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## Carbohydrate Composition of Amphiphilic Macromolecules Influences Physicochemical Properties and Binding to Atherogenic Scavenger Receptor A

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### Abstract

Amphiphilic macromolecules (AMs) based on carbohydrate domains functionalized with poly(ethylene glycol) can inhibit the uptake of oxidized low density lipoprotein (oxLDL) mediated by scavenger receptor A (SR-A) and counteract foam cell formation, the characteristic “atherosclerotic” phenotype. A series of AMs were prepared by altering the carbohydrate chemistry to evaluate the influence of backbone architecture on the physicochemical and biological properties. Upon evaluating the degree of polymer-based inhibition of oxLDL uptake in human embryonic kidney cells expressing SR-A, two AMs (**2a** and **2c**) were found to have the most efficacy. Molecular modeling and docking studies show that these same AMs have the most favorable binding energies and most close interactions with the molecular model of SR-A collagen-like domain. Thus, minor changes in the AMs architecture can significantly affect the physicochemical properties and inhibition of oxLDL uptake. These insights can be critical for designing optimal AM-based therapeutics for management of cardiovascular disease.

### Keywords

Amphiphilic polymer; atherosclerosis; self-assembled micelle; oxLDL inhibition

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### Supplementary Information Available

Molecular modeling was performed to demonstrate how the polymers interact with the SR-A homology model. This data is a qualitative representation of the AMs binding to SR-A. This material is available free of charge via the Internet at [www.sciencedirect.com](http://www.sciencedirect.com).

## 1. INTRODUCTION

Atherosclerosis, the occlusive artery disease, is triggered by the build-up of oxidized low density lipoprotein (oxLDL) in blood vessel walls.[1] The oxLDL accumulation generates an inflammatory response which results in the recruitment of circulating monocytes followed by their differentiation to macrophages and upregulation of macrophage membrane-based scavenger receptors.[2] The major scavenger receptors, namely scavenger receptor A (SR-A) and CD36, mediate the uptake of oxLDL,[3–5] leading to unregulated cholesterol accumulation and foam cell formation, a key characteristic of the onset of atherogenesis.[6,7]

Aside from lifestyle changes, cholesterol lowering therapies (i.e., statins) are the most common methods for atherosclerosis treatment. Such drugs, however, indirectly mediate atherosclerosis by decreasing cholesterol synthesis which ultimately results in decreased oxLDL in the blood vessel walls. A more direct and promising approach in the treatment and prevention of atherosclerosis involves designing functional inhibitors against scavenger receptors to abrogate uncontrolled oxLDL uptake.[8–11] Our research group has previously presented carbohydrate-based, nanoscale amphiphilic macromolecules (AMs) that competitively inhibit scavenger receptor-mediated oxLDL uptake.[12,13] Comprised of a hydrophobic domain based on the sugar, mucic acid, acylated with four aliphatic chains, and a hydrophilic poly(ethylene glycol) (PEG) tail, the AMs exhibit high biocompatibility and stability.[14] Further, the AMs self-assemble into micelles in aqueous media at relatively low ( $10^{-7}$  M) concentrations, enabling drug encapsulation within the micellar core. To understand the key structural features relevant to oxLDL inhibition, systematic variations to the AM structure were performed, including the PEG chain length, PEG architecture and aliphatic chain length as well as type, charge, number, and rotational motion of anionic charges. These studies also tested whether amphiphilicity was a significant attribute in preventing oxLDL internalization.[15,16] The structure-function studies demonstrated that while amphiphilicity was necessary, it alone was insufficient; charge and hydrophobicity better contributed to the polymer's ability to inhibit oxLDL uptake. Overall, anionic charge and a rigid, hydrophobic carboxylic acid presentation significantly enhanced the inhibitory activity of AMs.

The AM hydrophobic domain compositions have not been optimized to date, however, and the impact of carbohydrate stereochemistry and conformation remain to be elucidated. Minute changes in carbohydrate stereochemistry and conformation can potentially have far-reaching consequences on polymer physicochemical and biological properties.[17–20] For example, Wang *et al.* demonstrated that the L configuration of poly(N-acryloyl-L(D)-valine) films promote better cytocompatibility (i.e., adhesion, growth, assembly and proliferation) compared to the D configuration.[19] Additionally, the introduction of aromaticity can significantly affect the properties of the resultant polymers.[21] To this end, several AMs were synthesized with incremental backbone modifications to investigate the influence of carbohydrate stereochemistry, aromatic groups and cyclic carbohydrates that are incorporated within the polymer backbone on biological activity, specifically, oxLDL inhibition (Figure 1). Molecular modeling and docking studies were employed to quantify the structure-activity relationship between the polymers and SR-A scavenger receptors. Previous experiments performed with macrophages (both mouse and human) indicated that the polymers were acting, in part, through interactions with SR-A.[12, 15,16,22,23] Although polymer interactions with CD36 and other surface proteins were observed, the goal of the studies outlined in this manuscript was to isolate the interactions of the polymers with SR-A *only*. To confirm the extent of backbone influence on scavenger receptor binding, a functional assay was developed. The AMs were evaluated for their ability to inhibit SR-A mediated oxLDL uptake in human embryonic kidney cells expressing the SR-

A scavenger receptor (HEK-SRA). These cells have a well-established history of use for studying various SR-A functions due to their over expression of the SR-A construct.[23–26] AM binding to the cells was then determined by quantifying the amount of a competitor molecule (oxLDL) bound and internalized by the cells after 24 hours incubation in the presence of the polymers.

## 2. MATERIAL AND METHODS

### 2.1. Synthetic Materials

All reagents and solvents were purchased from Sigma-Aldrich and used as received unless otherwise noted. 4-(dimethylamino)pyridinium p-toluene-sulfonate (DPTS) was prepared as described by Moore and Stupp.[27] Monomethoxy-poly(ethylene glycol) (mPEG, Mn = 5000 Da) was azeotropically distilled with toluene prior to use. The synthesis of **2a** and **2d** have been previously reported.[14,28]

### 2.2 Instrumentation

<sup>1</sup>H-NMR spectra were obtained using a Varian 400 MHz or 500 MHz spectrophotometer with TMS as internal reference. Samples were dissolved in CDCl<sub>3</sub>, or CDCl<sub>3</sub> with a few drops of DMSO-d<sub>6</sub> if necessary. Molecular weights (MW) were determined using gel permeation chromatography (GPC) with respect to PEG standards (Sigma-Aldrich) on a Waters Stryagel® HR 3 THF column (7.8 × 300 mm). The Waters LC system (Milford, MA) was equipped with a 2414 refractive index detector, a 1515 isocratic HPLC pump, and 717plus autosampler. Samples (10 mg/mL) were dissolved in THF and filtered using 0.45 μm pore size nylon or PTFE syringe filters (Fisher Scientific). Dynamic light scattering (DLS) analysis was carried out on a Zetasizer nanoseries ZS90 (Malvern instruments). CMC studies were carried out on a Spex fluoromax-3 spectrofluorometer (Jobin Yvon Horiba) at 25 °C. Melting points were determined by DSC on a TA DSC Q200. TA universal analysis 2000 software was used for data collection on a Dell dimension 3000 computer. Samples (4–8 mg) were heated under dry nitrogen gas. Data were collected at heating and cooling rates of 10 °C min<sup>-1</sup> with a two-cycle minimum.

### 2.3. Polymer synthesis (Figure 2)

#### 2.3.1. Synthesis of **2b**

- a. D-Saccharic acid (2.0 g, 8.1 mmol) and zinc chloride (0.26 g, 1.9 mmol) were stirred with lauroyl chloride (26 mL, 12 mmol). The solution was heated to 95 °C in a temperature-controlled oil bath and stirred under argon overnight. Diethyl ether (100 mL) and water (30 mL) were added and the solution was allowed to stir for 45 minutes. The organic phase was washed with water (5 × 100 mL) and then concentrated via rotary evaporation. The concentrated solution was precipitated from hexanes (2.5 L) yielding intermediate **1b** as a white solid (4.2 g, 56%) <sup>1</sup>H NMR (CDCl<sub>3</sub>): δ = 0.82–0.91 (t, 12H, CH<sub>3</sub>), 1.21–1.40 (m, 64H, CH<sub>2</sub>), 1.61–1.71 (m, 8H, CH<sub>2</sub>), 2.40–2.50 (m, 8H, CH<sub>2</sub>), 5.85 (s, 1H, CH), 5.10–5.12 (1H, d, CH), 5.35–5.39 (d, 1H, CH), 5.49–5.51 (d, 1H, CH); <sup>13</sup>C NMR (CDCl<sub>3</sub>): 13.09, 21.67, 23.58, 23.68, 27.90, 28.05, 28.13, 28.16, 28.23, 28.32, 28.36, 28.38, 28.43, 28.55, 28.57, 30.89, 32.12, 32.31, 66.12, 71.90, 71.93, 73.47, 77.88, 167.98, 168.12, 171.82, 172.32; IR (NaCl cm<sup>-1</sup>): 3440, 2918, 2849, 1803, 1755 (C=O), 1298, 1196; T<sub>m</sub> = 75 °C.
- b. Intermediate **1b** (1.5 g, 1.6 mmol) and DPTS (0.15 g 0.48 mmol) were dissolved in anhydrous CH<sub>2</sub>Cl<sub>2</sub> (15 mL). mPEG was cooled to room temperature under argon and added to the **1b**/DPTS solution. Once the PEG had dissolved, N,N'-dicyclohexylcarbodiimide (DCC) (1.7 mL, 1.7 mmol) was added dropwise. The

reaction mixture was stirred under argon for 48 hours, cooled and the resulting white solid precipitate (dicyclohexylurea) was removed by vacuum filtration. The filtrate was washed with 0.1N HCl (20 mL), followed by 50:50 brine:water ( $2 \times 20$  mL), dried over  $\text{MgSO}_4$  and concentrated via rotary evaporation. The product was precipitated from diethyl ether yielding **2b** as a white solid (2.6 g, 83%).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta = 0.88$  (t, 12H,  $\text{CH}_3$ ), 1.30 (m, 64H,  $\text{CH}_2$ ), 1.65 (m, 8H,  $\text{CH}_2$ ), 2.38 (m, 4H,  $\text{CH}_2$ ), 2.60 (m, 4H,  $\text{CH}_2$ ), 3.63 (m,  $\sim 0.4\text{kHz}$ ,  $\text{CH}_2$ ), 5.42 (m, 1H,  $\text{CH}$ ), 5.50 (m, 1H,  $\text{CH}$ ), 5.59 (m, 2H,  $\text{CH}$ ); IR ( $\text{NaCl cm}^{-1}$ ): 3478 (O-H), 2883 (C-H), 1749 (C=O), 1241, 1113 (C-O);  $T_m = 54^\circ\text{C}$ ; GPC:  $M_w = 6.3$  kDa; PDI = 1.09.

### 2.3.2. Synthesis of 2c

- a. Procedure was same as that for 1b (2b part a) using 2,5-dihydroxyterephthalic acid (5.0 g, 25 mmol), zinc chloride (0.81 g, 5.9 mmol), and lauroyl chloride (87 mL, 378 mmol), yielding intermediate **1c** as a yellow solid (5.1 g, 68%).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta = 0.80$ – $0.93$  (t, 6H,  $\text{CH}_3$ ), 1.20–1.45 (m, 32H,  $\text{CH}_2$ ), 1.70–1.81 (m, 4H,  $\text{CH}_2$ ), 2.51–2.65 (m, 4H,  $\text{CH}_2$ ), 7.70 (s, 2H, Ar- $\text{CH}$ );  $^{13}\text{C}$  NMR ( $\text{CDCl}_3$ ): 14.32, 22.87, 29.34, 29.53, 29.67, 29.80, 29.82, 32.10, 34.37, 127.37, 128.74, 147.84, 165.27, 172.42; IR ( $\text{NaCl cm}^{-1}$ ): 2847, 1768, 1690 (C=O), 1268, 1177, 935, 897;  $T_m = 71^\circ\text{C}$ .
- b. Procedure same as that for 2b (part b) using intermediate 1c (0.4 g, 0.7 mmol), mPEG (1.0 g, 0.2 mmol), DPTS (0.1 g, 0.3 mmol), and DCC (1 mL, 1.0 mmol), yielding **2c** as a white solid (1.0 g, 91%).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta = 0.85$  (t, 6H,  $\text{CH}_3$ ), 1.30 (m, 32H,  $\text{CH}_2$ ), 1.71 (m, 4H,  $\text{CH}_2$ ), 2.21 (m, 4H,  $\text{CH}_2$ ), 2.36 (m, 4H,  $\text{CH}_2$ ), 3.63 (m,  $\sim 0.4\text{-kHz}$ ,  $\text{CH}_2$ ), 7.65 (d, 1H, Ar-H), 8.10 (s, 1H, Ar-H); IR ( $\text{NaCl cm}^{-1}$ ): 3426, 2879, 1650 (C=O), 1108, 949, 842;  $T_m = 56^\circ\text{C}$ ; GPC:  $M_w = 6.3$  kDa; PDI = 1.07.

### 2.3.3. Synthesis of 2e

- a. Procedure same as that for 1b (2b part a) using D-galacturonic acid (2.0 g, 9.2 mmol), zinc chloride (0.30 g, 2.2 mmol), and lauroyl chloride (10 mL, 46 mmol), yielding intermediate **1e** as a white solid (1.2 g, 13%).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta = 0.80$ – $0.86$  (t, 12H,  $\text{CH}_3$ ), 1.21–1.38 (m, 64H,  $\text{CH}_2$ ), 1.50–1.65 (m, 8H,  $\text{CH}_2$ ), 2.20–2.28 (m, 8H,  $\text{CH}_2$ ), 4.61 (d, 1H,  $H$ ), 5.01–5.08 (dxd, 1H,  $H$ ), 5.20–5.29 (t, 1H,  $H$ ), 5.60–5.69 (t, 1H,  $H$ ), 6.35 (d, 1H, anomeric- $H$ );  $^{13}\text{C}$  NMR ( $\text{CDCl}_3$ ): 14.33, 22.91, 22.91, 24.88, 24.92, 24.98, 25.14, 29.28, 29.32, 29.35, 29.50, 29.58, 29.71, 29.85, 32.15, 34.04, 34.16, 34.26, 34.30, 68.81, 68.92, 68.97, 70.15, 88.72, 171.58, 172.51, 172.84; IR ( $\text{NaCl cm}^{-1}$ ): 3492, 2920, 2851, 1740, 1715 (C=O), 1238, 1147;  $T_m = 51^\circ\text{C}$ .
- b. Procedure same as that for 2b (part b) using intermediate 1e (1.4 g, 1.5 mmol), mPEG (2.5 g, 0.5 mmol), DPTS (0.13 g, 0.4 mmol), and DCC (0.27 mL, 1.5 mmol) yielding **2e** as a white solid (1.9g, 65%).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta = 0.80$ – $0.86$  (t, 12H,  $\text{CH}_3$ ), 1.21–1.38 (m, 64H,  $\text{CH}_2$ ), 1.50–1.65 (m, 8H,  $\text{CH}_2$ ), 2.21–2.26 (m, 8H,  $\text{CH}_2$ ), 3.40–3.78 (m,  $\sim 0.4\text{-kHz}$ ,  $\text{CH}_2$ ), 4.61 (d, 1H,  $H$ ), 5.01–5.08 (dxd, 1H,  $H$ ), 5.20–5.29 (t, 1H,  $H$ ), 5.60–5.69 (t, 1H,  $H$ ), 6.35 (d, 1H, anomeric- $H$ ); IR ( $\text{NaCl cm}^{-1}$ ): 2887 (C-H), 1742 (C=O), 1280, 1148, 1113 (C-O);  $T_m = 57^\circ\text{C}$ ; GPC:  $M_w = 6.3$  kDa; PDI = 1.07.

### 2.3.4. Synthesis of 2f

- a. Procedure same as that for 1b (2b part a) using D-glucuronic acid (2.0 g, 10 mmol), zinc chloride (0.33 g, 2.4 mmol), and lauroyl chloride (12 mL, 51 mmol), yielding

If as a white solid (4.7 g, 51%).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta$  = 0.80–0.86 (t, 12H,  $\text{CH}_3$ ), 1.20–1.39 (m, 64H,  $\text{CH}_2$ ), 1.51–1.71 (m, 8H,  $\text{CH}_2$ ), 2.19–2.30 (m, 4H,  $\text{CH}_2$ ), 2.31–2.42 (m, 4H,  $\text{CH}_2$ ), 4.75 (s, 1H,  $H$ ), 5.30–5.42 (qxd, 2H,  $H$ ), 5.89 (s, 1H,  $H$ ), 6.56 (d, 1H, anomeric- $H$ )  $^{13}\text{C}$  NMR ( $\text{CDCl}_3$ ): 14.22, 22.90, 24.79, 24.98, 25.20, 29.26–29.88, 32.13, 34.09, 34.14, 34.16, 34.27, 65.98, 67.01, 68.28, 70.85, 89.37, 171.59, 172.56, 172.65, 172.84 (C=O); IR (NaCl  $\text{cm}^{-1}$ ): 3601, 2919, 2850, 1752, 1701 (C=O), 1299, 1172;  $T_m$  = 68 °C.

- b. Procedure same as that for 2b (part a) using intermediate 1f (1.4 g, 1.5 mmol), mPEG (2.5g, 0.5 mmol), DPTS (0.13g, 0.4mmol), and DCC (0.27 mL, 1.5 mmol) yielding 2f as a white solid (1.5g, 51%).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta$  = 0.88 (t, 12H,  $\text{CH}_3$ ), 1.28 (m, 64H,  $\text{CH}_2$ ), 1.60 (m, 8H,  $\text{CH}_2$ ), 2.21 (m, 4H,  $\text{CH}_2$ ), 2.36 (m, 4H,  $\text{CH}_2$ ), 3.63 (m, ~0.4-kH,  $\text{CH}_2$ ), 4.78 (s, 1H,  $\text{CH}$ ), 5.40 (s, 2H,  $\text{CH}$ ), 5.84 (s, 2H,  $\text{CH}$ ), 6.53 (s, 1H,  $\text{CH}$ ); IR (NaCl  $\text{cm}^{-1}$ ): 2883 (C-H), 1752 (C=O), 1280, 1148, 1114 (C-O);  $T$  = 55 °C; GPC:  $M_w$  = 6.3 kDa; PDI = 1.07.

#### 2.4. Critical Micelle Concentration (CMC) Measurements

A solution of pyrene, the fluorescence probe molecule, was made up to a concentration of  $5 \times 10^{-6}$  M in acetone. Samples were prepared by adding 1 mL of pyrene solution to a series of vials and allowing the acetone to evaporate so that the final concentration of pyrene in all of the samples was  $5 \times 10^{-7}$  M. AMs were dissolved in HPLC grade water and diluted to a series of concentrations from  $1 \times 10^{-3}$  M to  $1 \times 10^{-10}$  M. AM-pyrene solutions (10 mL) were shaken overnight at 37 °C to allow partition of the pyrene into the micelles. Emission was performed from 300 to 360 nm, with 390 nm as the excitation wavelength. The maximum absorption of pyrene shifted from 332 to 334.5 nm on micelle formation.[29–31] The ratio of absorption of encapsulated pyrene (334.5 nm) to pyrene in water (332 nm) was plotted as the logarithm of polymer concentrations. The inflection point of the curve was taken as the CMC.

#### 2.5. Cell Culture

Studies of polymer interactions were conducted using a tet-inducible cell line with controlled expression of scavenger receptor A (SR-A), human embryonic kidney (HEK) cells stably transfected with human SR-A (gift from Dr. Steven R. Post), which are referred to as HEK-SRA.[24,25] Cells were propagated in high glucose Dulbecco's Modified Eagle Medium (DMEM) (Invitrogen) supplemented with 10% fetal bovine serum (FBS), 1% penicillin/streptomycin, 15  $\mu\text{g}/\text{mL}$  Blasticidin and 100  $\mu\text{g}/\text{mL}$  HygromycinB at 37 °C in 5%  $\text{CO}_2$ . SR-A expression was induced with addition of 0.5  $\mu\text{g}/\text{mL}$  tetracycline overnight and throughout the experiment. SR-A expression was confirmed via antibody binding assays (results not shown).

#### 2.6. LDL Oxidation

Oxidized low density lipoprotein (oxLDL) was generated by incubating 50  $\mu\text{g}/\text{mL}$  LDL purified from human plasma (Molecular Probes Eugene, OR) with 10  $\mu\text{M}$   $\text{CuSO}_4$  at 37 °C for 18 hr exposed to air.[32,33] Oxidation was terminated with 0.01% w/v EDTA (Sigma, St. Louis, MO).

#### 2.7. OxLDL Accumulation in HEK-SRA Cells

The internalization of oxLDL by HEK-SRA cells was assayed by incubating boron-dipyrromethene (BODIPY)-labeled oxLDL (10  $\mu\text{g}/\text{mL}$ ) and  $10^{-6}$  M polymers with cells for 24 hr at 37 °C and 5%  $\text{CO}_2$  in serum containing DMEM. Conditions included a control of medium alone without polymer intervention, and non-induced cells. Cells were washed with  $1 \times$  PBS, fixed with 4% formaldehyde and imaged on a Nikon Eclipse TE2000-S fluorescent

microscope to determine fluorescently tagged oxLDL accumulation. The images were analyzed with ImageJ 1.42q (NIH) and fluorescence data was normalized to cell count. The levels of oxLDL uptake were normalized to those obtained in the absence of polymers.

## 2.8. Statistical Analysis

Each *in vitro* experiment was performed at least twice and three replicate samples were investigated in each experiment. Five images per well were captured and analyzed. The results were then evaluated using analysis of variance (ANOVA). Significance criteria assumed a 95% confidence level ( $P < 0.05$ ). Standard error of the mean is reported in the form of error bars on the graphs of the final data.

## 2.9. Polymer and SR-A Modeling

The polymers were modeled according to their chemical structures illustrated in Figure 1 using the build module in molecular operating environment (MOE) (Chemical Computing Group, Inc., Montreal, Canada). The model polymer molecules were parameterized for Amber99[34] force field and energy minimized until convergence ( $\text{grad} = 0.001$ ) was attained. The creation of the SR-A homology model was previously described.[15] Briefly, the 3D homology model of the SR-A collagen-like domain was generated using the program MODELLER35] with collagen type I chain A as template.

## 2.10. Docking and Scoring

Selected polymer models were docked to the collagen-like domain of SR-A using GOLD v3.2.[36] The GOLD program employs a genetic algorithm for docking flexible ligands into partially flexible receptor sites. The binding cavity was defined as residues Arg45 – Ser68 with an active site radius of 15 Å such that all major residues thought to be necessary for oxLDL binding were included. Dockings were performed with standard default settings; population size of 100, selection pressure of 1.1, number of operations at 100,000, number of islands at 5, and a niche size of 2. Twenty independent docking runs were performed for each polymer, which optimized the computational time required to dock and score non-redundant conformations. The docked pairs were ranked based on each GoldScore. The best ranking conformation of the polymer illustrated the most preferred conformation to interact with scavenger receptor and the binding energy was computed for the refined complexes using Equation 1.

$$\Delta E_{\text{binding}} = \Delta E_{\text{complex}} - \Delta E_{\text{SR-A}} - \Delta E_{\text{polymer}} \quad \text{Equation 1}$$

Where  $\Delta E_{\text{complex}}$  is the energy of the polymers docked to collagen-like domain of SR-A,  $\Delta E_{\text{SR-A}}$  is the energy of the homology model of the scavenger receptor collagen-like domain, and  $\Delta E_{\text{polymer}}$  is the energy of the polymer. Each structure (polymer model, homology model of the SR-A collagen-like domain, and the docked conformation of the pair) was parameterized using Amber99[34] force field and energy minimized until convergence ( $\text{grad} = 0.001$ ) was attained. These minimized energies were used to estimate the binding energy from Equation 1.

## 3. RESULTS AND DISCUSSION

A series of AMs (Figure 1) were designed and synthesized to investigate the influence of altering the hydrophobic backbone on the resultant physicochemical properties. The hydrophobic domain was varied to investigate the influence of stereochemistry, rigidity and shape. The hypothesis was that specific stereochemical and conformational variations would

alter polymer binding affinity to scavenger receptors and the corresponding capacity for inhibition of scavenger receptor-mediated oxLDL uptake.

Stereochemistry is widely recognized as an important design consideration for therapeutic molecules.[37–39] The hydroxyl stereochemistry of the polymers was altered by one carbon position to address the hypothesis that stereochemistry governs the spatial interactions of the carbohydrate with the receptor domains and serving to further enhance binding properties. Saccharic acid is a commercially available carbohydrate stereoisomer of mucic acid, differing in only one hydroxyl group position, and was chosen as the building block for the new AM, **2b**, with altered hydroxyl stereochemistry (Figure 1). **2b** was synthesized by modifying a previously published method for the preparation of **2a**, which comprises mucic acid.[14] Briefly, the two-step procedure involves acylating the carbohydrate with lauroyl groups followed by coupling to PEG (Figure 2). In the initial acylation step, some modifications were required when saccharic acid was used in place of mucic acid. For example, to achieve an acceptable yield (56%) of **1b**, the number of equivalents of acylating agent (lauroyl chloride) was significantly reduced (from 15 with mucic acid to 5 with saccharic acid) as isolation and purification proved problematic with a large excess of lauroyl chloride. Coupling of the PEG and **1b** using DCC as the coupling agent and DPTS as the catalyst proceeded in excellent yields irrespective of hydroxyl stereochemistry (98% for **2a** and 83% for **2b**).

Because increased rigidity previously demonstrated efficient oxLDL inhibition,[15, 16] an aromatic component was introduced to further increase the rigidity. Thus, 2,5-dihydroxyterephthalic acid was chosen as the building block as it possesses both two carboxylic acids and two hydroxyl groups for functionalization. The synthesis of **2c** was based on the same two-step procedure described above for **2b**; no modifications were necessary and the overall yield was 62%. While **2c** does not contain a sugar and is thus technically not a ‘carbohydrate-based’ polymer, it was included to ascertain the role of cyclic structures and flexibility in the five true, sugar-based AMs.

The effect of conformation on the resultant properties was further explored by preparing galacturonic and glucuronic acid-based AMs, **2e** and **2f**, respectively. Increasing the hydrophobic domain size was hypothesized to increase binding to the scavenger receptor pocket. The cyclic carbohydrates (galacturonic and glucuronic acid) were commercially available with carboxylic acid groups for PEG coupling and multiple hydroxyl groups for acylation. Following functionalization with aliphatic chains and PEG, both AMs based on these sugars (**2e** and **2f**) are uncharged; unlike the other polymers (**2a–2c**), these two polymers lack a free carboxylic acid. As stereoisomers, the cyclic AMs **2e** and **2f** also provide insight into how hydroxyl stereochemistry influences bioactivity. Acylation of galacturonic and glucuronic acid proved more difficult than the linear carbohydrates due to the  $\alpha$  and  $\beta$  anomeric forms. The acylation reaction was optimized (95 °C for 4–5 hours) to achieve the highest possible yield (13% for **1e** and 51% for **1f**). To prepare the AMs, the acylated cyclic saccharides, **1e** and **1f**, were coupled to PEG, resulting in **2e** and **2f** in 65% and 51% yield, respectively.

With this unique series of AMs, the influence of backbone architecture on the resultant physicochemical properties (hydrodynamic diameter, melting temperature ( $T_m$ ) and critical micelle concentration (CMC)) was evaluated. The  $T_m$  values of all synthesized polymers were ~60 °C, demonstrating that the PEG domain dominates the polymer’s thermal behavior. Previous work indicated that the AM biological activity was dependent on their ability to form stable micelles.[15,16] The AMs with the lowest CMC values are extremely efficacious and the change in CMC may have a significant biological impact for the newly synthesized AMs. CMC values were measured using a previously reported fluorimetry

technique using pyrene as the fluorescence probe.[29–31] Polymer **2b** formed ~19 nm micelles based on dynamic light scattering (DLS) with a CMC in the same range ( $10^{-7}$  M) as the previously described **2a** (Table 1). The similar hydrodynamic diameter and CMC values indicate that altering the hydroxyl stereochemistry by one position has little effect on polymer self-assembly into micelles. Incorporation of the rigid, aromatic 2, 5-dihydroxyterephthalic acid backbone of polymer **2c** yielded micelles almost twice the size (~35 nm) of the linear analogs (**2a** and **2b**), but with a comparable CMC value ( $10^{-7}$  M). The increase in size was anticipated as terephthalic acid is more sterically bulky than the linear carbohydrates. Similarly, the physicochemical properties of the cyclic carbohydrate-based polymers, **2e** and **2f**, differed from **2d**, the uncharged linear analog of **2a**. Compared to **2d**, **2e** and **2f** have slightly larger hydrodynamic diameters (38 nm and 40 nm, respectively) and CMCs on the order of  $10^{-5}$  M, an order of magnitude higher than that of **2d**. Both results are likely due to the steric bulk of the cyclic carbohydrates as compared to their linear counterparts. This increased CMC suggests that micelles of the cyclic analogs (**2e** and **2f**) are less stable under physiological conditions and may demonstrate decreased biological activity compared to micelles of **2d**. These combined results suggest that although the AM structure is dominated by the mass of PEG, small changes to the hydrophobic domain have a significant impact on the micellar properties.

The AMs were then evaluated for their ability to inhibit oxLDL internalization through the combination of molecular modeling and *in vitro* structure-activity relationship studies with HEK-SRA cells. For molecular modeling, the polymers were modeled according to their chemical structures and scaled to contain only 20 ethylene glycol repeat units instead of 110. We have seen previously that the PEG chain has little interaction with the collagen-like domain, and scaling allows for optimization of computational time. The aliphatic arms were modeled at full alkyl chain length to fully assess the role of the hydrophobic domain. The modeled polymers were docked to an SR-A collagen-like domain homology model using GOLD v.3.2[36] and ranked based on its GoldScore. Previous receptor blocking studies have indicated that the AMs bind specifically to HEK-expressed SR-A1.[26] As such, *in vitro* experiments were carried out via incubation of the HEK-SRA cells with  $10^{-6}$  M polymers and fluorescently labeled oxLDL for 24 hr at 37 °C. To maintain a relatively constant concentration and to ensure we evaluated polymers in the same state (i.e., micellar form), these experiments were performed at or near the critical micelle concentration of the various polymers. The 24-hr time point allowed for saturation of the cells with oxLDL and showed maximal differences between the polymers. As controls, the basal uptake of oxLDL when SRA-expression was not induced and the basal uptake of oxLDL when no polymer was present were both evaluated. The previously synthesized **2a**[14] and **2d**[28], the uncharged analog of **2a**, were compared to the newly synthesized anionic and neutral polymers. Altering the position of one hydroxyl group in the hydrophobic domain had a significant impact on the AMs ability to inhibit oxLDL uptake. **2b** showed only ~10% oxLDL inhibition compared to the ~60% inhibition of **2a** (Figure 3a). This data correlated well with the modeling studies which revealed that **2b** has minimal hydrophobic interactions with SR-A, resulting in unfavorable binding (40 kcal/mol) which may be due to the significant degree of free space between the aliphatic arms of **2b** and the protein binding pocket (see Figure S1). **2a**, on the other hand, possesses a favorable binding energy (–29 kcal/mol) as the aliphatic arms remain in close contact with the SR-A model indicating a significant amount of hydrophobic interactions.

Incorporation of a rigid, aromatic hydrophobic component (**2c**) resulted in binding comparable to **2a**. Although it has only half the number of aliphatic arms, **2c** demonstrated comparable binding in both *in vitro* and molecular modeling experiments, inhibiting ~60% oxLDL uptake in HEK-SRA cells while showing a favorable binding energy (–26 kcal/mol) in modeling studies. This data correlates with previous findings that suggest increased



rigidity positively affects oxLDL inhibition.[15,16] The incorporation of a cyclic aliphatic component showed slight improvements over **2d**, with **2e** and **2f** inhibiting ~50% and ~40% oxLDL uptake, respectively. Given the lack of anionic groups in **2e** and **2f**, SR-A binding is primarily mediated through hydrophobic interactions as demonstrated by molecular modeling. Polymer **2f** displays slightly better oxLDL inhibition and favorable binding energy, in comparison to **2e**, suggesting that the aliphatic arms of **2f** encircle the SR-A protein (Figure S1). While not as significant as the linear analogs, it is noteworthy that although **2e** and **2f** differ by only one stereocenter, they show different binding affinities to SR-A. This result once again illustrates how minute changes in the AM hydrophobic domains can significantly impact the polymer properties.

Notably, the ability to bind to HEK-SRA cells does not correlate to the solution stability of the AMs. The CMC value of **2b** is comparable to that of **2a** ( $10^{-7}$  M), yet was much less effective at inhibiting oxLDL uptake. **2e** and **2f**, however, formed less stable micelles when compared to their linear analog, **2d**, but were much more efficacious in regards to oxLDL inhibition. This data suggests that the polymer behavior is dominated by the unimer, not the micelle, such that the individual polymer chains govern binding to scavenger receptors.

## 4. CONCLUSIONS

A series of AMs were designed to investigate the influence of hydrophobic domain stereochemistry and composition on aggregation and biological properties. While seemingly small with respect to the overall polymer compositions, minor alterations in backbone architecture resulted in significant differences in the physicochemical properties of the AMs, particularly in micelle size and solution stability. *In vitro* and molecular modeling experiments were implemented to examine AM binding to SR-A. These studies also demonstrated that minute changes in the polymer structure significantly affect SR-A binding affinities and consequently modulate the competitive inhibition of oxLDL uptake. Changing the stereochemistry of one hydroxyl group in the backbone (**2b**) significantly reduces oxLDL inhibition while incorporating an aromatic backbone (**2c**) results in oxLDL inhibition comparable to that of the previously published “gold standard”, **2a**. Incorporation of a cyclic carbohydrate (**2e** and **2f**) demonstrated increased oxLDL inhibition and more favorable binding energies than AMs based on the neutrally charged aliphatic analog (**2d**). These findings establish that the composition of the hydrophobic domain is a critical design factor influencing the biological and physicochemical properties of these polymers. Previous *in vivo* studies have demonstrated that the polymers, when loaded in a collagen matrix and implanted around the carotid artery, have been found to be effective at reducing plaque in rats fed a high fat diet.[23] These results, in conjunction with our ability to manipulate polymer carbohydrate compositions, suggest AMs as candidates for cardiovascular therapeutics.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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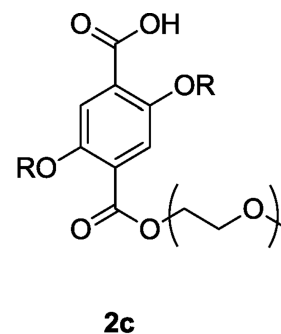
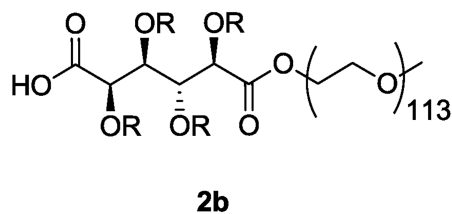
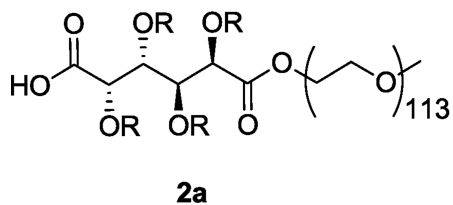
cells stably transfected with human scavenger receptor A for these studies. Additionally, we would like to thank Dr. Bryan Langowski for his insight and critical review of this manuscript.

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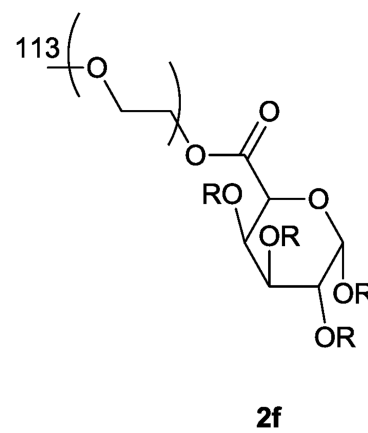
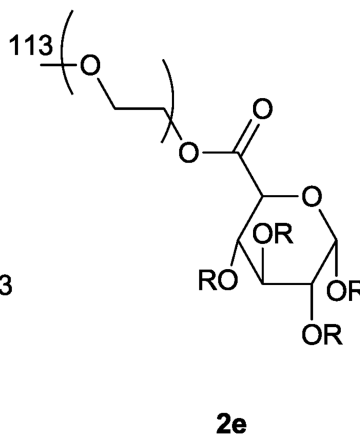
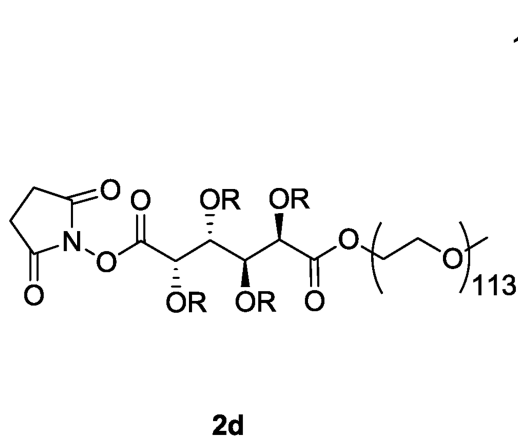
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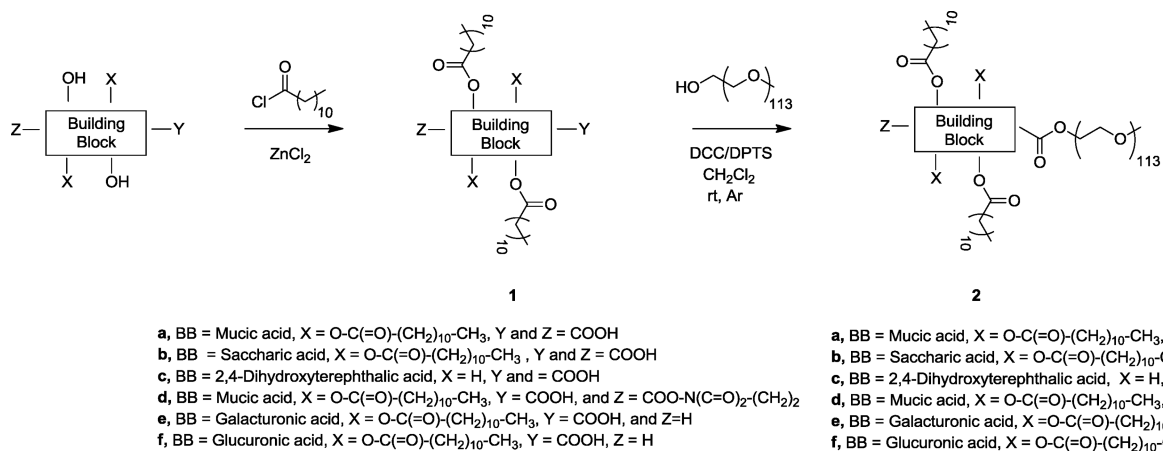
### Charged AMs



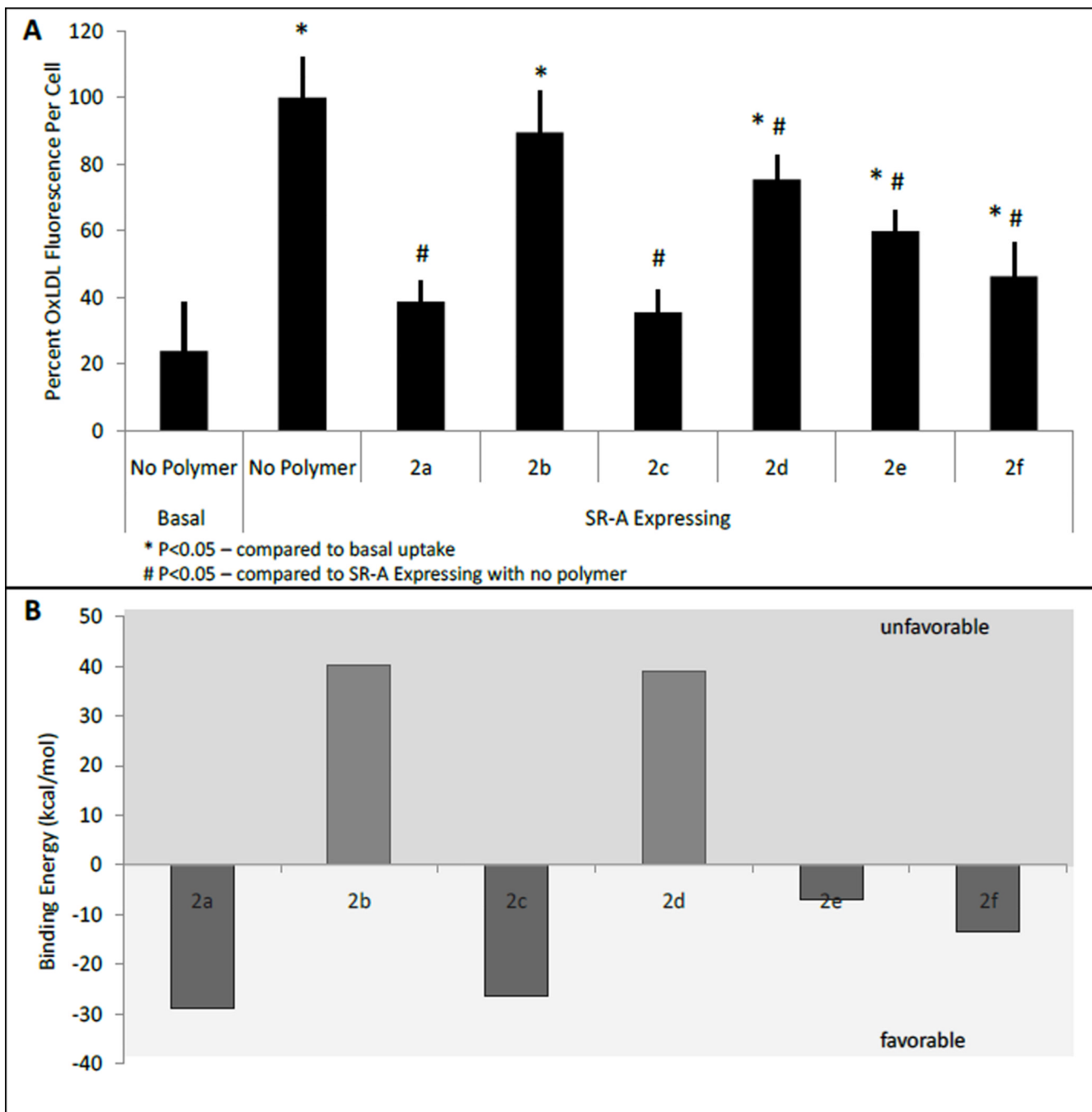
### Uncharged AMs

**Figure 1.**

Structures of charged and uncharged AMs where  $R = \text{---}(\text{CH}_2\text{CH}_2\text{O})_{113}\text{---}$ .

**Figure 2.**

General scheme for the AM synthesis where BB represents the “building block” of the hydrophobic domain.



**Figure 3.** Evaluation of AMs through a) *in vitro* inhibition of oxLDL uptake over 24 hr in HEK cells induced to express SR-A and b) energetics of AMs binding with the SR-A collagen-like domain via molecular modeling.

**Table 1**

Physicochemical properties (hydrodynamic size, CMC and  $T_m$ ) of AMs: as micelles(size and CMC) and in the solid state ( $T_m$ ).

Polymer	Size <sup>a</sup> (nm)	CMC (M)	$T_m$ (°C)
2a	20	$1.20 \times 10^{-7}$	56
2b	19	$1.71 \times 10^{-7}$	59
2c	35	$1.58 \times 10^{-7}$	56
2d	27	$1.67 \times 10^{-6}$	59
2e	38	$1.00 \times 10^{-5}$	55
2f	40	$1.30 \times 10^{-5}$	57

<sup>a</sup>Z-average size