stability of the PAM-PADT micelles on their $H₂S$ release profiles was evaluated. In our previous report, we showed that $H₂S$ release from ADT was induced by cell lysate from mouse macrophages¹⁶ and rat cardiomoycytes.¹⁷ In this study, we measured H₂S release from the micelles in cell lysate from HUVECs using the WSP-1 $H₂S$ detection probe.²⁸ Interestingly, the $H₂S$ release profiles differed significantly between the micelles. As shown in Figure 2, the order is **8a < 8b < 8c**, suggesting that the micelles with a higher stability (i.e., lower CMC) show the slowest H_2S release.

We previously showed that ADT-based H_2S donors release H_2S in cell lysate but not fetal bovine serum or glutathione-containing PBS.¹⁴ This implies that H_2S release is induced by intracellular components, presumably enzymes as was recently reported for rat liver microsomes.29 Since the ADT groups, which are located within the micellar core, are unlikely to be accessible for enzymes, it is reasonable to assume that micelles partially dissociate in the presence of cell lysate, resulting in H2S release. This hypothesis is not contradicted by the result in Figure 2 showing that the micelles with a higher CMC (**8c**), i.e. low thermodynamic stability, exhibits faster H_2S release.

H2S release from the micelles in HUVECs

We compared the H2S release profiles of the micelles **8a**, **8b** and **8c** in live cells. HUVECs were pretreated with the WSP-1 $H₂S$ detection probe and cultured in the presence of micelles. The fluorescence intensity within the cells was measured by a plate reader using the bottom reading mode for real time monitoring of H_2S release in HUVECs. As shown in Figure 3, the H₂S release follows the same trend as that observed in cell lysate.

To show that H_2S release occurred intracellularly we used the WSP-1 H_2S detection probe to visualize intracellular H2S by confocal laser scanning fluorescence microscopy (CLSFM). HUVECs were treated with the different H₂S donors (100 μ M) for 6 h at 37°C and observed by CLSFM. We observed increased fluorescence within cells treated with the small H2S donor ADT (Figure 4B) as well as the micelles (Figure 4C-E) compared to the non-treated cells (Figure 4A). These CLSFM images clearly show that ADT as well as the micelles can release H2S inside HUVECs. Furthermore, in the case of ADT, due to its low water solubility, small crystallites of ADT appeared, as indicated by the black spots in the DIC image in Figure 4B.

Furthermore, we investigated whether the micelles can be internalized by HUVECs. The micelles were prepared from cyanine3 (Cy3)-labeled polymers **8b** and **8c** (Figure S7, supporting information). As shown in Figure S8 (supporting information), HUVECs treated with the micelles showed bright fluorescence, supporting that the micelles were taken up by cells and thereafter released H2S intracellularly.

Pro-angiogenic effects of the micelles in the HUVEC migration and tube formation assays

H2S is known to induce angiogenesis by stimulating endothelial cell migration and tubulogenesis.^{5,8} We therefore tested the proangiogenic activity of the micelles by the gap closure migration and tube formation assays using HUVECs.

The effect of ADT and the micelles on migration of HUVECs was evaluated in the gap closure migration assay where a confluent cell monolayer with an artificial cell free gap (width: 500 μm) was first prepared and cell migration into the gap was monitored. As shown in Figure 5 and S9A-E in the supporting information, the micelles enhanced migration into the cell free gap and about 75% of the gap surface was covered over the course of 9 h. Note that the effect on cell migration is very similar with that of the growth factor $VEGF_{121}$.

Compared to the micelles, ADT showed less migratory activity. This may be related to its higher toxicity compared to micelles in HUVECs as shown in Figure S10 in the supporting information. Furthermore, the images obtained at different time points were analysed to determine the rate of cell migration (Figure S9F in the supporting information). Compared to non-treatment (NT) and ADT, the micelles showed a higher cell migration rate. These results show that micelles efficiently stimulate migration of HUVECs compared to ADT. There was no obvious effect of the difference in H_2S release profiles of the micelles on cell migration.

We then tested tubulogenic activity of the micelles in the *in vitro* tube formation assay. Cells were seeded on fibrin gels and cultured for 24 h in medium containing $VEGF₁₂₁$, ADT and

micelles. As shown in Figure $6B$, in the presence of $VEGF₁₂₁$, cells have migrated and aligned themselves to form tubular structures, as indicated by the white arrows, which were not observed for the non-treated cells (Figure 6A). The micelles showed formation of capillary tubes (Figures 6D-F) indicating the tubulogenic effect of H_2S released from the micelles. Interestingly, the number of capillary-like tubes formed by the micelles was in the order of **8a < 8b < 8c** as shown by a quantitative analysis (Figure S11 supporting information). This suggests that the tubulogenesis of HUVECs is affected by the $H₂S$ release rate. The micelles with a faster H2S release rate stimulate endothelial cell tube formation more efficiently.

Pro-angiogenic effect of the micelles in the in ovo chick chorioallantoic membrane assay

To evaluate the pro-angiogenic effects of the micelles in a more realistic model, we used the in ovo chick chorioallantoic membrane (CAM) assay.³⁰ The CAM is a vascularized membrane that forms around day 4 to 5 of embryonic development that covers the chick embryo. During embryo development, the CAM capillary endothelium develops into a dense network of arteries and veins in the mesodermal layer. From this network, the capillary plexus is formed, i.e., branching network of blood vessels, that serves as the respiratory organ until the time of hatching.^{31,32} Due to its unique capillary plexus formation, CAM has been used to study the pro-angiogenic effect of drugs including $Na₂S^{4,33,34}$

We placed growth factor reduced Matrigels containing $VEGF₁₂₁$ and the micelles on the CAMs on embryonic day 9. The Matrigels containing ADT could not be prepared due to its low aqueous solubility. Instead, we used a PEG-Gly-ester-ADT conjugate having ADT linked via a hydrolysable ester bond as a prodrug of ADT-OH.¹⁴ As shown in our previous report, this conjugate is rapidly hydrolyzed in aqueous solutions to release ADT-OH. Although ADT-OH lacks the methyl group, this compound is also known to release H_2S similar to ADT. On embryonic day 12, the CAM area around the Matrigels was taken out and observed under a macro zoom microscope (Figure 7A-F).

Compared to the non-treated CAMs (NT), the CAM treated with all formulations increased the number of blood vessels that are sprouting out from the area where Matrigels were placed. We further used the reported semi-quantitative scoring system to compare the proangiogenic effect of the different formulations.²⁶ As can be seen in Figure 7G, the micelles showed significantly stronger pro-angiogenic effects compared to NT ($p < 0.001$) while no statistically significant effects was observed for ADT-OH. All micelles were more effective in inducing angiogenesis compared to ADT-OH, and the micelles **8b** and **8c** showed a significantly stronger effect than $VEGF_{121}$. These data clearly show the advantage of using the micelles for promoting vascularization on the CAMs. Furthermore, although the difference between the micelles **8b** and **8c** was not significant, the proangiogenic activities of both micelles **8b** and **8c** were significantly stronger compared to **8a** (p < 0.05) showing that the H₂S-release profile affects vascularization in the CAM assay (Figure 7).

Conclusions

We prepared amphiphilic block copolymers composed of a hydrophilic poly $(N$ -acryloyl morpholine) segment and a hydrophobic segment bearing different numbers of ADT groups,

using the RAFT polymerization technique. The polymers assembled in water to form spherical micelles. The thermodynamic stability of the micelles, which was determined by critical micelle concentration, was enhanced by increasing the number of ADT groups. The micelles with higher thermodynamic stability showed slower H_2S release in cell lysate of HUVECs. The micelles showed stronger pro-angiogenic effects in the in vitro migration and tube formation assay compared to the small H_2S donor ADT. Furthermore, the micelles efficiently induced vascularization in the *in ovo* CAM assay. The micelle approach would have promise in promoting angiogenesis by delivering H2S in a controlled manner.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgements

This work was supported by Center for Molecular Analysis of Disease Pathways (CMADP), National Institute of General Medical Sciences of the National Institutes of Health, Award Number P20GM103638.

Notes and references

- 1. Kimura H, Molecules, 2014, 19, 16146–16157. [PubMed: 25302704]
- 2. Mistry RK and Brewer AC, Free Radical Biology and Medicine, 2017, 108, 500–516. [PubMed: 28433660]
- 3. Carmeliet P, Nature Medicine, 2003, 9, 653–660.
- 4. Papapetropoulos A, Pyriochou A, Altaany Z, Yang G, Marazioti A, Zhou Z, Jeschke MG, Branski LK, Herndon DN, Wang R and Szabó C, Proceedings of the National Academy of Sciences, 2009, 106, 21972–21977.
- 5. Kan J, Guo W, Huang C, Bao G, Zhu Y and Zhu YZ, Antioxidants & Redox Signaling, 2014, 20, 2303–2316. [PubMed: 24180631]
- 6. Wang MJ, Cai WJ, Li N, Ding YJ, Chen Y and Zhu YC, Antioxidants & Redox Signaling, 2010, 12, 1065–1077. [PubMed: 19842913]
- 7. Liu F, Chen D-D, Sun X, Xie H-H, Yuan H, Jia W and Chen AF, Diabetes, 2014, 63, 1763–1778. [PubMed: 24487028]
- 8. Cai W-J, Wang M-J, Moore PK, Jin H-M, Yao T and Zhu Y-C, Cardiovascular Research, 2007, 76, 29–40. [PubMed: 17631873]
- 9. Mustafa AK, Sikka G, Gazi SK, Steppan J, Jung SM, Bhunia AK, Barodka VM, Gazi FK, Barrow RK, Wang R, Amzel LM, Berkowitz DE and Snyder SH, Circulation Research, 2011, 109, 1259– 1268. [PubMed: 21980127]
- 10. Yuan S, Pardue S, Shen X, Alexander JS, Orr AW and Kevil CG, Redox Biology, 2016, 9, 157– 166. [PubMed: 27552214]
- 11. Zheng Y, Ji X, Ji K and Wang B, Acta Pharmaceutica Sinica B, 2015, 5, 367–377. [PubMed: 26579468]
- 12. Szabo C and Papapetropoulos A, Pharmacological Reviews, 2017, 69, 497–564. [PubMed: 28978633]
- 13. Powell CR, Dillon KM and Matson JB, Biochemical Pharmacology, 2018, 149, 110–123. [PubMed: 29175421]
- 14. Hasegawa U and van der Vlies AJ, Bioconjugate Chemistry, 2014, 25, 1290–1300. [PubMed: 24942989]
- 15. Yuan S, Patel RP and Kevil CG, American Journal of Physiology-Lung Cellular and Molecular Physiology, 2015, 308, L403–L415. [PubMed: 25550314]
- 16. Hasegawa U and van der Vlies AJ, MedChemComm, 2015, 6, 273–276.

- 17. Takatani-Nakase T, Katayama M, Matsui C, Hanaoka K, van der Vlies AJ, Takahashi K, Nakase I and Hasegawa U, Molecular BioSystems, 2017, 13, 1705–1708. [PubMed: 28681875]
- 18. Connal LA, Journal of Materials Chemistry B, 2018, 6, 7122–7128. [PubMed: 32254628]
- 19. Urquhart MC, Ercole F, Whittaker MR, Boyd BJ, Davis TP and Quinn JF, Polymer Chemistry, 2018, 9, 4431–4439.
- 20. Foster JC, Carrazzone RJ, Spear NJ, Radzinski SC, Arrington KJ and Matson JB, Macromolecules, 2019, 52, 1104–1111. [PubMed: 31354172]
- 21. Yu SH, Esser L, Khor SY, Senyschyn D, Veldhuis NA, Whittaker MR, Ercole F, Davis TP and Quinn JF, Journal of Polymer Science Part A: Polymer Chemistry, 2019, 57, 1982–1993.
- 22. Li L, Rossoni G, Sparatore A, Lee LC, Del Soldato P and Moore PK, Free Radical Biology and Medicine, 2007, 42, 706–719. [PubMed: 17291994]
- 23. Hasegawa U, Tateishi N, Uyama H and van der Vlies AJ, Macromolecular Bioscience, 2015, 15, 1512–1522. [PubMed: 26102371]
- 24. van der Vlies AJ, Morisaki M, Neng HI, Hansen EM and Hasegawa U, Bioconjugate Chemistry, 2019, 30, 861–870. [PubMed: 30676733]
- 25. Samkoe KS, Clancy AA, Karotki A, Wilson BC and Cramb DT, Journal of Biomedical Optics, 2007, 12, 1–14.
- 26. Ribatti D, Nico B, Vacca A and Presta M, Nature Protocols, 2006, 1, 85–91. [PubMed: 17406216]
- 27. Hasegawa U, van der Vlies AJ, Simeoni E, Wandrey C and Hubbell JA, Journal of the American Chemical Society, 2010, 132, 18273–18280. [PubMed: 21128648]
- 28. Peng B, Chen W, Liu C, Rosser EW, Pacheco A, Zhao Y, Aguilar HC and Xian M, Chemistry A European Journal, 2014, 20, 1010–1016.
- 29. Dulac M, Nagarathinan C, Dansette PM, Mansuy D and Boucher J-L, Drug Metabolism and Disposition, 2018, 46, 1390–1395. [PubMed: 30018103]
- 30. Mary R and Gurmit S, Current Drug Targets Cardiovascular & Hematological Disorders, 2003, 3, 155–185. [PubMed: 12769641]
- 31. Ribatti D, Nico B, Vacca A, Roncali L, Burri PH and Djonov V, The Anatomical Record, 2001, 264, 317–324. [PubMed: 11745087]
- 32. Dimitropoulou C, Malkusch W, Fait E, Maragoudakis ME and Konerding MA, Angiogenesis, 1998, 2, 255–263. [PubMed: 14517465]
- 33. Melkonian G, Munoz N, Chung J, Tong C, Marr R and Talbot P, Journal of Experimental Zoology, 2002, 292, 241–254.
- 34. Vázquez F, Hastings G, Ortega M-A, Lane TF, Oikemus S, Lombardo M and Iruela-Arispe ML, Journal of Biological Chemistry, 1999, 274, 23349–23357

Figure 1.

Morphology of the micelles (**A**) **8a**, (**B**) **8b** and (**B**) **8c** by atomic force microscopy (AFM). The micelles were adsorbed onto a fresh mica surface and observed by AFM after removing water by air blower. The images show the height image with the height profile from 0 to 4.87 nm indicated below each image.

Figure 2.

H2S release profiles of the micelles **8a**, **8b** and **8c**. The micelles (ADT concentration: 100 μ M) were added to HUVEC cell lysate containing the WSP-1 H₂S detection fluorescent probe (10 μM). $λ_{ex}$ = 465 nm, $λ_{em}$ = 515 nm. n=3.

Figure 3.

 $H₂S$ release in HUVECs. Cells in a 96 well plate were pretreated with the WSP-1 $H₂S$ detection probe in PBS for 30 min and cultured in medium without (NT, squares) or with micelles **8a** (rhombus), **8b** (triangles) and **8c** (circles). ADT concentration: 50 μM. The fluorescence intensity was monitored using the bottom reading mode (λ_{ex} =465 nm, λ_{em} =515 nm). * $p < 0.05$, ** $p < 0.01$,*** $p < 0.001$ (n=3).

Figure 4.

H2S release from the micelles in HUVECs. The cells were treated with (**A**) NT (PBS), (**B**) ADT, (**C**) micelles **8a**, (**D**) **8b** and (**E**) **8c** for 6 h. Cells were washed with PBS and treated with the WSP-1 probe to detect H_2S within cells. ADT concentration: 100 μM. Scale bars: 50 μm.

Figure 5.

The gap closure migration assay. HUVECs were seeded in the ibidi Culture-Insert 2 Well to create a confluent cell monolayer with an artificial gap. After removing the insert, cells were cultured in culture medium containing (A) NT (water), (B) VEGF₁₂₁ (100 ng/mL), (C) ADT, (**D**) micelles **8a**, (**E**) **8b** and (**F**) **8c**. ADT concentration 100 μM. Top panels: the cell free gap at 0 h, Bottom panels: the same area after 9 h. Scale bar: 500 μm.

Figure 6.

Tube formation assay. Representative images of HUVECs treated with (**A**) NT (water), (**B**) VEGF121, (**C**) ADT, (**D**) micelles **8a**, (**E**) **8b** and (**F**) **8c**. HUVECs were seeded on fibrin gel and treated with 100 ng/mL VEGF₁₂₁, ADT and micelles (ADT concentration: 330 μ M) for 24 h. The cells were stained with calcein AM and observed under a macro zoom fluorescence microscope. Arrows indicate capillary tubes. Scale bar: 500 μm.

Figure 7.

Pro-angiogenic effect in the CAM assay. Photographs of CAMs treated with Matrigel containing (**A**) NT (PBS), (**B**) VEGF₁₂₁ (11 μg/mL), (**C**) ADT-OH (PEG-Gly-ester-ADT), (**D**) micelles **8a,** (**E**) **8b** and (**F**) **8c**. ADT concentration: 400 μM. The location where the gel was applied has been indicated by the circled area. Scale bar: 2 mm. (**G**) Semi-quantitative scoring. * $p < 0.001$ versus NT, $^{#}p < 0.001$, $^{#}p < 0.05$ versus VEGF, $^{*}p < 0.001$, $^{*}p < 0.05$ versus ADT-OH. $\frac{4}{7}$ p < 0.05 versus **8a**. n= 8-10.

Scheme 1.

Structures of (**A**) ADT, (**B**) PEG-ADT conjugate and (**C**) micelle-forming PEG-PADT block copolymer.

Scheme 2.

Synthesis of the PAM-PADT diblock copolymers.

(a) AIBN, 1,4-dioxane, 70°C, (b) acryloyl morpholine, AIBN, 1,4-dioxane, 70°C, (c) EPHP, AIBN, 1,4-dioxane, 95°C, (d) TFA/H₂O, (e) NHS, DMAP, DMF, 25°C, (f) Et₃N, DMF, 25°C. For explanation of the used abbreviations see the materials section.

Table 1.

PAM-PADT polymers and their micelles.

 α ²Number of ADT groups per polymer as determined by ¹H NMR

 b _{Z-average hydrodynamic diameter} (D_h) and polydispersity index (PDI=μ₂Γ⁻²) of the micelles as determined by dynamic light scattering (DLS) after fitting the correlation function using the cumulant method.

 c Critical micelle concentration (CMC) as determined by the surface tension method.

 $d_{p=0.0031}^{d}$ vs. **8a**.

 $e^b_p = 0.035$ vs. **8b**.