VPO1 Mediates ApoE Oxidation and Impairs the Clearance of Plasma Lipids

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Abstract

Objective: ApoE is an abundant component of chylomicron, VLDL, IDL, and HDL. It binds to multiple types of lipids and is implicated in cholesterol and triglyceride homeostasis. Oxidation of ApoE plays a crucial role in the genesis of atherosclerosis. It is proposed that heme-containing peroxidases (hPx) are major mediators of lipoprotein oxidation. Vascular peroxidase 1 (VPO1) is a recently-discovered hPx, which is expressed in cardiovascular system, lung, liver etc. and secreted into plasma. Its plasma concentration is three orders of magnitude of that of myeloperoxidase. If VPO1 mediates ApoE oxidation and affects the lipid metabolism remains to be elucidated.

Methods: Recombinant ApoE and VPO1 were expressed and purified from stably-expressing cell lines derived from HEK293 cells. ApoE oxidation was carried out by VPO1 in the presence of H₂O₂ and chloride. ApoE oxidation was verified by a variety of approaches including immunoblot and amino acid analyses. To evaluate the functional changes in VPO1-oxidized ApoE, lipid emulsion particle binding assays were employed.

Results: Oxidized ApoE binds weaker to lipid emulsion particles, which mimic the large lipid complexes in vivo. In lipid efflux assay, oxidized ApoE showed reduced capability in efflux of lipids from foam cells. Mice administrated with oxidized ApoE via blood exhibited weaker clearance ability of plasma lipids.

Conclusions: Our data suggest that VPO1 is a new mediator regulating lipid homeostasis, implying a role in genesis and development of atherosclerosis.

Introduction

The physiological functions of lipoproteins are to transport lipids in blood and facilitate lipids through the cellular membrane into cells. Liver, intestine, and adipose tissue are the major tissues for metabolism and storage of lipids [1]. Blood lipoproteins form five types of lipoprotein particles based on electrophoresis and ultracentrifugation, named high density lipoprotein (HDL), low density lipoprotein (LDL), intermediate-density lipoprotein (IDL), very low density lipoprotein (VLDL), and chylomicrons [2]. These particles have distinct functions in homeostasis and transportation of lipids.

Apolipoprotein E (ApoE) is a glycoprotein of 299 amino acids with a molecular mass of ~34 kDa. It is a major component of chylomicron, VLDL, IDL and HDL [3], [4]. It is expressed in multiple tissues including liver, brain, kidneys, and spleen, and secreted into plasma. The major function of ApoE is proposed to facilitate lipid transportation [5]. Pathologically, ApoE plays similar roles in genesis of atherosclerosis in both mouse and humans. ApoE-deficient mice showed severe hypercholesterolemia and advanced atherosclerotic lesion [6], [7]. In this animal model, formation of foam cells and development of fibroproliferative lesions are histochemically similar to that of human atherosclerosis [8], [9]. Formation of foam cells is the benchmark of atherosclerosis. On the other hand, elevated plasma ApoE levels reduced hyperlipidemia and atherosclerosis such as in the transgenic mouse models with macrophage-specific ApoE expression. ApoE-deficient mice transplanted with bone marrow cells containing human ApoE gene revealed elevated plasma ApoE levels, decreased blood lipid levels, and less atherosclerotic lesions [10], [11], [12].

Atherosclerosis is a metabolic and inflammatory disorder which characterized by reduction in the luminal diameter of the main arteries [13]. The first stage in the development of atherosclerosis is the formation of foam cells on the wall of blood vessel [14]. Foam cells derived from blood monocytes by the uptake and accumulation of lipids, mainly cholesterol and triglycerides, resulting in impaired transportation of lipids. It is assumed that
the impaired transportation and accumulation of lipids are caused by the modification and dysfunction of lipoproteins, e.g. ApoE. A line of evidence shows that animal heme-containing peroxidases (hPxes) play an important role in oxidation of lipids and proteins, and development of atherosclerosis [15]. Myeloperoxidase (MPO) is proto-enzyme of the family and is extensively studied. MPO is restricted expressed in neutrophils and monocytes. Only a small fraction of MPO is secreted into plasma by a process called “leaking”. It is proposed that monocytes migrate into the intermediate of the vessel walls at inflammatory sites. Macrophages, which are derived from monocytes, then oxidize LDL and transform foam cells [16]. MPO mediates the oxidation of proteins, lipids and DNA by generating potent hypochlorous acid (HOCI), a key mediator in the genesis of atherosclerosis [16], [17].

Recently, our group identified and characterized a novel hPx, named vascular peroxidase 1 (VPO1) [18]. It is mainly expressed in the cardiovascular system, lung, liver, and spleen, and most of VPO1 proteins are secreted into blood stream with the concentration of three orders of magnitude of that of MPO in plasma [19]. Like MPO, VPO1 is able to oxidize chloride to generate HOCl in the presence of hydrogen peroxide (H2O2) [20], [21]. However, the pathophysiological role of VPO1 in genesis and development of atherosclerosis remains unclear. In the present study, we showed that ApoE and VPO1 are localized in the same area of atherosclerotic plaques. ApoE is susceptible to oxidation by VPO1. Moreover, oxidized ApoE (ox-ApoE) has decreased functions in promoting triglyceride and cholesterol efflux from foam cells, and in plasma lipid clearance. These data suggest that VPO1 participates in the dysfunction of ApoE and the formation of atherosclerosis.

### Materials and Methods

#### Ethics Statement

The experiments on mice were approved by the University of Alabama at Birmingham Animal Care and Use Committee (Animal Project Number: 120309582).

#### Cells and Reagents

HEK293H (Invitrogen, Carlsbad, CA) cells were maintained in DMEM medium with 10% fetal bovine serum, 100 U/ml penicillin, and 100 µg/ml streptomycin. THP-1 cells (human acute monocytic leukemia cell line, American Type Culture Collection, Manassas, VA) were maintained in RPMI 1640 media with 10% fetal bovine serum, 100 U/ml penicillin, and 100 µg/ml streptomycin. Trinitrobenzene sulfonic acid, sodium hypochlorite solution, triolein, egg phosphatidylcholine (PC), PC assay kit, and NaCl were purchased from Sigma-Aldrich (St. Louis, MO); chemiluminescent substrate for immunoblots was from Pierce Biotechnology (Rockford, IL); goat polyclonal antibody against human ApoE was from Millipore (Billerica, MA); cholesterol assay kit was from Wako Chemicals (Richmond, VA); triglycerides assay kit was from Pointe Scientific (Lincoln Park, MI); Fluorescence-labeled second antibodies were from Invitrogen (Carlsbad, CA). Recombinant VPO1 proteins were expressed and purified as described in the article [18].

#### Mice

C57BL/6j mice were originally obtained from the Jackson Laboratory and maintained in micro-isolator cages. The experiments on mice were carried out according to the guidelines of the University of Alabama at Birmingham Animal Care and Use Committee. Unless otherwise stated, mice were fed with normal chow diet. Mice at 8–10 weeks were used in the experiments.

#### Immunofluorescence Assay

Glycercosctions of atherosclerotic aorta from low-density lipoprotein receptor (LDLR)−/−deficient mice (generously provided by Dr. Janusz H. Kabarowski at Department of Microbiology, University of Alabama at Birmingham) were fixed by incubation in cold acetone (−20°C) for 2 minutes and rehydrated in phosphate buffered saline (PBS). Standard immunofluorescence staining was carried out. The primary antibodies were polyclonal goat anti-ApoE antibody (Millipore, Billerica, MA) and affinity-purified rabbit anti-VPO1 antibody as described in the article [18]. The secondary antibodies were Alexa Fluor 488-conjugated anti-goat antibody and Alexa Fluor 555-conjugated anti-rabbit antibody. Images were taken and analyzed by Zeiss Axiocamt fluorescence microscope (Carl Zeiss, Heidelberg, Germany).

#### Establish ApoE3-stably Expressing Cell Lines

Full-length human ApoE3 gene was constructed in plasmid pcDNA3. The plasmid containing ApoE3 was transfection into HEK293 cells using FuGENE 6 (Roche, Indianapolis, IN) following the manufacturer’s instruction. After 48 hrs, cells were re-plated and G418 was added into medium at 500 µg/ml. Media containing G418 were changed every 3 days for 3 weeks. The positive colonies of ApoE3-stably expressing cell lines were screened by immunoblot analysis using anti-ApoE antibody.

#### Expression and Purification of ApoE3

For expression and purification of recombinant ApoE3 (rApoE), ApoE3-stably expressing cells were cultured in spinner bottle in serum-free medium [DMEM medium supplemented with 0.1% (w/v) Phuronic® F68 and 3.6 g/L CDM-HD (FiberCell Systems Inc. Frederic, MD)]. In brief, ~0.5 x 10⁶ cells/ml were grown in 100 ml spinner bottles with 40 ml of the serum-free medium at 37°C and 8% CO2 atmosphere with stirring at 100 rpm. After three days, 60 ml of fresh serum-free medium was added and the culture was continued for four days. Subsequently, the culture was transferred to a 1000 ml spinner bottle and 400 ml fresh media was added. When cell density reached 3 x 10⁶/ml, 400 ml fresh media was added and culture was continued for another 3–4 days. The supernatant was harvested by centrifugation at 1000 x g at 4°C for 10 min. The conventional procedures of chromatography were carried out for purification of rApoE. The supernatant was loading into a Q-Sepharose column (1 x 8 cm) with speed at 0.5 ml/min. The column was washed with 100 ml of the serum-free medium at 37°C and 8% CO2 atmosphere with stirring at 100 rpm. After three days, 60 ml of fresh serum-free medium was added and the culture was continued for four days. Subsequently, the culture was transferred to a 1000 ml spinner bottle and 400 ml fresh media was added. When cell density reached 3 x 10⁶/ml, 400 ml fresh media was added and culture was continued for another 3–4 days. The supernatant was harvested by centrifugation at 1000 x g at 4°C for 10 min. The conventional procedures of chromatography were carried out for purification of rApoE. The supernatant was loading into a Q-Sepharose column (1 x 8 cm) with speed at 0.5 ml/min. The column was washed with 100 ml of 20 mM phosphate buffer, pH 8.0, and gradually eluted by 60 ml of 20 mM phosphate buffer containing 0–0.6 M NaCl. rApoE was detected by SDS-PAGE following Coomassie Blue staining and immunoblot analysis. The elution containing rApoE and ~0.4 M NaCl was subject to be concentrated by using GE Vivaspin® 20 (10 kDa MWCO) (GE Healthcare Biosciences, Pittsburgh, PA). The concentrated sample was loaded onto a Sephacryl S-300 column (1 x 100 cm) for further purification. The column was eluted with phosphate buffer at 0.3 ml/min and samples were collected at 4 ml/tube. A280 was monitored. The purity of rApoE was determined by SDS-PAGE. Single band was observed. Protein concentration was measured by Bradford dye-binding procedure using protein assay kit (Bio-Rad, Hercules, CA).

#### Preparation of Lipoproteins

VLDL was isolated from human plasma by ultracentrifugation as described previously [22]. Briefly, VLDL was isolated by ultracentrifugation at plasma density (d = 1.06 g/cm³), at 100,000 × g for 18 hrs at 12°C. VLDL was dialyzed against 20 mM phosphate buffer, pH 7.4, before use.
Oxidation of rApoE and VLDL
rApoE or VLDL was incubated in 800 μM HOCl, or in the presence of 50 μM H₂O₂, 140 mM NaCl, 750 mM/heme VPO1 or 50 nM/heme MPO at 37 °C for 24 hrs. The reactions were terminated by dialysis in 20 mM phosphate buffer, pH 7.4.

Western Blot Analysis
Protein samples were resolved by 12% SDS-PAGE and transferred to polyvinylidene difluoride (PVDF) membranes using a Mini-Tank Transfer System (Bio-Rad Laboratories, Hercules, CA). Proteins were detected by chemiluminescence (Pierce, SuperSignal West Pico Chemiluminescent Substrate) with the primary antibody as indicated.

Measurement of Free Amino Groups
Free Amino acid groups of apolipoprotein were quantified as described previously [23]. Briefly, samples were mixed with 1 ml of 4% NaHCO₃ (w/v; pH 8.4) and 50 μl of 0.1% (v/v) trinitrobenzene sulfonic acid. After incubation at 37 °C for 1 hr, 100 μl of 1 M HCl and 100 μl of 10% SDS were added. Absorbance at 340 nm was recorded.

Preparation of Emulsion Particles
Emulsion particles were prepared by sonication and ultracentrifugation as described in [25]. In brief, triolein and egg PC emulsion particles were prepared by mixing triolein and PC with the weight ratio of 3.5:1 while PC emulsion particles contained only PC. To prepare triolein and egg PC emulsion particles, 5 μl of 0.1 mg/μl egg PC and 1.75 μl of 1 mg/μl triolein in chloroform/methanol solution were mixed in a glass tube and dried under a stream of nitrogen. The tube containing triolein and egg PC was placed into a vacuum desiccator overnight. The dry lipids were suspended in Tris buffer (10 mM Tris-HCl, pH 7.4, 150 mM NaCl, 0.02% NaN₃, 1 mM EDTA) and sonicated for 30 min at 50–60 °C under a stream of nitrogen. Titanium was removed from the crude emulsions by centrifuging at 3000 rpm for 15 min. The samples were then centrifuged at 15,000 rpm for 15 min at 20°C to remove creamy layer and washed twice with the buffer. Additional spinning at 17,000 rpm for 1 hr at 20°C min to remove any contaminated liposome. The resulting creamy top layer was collected as emulsion particles. After purification, the final PC concentration in solution is ~0.3 mg/ml while triolein is ~1 mg/ml.

Binding of ApoE or ox-ApoE to Emulsion Particles
Binding of ApoE or ox-ApoE to emulsion particles was assayed with a centrifugation method as described in [25]. The emulsion particles were incubated with ApoE or ox-ApoE for 1 hr at room temperature with gentle shaking in Tris buffer (pH 7.4) containing 0.25 M sucrose. The final concentrations of ApoE, PC and triolein were 20 μg/ml, ~0.03 mg/ml and ~0.1 mg/ml, respectively. After incubation, the samples were centrifuged in a Beckman ultracentrifuge with a 50 Ti rotor at 30,000 rpm for 30 min. All of lipids were found in the top fraction and free ApoE was in the down layer. The top and lower layers were collected. Equal amount of samples from top or lower layers were carried out for Western blot analysis. The ratio was calculated as emulsion particle-bound ApoE/free ApoE.

Triglyceride, cholesterol and PC assay
To load lipids, THP-1 cells were treated with 100 nM phorbol myristate acetate for 24 hrs, rested another 24 hrs, and then incubated with acetylated-LDL (Ac-LDL) for 24 hrs. LDL enriched cells were washed with PBS and incubated with medium containing 50 mg/L native or ox-ApoE. After 24 hrs, cells were extracted using lysis buffer (CelLytic M, Sigma-Aldrich St. Louis, MO). The cellular triglycerides, cholesterol and PC were measured by Pointe Scientific triglycerides assay kit, Wako cholesterol assay kit, and Sigma–Aldrich PC assay kit, respectively, following the manufacturers’ instructions. For detection lipids in supernatant, LDL-enriched cells were washed with PBS and incubated with medium containing 100 mg/L native or ox-ApoE. After 9 hrs, triglycerides, cholesterol and PC in supernatant were measured by the relative kit. As foam cells are full of engulfed lipid vesicles, the levels of free glycerol/glycerol-3-phosphate are negligible as our previous report [26].

Assays for Mouse Plasma Triglyceride and Cholesterol Concentrations
C57BL/6j mice were deprived of food for 16 hrs before the administration of native or ox-ApoE proteins. Blood volume of mouse is calculated by 38.5 ml per kilogram mice body weight [27], and the endogenous ApoE concentration in mouse serum is 38.7 mg/L [27]. The final concentration of administered rApoE is 60 mg/L. Venous blood was drawn for analysis before and after 1 hr of the injection. Mouse plasma was collected immediately and the triglyceride and cholesterol concentrations were measured as described above. This study was carried out in strict accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. The protocol was approved by the Institutional Animal Care and Use Committee of the University of Alabama at Birmingham (Animal Project Number: 120309582).

Statistical Analysis
Data are shown as means ± SEM, unless otherwise indicated. Quantitative variables were compared by means of ANOVA followed by Tukey’s multiple comparison tests. A value of P<0.05 was considered significant.

Results
VPO1 and ApoE are Localized at Atherosclerotic Lesions
Previous reports demonstrated that ApoE localized at lipid streaks and atherosclerotic plaques [28], [29]. Our immunofluorescence assay confirmed the specific localization of ApoE in aortic atherosclerotic plaques using LDLR-deficient mice (Fig. 1A). VPO1 is a new hPx expressed in vascular endothelial cells and small muscle cells. We evaluated if VPO1 is co-localized with ApoE in atherosclerotic plaques. Immunofluorescence assay showed that VPO1 was also located in the atherosclerotic plaque and was close to ApoE (Fig. 1H). H₂O₂ and chloride are the co-substrate of hPxes and virtually, all cells in vessel walls produce H₂O₂ in varying amounts and in response to diverse stimuli [30] while chloride readily exists in cells with physiological concentration of 100–140 mM. Thus, our data support that VPO1 is proximal to ApoE and potentially mediates oxidation of ApoE.

Oxidation of ApoE by VPO1/H₂O₂/Cl⁻ System
An array of evidence shows that ApoE is highly susceptible to oxidation by MPO [31]. Treatment of VLDL with MPO lead to a variety of oxidations of ApoE proteins including the formation of...
intra- and intermolecular crosslinks with the concomitant formation of high molecular mass aggregates [32], [33]. The peroxidase domain of VPO1 is highly similar to MPO and is able to generate HOCl in the presence of H$_2$O$_2$ [18], [20], [21]. To evaluate the oxidation of ApoE by VPO1, we incubated VLDL or rApoE with VPO1/H$_2$O$_2$/Cl$^-$, comparing with MPO/H$_2$O$_2$/Cl$^-$ system. Oxidation of ApoE was examined by immunoblot analysis. The membranes were blotted by anti-ApoE antibody. It is expected that the migration rate of ox-ApoE is different with the native ApoE. Indeed, ApoE in VLDL oxidized by VPO1/H$_2$O$_2$/Cl$^-$ as well as by MPO/H$_2$O$_2$/Cl$^-$ revealed distinct migration rates comparing with native ApoE controls. In addition to the 34 kDa band, which represents the native ApoE, the immunoreactive bands of ox-ApoE in VLDL showed additional molecular masses of 42 and 80 kDa (Fig. 2A). Some degraded bands in ox-ApoE were also observed (Fig. 2A). Similarly, rApoE oxidized by VPO1/H$_2$O$_2$/Cl$^-$ or by MPO/H$_2$O$_2$/Cl$^-$ revealed the patterns of immunoreactive bands as ox-ApoE in VLDL (Fig. 2A and 2C). Omitting H$_2$O$_2$ in the system, migration rates of rApoE did not change (Data not shown).

The oxidized VLDL was also separated using native PAGE and transferred to PVDF membrane for immunoblot analysis using anti-ApoE antibody (Fig. 2B). VLDL, which size is approximately 25–90 nanometers, stayed on top of the gel whereas the oxidized VLDL revealed smear bands below the main band in native VLDL (Fig. 2B). Taken together, our data suggest both ApoE in VLDL and rApoE are readily oxidized by VPO1/H$_2$O$_2$/Cl$^-$, resulting in a variety of oxidations including intra- and intermolecular crosslinks, degradation etc.

VPO1-mediated Oxidation of Amino Acid Residues in ApoE

MPO/H$_2$O$_2$/Cl$^-$ system generates potent oxidant, HOCl, and mediates oxidation of a variety of biomolecules including proteins, lipids and DNA. In atherosclerosis, lipoproteins are the common targets. Treatment of VLDL with reagent HOCl or MPO/H$_2$O$_2$/Cl$^-$ resulted in the modification of the protein moiety and impairment of protein functions [34]. Lysine-derived amino groups are highly susceptible to HOCl-mediated attack [35]. We incubated rApoE with VPO1/H$_2$O$_2$/Cl$^-$ and examined the status of its amino acid residues. Like MPO/H$_2$O$_2$/Cl$^-$ systems or reagent HOCl, VPO1/H$_2$O$_2$/Cl$^-$ mediated the loss of free amino groups (Fig. 3A). VPO1 showed weaker effects than MPO thought VPO1 concentration was 10 folds (500 nM) than that of MPO (Fig. 3A), consistent with our previous report that VPO1 activity was approximately one twentieth of that of MPO [18]. However, VPO1 is expressed in vessel walls and its concentration in plasma is 1000 folds than that of MPO [18], [19]. It is assumed that the ability of VPO1-mediated oxidation of free amino acids is higher than that of MPO in vessel walls. HOCl at (800 mM) showed a similar ability of oxidation as MPO at 50 nM. Tryptophan residues are also susceptible to oxidation by HOCl [36]. After incubation with VPO1/H$_2$O$_2$/Cl$^-$, MPO/H$_2$O$_2$/Cl$^-$ or reagent HOCl, the quantity of tryptophan residues in ApoE was measured by fluorescence assay (emission 335 nm/excitation 280 nm). We found approximately 6% of tryptophan residues was oxidized by VPO1 (p<0.05 vs native ApoE) (Fig. 3B) similar to that by MPO and HOCl. The data suggest that multiple types of amino acid residues may be oxidized by VPO1/H$_2$O$_2$/Cl$^-$ system like MPO/H$_2$O$_2$/Cl$^-$ and reagent HOCl.

Figure 1. Localization of VPO1 and ApoE in atherosclerotic lesion. Immunofluorescence staining was carried out for detection of ApoE (A) and VPO1 (B) using sections from mouse atherosclerotic lesions. C. Merged image of A and B. D. Nuclei were visualized with Hoechst staining (blue). E. Bright-field image. F. Merged image of A, B, D, and E. Scale bar, 20 μm. Magnification: x400.

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Ox-ApoE Binds Less Lipids in Emulsion Particles than Native ApoE

ApoE plays multiple roles in regulation of metabolism and transportation of lipids. Lipid molecules in the circulation system lie in two ways: as free lipids in very low concentration or in emulsion particles, such as the lipoprotein complexes VLDL, LDL, and HDL [21]. Most lipids in plasma exist in the lipoprotein complexes for facilitating transportation. Takahashi et al reported that rApoE enhanced the binding and internalization of triglyceride-rich lipoproteins to the VLDL receptor [37]. To determine the binding ability of ox-ApoE, emulsion particles from PC or Triolein-PC mixture were prepared by sonication and ultracentrifugation [25]. Emulsion particles were then incubated with ApoE or ox-ApoE. Emulsion particle-bound ApoE was then separated from unbound ApoE (free ApoE) by ultracentrifugation. The separated ApoE proteins were analyzed by immunoblot and quantified by using Image J software (The National Institutes of Health, Bethesda, MD). As shown in Fig. 4, ox-ApoE bound significantly less PC or Triolein-PC in emulsion particles than native ApoE.

Figure 2. Immunoblot analysis of ApoE oxidized by VPO1. A. VLDL was oxidized by MPO or VPO1 as described in Materials and Methods. The oxidized lipoproteins were separated by SDS-PAGE and transferred to PVDF membrane. The blot was analyzed by using anti-ApoE antibody and visualized using chemiluminescence. B. The same as in A, but the samples were separated by using native PAGE. C. rApoE was oxidized by VPO1 or reagent HOCl. Blot was analyzed as in A.
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Figure 3. Loss of free amino groups and tryptophan residues in oxidized ApoE. A. Loss of free amino groups in oxidized ApoE. ApoE was oxidized by VPO1/H2O2/Cl−, MPO/H2O2/Cl−, or HOCl for 3 hrs at 37°C (n = 3). Unmodified amino groups were quantified with the method using trinitrobenzene sulfonic acid as described in Materials and Methods. Absorbance was measured at 340 nm. B. Loss of tryptophan residues following ApoE oxidation. ApoE was incubated with VPO1/H2O2/Cl−, MPO/H2O2/Cl− or HOCl for 3 hrs at 37°C (n = 3) as in A. Fluorescence from free tryptophan residue (emission 335 nm/excitation 280 nm) was quantified with a BioTek Microplate Reader. Trp: tryptophan. Data are shown as means ± SEM; *p<0.05 vs. native ApoE; n = 3.
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Ox-ApoE Shows Less Efflux of Triglycerides, Cholesterol, and PC from Foam Cells

It is reported that ApoE increased free cholesterol efflux into the medium [38]. ApoE as an extracellular receptor promoted cholesterol efflux activity at a dose-dependent manner [39]. To investigate the effect of rApoE on transportation of triglycerides, cholesterol, and PC, we added native ApoE or ox-ApoE in cultured THP-1 foam cells. Ox-ApoE decreased function of THP-1 cells in efflux of triglycerides, cholesterol, and PC, comparing with native ApoE (Fig. 5A, B, and C). Consistently, less triglycerides, cholesterol, and PC in supernatant were detected in ox-ApoE groups than that of the native ApoE group (Fig. 5D, E, and F). Our data suggest that the oxidization of ApoE impairs its function of lipid transportation from foam cells.

Ox-ApoE Exhibits Less Function in Clearing Plasma Triglyceride

Previous studies on mice with transgenic expression of ApoE and mice transfected with virus containing ApoE indicated that ApoE facilitated plasma lipid clearance by liver [10], [11], [12]. In the present study, we ask the effect of ox-ApoE oxidized by VPO1 on transportation and clearance of triglyceride in plasma. We injected PBS, native ApoE or ox-ApoE (either by VPO1 or reagent HOCl) into the circulation via mouse tail vein. To minimize effects of the stress on triglyceride levels, plasma was taken 1 hr before injection. After 1 hr of injection, plasma was taken for analysis. The concentrations of triglycerides were measured. Triglyceride concentrations at pre- and post-injection were compared and the percentage changes were expressed. Decreased plasma triglyceride concentrations were observed after native ApoE administration, comparing with PBS groups (Fig. 6A). Significantly, ox-ApoE increased plasma triglyceride levels. Both reagent HOCl-oxidized ApoE and VPO1-oxidized ApoE had the similar effects (Fig. 6A). Similarly, plasma cholesterol concentrations were decreased by administrating native ApoE whereas slight increase was observed by administrating ox-ApoE (Fig. 6B).

Discussion

The milestone of genesis of atherosclerosis is the deposition of lipoprotein in vessel walls [40]. ApoE is an important protein component of triglyceride-rich lipoprotein complexes including chylomicron, VLDL, IDL, and HDL. These complexes play a
crucial role in transportation, distribution and catabolism of triglyceride-rich lipoproteins in plasma. Increased levels of ApoE facilitate lipid transportation and catabolism whereas decreased levels of ApoE lead to less transportation of lipids, resulting in elevated triglyceride levels in plasma and lipid deposition in vessel walls [10], [11], [12]. ApoE accomplishes its lipid transport and delivery function mainly through VLDL receptor. Therefore, abnormal expression or structural modifications of ApoE will cause its dysfunction. For example, ApoE-deficient mice spontaneously develop atherosclerotic lesions on a standard chow diet [41]. Ox-ApoE oxidized by MPO has impaired function for transporting lipids [32]. Our previous report demonstrated that VPO1 is the second member of hPx family generating potent oxidant, HOCl [20]. In the present study, we further define that VPO1 can, like MPO or reagent HOCl, oxidize ApoE, and impair the ability of ApoE to transport lipids, causing elevated plasma lipid levels.

Oxidative modification of ApoE leads to its structure changes and reduced transportation of lipids. The major lipid-binding domain of ApoE is at the C-terminal (amino acids 241–272) [42]. This region shows higher hydrophobicity on the surface. The present study reveals that amino acid residues of ApoE are widely oxidized, especially that of the hydroxyl group, carboxyl group, and imidazole group. The modification of these residues causes changes of the lipid-binding capacity of ApoE in two ways. One is the damage of its lipid binding domain, causing less lipid attachment [43]. Another is the oxidation of surface polar residues, turning ApoE more hydrophobic [44]. Our data show that the latter seems dominant. The ability of ox-ApoE for the transportation of lipids into cells is impaired, assuming that ox-ApoE has decreased ability to bind to VLDL receptor. Furthermore, ox-ApoE becomes unstable and prone to deposit in the vessel walls. In addition to oxidize ApoE, VPO1 may also oxidize other lipid proteins such as apolipoprotein B, apolipoprotein A-I, apolipoprotein A-II in a similar way since the potent oxidants generated by VPO1 oxidize a variety of proteins without selectivity.

Unlike MPO, which restrictedly expresses in monocytes and neutrophils, VPO1 highly expresses in a variety of tissues including cardiovascular system, lung and liver and is secreted into blood [18]. The plasma VPO1 concentration is approximately three orders of magnitude of that of MPO [19]. Interestingly, VPO1 and ApoE are co-located at the plaque of atherosclerotic aorta from LDLR−/− mice. These data implicate that VPO1 participates ApoE metabolism in plasma as well as in vessel walls at inflammatory sites.

It is well known that inflammatory factors play a key role in genesis and development of atherosclerosis. Lipopolysaccharides (LPS) induce the migration and activation of neutrophils [45]. Ox-LDL stimulates macrophages to release a variety of inflammatory factors including TNF-α and mediates migration and proliferation of smooth muscle cells [46], [47], [48]. The interplay of oxidized lipids and inflammatory factors improve the progress of atherosclerosis. MPO is currently considered as one of major mediators in the genesis and development of atherosclerosis. Like MPO, VPO1 expression is also regulated by inflammatory factors such as TNF-α and LPS [19]. VPO1 is located at the inflammatory sites in vessel walls. Thus, the inflammatory factors and increased expression of VPO1 may interplay in genesis and development of atherosclerosis. We propose that VPO1 is a new and important factor in the ox-ApoE oxidation and clearance of plasma lipids.
mediator for genesis and development of atherosclerosis. One question is which one plays a major role in genesis of atherosclerosis? Because MPO is distributed in inflammatory cells with approximately 5% of dry weight of the cell and has stronger peroxidase activity, we hypothesize that MPO may play a dominant role in the acute stage of atherosclerosis. However, the basal level of VPO1 is relative higher and it is also induced by a variety of inflammatory factors as our previous report that LPS and TNF-α induced VPO1 expression and secretion in HUVECs [19]. Indeed, increased levels of VPO1 were seen in the stable angina pectoris and acute coronary symptom (Unpublished observation). Thus, it is currently unclear if VPO1 plays a major role in atherosclerosis. It is important, but beyond our present study.

H₂O₂ is the co-substrate of VPO1. The major sources of H₂O₂ in vessel walls include mitochondrial electron transport chain, NADPH oxidases (Noxes) and uncoupled nitric oxide synthase [49], [50]. Noxes are professional enzymes for generation of reactive oxygen species and are considered to be regulated deliberately [25]. Former studies had proved that there were several Nox genes expressed in epithelial cells and atherosclerotic plaques [51], [52]. Of interest, at the inflammatory sites, a number of chemokines and cytokines induce the expression and activation of Noxes. For example, TNF-α induced Nox2 activation via PKC-mediated phosphorylation of p47phox in endothelial cells [53]. In addition, it is well known that hPxes are functionally associated with Noxes. MPO utilizes Nox2-generated H₂O₂ (derived from superoxide by dismutation) to produce HOCl in host defense.

Pathologically, synergic actions of MPO and Noxes produce HOCl and oxidize ApoE [18]. We previously found that VPO1 may regulate the proliferation of vascular smooth muscle cells via Nox/H₂O₂/VPO1/HOCl/ERK1/2 pathway and involve in ox-LDL-induced endothelial cell apoptosis through Nox/p38 MAPK/caspase-3 pathway [54], [55]. Thus, we hypothesize that Nox-derived H₂O₂ is the major source of VPO1 catalysis in vessel walls and plasma.

Conclusions

In the present study, we investigated the role of VPO1 in regulating ApoE-mediated lipid metabolism and transportation. We found that ApoE and VPO1 are co-localized at atherosclerotic plaques. VPO1-mediated oxidation of ApoE causes decreased efflux and impaired clearance of plasma lipids. We suggest that VPO1 is a new mediator of genesis and development of atherosclerosis.

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Author Contributions

Conceived and designed the experiments: GC YY LT. Performed the experiments: YY ZC LT WTG. Contributed reagents/materials/analysis tools: YY ZC LT WTG GC. Wrote the paper: GC YY LT WTG.

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Author Contributions

Conceived and designed the experiments: GC YY LT. Performed the experiments: YY ZC LT. Analyzed the data: GC YY LT WTG. Contributed reagents/materials/analysis tools: YY ZC LT WTG GC. Wrote the paper: GC YY LT WTG.
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