Exploration and application of nanomedicine in atherosclerotic disease

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Chapter 3

Inhibiting macrophage proliferation suppresses atherosclerotic plaque inflammation

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Abstract

Inflammation drives atherosclerotic plaque progression and rupture, and is a compelling therapeutic target. Consequently, attenuating inflammation by reducing local macrophage accumulation is an appealing approach. This can potentially be accomplished by either blocking blood monocyte recruitment to the plaque or increasing macrophage apoptosis and emigration. Because macrophage proliferation was recently shown to dominate macrophage accumulation in advanced plaques, locally inhibiting macrophage proliferation may reduce plaque inflammation and produce long-term therapeutic benefits. To test this hypothesis, we used nanoparticle-based delivery of simvastatin to inhibit plaque macrophage proliferation in apolipoprotein E–deficient mice (Apoe^{-/-}) with advanced atherosclerotic plaques. This resulted in the rapid reduction of plaque inflammation and favorable phenotype remodeling. We then combined this short-term nanoparticle intervention with an 8-week oral statin treatment, and this regimen rapidly reduced and continuously suppressed plaque inflammation. Our results demonstrate that pharmacologically inhibiting local macrophage proliferation can effectively treat inflammation in atherosclerosis.
Introduction

Atherosclerosis, primarily affecting large and midsized arteries, is a lipid-driven inflammatory disease and the underlying cause of most cardiovascular events. Macrophage accumulation plays a major role in atherosclerotic plaque progression, promotes maladaptive inflammation, and aggravates the disease. Therefore, effective therapeutic strategies to reduce plaque macrophage accumulation are a promising approach to dampening inflammation and treating the disease.

Local macrophage accumulation has long been believed to primarily associate with the recruitment rate of blood Ly-6C<sup>high</sup> monocytes, as well as apoptosis and possibly emigration of plaque macrophages. Therefore, anti-inflammatory therapeutic strategies currently being developed focus on blocking monocyte recruitment, promoting macrophage apoptosis, or increasing macrophage emigration. The established macrophage accumulation paradigm is now being revisited. Recent studies demonstrated that local macrophage proliferation can maintain macrophage accumulation in inflammatory and normal tissues. In the context of atherosclerosis, macrophage proliferation dominates in advanced atherosclerotic plaques, in which proliferation contributed more than 80% of local macrophage accumulation over a 1-month period. However, whether directly inhibiting macrophage proliferation can reduce plaque inflammation and produce therapeutic benefits for atherosclerosis remains unknown.

3-Hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase inhibitors, also known as statins, specifically act on the mevalonate pathway, which is indispensable for cell membrane attachment of many small guanosine triphosphatases (GTPases) and their function in promoting cell proliferation. In cell culture, statins effectively inhibit the proliferation of breast cancer cells, smooth muscle cells, and macrophages. We incorporated the HMG-CoA reductase inhibitor simvastatin in a high-density lipoprotein nanoparticle (S-HDL) and achieved specific delivery to plaque macrophages. Here, we show that S-HDL inhibits macrophage proliferation in advanced atherosclerotic plaques, reduces plaque inflammation, and, when combined with oral statin treatment, generates long-term therapeutic benefits.

Results

Study focuses on S-HDL’s therapeutic mechanism and efficacy. Our study focused on inhibiting macrophage proliferation to treat plaque inflammation in Apoe<sup>-/-</sup> mice with advanced atherosclerosis. Using a combination of flow cytometry, magnetic resonance imaging (MRI), near-infrared fluorescence (NIRF) imaging, laser capture microdissection, and mRNA profiling, we dissected the mechanism by which S-HDL exerts its anti-inflammatory effect (Fig. 1A). Having acquired a mechanistic understanding of S-HDL, we designed a two-step treatment regimen that consisted of a 1-week intravenous S-HDL treatment followed by an 8-week oral statin routine intervention (Fig. 1B). After nine weeks of treatment, we used immunostaining, laser capture microdissection, mRNA profiling, blood chemistry, and histology to study the regimen’s effects on advanced plaque inflammation and phenotype in Apoe<sup>-/-</sup> mice (Fig. 1B).

S-HDL does not affect Ly-6C<sup>high</sup> monocyte recruitment. S-HDL was labeled with <sup>89</sup>Zr and its biodistribution was evaluated in wild-type B6 and Apoe<sup>-/-</sup> mice (Fig. S1A). We observed a more than fourfold higher S-HDL aortic arch accumulation in atherosclerotic Apoe<sup>-/-</sup> mice compared to healthy B6 mice (Fig. S1C). Clearance occurs primarily through mononuclear phagocyte system organs (Fig. S1, D and E), with a favorable aorta-to-liver accumulation ratio in diseased mice (Fig. S1F). Blocking Ly-6C<sup>high</sup> monocyte recruitment reduces atherosclerotic plaque inflammation, whereas acute and chronic psychosocial stress aggravates vascular inflammation due to monocytosis. To investigate S-HDL’s effects on monocyte recruitment to the plaque, we first characterized Ly-6C<sup>high</sup> monocyte targeting dynamics after a single S-HDL intravenous injection to Apoe<sup>-/-</sup> mice with advanced atherosclerosis. In plasma and in association with Ly-6C<sup>high</sup> monocytes, S-HDL had about a 20-hour blood half-life, without affecting important recruitment proteins CCR2 and CD115 (Fig. S2). A 1-week nanotherapy treatment course consisting of four intravenous S-HDL injections at a simvastatin dose of 60 mg/kg, which corresponds to a simvastatin dose of 5 mg/kg in humans, did not affect the blood Ly-
6C<sup>high</sup> monocyte levels (Fig. 2, A and B). The aortas showed no change in total mRNA levels of the key genes involved leukocyte recruitment (Ccl2, Icam1, Vcam1, Ccl3, Ccl5, and Cxcl12) and macrophage inflammation state (Tnfa, Il1α, Il1β, and Spp1) after treatment (Fig. 2C). Flow cytometry revealed similar expression patterns of ICAM-1 and VCAM-1 (Fig. 2D), two critical adhesion molecules involved in monocyte recruitment. Finally, using flow cytometry, we quantified Ly-6C<sup>high</sup> monocytes in the whole aortas and found no difference between placebo and S-HDL–treated animals (Fig. 2, E and F). In line with previous studies<sup>14, 23</sup>, little Ly-6C<sup>low</sup> monocytes were observed in plaques (Fig. 2E).

**Figure 1.** Study summary. (A) First, we investigated the mechanisms by which statin-loaded HDL nanoparticles (S-HDL) reduce plaque inflammation. Blood Ly-6C<sup>high</sup> monocyte targeting, monocyte recruitment, plaque phenotype, macrophage proliferation, macrophage emigration, and macrophage inflammation were investigated by flow cytometry, MRI, latex bead–based in vivo cell tracking, laser capture microdissection, and mRNA profiling. (B) To evaluate S-HDL’s translational potential, we combined one-week S-HDL intervention with an eight-week oral statin treatment. Plaque macrophage accumulation, macrophage phenotype, plaque phenotype, and toxic effects on the liver were evaluated in Apoe<sup>−/−</sup> mice fed a high-cholesterol diet (HCD) for 26 weeks.

**S-HDL reduces local macrophage accumulation in plaques.** We previously found that S-HDL reduced macrophage levels in Apoe<sup>−/−</sup> mice that received an HCD for 26 weeks<sup>21</sup>. Here, we used an iHCD that induces established plaques within 12 weeks<sup>14</sup> and advanced plaques in aortic roots after 16 to 20 weeks. We developed a delayed-enhancement T1-weighted MRI (DE-MRI) protocol to evaluate changes in atherosclerotic plaque phenotypes<sup>25, 26</sup>. Similar to DE-MRI after myocardial infarction (MI)<sup>27</sup>, in aortic roots, this technique also reports on inflammation. Figure S3 outlines the procedure that was used to locate and measure signal enhancement in Apoe<sup>−/−</sup> mouse aortic roots and arches after intravenously administered Gd-DTPA (gadolinium–diethylenetriamine pentaacetic acid). We measured signal enhancement before and after one week of S-HDL treatment. We found that, in contrast with placebo-treated mice, a 1-week S-HDL treatment regimen decreased signal enhancement in aortic roots, which indicates a plaque phenotype change (Fig. 3, A and B). These in vivo MRI data were corroborated by NIRF imaging that showed decreased dye-labeled albumin accumulation in the same tissues (Fig. 3, C and D).

Immunofluorescence staining revealed that S-HDL reduced macrophage accumulation in aortic roots by 45% (P<0.01) in Apoe<sup>−/−</sup> mice receiving iHCD. Using laser capture microdissection, we isolated plaque macrophages and measured the expression of key inflammatory genes Tnfa, Ccl2, Icam1, Vcam1, Ccl3, Ccl5, and Cxcl12. This process revealed significantly reduced Tnfa, Ccl2, Vcam1, and Ccl3 expression in mice treated with S-HDL (Fig. 3G). In line with the reduced
macrophage burden, using a bead-based protocol\textsuperscript{28}, we observed lower numbers of emigrating macrophages (fig. S4) but no changes in macrophage apoptosis (fig. S5).

Figure 2. S-HDL does not affect Ly-6C\textsuperscript{high} monocyte recruitment to atherosclerotic plaques. (A and B) Apoe\textsuperscript{-/-} mice receiving a 20-week iHCD were treated with either 1-week S-HDL intravenous injection (n=9) or placebo [phosphate-buffered saline (PBS), n=9]. (A) Representative graph. (B) Quantification. (C) Tested gene expression was normalized to housekeeping gene Hprt1. Statistics were calculated using Student's t test between S-HDL (n=7) and placebo (n=7). (D) Endothelial cells (ECs) were separated from aortas, and the cells' ICAM-1 and VCAM-1 levels were measured by flow cytometry. Gray-filled graphs in (D) are isotype controls. (E and F) Cells from 18 aortas (n=9 for S-HDL and n=9 for placebo) were released by enzyme digestion, and they were identified and enumerated by flow cytometry. Lineage (Lin) included antibodies recognizing CD90, B220, CD49b, NK1.1, Ly-6G, and Ter-119. Quantification is shown in (F). All graphs are presented as means ± SEM. Differences between placebo and S-HDL were calculated by Mann-Whitney U tests if not particularly noted, and none of the comparisons showed significant differences.

\textbf{S-HDL inhibits plaque macrophage proliferation.} The above data demonstrate that S-HDL treatment significantly decreased macrophage accumulation but did not affect Ly-6C\textsuperscript{high} monocyte recruitment, macrophage emigration, or macrophage apoptosis. Recent research shows that cell proliferation is the primary contributor to macrophage accumulation in advanced atherosclerotic plaques\textsuperscript{14}. In light of this discovery, we evaluated S-HDL's ability to reduce macrophage proliferation. Twenty-four hours before the study end point, we injected 5-bromo-2'-deoxyuridine (BrdU) into the peritoneal cavity of mice that had received 1 week of S-HDL treatment (four intravenous injections; simvastatin, 60 mg/kg). BrdU's short systemic half-life renders this a pulse-chase experiment that labels actively proliferating cells for only a short period of time. We isolated aortic macrophages and monocytes for analysis by flow cytometry (Fig. 4, A to C). We found 25% fewer BrdU\textsuperscript{+} macrophages in the aortas of mice treated with S-HDL than the placebo group. In accordance with previous work\textsuperscript{14}, we did not observe robust monocyte proliferation (Fig. 4C). To have a complete view on plaque macrophage proliferation, we subsequently focused on aortic roots, in which atherosclerosis development is the most advanced and macrophage proliferation is dominant. Ki67 and macrophage co-staining revealed that S-HDL reduced macrophage proliferation by 67\% when normalized to the CD68-positive area (P < 0.01) (Fig. 4, D and E).

To unravel the mechanism by which S-HDL reduced macrophage proliferation, we performed in vitro experiments on a murine macrophage cell line (J774A.1). Cells were incubated with S-HDL
at 1.0 or 3.3 μM simvastatin, which effectively inhibited proliferation in a dose-dependent manner (Fig. 4, F and G). At 1.0 μM simvastatin S-HDL, we observed significantly diminished cell proliferation but no change in cell viability, indicating that the antiproliferative effect did not result from diminished cell viability. Finally, the inhibitive effects were largely reverted by the addition of mevalonate, which shows that S-HDL curtails proliferation by blocking the mevalonate pathway.

**Designing a two-step treatment regimen.** Patients hospitalized after MI or stroke have a high recurrence rate of up to 20% (ref 29). We envision that S-HDL, if translated for human clinical use, could be a short-term infusion therapy in patients with acute coronary syndrome, who benefited from early intervention of high-dose oral statin treatment but to a limited extent30. Used as an induction therapy, S-HDL would rapidly suppress plaque inflammation during the vulnerable period after an acute coronary syndrome31. Subsequently, this suppressed inflammation could be sustained by current standard-of-care oral statin therapy. To mimic a clinically relevant scenario, we designed a two-step treatment regimen consisting of a 1-week S-HDL intervention and a subsequent 8-week oral statin treatment. We hypothesize that in Apoe−/− mice with advanced atherosclerosis, a 1-week S-HDL treatment rapidly reduces plaque inflammation and, in conjunction with oral statin therapy, results in continuously reduced inflammation.

After developing advanced atherosclerosis, Apoe−/− mice were randomly assigned to five groups: (i) a control group (referred to as “Control”) received nine weeks of ongoing HCD; (ii) an oral statin group (“Oral Statin”) received nine weeks of oral statin to mimic the current standard of care; (iii) a two-step treatment regimen group (“Hi + Oral”) received a 1-week high-dose S-HDL followed by eight weeks of oral statin; (iv) an S-HDL only group (“Hi + No”) received a 1-week...
high-dose of S-HDL followed by eight weeks without treatment; and (v) a high-dose S-HDL plus low-dose S-HDL group (“Positive”), serving as a positive control, received one week of high-dose S-HDL followed by eight weeks of low-dose S-HDL (fig. S6A).

To measure the therapeutic effects, we used immunostaining for the macrophage marker CD68, histology, laser capture microdissection, and real-time PCR (fig. S6B). The immunostaining and histological data were analyzed by a Matlab-derived procedure that provided objective semi-automatic measurement (fig. S7). Potential toxic effects on the liver and blood cholesterol levels were assessed by standard blood chemistry procedures.

**Figure 4.** S-HDL inhibits macrophage proliferation in atherosclerotic plaques. (A to C) Macrophages and monocytes from whole aortas of Apoe^{-/-} mice treated with either 1-week S-HDL (n=8) or a placebo (n=8) were identified by flow cytometry, and BrdU\(^{+}\) cells were enumerated (A). The percentages of BrdU\(^{+}\) macrophages and monocytes are shown in (B) and (C), respectively. (D and E) Proliferating macrophages in aortic roots were identified as triple-positive with macrophage (MΦ) marker (CD68, green), Ki67 (red), and 4',6-diamidino-2-phenylindole (DAPI) (blue). See (D) for representative images and (E) for quantification. Placebo group has average 5.5 ± 3.0 proliferating macrophages per section; S-HDL group, 0.7 ± 0.5 per section. (F and G) A murine macrophage cell line (J774A.1) was cultured and treated with S-HDL (1.0 or 3.3 \(\mu\)M incorporated simvastatin) in the presence or absence of mevalonate (Mev) (100 \(\mu\)M) for 24 hours. BrdU was added in the last 45 min, and BrdU\(^{+}\) cells were enumerated with flow cytometry (F). See (G) for the percentage of BrdU\(^{+}\) cells and overall viability. All data are means ± SEM, except for (G), which is means ± SD. P values were calculated with Student’s t-tests. *P<0.05, **P<0.01, ***P<0.001, ****P<0.0001.

**Two-step regimen continuously suppresses plaque inflammation.** According to blood chemistry data, the two-step regimen (Hi + Oral) had no toxic effects on the liver and did not change blood cholesterol levels (fig. S8). Histology revealed that one week into the two-step regimen (Hi + Oral), plaque size was significantly reduced by 43% when compared to placebo (P<0.0001), whereas a 1-week oral statin treatment (Oral Statin) produced no significant changes (fig. S9). Further, the two-step regimen (Hi + Oral) maintained the plaque size reduction throughout the 9-week treatment course when compared to control groups. The two-step
regimen (Hi + Oral) also produced a favorable collagen-to-macrophage ratio as early as one week into treatment and maintained that ratio throughout the subsequent 8-week oral statin regimen (fig. S10).

We used anti-CD68 immunostaining to monitor macrophage levels throughout the 9-week treatment course (Fig. 5, A and B) and observed that 1-week S-HDL treatment (Hi S-HDL) led to 65% fewer macrophages than the control group (P<0.0001) and 60% fewer than the oral statin group (P<0.0001) (Fig. 5B). The subsequent 8-week oral statin treatment maintained macrophage reduction and resulted in 33% lower macrophage levels than the control group (P<0.05). Conversely, the data showed that oral statin treatment alone (Oral Statin) did not yet kick in at five weeks. Significantly lower macrophage levels, as compared to the control group, were achieved only after the full 9-week treatment (40% fewer macrophages than the control group, P<0.01). At the end of the 9-week treatment, the macrophage levels in the statin-only group were comparable to those in the two-step regimen group (Fig. 5B). Ki67 quantification corroborated that the initial (and very rapid) reduction of plaque inflammation is due to the inhibition of macrophage proliferation (fig. S11). In line with macrophage burden, eight weeks into the course of oral simvastatin treatment, no significant differences in proliferation levels were observed (fig. S11). Therapeutic anti-inflammatory benefits were realized as early as one week into the treatment for the two-step nanomedicine regimen, whereas nine weeks of treatment was required for the statin-only treated mice.

Using laser capture microdissection, we isolated macrophages from aortic roots and measured their expression of key inflammatory genes (Ccl2, Vcam1, Icam1, Ccl3, Cxcl12, Ccl5, Tnfα, Spp1, Il1α, and Il1β) by quantitative PCR (qPCR). As with macrophage levels, we found that 1-week S-HDL treatment significantly reduced the expression of most genes, including Ccl2, Vcam1, Icam1, Ccl3, Tnfα, Spp1, Il1α, and Il1β, as compared to controls (Fig. 5C and fig. S12A). A 1-week oral statin treatment did not result in statistically significant reductions in any of the genes (Fig. 5C, left panel, and fig. S12A). In the two-step regimen (Hi + Oral), inflammatory gene expression could be maintained by the 8-week oral statin treatment (Fig. 5C and fig. S12, B and C). Collectively, these results demonstrate that the two-step regimen not only reduces macrophage accumulation but also renders the remaining cells less inflammatory.

In mice treated with a 1-week S-HDL regimen that was not followed by oral statin therapy (Hi + No), the macrophage levels and inflammatory state were similar to those in the control group at the end of the 9-week treatment period (Fig. 5D and fig. S12C). These data suggest the 8-week oral statin component is essential to continuous suppression of plaque inflammation. In the positive control group, the 8-week oral statin treatment was replaced by an 8-week low-dose S-HDL infusion (Positive). This regimen achieved therapeutic benefits similar to those seen in the two-step (Hi + Oral) and statin-only (Oral Statin) groups (Fig. 5, C and D), albeit at the cost of two intravenous injections per 8 weeks.
**Figure 5.** The two-step treatment regimen continuously suppresses plaque inflammation. (A) Representative binary mask generated from a Matlab procedure identified areas positive for macrophages. (B) Macrophage levels were measured throughout the different 9-week treatment programs. (C) Inflammatory genes expressed by macrophages isolated with laser capture microdissection show differences between treated and control groups. Bar graph quantification of inflammatory genes’ expression is provided in fig. S12. (D) Macrophage levels at weeks one and nine. Data are means ± SEM. P values were calculated with Mann-Whitney U tests. *P<0.05, **P<0.01, ***P<0.001, ****P<0.0001. Statistics were calculated by comparing the Control group with nonparametric Mann-Whitney U tests.
Discussion

Here, we show that inhibiting local macrophage proliferation is an effective therapeutic strategy for suppressing atherosclerotic plaque inflammation. More specifically, we use a 1-week nanoparticle therapy (S-HDL) that specifically delivers statins to plaque macrophages, thereby inhibiting their proliferation and reducing inflammation. These therapeutic anti-inflammatory benefits can be maintained by a subsequent 8-week oral statin treatment. The combined 1-week nanotherapy and 8-week oral statin regimen produced no toxic effects on the liver.

Statins, the most widely used cholesterol-lowering drug class, specifically inhibit the mevalonate pathway. This pathway produces isoprenoids, a class of molecules used not only for cholesterol synthesis but also for membrane attachment of small GTPases such as Ras and Rho. Ras plays a central role in nuclear factor-κB (NF-κB)–mediated inflammatory responses, whereas Rho initiates proliferation of certain cells, including macrophages. Systemic statin delivery promotes macrophage emigration from plaques, whereas targeted statin delivery to blood monocytes inhibits monocyte recruitment to plaques, and both effects are probably related to statins’ inhibitory effects on NF-κB–mediated inflammatory function. For example, in atherosclerotic rabbits, it has been shown that lengthy periods of oral therapy are required to produce therapeutic benefits, which rely on statins’ cholesterol-lowering and systemic anti-inflammatory effects. Here, we observed S-HDL’s ability to reduce inflammatory gene expression, but our primary focus is statins’ inhibitory effects on cell proliferation, likely due to decreased production of isoprenoid intermediates in the inhibited mevalonate pathway. Neither Ly-6C<sup>hi</sup> blood monocyte recruitment and associated gene expression nor macrophage emigration was significantly changed by S-HDL therapy, whereas flow cytometry assays in vivo and in vitro indicated strong antiproliferative effects. Our approach positions nanomedicine as a novel therapeutic strategy that capitalizes on statins’ antiproliferative benefits in treating atherosclerosis.

Extensive technical advances in nanotechnology production and formulation chemistry now allow us to incorporate various diagnostic and therapeutic agents in HDL nanoparticles and deliver them to plaque macrophages. S-HDL’s capacity to inhibit local macrophage proliferation indicates the therapeutic potential of targeted delivery of compounds that are designed to specifically inhibit key processes important in macrophage plaque accumulation. These compounds’ HDL formulations can perhaps reveal the extent to which recruitment, migration, and proliferation can combat inflammation in atherosclerosis.

At the dose used in the current study, conventionally delivered HDL has no therapeutic benefits. In mice, HDL at 400 mg/kg ApoA1 dose has been shown to cause plaque regression, and several clinical trials have safely administered HDL at doses up to 80 mg/kg. However, the precise therapeutic benefits of HDL remain a topic of ongoing research. Our S-HDL nanotherapy was based on the aforementioned clinically validated, reconstituted HDL injectables. Because our nanomaterial produces excellent therapeutic benefits at low toxicity, it has significant translational potential for more effectively treating atherosclerosis and its related conditions.

Our nanotherapy treatment strategy may be particularly useful immediately after atherosclerosis-related events, such as MI or stroke, which have an up to 20% recurrence rate within three years. Recent studies in both mice and humans suggest that MI itself elevates the inflammatory state of high-risk vulnerable plaques, possibly contributing to recurring events. Additionally, high levels of macrophage proliferation have been found in high-risk human plaques. In this context, we foresee a scenario in which a hospitalized patient receives intravenous nanoparticle therapy to rapidly reduce plaque inflammation and then subsequent oral statin therapy to maintain the reduced level of plaque inflammation. We demonstrated the potential of a two-step regimen in mice that received a 1-week short-term nanoparticle intervention followed by an 8-week oral statin treatment. This approach reduced plaque inflammation far more quickly than oral statin treatment, which required nine weeks to induce the desired therapeutic effects. The two-step approach, on the other hand, produced comparable therapeutic benefits within one week via S-HDL nanotherapy and maintained them throughout the study period using oral statin therapy. This approach produced similar effects to the positive control protocol of 1-week high-dose S-HDL followed by eight weeks of low-dose S-HDL treatment. The latter low-dose nanoparticle treatment was previously shown to effectively prevent plaque formation.

This study demonstrates that pharmacologically inhibiting plaque macrophage proliferation by
nanotherapy suppresses plaque inflammation and alleviates atherosclerosis. In conjunction with oral statin therapy, rapid and continuous suppression of plaque inflammation could be achieved with minimal toxic effects. This study opens a therapeutic avenue for tackling inflammation in atherosclerosis using a macrophage-targeted antiproliferative strategy.

**Materials**

**Synthesizing S-HDL nanoparticles.** Briefly, simvastatin (AK Scientific), 1-myristoyl-2-hydroxy-sn-glycero-phosphocholine (MHPC), and 1,2-dimyristoyl-sn-glycero-3-phosphatidylcholine (DMPC) (Avanti Polar Lipids) were dissolved in chloroform/methanol (4:1 by volume) solvent and dried to form a thin film. Human apolipoprotein A1 (ApoA1) proteins, separated from human plasma, were added to the film, and the solution was incubated at 37°C until the film was hydrated and a homogeneous solution was formed. The solution was sonicated, and aggregates were removed by centrifugation to yield small simvastatin-loaded nanoparticles (S-HDL). For fluorescence detection purposes, either DiR or DiO (Invitrogen) was incorporated when fluorescence imaging techniques were applied. Nanoparticle solution was washed extensively using 100-kD filter (Vivaspin, Vivaproducts) and filtered through a 0.22-mm nylon filter before being administered to animals. Simvastatin incorporation efficiency was determined by high-performance liquid chromatography (Shimadzu).

Preparation of ⁴²⁴⁴Tc-labeled S-HDL is detailed in Supplementary Materials and Methods and used a method previously reported.**

**Animal protocol, diet, and immunostaining.** More than 300 Apoe⁻⁻ (B6.129P2-Apoemtm1Unc) and 5 B6 wild-type (C57BL/6) mice were used for this study. All animal care and procedures were based on an approved institutional protocol from Icahn School of Medicine at Mount Sinai. Five-week-old Apoe⁻⁻ male and female mice were purchased from The Jackson Laboratory. In a mechanistic study, the mice were fed an intensively high-cholesterol diet (0.2% weight cholesterol; 15.2% kcal protein, 42.7% kcal carbohydrate, 42.0% kcal fat; Harlan TD88137) for 16 to 20 weeks (referred to as iHCD). In the therapeutic study, mice were fed a more moderate high-cholesterol diet (0.2% weight cholesterol; 20% kcal protein, 40% kcal carbohydrate, 40% kcal fat; Research Diets Inc.) that needed 26 weeks to induce advanced atherosclerosis in Apoe⁻⁻ mice (referred to as HCD). The more moderate diet allowed the mice to reach an average age of 6 to 9 months before receiving treatment to more closely replicate the fact that atherosclerosis mainly affects adult and senior patients. Aortic roots were dissected and used for staining with CD68 (clone 557, Serotec), activated caspase-3 (ab13847, Abcam), or Ki67 (ab15580, Abcam).

**Flow cytometry.** To evaluate S-HDL’s effects on blood monocytes, 20 Apoe⁻⁻ mice received a single intravenous infusion of high-dose S-HDL loaded with a trace amount of DiR. About 40 ml of blood was collected by retro-orbital bleeding before the infusion and at days 0.25, 0.5, 1, 2, 3, 4, 5, and 6 after. Red blood cells in these samples were removed using a red blood cell lysis buffer (BD Biosciences). White blood cells were identified as DAPI CD45⁺. Blood monocytes were identified as DAPI CD45⁺ CD115⁺ SSC⁻. Monocytes were further identified as Ly-6C⁻ or Ly-6C⁺ subsets on the basis of the expression of Gr-1. Fluorescence intensity of DiR from the cells was used to measure S-HDL uptake levels.

To evaluate the expression levels of adhesion molecules VCAM-1 and ICAM-1, whole aortas were digested using a cocktail of liberase TH (4 U/ml) (Roche), deoxyribonuclease (DNase) I (0.1 mg/ml) (Sigma-Aldrich), and hyaluronidase (1 mg/ml) (Sigma-Aldrich). Cells were digested with trypsin/EDTA (100 units/ml) and then filtered through a 40-μm nylon filter (BD Biosciences). The mixture was incubated on an orbital shaker for 10 minutes at 37°C. Monocytes were stained and fixed by following the protocol provided by the manufacturer (BD APC-BrdU Kit, 552598), and detected and digested 24 hours later. Cells from aortas were isolated and stained as described above, and then fixed and permeated using a red blood cell lysis buffer (BD Biosciences). White blood cells were identified as DAPI CD45⁺. Neutrophils were identified as Ly-6G⁻. Neutrophils were defined as Ly-6C⁺ CD11b⁻ cells identified by flow cytometry.

Aortic roots were dissected and used for staining with CD68 (clone 557, Serotec), activated caspase-3 (ab13847, Abcam), or Ki67 (ab15580, Abcam). Aortic roots were dissected and used for staining with CD68 (clone 557, Serotec), activated caspase-3 (ab13847, Abcam), or Ki67 (ab15580, Abcam).

**In vivo and in vitro BrdU labeling.** BrdU (1 mg) was injected into the peritoneal cavity, and aortas were harvested and digested 24 hours later. Cells from aortas were isolated and stained as described above, and then fixed and permeated according to the manufacturer's protocol (BD APC-BrdU Kit, 55298). BrdU⁻⁻ cells were identified by flow cytometry.

In vitro labeling was done with J774A.1 cells cultured with Dulbecco's modified Eagle's medium with 10% fetal bovine serum (Sigma-Aldrich) and labeled with BrdU for 45 min before being harvested. Cells were stained and fixed by following the protocol provided by the manufacturer (BD APC-BrdU Kit, 55298), and detected by flow cytometry.

**In vivo MRI and ex vivo NIRM.** Twenty-two Apoe⁻⁻ mice with advanced atherosclerosis were scanned using an electrocardiography (ECG)-gated delayed-enhancement T1-weighted protocol on a 7 T MRI scanner equipped with a 25 mm Quad H1 coil (Bruker). The ECG trigger was derived from two electrodes inserted into the animals' front limbs. Briefly, the aortic root regions were imaged before and after Gd-DTPA (0.3 mmol/kg, Magnevist, Bayer) injection through the tail vein. A T1-weighted spin echo (repetition time = 800 ms, echo time = 7.543 ms, spatial resolution = 0.117 mm², field of view = 30 mm × 30 mm, slice thickness = 0.5 mm, number of acquisitions = 4, scan time = 15 min) sequence with a black-blood saturation band was used to scan the aortic roots.

Signal enhancement was determined from the SNR of the vessel wall before (SNRpre) and after (SNRpost) Gd-DTPA injection. Seven animals were sacrificed after the first scan and used to determine the baseline (n = 7). The remaining animals received either 1-week S-HDL (n = 8) or placebo (n = 7) and were subjected to a second scan. SNR change was calculated using the following formula: SNR change = [SNRpost/SNRpre] – 1. Thirty minutes before sacrifice, the animals received a Cy5.5-albumin injection (Cy5.5, 1 mg/kg) to measure endothelial permeability of the same region.
Aortic roots and arches were collected and imaged by NIRF imaging using the IVIS200 spectrum optical imaging system (PerkinElmer). Total fluorescence signal from aortic roots was quantified using Living Imaging (PerkinElmer).

**Design for therapeutic study and ex vivo analyses.** One hundred twenty-six Apoe−/− mice with advanced atherosclerosis were randomly assigned to 14 groups with 9 animals per group. Thirty-six animals received a 1-week placebo treatment (intravenous PBS infusion) and no treatment in the subsequent eight weeks. Atherosclerosis progression was monitored by sacrificing animals at weeks zero (n=9), 1 (n=9), and nine (n=9) during the treatment period (referred to as Control). Twenty-seven animals received oral simvastatin from diet (15 mg/kg per day), and animals were sacrificed for analysis at weeks 1 (n=9), 5 (n=9), and 9 (n=9) (Oral Statin). Twenty-seven animals first received a 1-week high-dose intravenous treatment with S-HDL (simvastatin, 60 mg/kg; ApoA1, 40 mg/kg; four infusions per week) and a subsequent 8-week oral statin treatment (simvastatin, 15 mg/kg per day). Animals were sacrificed at weeks one, five, and nine (Hi + Oral). Eighteen animals received the same 1-week high-dose intravenous infusion of S-HDL but received no treatment in the following eight weeks. Nine animals were sacrificed in weeks five and nine (Hi + No). Eighteen animals received the same 1-week high-dose S-HDL and a subsequent 8-week intravenous infusion of low-dose S-rHDL (simvastatin, 15 mg/kg; ApoA1, 10 mg/kg; two times per week), and nine animals were sacrificed at weeks five and nine (Positive). This treatment schedule is summarized in fig. S6A.

For each animal, 72 aortic root sections (6 μm thick) were prepared. Nine sections per root were stained with hematoxylin phloxine saffron; nine sections, for trichrome stain; nine sections, for anti-CD68 stain (Serotec, clone MCA1957); and three sections, for anti-Ki67 immunofluorescence staining. A custom-made Matlab (MathWorks) procedure was used to quantify the stained areas on the histological sections. Laser capture microdissection was used to extract macrophage mRNA from the remaining sections. These cells’ inflammatory gene expression levels were quantified by real-time PCR. Figure S6B provides an overview. In the therapeutic study, 3522 histology sections were analyzed. The Matlab-based procedures and information for tested genes are described in the Supplementary Materials. Blood samples were collected upon sacrifice and used for blood chemistry analyses.

**Statistical analysis.** Results are expressed as means ± SEM. Significance of differences was calculated using Student's t test and nonparametric Mann-Whitney U test. GraphPad Prism 5.0 for PC (GraphPad Software Inc.) was used for statistical analysis. P < 0.05 was regarded as significant as denoted by “∗” if not specifically noted.
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Supplementary results

Supplementary material for this article is available at http://advances.sciencemag.org/cgi/content/full/1/3/e1400223/DC1.
Supplementary figures

Fig. S1. S-HDL’s biodistribution in atherosclerotic and wild type mice. (A) Chromatograms show the same eluting profiles of S-HDL and $^{89}$Zr radioactivity. (B) Autoradiographs of Apoe$^{-/-}$ or B6 mice aortas (left panel, n=5) and other organs (right panel, n=1). (C) Mean radioactivity in aortic arches, normalized to arch radioactivity of B6 mice, is 4.33-fold higher in Apoe$^{-/-}$ than in B6 mice. Radioactivity in whole aortas and selected tissues (D) expressed as % ID/g and (E) normalized to that of B6 mice. (F) Radioactivity (%ID/g) aorta-to-liver ratio is 4.53-fold higher ratio in Apoe$^{-/-}$ mice than in B6 mice. A student’s t-test with Welch’s correction was used to calculate statistical significance. ** P<0.01. * P<0.05, *** P<0.001, and **** P<0.0001.

Fig. S2. Targeting dynamics of S-HDL to blood Ly-6C$^{hi}$ monocytes and their expression of key recruitment proteins (A) One dose of DiR-labeled S-HDL (n=7) or placebo (n=6) was intravenously injected to Apoe$^{-/-}$ mice that had been fed a intensive high-cholesterol diet (iHCD) for 16 weeks. Mice were bled at different time points (day 0, 0.25, 0.5, 1, 2, 3, 4, 5, and 6). Sample plasma and cellular components were separated by centrifuge. Ly-6C$^{hi}$ blood monocytes were identified as CD45$^+$, CD115$^+$, and Gr1$^{hi}$. (B) By measuring DiR fluorescence intensity, the relative levels of free S-HDL in blood plasma and ingested particles in Ly-6C$^{hi}$ cells were normalized. Here, those levels are presented as functions of time. (C) Gating procedure to identify Ly-6C$^{hi}$ monocytes 24 hours after administering S-HDL (n=5) or placebo (n=5). In the two right panels, representative expression of CCR2 and CD115 are shown relative to isotype controls (gray histograms). (D) Quantification of CCR2 and CD115 on Ly-6C$^{hi}$ monocytes from mice treated with S-HDL or placebo. A student’s t-test was used to calculate significance. ** P<0.01.

Fig. S3. Procedure for in vivo delayed-enhancement MRI (DE-MRI). (A) Mice first underwent a pilot scan
to identify ascending and descending aortas, and the results were used to place new sagittal scans (yellow line indicates the plane) that covered the aortic roots and arches. (B) In the sagittal view of aortic roots and arches, the subsequent T1-weighted spin-echo scan was placed parallel to the aortic root (yellow line). This scan had 8 slices (0.5 mm thick) above the yellow line to cover the whole aortic arch. (C) Representative image shows ascending aorta (AA), descending aorta (DA), and pulmonary trunk (PT). (D) Signal enhancement accumulated in aortic root area (AR) vessel walls after Gd-DTPA injection. Post-scan started immediately after Gd-DTPA injection and lasted for 15 minutes.

**Fig. S4.** Latex bead-based method for tracking macrophage emigration. (A) Representative immunofluorescent image of a section stained with anti-CD68 macrophage marker (green), nuclei (blue), and latex beads (red spots denoted by white arrow). (B) The number of remaining beads inversely correlates to the number of emigrating macrophages, demonstrating S-HDL-treated mice (n=10) had fewer emigrating macrophages than placebo (n=10). Data are mean ± SEM. * p<0.05.

**Fig. S5.** Quantification of apoptosis in aortic roots after S-HDL treatment. Ten Apoe−/− mice received a one-week S-HDL treatment (60 mg/kg simvastatin, 4 intravenous injections / week). Three sections per aortic root from each animal were dual-stained against CD68 (green) and activated Caspase 3 (red). The area filled with apoptotic macrophages was yellow. (A) Representative images from either placebo- or S-HDL-treated animals. (B) Quantification of apoptotic macrophages was achieved by measuring the yellow area within dash-line masks (A). Three plaques were measured per aortic root section. Data are mean ± SEM. N.S. means not statistically significant.

**Fig. S6.** Designing a two-step treatment regimen. (A) 126 Apoe−/− mice were used for this study. Mice were fed a mildly high-cholesterol diet (HCD) for 26 weeks before receiving different nine-week treatments. In each treatment group, nine mice were sacrificed at weeks 0, 1, 5, and 9. (B) Each mouse aortic root generated 72 6-μm thick frozen sections. Plaque size was measured by adding macrophage stain (CD68) to connective tissue stain (hematoxylin phloxine saffron, HPS). Collagen levels were determined by trichrome stain (Col.). Macrophage levels were determined by anti-CD68 staining (CD68). Proliferating macrophages were determined by anti-Ki67, anti-CD68, and DAPI triple stain. Thirty sections per animal were used for histology. The remaining sections underwent laser capture microdissection to extract mRNA for profiling experiment.
**Fig. S7.** Representative binary masks of histological images. Three consecutive sections were stained with anti-CD68 (A), trichrome (B), or hematoxylin phloxine saffron (HPS) (C). The upper panels show the original images taken under a 4 X magnification in bright field. The lower panels show the binary masks generated by programs written in MATLAB. Plaque size was calculated by adding HPS positive area to CD68 positive area. Collagen/macrophage ratio was calculated by dividing collagen levels by CD68 levels.

**Fig. S8.** Blood chemistry results. The different treatments’ toxic effects on the liver were determined by measuring the levels of alanine transaminase (ALT) and aspartate transaminase (AST) in the blood at weeks 1 and 9. (A), (B), and (C) show ALT, AST, and cholesterol levels at week 1, respectively. (D), (E), and (F) show ALT, AST, and cholesterol levels at week 9, respectively. Data are mean ± SEM. Student’s t-tests were used to measure statistical significance between control and other groups. No significance was found.

**Fig. S9.** Plaque size changes during the nine-week treatment. (A) Plaque size change is plotted as a function of time. (B) Plaque sizes are presented in group-specific graphs. Data are mean ± SEM. * p<0.05, ** p<0.01, *** p<0.001, **** p<0.0001. Statistics are calculated by comparing to the control group using student’s t-tests unless otherwise noted.
**Fig. S10.** Changes in macrophage-to-collagen ratio during the nine-week treatment. (A) Collagen-to-macrophage ratio is plotted as a function of time. (B) All groups’ ratios are presented in individual graphs. Data are mean ± SEM. *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001. Statistics were calculated by comparing to the control group unless otherwise noted. Student’s t-tests were used.

**Fig. S11.** Quantification of proliferating macrophages in week 1 and 9 of therapy. (A) Representative images to identify proliferating macrophages (Ki67+, DAPI+, and CD68+ cells). Arrows in the right panel indicate proliferating macrophages. (B) Quantification of proliferating macrophages in different treatment groups (n=5 per group; 3 sections per aortic roots). Statistics were calculated by comparing tested groups to the control group, unless otherwise noted, with student’s t-test. On week 9 no significant differences were observed. * P<0.05 and ** P<0.01.

**Fig. S12.** Inflammatory gene expression during the nine-week treatment. Gene expression at weeks 1 (A), 5 (B), and 9 (C). Data are mean ± SEM. * p<0.05. Statistics were calculated by comparing to the control group using student’s t-tests.