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Single Step Reconstitution of Multifunctional High-Density Lipoprotein-Derived Nanomaterials Using Microfluidics

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Abstract

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Supporting Information Available: Supporting information with 5 figures includes confocal microscopy images and flow cytometry data. This material is available free of charge via the Internet at http://pubs.acs.org.
High-density lipoprotein (HDL) is a natural nanoparticle that transports peripheral cholesterol to the liver. Reconstituted high-density lipoprotein (rHDL) exhibits antiatherothrombotic properties and is being considered as a natural treatment for cardiovascular diseases. Furthermore, HDL nanoparticle platforms have been created for targeted delivery of therapeutic and diagnostic agents. The current methods for HDL reconstitution involve lengthy procedures that are challenging to scale up. A central need in the synthesis of rHDL, and multifunctional nanomaterials in general, is to establish large-scale production of reproducible and homogeneous batches in a simple and efficient fashion. Here, we present a large-scale microfluidics-based manufacturing method for single-step synthesis of HDL-mimicking nanomaterials (µHDL). µHDL is shown to have the same properties (e.g., size, morphology, bioactivity) as conventionally reconstituted HDL and native HDL. In addition, we were able to incorporate simvastatin (a hydrophobic drug) into µHDL, as well as gold, iron oxide, quantum dot nanocrystals or fluorophores to enable its detection by computed tomography (CT), magnetic resonance imaging (MRI), or fluorescence microscopy, respectively. Our approach may contribute to effective development and optimization of lipoprotein-based nanomaterials for medical imaging and drug delivery.

Keywords
HDL; multifunctional; nanoparticle; reconstitution; microfluidics; high-throughput

Lipoproteins are natural nanoparticles that transport cholesterol and triglycerides throughout the body and have important roles in a variety of pathological processes,\(^1\) most prominently atherosclerosis. Atherosclerosis is a progressive disease in which the buildup of cholesterol and inflammatory cells in vessel wall of arteries can eventually lead to myocardial infarction or stroke.\(^2\) High levels of serum low-density lipoprotein (LDL) increase the progression of atherosclerosis and the risk of coronary artery disease.\(^3\) On the other hand, high-density lipoprotein (HDL) is believed to exhibit athero-protective properties. In vitro, in a number of preclinical studies and in limited clinical studies,\(^4\) the mechanistic basis for a high level of HDL and a decreased risk of coronary artery disease is suggested.\(^5\) These athero-protective properties include HDL’s ability to remove and transport cholesterol from atherosclerotic plaques to the liver via a process known as reverse cholesterol transport.\(^6,7\) Traditional antiathero-sclerotic therapies such as statins lower LDL levels systemically, but elevating
HDL levels is believed to hold great promise as an alternative strategy. Among different approaches, including the application of cholesteryl ester transfer protein inhibitors, direct infusion of rHDL is an emerging treatment for cardiovascular disease.

For example, HDL infusions have been reported to modulate fatty acid metabolism and support cholesterol efflux, which therefore reduces myocardial lesions in a rat model and the size of human atherosclerotic plaques or their inflammatory state.

Moreover, HDL’s endogenous character makes it well suited as a vehicle for targeted delivery of diagnostic and therapeutic agents. For example, HDL nanoparticles have been reconstituted to carry inorganic nanocrystals as contrast agents for medical imaging as well as to serve as delivery vehicles for siRNAs or therapeutic molecules. The reconstitution of such HDL nanoparticles involves multistep processes, which are highly dependent on synthetic conditions, difficult to scale up, and laborious. For example, the cholate, sonication, and vesicle insertion methods are time-consuming, requiring at least 24 h to perform. A central challenge in the synthesis of therapeutic and diagnostic HDL-based nanomaterials is to establish large-scale and continuous production methods with high reproducibility, yield, and homogeneity, while simultaneously decreasing the number of formulation steps.

Microfluidic technologies using diffusion, emulsification, or mixing have recently emerged for continuous formation of a variety of nanoparticles including liposomes, polymeric nanoparticles, and lipidpolymer hybrid nanoparticles. Thanks to their ability to tune nano- and microscale interactions between precursors, microfluidic formulation processes offer effective control of the formation and characteristics of produced nanomaterials leading to a narrow size distribution and high batch-to-batch reproducibility.

In the current study, we apply the above microfluidic methodology for the synthesis of biologically active HDL-mimicking nanomaterials (µHDL) that can be loaded with hydrophobic molecules. The microfluidic approach enables us to tailor µHDL lipid composition and encapsulate compound such as simvastatin (S), fluorophores, or inorganic nanocrystal cores such as gold (Au), iron oxides (FeO) and quantum dots (QD) using a single-step production process that may easily be adapted for large-scale production.

In this work, we show that the physicochemical properties of µHDL can be readily varied and optimized by manipulating mixing speeds and the lipid to protein ratios. We demonstrate that µHDL has similar morphological and compositional properties to native HDL and conventionally reconstituted HDL (rHDL). We also validate the biological properties of µHDL by studying its interaction with macrophages and comparing its cholesterol efflux capacity with native HDL. Finally, we demonstrate the diagnostic properties of nanocrystal loaded µHDL.
RESULTS AND DISCUSSION

Microfluidic Platform for Single Step Assembly of HDL-Derived Nanomaterials

Multifunctional HDL-mimicking nanomaterials (µHDL, DiO-µHDL, [S]-µHDL, Au-µHDL, FeO-µHDL, and QD-µHDL) were reconstituted using a single-step, self-assembly method in a single layer, 3-inlet microfluidic device (Figure 1a and Table 1). This large-scale microfluidic device (2 mm wide and 400 µm high) generates tunable dual microvortices and a focusing pattern at Reynolds number \((Re) \approx 150\) (Figure 1b – d), thereby allowing rapid and effective mixing of the solutions in the central inlet and the two outer inlets. We adjusted the type or composition of the phospholipids \(\text{e.g., 1,2-dimyristoyl-sn-glycero-3-phosphocholine (DMPC) and 1-myristoyl-2-hydroxy-sn-glycero-3-phosphocholine (MHPC)}\) and its mixing conditions with apolipo-protein A–I (apoA–I) by tuning the flow patterns in the microfluidic device. The solvent of the central inlet was determined by the solubility of the individual imaging agent payloads (Table 2). In this microfluidic approach, the self-assembly process occurs due to: (1) the transition of the lipid/payload from an organic solution to an aqueous one, which initiates the formation of lipid aggregates, while (2) the microvortices cause apoA-I to swiftly incorporate in the nascent aggregates, resulting in instantaneous formation of small µHDL nanoparticles. The conventional multistep synthesis of a 120 mg (total weight) batch of rHDL would normally first require lipid film formation (2 h), followed by lipid film hydration (3 h), 1 h of sonication and finally 16 h incubation with apoA–I.\(^\text{38}\) Remarkably, in this microfluidic approach, HDL was formed instantaneously, as determined by DLS and TEM measurements (\textit{vide infra}). This raises the prospect of on site or at point-of-care synthesis of these nanomedicines. Furthermore, this new microfluidic approach can result in the continuous production of µHDL at a rate of 420 mg/h (total weight), with minimal variability in the product while maintaining the same yield to that of conventional synthesis (\(~57\%\) of apoA–I) (Table 3). The microfluidic approach also has the ability for further scale up by running multiple chips in parallel. While conventional methods can also be scaled up, they would suffer from greater challenges in ensuring the desired nanoparticle properties are maintained (Tables 1–3).

Physicochemical Properties of µHDL

We investigated the size of µHDL as a function of the flow pattern in the microfluidic device. By varying the flow rate of each inlet (\(i.e.,\) Reynolds number), we were able to judiciously adjust the µHDL diameter at a given lipid-to-protein ratio (\(i.e.,\) DMPC:apoA–I = 2.5) (Figure 2a). By increasing the Reynolds number \((Re)\) from approximately 30 to 150, we were able to decrease the average size of µHDL from 30.0 to 8.1 nm. We noted that the solutions of DMPC and apoA-I were not well mixed at \(Re \approx 30\), whereas they were strongly mixed at \(Re \approx 150\). In Figure 2a the size of µHDL before and after purification indicates that for most synthesis conditions, \(i.e.,\) changes in \(Re\), the size remained the same. This is indicative of a high yield of homogenously sized µHDL nanoparticles. We then tested how the average size changes as a function of the DMPC:apoA–I ratio at \(Re \approx 150\). Results obtained showed that the average size of µHDL remained 7.6–8.5 nm as the DMPC:apoA–I ratio increased from 0.625 to 2.5 but increased to approximately 30.6 nm with a 12.5 ratio (Figure 2b). This increase is likely the result of the formation of larger lipid aggregates that do not incorporate sufficient apoA-I. Additionally, as the Reynolds number increased, the
polydispersity of µHDL gradually decreased to approximately 0.1 (Figure 2c). We note that an excessive increase of the DMPC:apoA–I ratio led to an increase of the polydispersity to 0.218 (Figure 2d).

We then compared the structure of rHDL reconstituted using conventional multistep methods27 (Figure 2e) with that of µHDL (Figure 2f) using transmission electron microscopy (TEM). The images revealed that both formulations adopted discoidal shape and characteristic rouleaux structures (stacks of discs on their edge), characteristic of native nonspherical HDL particles. Furthermore, dynamic light scattering confirmed that both methods resulted in HDL nanoparticles with a similar size around 8–9 nm (Figure 2g).

**Biological Properties of µHDL**

HDL has been shown to bind to macrophage cells *via* several cell surface proteins such as SR-B1, ABCA1, and ABCG1.39,40 We have previously shown HDL to be taken up by macrophages *via* such cell surface proteins.27,41 We incubated murine macrophages with rhodamine labeled µHDL to probe the nanoparticle uptake and observed that the µHDL was taken up by macrophages in a time-dependent fashion (Figure 3a–c). Next, we performed a competitive-inhibition experiment, where µHDL was co-incubated with different concentrations of native HDL purified from serum. We observed a significant decrease of µHDL uptake in the cells with increasing concentrations of native HDL (Figure 3d), indicating that the µHDL was taken up by macrophages through the same pathway as native HDL. Last, we compared the macrophage cholesterol efflux capacity of µHDL with that of native HDL and rHDL. We found that the three HDL types (µ-HDL, native HDL, and rHDL) produced significant and comparable efflux in both bone marrow derived macrophages (BMDM) and J774A.1 (Figure 3e). These assays demonstrated that, similar to the rHDL we previously reconstituted using conventional multistep methods,27 µHDL has similar bioactivity to native HDL.

**Incorporation of Hydrophobic Molecules into µHDL**

Using the single-step microfluidic approach, we were able to incorporate a fluorescent hydrophobic agent 3,3′-diodicycloxocarbocyanine perchlorate (DiO) or an anti-inflammatory hydrophobic drug simvastatin [S] into µHDL (Table 2). Nanoparticle loading of simvastatin is of interest as, when delivered orally, it is mainly confined to the liver, whereas nanoparticle formulation may allow it to be delivered to other tissues. The mean sizes of DiO-µHDL and [S]-µHDL, respectively, were 7.3 ± 1.1 and 32 ± 1.3 nm with a polydispersity of less than 0.1. The entrapment efficiencies were 94.2 ± 9.6% for DiO (DiO (mg)/apoA–I (mg) = 0.07) and 70.1 ± 7.0% for simvastatin (simvastatin (mg)/apoA–I (mg) = 1.4). We studied the uptake of DiO-µHDL in murine macrophages using confocal microscopy (Figure 4a,b; see also Figure S1). Incubation of [S]-µHDL with lipopolysaccharide (LPS) activated macrophages resulted in a 30% decrease in the secretion of tumor necrosis factor-alpha (TNF-α), a pro-inflammatory cytokine (Figure 4c). Co-incubation of [S]-µHDL with 100 µM of mevalonate resulted in no decrease of TNF-α secretion, demonstrating that the effects of [S]-µHDL are due to inhibition of the HMG-CoA reductase enzyme, as mevalonate is the product of that enzyme (Figure S2).42
**Gold Nanocrystal Core µHDL (Au-µHDL)**

We next incorporated gold nanocrystals into the core of µHDL (Table 2). The central stream contained gold nanoparticles mixed with phospholipids dissolved in a solution of ethanol, chloroform, and methanol at a ratio of 35:4:1, while apoA–I in a PBS buffer was infused in the outer channels at a final ratio of lipid:apoA–I ratio of 2.5:1. The crude product contained both Au-µHDL and ‘empty’ µHDL (Figure 5a and Figure 3). Subsequently, Au-µHDL was isolated and purified (Figure 5b) using a gradient density purification method derived from Havel’s ultracentrifugation lipoprotein separation protocol.25,43 Next, we tested the computed tomography (CT) detectability of Au-µHDL phantoms (Figure 5c). We then compared the uptake of Au-µHDL or Au-PEG (gold nanocrystals shielded by a nonspecific phospholipid-polyethylene glycol) by macrophages in vitro by imaging the resultant cell pellets with CT. The contrast in CT images revealed that Au-µHDL was taken up more avidly by the cells as compared to Au-PEG (Figure 5d). Using this microfluidic method we were able to decrease the weight ratio of phospholipid:gold needed in the formation process from 5:1 for the previously reported method to 1:1 in the microfluidic method, leading to a better yield and purer product. Furthermore, we were able to form Au-µHDL from a range of phospholipids and their mixtures with the possibility to decrease the ratio of, or even exclude, lysolipids in the formulation (Figure S3b), which are prone to destabilize cell membranes at higher concentrations. Our single step microfluidic method not only enabled scale-up production of Au-µHDL but also eliminated the need for long incubation steps, heating and extensive purification that were necessary in our previous methods.25

**FeO and QD Core µHDL (FeO-µHDL and QD-µHDL)**

FeO nanoparticles and quantum dots were first co-dissolved with the phospholipids in a solution of chloroform and methanol before being infused into the microfluidic device. We were able to produce a homogeneous population of single nanocrystal core µHDL after evaporating the chloroform in the produced nanoparticle solution and purifying the solution by gradient density separation (Figure 6a,b). We observed that the nanoparticle size was dependent on the size of the quantum dots or iron oxide nanocrystals injected into the microfluidic device. Confocal images of macrophages incubated with QD-µHDL (Figure 6c; see also Figure S5) revealed its cellular uptake. In addition, magnetic resonance images of macrophages incubated with FeO-PEG and FeO-µHDL (Fe: 40 µg/mL) revealed a decrease in signal intensity originating from the cell pellets incubated with FeO-µHDL when compared to nontreated cells or cells incubated with FeO-PEG. This signal decay indicated preferential internalization of FeO-µHDL by macrophages.

**CONCLUSIONS**

The present studies demonstrated that multifunctional HDL-mimicking nanomaterials (µHDL) can be reconstituted by a microfluidic production process using rapid mixing of precursors in microvortices. We identified an optimal ratio of lipid:apoA–I in the microfluidic platform and validated that µHDL has similar bioactivity properties to native HDL and rHDL synthesized using a conventional multistep bulk method. Using this microfluidic approach, we were able to incorporate various hydrophobic molecules and inorganic nanocrystals into µHDL and image these nanomaterials using computed tomography (CT),
magnetic resonance imaging (MRI) and fluorescent techniques. Our new microfluidic based formulation process enables the continuous and reproducible manufacture of homogeneous HDL with the ability to readily control the properties by varying concentrations of different combinations of the precursors in a single-step procedure. This new approach could greatly contribute to effective development and optimization of HDL-based multifunctional nanomaterials for medical imaging and drug delivery. Furthermore, this microfluidic technology, once operated in parallel, has potential to facilitate reproducible and controlled synthesis of HDL nanoparticles at scales suitable for rapid clinical development and commercialization.

METHODS

Materials

Myristoyl hydroxy phosphatidylcholine (MHPC), 1,2-dimyristoyl-sn-glycero-3-phosphocholine (DMPC), 1,2-distearoyl-sn-glycero-3-phosphoethanolamine-N-[methoxy(poly-ethylene glycol)-2000] (PEG2000-DSPE) and 1,2-dimyristoyl-sn-glycero-3-phosphoethanolamine-N-(lissamine rhodamine B sulfonyl) (Rhodamine-DMPE) were purchased from Avanti Polar Lipids, Inc. (Alabaster, AL).3,3’-Dioctadecyloxacarbocyanineper-chlorate (DiO) was purchased from Invitrogen. Oleic acid coated iron oxide nanoparticles were purchased from NN Laboratories (Fayetteville, AR). Dodecanethiol coated gold nanoparticles were synthesized as described previously.25 The 650 nm emitting CdSe/CdS/CdZnS/ZnS core-shell-shell quantum dots were synthesized as described previously.28 Apolipoprotein A-1 was a kind gift of CSL Ltd. (Parkville, Australia).

Microfluidic Device Design and Fabrication

Our microfluidic device had three inlet channels (with rectangular cross sections (200 × 400 \( \mu \text{m}^2 \)) converged to create a single mixing zone with dimensions 2 mm wide, 400 \( \mu \text{m} \) high, and 20 mm long. The device was made of polydimethylsiloxane (PDMS) (SYLGARD 184, Dow Corning, Midland, MI) using standard soft-lithography techniques.44

Flow Visualization

Flow patterns in the single mixing zone of the microfluidic device were visualized using a stereo microscope (Leica M125, Leica Microsystems, Bannockburn, IL). In the visualization, the central stream has a 10:1 ratio of deionized water and black ink and the outer streams have deionized water.

Reynolds Number and Flow Rate

Syringe pumps (NE-1010-U, Kats Scientific) were used to regulate flow rates in the microfluidic device. The flow rate in the outer streams was set to 5 mL/min (total 10 mL/min for both outer channels), while the flow rate in the central stream was 1 mL/min. Reynolds number (Re) was calculated using the following equation.45
where \( Q \) is the flow rate; \( \mu \) represents the fluid’s viscosity; \( w \) and \( h \) represent the channel width (2000 \( \mu \)m) and height (400 \( \mu \)m); \( \rho \) represents the fluid’s density; \( U \) represents the fluid’s average velocity.

**Numerical Simulations**

Computational fluid dynamics simulations predicted the flow field using the commercial CFD solver (SC/Tetra, CRADLE, Beavercreek, OH). We assumed a Newtonian fluid with the properties of water at room temperature and no-slip boundary conditions on all the walls. Mesh independence study was conducted. The convergence limit of the velocity was set to 0.1\%, in which the mass fraction reached the asymptotic value within 0.01\%.

**\( \mu \)HDL Synthesis**

Organic solution containing 5 mg/mL of phospholipids (DMPC) was injected into the middle inlet channel of the microfluidics device at a rate of 1 mL/min using a programmable syringe pump, while aqueous solutions of apoA-I (0.2 mg/mL) in PBS were injected in both the outer channels at a rate of 5 mL/min. \( \mu \)HDL produced through controlled mixing of the solutions was collected at the outlet of the device. \( \mu \)HDL was then washed 3 times in PBS and concentrated to desired volumes using 10000 MWCO centrifugal concentrators.

**Production Rate**

The production rate was calculated using the following formula:

\[
\text{Production rate (total wt./h)} = [C_1 \cdot (fr_1 + fr_1) + C_2 \cdot fr_2]
\]

where \( C_1 \) represents the concentration of the apoA-I in the aqueous solution (mg/mL); \( C_2 \) represents the concentration of the phospholipid in the organic solution (mg/mL); \( fr_1 \) represents the flow rate of the aqueous solutions (mL/h) and \( fr_2 \) represents the flow rate of the organic solution (mL/h). To load \( \mu \)HDL with either hydrophobic molecules or nanocrystals, we used the above method with slight alterations as indicated in Table 2.

**Gradient Density Purification**

Nanocrystal loaded HDL was separated from empty \( \mu \)HDL using a dual density based centrifugation method adapted from Havel’s lipoprotein separation method. Briefly, 200 \( \mu \)L of \( \mu \)HDL mixture in PBS was carefully added on top of 1 mL of a 30\% (w/w) KBr solution and subsequently centrifuged for 1 h at 14100g. During centrifugation, \( \mu \)HDL remained within the top layer, while denser nanocrystals loaded HDL pelleted into the KBr layer. After centrifugation, the top layer was discarded, while the remaining solution was collected and its content washed three times with PBS. This above-mentioned procedure was performed twice before the particles were concentrated to the desired volume.
Dynamic Light Scattering
The nanoparticle samples were diluted in deionized filtrated water (18 MΩ cm) and nanoparticle size was analyzed a minimum of three times for each sample by dynamic light scattering (DLS) using a ZetaPALS system (Brookhaven Instruments Corporation).

Transmission Electron Microscopy
HDL sample solutions were transferred to an ammonium acetate buffer solution, then negatively stained using an ammonium acetate buffer containing 2% sodium phosphotungstate. Drops of samples were then placed onto 100 mesh Formvar coated nickel grids (Electron Microscopy Sciences) and allowed to air-dry. Grids were imaged using a Hitachi 7650 microscope operated at 80 kV and connected to a digital camera (Scientific Instruments and Applications) controlled by Maxim CCD software.

DiO Loading Measurements
HDL solution samples were completely dried for 16 h in a desiccator, then solubilized in ethanol. Solutions were then centrifuged and the supernatants were collected. DiO concentrations were then quantified by measuring sample absorbance at 480 nm using a Spectramax m5 plate reader (Molecular Devices, Sunnyvale, CA).

Simvastatin Loading Measurements
HDL solution samples were dried for 16 h in a desiccator then solubilized in acetonitrile. Acetonitrile solutions were then centrifuged and the supernatants collected. Simvastatin concentrations were then measured using reverse phase C18 column connected to a Shimadzu HPLC system.

Cell Propagation
Murine sarcoma macrophage cells J774A.1 (ATCC, Manassas, VA) were propagated in Dulbecco’s Modified Eagle Medium supplemented with 1% streptomycin/penicillin (Cellgro Mediatech Manassas, VA) and 10% fetal bovine serum (Sigma-Aldrich, St. Louis, MO). Before experiments, cells were detached, washed in PBS, and counted. Defined numbers of cells were then seeded and left overnight to adhere. Immediately before experiments were initiated, the media was replaced with fresh media.

Rhodamine Assay
Rhodamine labeled µHDL was incubated with murine macrophages and subsequently washed with PBS to discard nonbound/internalized particles. In competition-inhibition experiments, macrophages were co-incubated with a fixed concentration of the rhodamine labeled µHDL (21.4 µg/mL apoA–I) and increasing amounts of native murine HDL extracted from serum.

Fluorescence/Confocal Microscopy
Macrophages were seeded on microscope chamber slides and allowed to adhere overnight. The cell media was then refreshed and cells incubated for 2 h with various nanoparticle solutions. After incubation, slides were washed three times with PBS and cells were fixed.
with 4% paraformaldehyde (PFA) for 30 min at 37 °C. Slides were then washed again three times with PBS and mounted with mounting media containing DAPI (Vectashield, Vector Laboratories). Epifluorescence images were taken using an Olympus 1 x 71 fluorescence Inverted Microscope. Confocal images were taken using a Leica TCS SP5 DM confocal using a sequential setup and excitation at 405 nm.

**Cholesterol Efflux Assay**

J774A.1 macrophages and murine bone marrow derived macrophages (BMDM) were first incubated with media containing 0.5 µCi/mL [3H]-cholesterol and 0.34 µM phorbol 12-myristate 13-acetate for 24 h. Cells were subsequently washed twice with PBS and incubated with media containing 0.34 µM phorbol 12-myristate 13-acetate and 2 mg/mL fatty acid-free albumin. After washing, cells were incubated for 16 h with different HDL solutions (50 µg/mL by protein content). Cell media was then removed and the cells were lysed in 0.1 N NaOH solution. [3H]-Cholesterol contents of media and cell lysates were measured by liquid scintillation counting.

**TNF-α Secretion Assay**

J774A.1 cells were seeded in 96-well plates (25 000 cells/well) and allowed to adhere overnight. Cells were then activated for 6 h with 300 units/mL of interferon-γ, then 16 h with 300 units/mL of interferon-γ and 25 ng/mL of LPS. Cells were then incubated for 24 h with serum free media containing either 10µM of free simvastatin, [S]-µHDL equivalent to 10 µM simvastatin or an equal amount of µHDL. The resulting TNF-α concentration in the media was measured using a TNF-α ELISA kit (Invitrogen, CA).

**Au-µHDL Incubation with Macrophages**

Three million cells/dish were seeded in 100 nm tissue culture dishes and allowed to adhere overnight. Cells were then incubated for 8 h with 10 mL of media containing 1.25 mg (gold content) of Au nanoparticles. Cells were then washed three times with PBS, collected using cell scrapers and centrifuged for 5 min at 500g. Cells were resuspended in 4% PFA and centrifuged to form a pellet.

**FeO-µHDL Incubation with Macrophages**

A total of 500 000 cells/ well were seeded in 6-well plates and allowed to adhere overnight. The cells were then incubated with 2 mL of media containing FeO nanoparticles at a concentration of 40 µg Fe/mL for 7 h. Cells were then washed three times with PBS, collected using scrapers and centrifuged for 5 min at 500g. Cells were resuspended in 4% PFA and allowed to form a pellet.

**CT Scanning**

Particle solutions and cell pellets were imaged on 256-slice Brilliance iCT scanner (Philips Medical Systems, Nederland B.V., The Netherlands). Gold attenuation values (expressed in Hounsfield units (HU)) were the average obtained from 3 selected regions of interest using the instrument’s software and converted to Au concentration using a predetermined
The CT images presented were prepared using OsiriX 32-bit (Geneva, Switzerland).

**MR Scanning**

T\(_2\)-weighted cell pellet images were obtained using a 7T MRI system (Bruker Instruments) with a gradient echo sequence: echo time, 2 ms; repetition time, 2000 ms; field of view, 20 × 20 × 10 mm; imaging matrix, 128 × 128 × 16; flip angle, 3°. The images presented in the manuscript were prepared using the Osirix software mentioned above.

**Relaxometry**

Iron oxide HDL samples were completely dried for 16 h in a desiccator then solubilized in ethanol. Samples were then incubated with 1 N HCl at 40°C until total dissolution. \(T_1\) measurements of the nanoparticle solutions were performed on a 60 MHz Bruker Minispec (Bruker Medical BmbH, Ettingen) operating at 40 °C. Iron concentration of the samples was then calculated using the calibration curve of \(1/T_1\) measurements of samples containing a known quantity of iron oxide.

**Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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Figure 1.
Microfluidic reconstitution of HDL-derived nanomaterials (µHDL). (a) A schematic depiction of a microfluidic platform that allows single-step and large-scale production of the HDL: reconstituted HDL by microfluidics (µHDL); DiO-µHDL for fluorescence; [S]-µHDL for a therapeutic platform; Au-µHDL for CT imaging; FeO-µHDL for MRI; and QD-µHDL for fluorescence. (b) Photograph of the microfluidic device. (c) Microscope image of dual microvortices at Re = 150. Scale bar is 500 µm. (d) Computational fluid dynamics simulation showing the microvortices at Re = 150.
Figure 2.

μHDL is controlled by Reynolds number as well as lipid-apoA-I compositions. (a) Size of μHDL with respect to Reynolds number, i.e., mixing speed (DMPC:apoA–I = 2.5). White column bars before purification; black bars after purification. (b) Size of μHDL with respect to lipid-apoA-I ratio (Re = 150). White column bars before purification; black bars after purification. (c) Polydispersity of μHDL with respect to Reynolds number. (DMPC:apoA–I = 2.5). White column bars before purification; black bars after purification. (d) Polydispersity of μHDL with respect to lipid:apoA-I ratio (Re = 150). White column bars before purification; black bars after purification.
before purification; black bars after purification. (e) TEM image of rHDL synthesized by sonication. Scale bar is 20 nm. (f) TEM image of μHDL produced using our microfluidic approach. Scale bar is 20 nm. (g) Size and polydispersity of rHDL and μHDL. Error bars are standard deviation. $N = 3$. 
Figure 3.
µHDL is bioactive. (a) Microscope image of DAPI-stained macrophages. Scale bar is 50 µm. (b) Microscope image of DAPI-stained macrophages incubated with rhodamine-conjugated µHDL (shown in red). Scale bar is 50 µm. (c) µHDL uptake by macrophages over time. Error bars are standard deviation. N = 4. (d) Competition assay that shows µHDL uptake by macrophages with respect to increasing concentrations of native HDL extracted from human plasma. Error bar is standard deviation. N = 5 (***P<0.0001). (e) Cholesterol efflux of BSA, µHDL, native HDL, and rHDL (Black column bars BMDM; white bars J774A.1). Error bar are standard deviation. N = 4. No statistical significances were shown between HDL cases except for between µHDL and native HDL in J774A.1 (P < 0.0001), while the comparisons between BSA and other HDLs showed statistical significance (P < 0.0001).
Figure 4.
µHDL as hydrophobic molecules delivery agent. (a) Microscope images of DAPI-stained macrophages. Scale bar is 50 µm. (b) Microscope images of DAPI-stained macrophages incubated with DiO-µHDL (shown in green). Scale bar is 50 µm. (c) TNF-α secretion for 24 h from preactivated macrophages incubated with [S]-µHDL solution. TNF-α secretion was measured using ELISA. Error bars are standard deviation. N = 4 (**P < 0.01).
Figure 5.
Au-µHDL. (a and b) TEM images of Au-µHDL (a) before and (b) after purification. Scale bar is 100 nm. (c) Phantom images of PBS and Au-µHDL solutions using CT imaging. (d) CT images of macrophage cell pellets: non-treated cells, cells incubated with Au-PEG, and cells incubated with Au-µHDL (Figure S4).
Figure 6.
QD-µHDL and FeO-µHDL. (a and b) TEM images of (a) QD-µHDL and (b) FeO-µHDL. Scale bar is 100 nm. (c) Confocal microscopy image of macrophages incubated with QD-µHDL. Scale bar is 100 µm. (d) T2-weighted magnetic resonance images of macrophage cell pellets: control (nontreated cells), cells incubated with FeO-PEG, and cells incubated with FeO-µHDL (Figure S4).
### TABLE 1
Experimental Setup in Microfluidics for µHDL Syntheses

| inlet | precursor | solvent          | flow rate (mL/min) | Re   |
|-------|-----------|------------------|--------------------|------|
| Left  | apoA-I (0.2 mg/mL) | PBS               | 5                  | 152.8|
| Center| DMPC, MHPC, and imaging agents/drugs (see Table 2) | ethanol, methanol, chloroform | 1      |      |
| Right | apoA-I (0.2 mg/mL) | PBS               | 5                  |      |
| HDL type | lipid/imaging agent | solvent (mL) |
|----------|---------------------|--------------|
|          | DMPC                | ethanol      | methanol | chloroform |
| µHDL     | 10 mg               | 2            |          |            |
| DiO-µHDL | 10 mg               | DIO: 0.3 mg  | 2         |            |
| [S]-µHDL | 20 mg               | Simvastatin: 4 mg | 2         |            |
| Au-µHDL  | 3.5 mg              | Au: 10 mg    | 1.75     | 0.05       | 0.2 |
| FeO-µHDL | 80 mg               | FeO: 5 mg    | 0.2      | 1.8        |
| QD-µHDL  | 50 mg               | QD: 3 nmol   | 0.2      | 1.8        |

DiO is 3,3′-dioctadecyloxacarbocyanine perchlorate. [S] is simvastatin.
### TABLE 3
Yield of the Final Product in Conventional Process (rHDL) and Microfluidics Approach (µHDL)

| precursor   | conventional process ($n = 3$) | microfluidics approach ($n = 4$) |
|-------------|---------------------------------|----------------------------------|
| apoA-I      | 59% ± 6                         | 57% ± 11                         |
| phospholipid| 63% ± 9                         | 53% ± 3.7                        |