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Neutrophil effector responses are suppressed by secretory phospholipase A₂ modified HDL

Sanja Curcic, MSc, Michael Holzer, PhD, Robert Frei, MSc, Lisa Pasterk, MSc, Rudolf Schicho, PhD, Akos Heinemann, MD, and Gunther Marsche, PhD

Institute of Experimental and Clinical Pharmacology, Medical University of Graz, Austria

Abstract

Secretory phospholipase A₂ (sPLA₂) generates bioactive lysophospholipids implicated in acute and chronic inflammation, but the pathophysiologic role of sPLA₂ is poorly understood. Given that high-density lipoprotein (HDL) is the major substrate for sPLA₂ in plasma, we investigated the effects of sPLA₂-mediated modification of HDL (sPLA₂-HDL) on neutrophil function, an essential arm of the innate immune response and atherosclerosis.

Treatment of neutrophils with sPLA₂-HDL rapidly prevented agonist induced neutrophil activation, including shape change, neutrophil extracellular trap formation, CD11b activation, adhesion under flow and migration of neutrophils. The cholesterol-mobilizing activity of sPLA₂-HDL was markedly increased when compared to native HDL, promoting a significant reduction of cholesterol-rich signaling microdomains integral to cellular signaling pathways. Moreover, sPLA₂-HDL effectively suppressed agonist-induced rise in intracellular Ca²⁺ levels. Native HDL showed no significant effects and removing lysophospholipids from sPLA₂-HDL abolished all anti-inflammatory activities.

Overall, our studies suggest that the increased cholesterol-mobilizing activity of sPLA₂-HDL and suppression of rise in intracellular Ca²⁺ levels are likely mechanisms that counteract agonist induced-activation of neutrophils. These counterintuitive findings imply that neutrophil trafficking and effector responses are altered by sPLA₂-HDL during inflammatory conditions.

Keywords

HDL; secretory phospholipase A₂; lysophospholipid; neutrophil

1. Introduction

Concentrations of some secretory phospholipase A₂ (sPLA₂) subspecies can increase hundred-folds in plasma during acute inflammation [1]. More than one third of the mammalian PLA₂ enzymes belong to the sPLA₂ family, which consists of low molecular mass, Ca²⁺ dependent enzymes with a His–Asp catalytic dyad [2]. sPLA₂ enzymes,

Corresponding author: Gunther Marsche, PhD, Institute of Experimental and Clinical Pharmacology, Medical University of Graz, Austria Universitätsplatz 4, 8010 Graz Tel.: +43 316 380 4513 Fax: +43 316 380 9645 gunther.marsche@medunigraz.at.

Conflict of Interest

None declared.

including groups IIA, III, V and X, catalyze the hydrolysis of glycerophospholipids to lysophospholipids and free fatty acids at the sn-2 position. Considerable evidence implicates a potential role for groups IIA and V sPLA₂ in cardiovascular disease and in patients with sepsis and septic shock, and increased plasma levels of sPLA₂ have been observed in acute inflammatory conditions such as sepsis as well as in chronic inflammatory diseases [3-5].

Circulating levels of sPLA₂-IIA and increased sPLA₂ activities associate with cardiovascular risk in patients with sepsis and coronary artery disease [6]. Pathologic studies demonstrated the presence of sPLA₂ isoforms in atherosclerotic lesions and myocardial regions that have sustained ischemic injury [7]. Unexpectedly, inhibition of sPLA₂ in patients with acute coronary syndromes resulted in an excess rate of myocardial infarction and stroke [8] and showed no overall survival benefit in patients with severe sepsis [9]. These studies clearly suggest that the pathophysiologic roles of sPLA₂ are poorly understood. Most sPLA₂ isoforms have the unique capacity to bind and hydrolyze lipoproteins in distinct and specific manners, thereby producing various lipid mediators and modifying the lipid particles. In the acute phase of the inflammatory response sPLA₂ is mostly associated with high-density lipoproteins (HDL) [10], which are the major source of phospholipids in plasma [11] and are important players in the innate and adaptive immunity and cardiovascular disease [12-14]. sPLA₂ overexpression in apoA-I transgenic mice results in a dramatic shift of the HDL particle size toward smaller particles and virtually all plasma sPLA₂ is found in the HDL fraction [15]. Surprisingly, direct effects of sPLA₂ modified HDL on innate immune cells have not been assessed so far. Neutrophils are classically considered as short-lived phagocytes with the ability to release vast amounts of proteolytic enzymes and reactive oxygen species, both of which are important during bacterial infections. Thus, both exuberant and/or diminished responses by the innate immune system may worsen clinical outcomes in severely ill patients with acute and chronic liver diseases [16], airway diseases [17] or acute coronary syndrome [18]. Neutrophils play key roles in sepsis [19] and atherosclerosis [20] and were shown to aggravate endothelial dysfunction, to activate macrophages, promote foam cell formation and to contribute to weakening of the fibrous cap [20]. We therefore thought to assess whether sPLA₂-treated HDL modulates neutrophil function, an essential arm of the innate immune response.

2. Materials and Methods

2.1. Materials

All laboratory reagents were from Sigma (Vienna, Austria), unless otherwise specified. Interleukin 8 (IL-8) and intracellular adhesion molecule-1 (ICAM-1) were purchased from Peprotech (London, UK). Free fatty acids (FFA), lysophosphatidylcholines (LPC) and lysophosphatidylserine 16:0 were from Avanti Polar Lipids (Birmingham, AL, USA). FFAs were dissolved in ethanol and LPCs in chloroform/methanol and stored at -20°C under argon atmosphere. Required amounts of LPC were dried under a stream of nitrogen and redissolved in PBS (pH 7.4). CellFix and FACS-Flow were from BD Bioscience (Vienna, Austria). Secretory phospholipase A₂ (sPLA₂) type III from bee venom and human recombinant secretory phospholipase A₂ type V were purchased from Cayman Europe (Tallin, Estonia). Varespladib was purchased from Eubio (Vienna, Austria). Anti-human

CD11b-FITC and CD16-PE antibodies were obtained from Biozym-Biotech (Vienna, Austria). Fluo-3-AM and SYTOX green were from Life Technologies (Vienna, Austria). Fixative solution was prepared by adding 9 ml distilled water and 30 ml FACS-Flow to 1 ml CellFix.

2.2. Isolation of HDL

HDL was isolated by density gradient ultracentrifugation as described [21-23]. Plasma density was adjusted with potassium bromide (Sigma, Vienna, Austria) to 1.24 g/ml and a two-step density gradient was generated in centrifuge tubes (16 × 76 mm, Beckman) by layering the density-adjusted plasma (1.24 g/ml) underneath a NaCl-density solution (1.006 g/ml). Tubes were sealed and centrifuged at 90.000 rpm for 4 hours in a 90Ti fixed angle rotor (Beckman Instruments, Krefeld, Germany). After centrifugation, the HDL-containing band was collected and desalted via PD10 columns (GE Healthcare, Vienna, Austria) and immediately used for experiments.

2.3. sPLA₂ treatment of HDL

HDL was incubated in the presence of 200 ng/ml sPLA₂ type III from bee venom or 400 ng/ml human recombinant type V sPLA₂ in PBS containing Ca²⁺ and Mg²⁺, overnight at 37°C, in order to hydrolyse HDL-associated phospholipids.

2.4. Lysophosphatidylcholine (LPC), free-fatty acid (FFA) and lysophosphatidylserine enrichment/depletion of HDL

In order to generate LPC-, FFA-, or lysophosphatidylserine-enriched HDL, 1 mg/ml HDL was incubated with 0.6 mmol/L 16:0, 18:1, 18:2 or 20:4 FFA, with 0.6 mmol/L 16:0, 18:1, 18:2 or 20:4 LPC or with 0.6 mmol/L lysophosphatidylserine 16:0 for 2 hours at 37°C. Unbound LPCs and FFAs were removed by gel filtration and HDL-associated LPC and FFA contents were determined as described below. In some experiments, sPLA₂-treated HDL was incubated in the presence or absence of 50 mg/ml albumin (1 h, 37°C) and sPLA₂-HDL was re-isolated by density gradient ultracentrifugation in order to remove LPCs and FFAs. LPC and FFA contents were assessed as described below.

2.5. Determination of FFAs and LPCs

HDL-associated lipids were extracted according to Bligh and Dyer [24] and dried under a stream of nitrogen. Dried lipid extracts were resuspended in 200 µl CHCl₃/MeOH (1:1, v/v) containing 1 pmol/µl of LPC 17:1 serving as internal standard. Chromatographic separation of lipids was performed by an Accela HPLC (Thermo Scientific) on a Thermo Hypersil GOLD C18, 100 × 1 mm, 1.9 µm column. Solvent A was a water solution of 1% ammonium acetate (v/v) and 0.1% formic acid (v/v) and solvent B was acetonitrile/2-propanol (5:2, v/v) supplemented with 1% ammonium acetate (v/v) and 0.1% formic acid (v/v), respectively. The gradient was run from 35% to 70% B for 4 min, then to 100% B in additional 16 min with subsequent hold at 100% for 10 min. The flow rate was 250 µl/min. Phospholipid species were determined by a TSQ Quantum ultra (Thermo Scientific) triple quadrupole instrument in positive ESI mode. The spray voltage was set to 4500 V and capillary voltage to 35 V. The LPC species were detected in a precursor ion scan on m/z 184 at 34 eV. Peak

areas were calculated by QuanBrowser for all lipid species and the calculated peak areas for each species were expressed as a % of internal standard. Results are expressed as nmol/mg HDL protein. FFA content was assessed using a non-esterified fatty acids kit (Diasys, Holzheim, Germany). In some experiments the Azwell LPC Assay Kit (Hözl Diagnostika) was used to assess the LPC content of sPLA₂-HDL.

2.6. Neutrophil isolation

Human polymorphonuclear leukocytes (PMNL) were isolated as previously described [25;26], from peripheral blood of healthy volunteers according to a protocol approved by the Ethics Committee of the Medical University of Graz. Prior to blood sampling from healthy volunteers, all donors signed an informed consent form. Platelet-rich plasma was removed by centrifugation of citrated whole blood. Erythrocytes were removed by dextran sedimentation. PMNL were isolated by Histopaque gradient centrifugation. Any erythrocyte contamination of the PMNL pellet was removed by hypotonic shock lysis. The purity and viability of neutrophil preparation was greater than 95%. All functional assays of neutrophils were performed in assay buffer (PBS with Ca²⁺ and Mg²⁺, HEPES 10 mmol/L, glucose 10 mmol/L, bovine serum albumin 0.1%, pH 7.4).

2.7. Neutrophil shape change assay

Neutrophil shape change was measured as previously described [27;28]. Isolated PMNL were resuspended in assay buffer and aliquots of cells (about 3×10^5 cells per sample) were preincubated with HDL samples and then stimulated in 37°C shaking water bath with interleukin-8 (IL-8), N-formyl-methionyl-leucyl-phenylalanin (fMLP) or complement component 5a (C5a) for 4 min, with lipopolysaccharide (LPS) in the presence of 2 % serum for 90 min or with *E.coli* bacteria for 60 min at a final volume of 100 µl. Cells were transferred to ice and 150 µl of ice-cold fixative solution was added to terminate the reaction and maintain the change in cell shape until analysis. The samples were then analyzed on a FACScalibur flow cytometer (BD Biosciences). Eosinophils were distinguished from neutrophils according to granularity (side scatter) and by their autofluorescence in the FL-2 channel. Shape change was determined as the increase in the forward scatter property of the cell compared with vehicle stimulation.

2.8. CD11b activation

Isolated PMNL were preincubated with HDL samples and stimulated with IL-8 (3 nmol/L), fMLP (5 nmol/L) or C5a (30 nmol/L) for 4 min at 37°C in shaking water bath in the presence of FITC-conjugated Ab to the active epitope of CD11b. Cells were fixed and then analyzed by flow cytometry [29].

2.9. Neutrophil adhesion under flow conditions

Vena8 biochips (Cellix Ltd, Dublin, Ireland) were coated with 10 µg/ml intracellular adhesion molecule-1 (ICAM-1) at 4°C overnight in a humidified box. On the next day, the chips were washed twice with distilled water, blocked with 0.1 % bovine serum albumin for 30 minutes and rinsed with distilled water. Isolated PMNL were resuspended in assay buffer (containing Ca²⁺ and Mg²⁺) and treated with vehicle, HDL, or sPLA₂-HDL for 15 min at

37°C. Cells (3×10^6 /ml) were then perfused over the ICAM-1 coated channels at constant shear stress of 0.5 dyne cm^{-2} for 5 minutes using the Mirus nanopump (Cellix). Neutrophil adhesion was recorded on an Olympus IX70 fluorescence microscope and an Olympus UPIanFI-X20/0.40 lens, using a Hamamatsu ORCA-ER digital camera and the Olympus CellP software. Cell images were taken 5 minutes after the start of perfusion and adherent neutrophils were analyzed using ImageJ software (National Institutes of Health) as described before [30].

2.10. Migration assay

Migration of freshly isolated human neutrophils (1.5×10^5 cells per well) was assessed using 96-well transwell plates with a pore size of $8 \mu\text{m}$ (Corning) as described [31]. Neutrophils were preincubated with vehicle, sPLA₂-HDL, HDL or PLA₂ for 15 minutes at 37°C and seeded into the upper wells. Cells were allowed to migrate towards IL-8 (10 nmol/L) for 1 hour at 37°C. Cells that had migrated to the lower compartment were enumerated by flow cytometric counting for 30 s. Spontaneous migration was determined in wells containing only assay buffer. To calculate the chemotactic index, the number of cells migrated in response to IL-8 was divided by the number of spontaneously migrated cells.

2.11. Ca²⁺ Flux

Changes in intracellular Ca²⁺ levels in neutrophils were analyzed by flow cytometry as described previously [32]. PMNL were loaded with the cell membrane permeable Ca²⁺-sensitive dye Fluo-3-AM ($2 \mu\text{mol/L}$) in the presence of 0.02 % F-127 pluronic acid for 60 min at room temperature. Cells were then washed, stained with anti-CD16 (PE) and resuspended in assay buffer. Neutrophils were identified as CD16-positive cells.

2.12. Neutrophil extracellular traps (NET)

The kinetics of NET formation was measured as previously described [33]. Isolated human neutrophils (5×10^4 cells per well) were treated in 96-well black plates in a final volume of $200 \mu\text{l}$ in the presence of SYTOX green ($5 \mu\text{mol/L}$), a cell-impermeable nucleic acid dye. 10 nmol/L PMA was used to induce NET formation. NET formation was observed by measuring mean fluorescence (Ex 488 nm, Em 523 nm) every 10 min for 5 hours at 37°C (FlexStation II; Molecular Devices).

2.13. Cholesterol-rich microdomain (lipid raft) assessment

Isolated human neutrophils were incubated with different HDL preparations for 5 to 120 min at 37°C; subsequently cells were washed with PBS and incubated with $1 \mu\text{g/ml}$ FITC-cholera toxin B for 1 hour at room temperature. Neutrophils were fixed and lipid raft abundance was measured by flow cytometry. For fluorescence microscopy, cells were spun on glass coverslips using a Cytospin 3 centrifuge (Shandon). The cells were mounted with Vectashield mounting medium including DAPI (Szabo Scandic, Vienna, Austria) and analyzed using OLYMPUS fluorescence microscope equipped with a Hamamatsu ORCA CCD camera.

2.14. Cholesterol efflux measurement

Isolated neutrophils were labeled for 3 hours with [³H]-cholesterol (1 μCi/ml) at 37°C in the presence of 2 μg/ml Acyl-CoA:cholesterol acyltransferase (ACAT) inhibitor Sandoz 58-035. The cells were then washed three times with PBS containing 1 mg/ml bovine serum albumin by centrifugation. Efflux to the respective cholesterol acceptors was allowed to occur for 1 hour at 37°C in a total volume of 400 μl. After the efflux period, cells were pelleted and the radioactivity in cells and supernatant was measured.

2.15. Filipin staining of unesterified cholesterol

Neutrophils were incubated with HDL samples, cells were then washed with ice cold PBS, fixed on ice for 20 min, washed with PBS and stained for 1 h with filipin (100 μg/ml). Cells were washed before quantification by flow cytometry (LSR II, BD Biosciences).

2.16. Statistical analysis

Data are shown as mean ± SD for n observations. Experiments were repeated three to six times with neutrophils from different donors. Statistical analyses were performed using GraphPad Prism Version 4.03. Student's t-test was used for experiments comparing two groups and one-way ANOVA with Tukey multiple-comparison post hoc test was used for experiments comparing three or more groups. Significance was accepted at p<0.05.

3. Results

3.1. sPLA₂ modification of HDL generates particles with potent shape change inhibitory activity

Stimulation of PMNL with chemoattractants results in rapid reorganization of the cytoskeleton and changes of their shape, which can be assessed by flow cytometry (Supplementary Figure IA to ID) [34]. These responses are important prerequisites for granulocyte adhesion, polarization, locomotion, and degranulation. To elucidate the role of sPLA₂ in chemoattractant-induced shape change responses, neutrophils were added to 20 % serum pretreated for 20 hours in the absence or presence of sPLA₂ type III. Cells were subsequently stimulated with interleukin 8 (IL-8) and increase in the forward scatter (shape change response) was assessed by means of flow cytometry. Surprisingly, sPLA₂-treated serum almost completely reversed IL-8 induced shape change (Figure 1A). Given that HDL is the major carrier of phospholipids in plasma and the major substrate for sPLA₂ [10;15], we isolated HDL from sPLA₂-treated plasma by ultracentrifugation and assessed whether the HDL containing fraction shows similar effects. Indeed, HDL isolated from sPLA₂ treated plasma effectively suppressed neutrophil shape change, whereas HDL isolated from control plasma showed no effect (Figure 1B). We next assessed whether direct modification of HDL with sPLA₂ (sPLA₂-HDL) generates particles with shape change inhibitory activity. The lysophosphatidylcholine (LPC) content of sPLA₂-HDL and control HDL is shown in Supplementary Table I. We observed that sPLA₂-HDL effectively inhibited the activation of neutrophils induced by various agonists (IL-8, formyl-methionyl-leucyl-phenylalanine (fMLP), lipopolysaccharide (LPS), complement 5a (C5a) and *E. coli* bacteria) (Figure 2A to 2E and Supplementary Figure ID), whereas native HDL or addition of sPLA₂ alone showed

no effects. Importantly, sPLA₂-HDL-induced inhibition of shape change was seen in neutrophils from all donors. Several HDL preparations from different donors showed comparable inhibitory effects after sPLA₂ treatment. Notably, also low-density lipoprotein treated with sPLA₂ suppressed agonist induced shape change (Supplementary Figure II). sPLA₂-HDL did not impact neutrophil viability (Supplementary Figure III). These results suggest that sPLA₂ induced hydrolysis of HDL-phospholipids is required for the observed effects. The inhibitory activity of sPLA₂-HDL on neutrophils was dose dependent (Figure 2F).

Notably, addition of sPLA₂-HDL after stimulation of neutrophils (with LPS or IL-8) rapidly reversed shape change (Supplementary Figure IV).

3.2. sPLA₂-generated lysophosphatidylcholines are the active moiety in HDL

Next, we assessed whether the potent effects of sPLA₂-HDL on neutrophils result from phospholipid hydrolysis products that are formed and enriched in sPLA₂-HDL. Therefore, we added an excess of fatty-acid free albumin to sPLA₂-HDL to remove lysophospholipids and free fatty acids and re-isolated HDL using density gradient ultracentrifugation. The lysophospholipid and FFA content following albumin incubation was effectively reduced as shown in Supplementary Figure V. We observed that the activity of sPLA₂-HDL was not due to reduced particle size or other structural alterations, but rather depended on the presence of lysophospholipids and/or free fatty acids in HDL (Figure 3A and 3B). In order to investigate whether specific free fatty acids and/or lysophospholipids mediate the effects on neutrophils, we enriched HDL with different free fatty acids and lysophospholipids. We observed that the major sPLA₂-HDL associated lysophospholipid LPC 16:0 was the most effective, whereas other LPCs and free fatty acids showed weak or no inhibitory effects (Figure 3C and 3D). Interestingly, besides LPC 16:0 also lysophosphatidylserine 16:0 showed potent inhibitory activity (Supplementary Figure VI).

3.3. sPLA₂-HDL suppresses CD11b activation, adhesion, migration and extracellular trap (NET) formation of neutrophils

When stimulated, neutrophils display an increased activation of CD11b, which is an important integrin for leukocyte adhesion and migration. In order to further investigate the effects of sPLA₂-HDL on neutrophil function, we measured CD11b activation. sPLA₂-HDL potently inhibited CD11b activation induced by IL-8, fMLP and C5a (Figure 4A to 4C).

Interestingly, LPS induced IL-8 production of neutrophils was not inhibited by sPLA₂-HDL (Supplementary Figure VII). Neutrophils, along with other inflammatory cells, infiltrate atherosclerotic plaques. An intriguing role has recently been suggested for neutrophil extracellular trap (NET) formation in this process [35;36]. NETs are pro-inflammatory, antimicrobial structures consisting of extracellular chromatin decorated with granular and cytoplasmic proteins, such as myeloperoxidase and neutrophil elastase. Of particular interest, stimulation of NET formation with the protein kinase C activator phorbol 12-myristate 13-acetate was effectively suppressed by sPLA₂-HDL (Figure 4D).

To show that the inhibition of neutrophil activation was dependent on the enzymatic activity of sPLA₂, varespladib (an inhibitor of sPLA₂ types IIA, V and X but not sPLA₂ type III)

was added during incubation of sPLA₂ with HDL. In agreement with results above, varespladib prevented the effects of sPLA₂ type V-treated HDL on neutrophil shape change and CD11b activation induced with IL-8 (Figure 5) and other agonists (Supplementary Figure VIII). To assess whether the sPLA₂-HDL-induced inhibition of CD11b activation is associated with functional alterations, we examined neutrophil adhesion to intercellular adhesion molecule 1 coated surfaces under shear stress. sPLA₂-HDL significantly reduced neutrophil adhesion under flow conditions (Figure 6A and 6B). Moreover, in accordance with these data, incubation of neutrophils with sPLA₂-HDL potently inhibited neutrophil migratory response towards IL-8 (Figure 6C).

3.4. sPLA₂-HDL suppresses agonist induced Ca²⁺ flux and reduces lipid raft assembly

Cytoplasmic Ca²⁺ changes in neutrophils play a key role in a series of pathways leading to activation. Stimulation with IL-8 resulted in a marked rise in the intracellular Ca²⁺, which was effectively suppressed by sPLA₂-HDL, whereas control HDL showed no effect (Figure 7A). Interestingly, we observed that sPLA₂-HDL on its own moderately induced Ca²⁺ flux in unstimulated (resting) cells. To gain insights into the potential mechanism(s) involved in sPLA₂-HDL effects on neutrophils, we next examined the involvement of cAMP, as previous investigations showed that lysophospholipids increase intracellular cAMP in neutrophils. However, we did not observe altered cAMP levels in neutrophils treated with sPLA₂-HDL (Supplementary Figure IX). Native HDL differs from sPLA₂-HDL in its lysophospholipid content, raising the possibility that the variant effects of these particles on neutrophil function may be related to their cholesterol efflux activity. Interestingly, cholesterol efflux capability of sPLA₂-HDL was increased when compared to control HDL (Figure 7B), reflected by a significantly decreased membrane free cholesterol content in neutrophils as shown by filipin fluorescence staining (Figure 7C).

Given that cholesterol-rich lipid rafts are known to play an important role in leukocyte activation, we investigated whether increased cholesterol efflux capability of sPLA₂-HDL alters lipid raft abundance in neutrophils. For that purpose, FITC-labeled cholera toxin B, which binds to the raft constituent ganglioside GM1, was used to visualize lipid rafts. Indeed, we observed that sPLA₂-HDL caused rapid depletion of cholera-toxin positive lipid rafts in neutrophils (Figure 7D, 7E). ABCA1 and SR-BI have been reported to localize selectively in lipid rafts or caveolae [37;38] and to be involved in cholesterol efflux. However, neither the ABCA1-inhibitor probucol [39] nor the SR-BI inhibitor BLT-1 [40] diminished sPLA₂-HDL and native HDL mediated efflux from neutrophils, suggesting that the increased cholesterol efflux capability of sPLA₂-HDL is independent of these receptors (Supplementary Figure X). Moreover, when cholesterol efflux experiments were performed with J774 macrophages, cholesterol efflux capability of sPLA₂-HDL was not different compared with native HDL, suggesting a cell type specific effect (Supplementary Figure XI).

4. Discussion

The studies presented here demonstrate that sPLA₂ modification of HDL generates an anti-inflammatory particle that has the ability to effectively alter neutrophil function. Of

particular interest is the observation that sPLA₂-HDL, but not control HDL, effectively depleted the free cholesterol content of neutrophils and suppressed agonist-induced rise in intracellular Ca²⁺ levels. It is now well established that alterations in membrane cholesterol content cause disruption of cholesterol-rich signaling microdomains integral to cellular signaling pathways known as lipid rafts [41]. Several lines of evidence suggest that HDL regulates the plasma membrane cholesterol content through the ability to remove cholesterol and other lipid species from cells, thereby dampening plasma membrane receptor signaling [42]. Our studies suggest that the increased cholesterol-mobilizing activity of sPLA₂-HDL as well as the suppression of agonist-induced rise in intracellular Ca²⁺ levels are likely mechanism that counteracts activation of neutrophils, resulting in an effective inhibition of CD11b activation and NET formation. CD11b mediates inflammation by regulating leukocyte adhesion and migration and has been implicated in several immune processes such as phagocytosis, cell-mediated cytotoxicity, chemotaxis and cellular activation [43]. Importantly, sPLA₂-HDL efficiently inhibited neutrophil adhesion to ICAM-1 coated surfaces under shear stress and potently inhibited neutrophil migratory response towards IL-8. Although NETs are able to trap and kill bacteria and their formation is an important first-line defense against various pathogens, excessive NET production may strongly promote inflammation, leading to endothelial damage, infiltration of further neutrophils into atheromatous lesions, thrombus formation and autoimmunity [35;36].

Of particular interest, we observed that the major sPLA₂-HDL associated lysophospholipid, LPC 16:0, was the most effective, whereas other LPCs and free fatty acids showed weak or no inhibitory effects. Given that groups IIA, V and X show a differential hydrolysis of molecular species of phosphatidylcholine [44], sPLA₂ modified HDLs with various activities are expected to be formed under inflammatory conditions.

Notably, treatment with native HDL showed only minimal effects on neutrophil function despite its ability to efflux cholesterol. Our results suggest that bidirectional lipid flux (efflux of cholesterol is accompanied by influx of HDL-cholesterylester) mediated by control HDL results in minimal net change in the cholesterol content of the neutrophil membrane as observed by filipin staining. We assume that lysophospholipid-enrichment of HDL generates a particle that more efficiently depletes neutrophil cholesterol. In line with our finding is the observation that that lysophospholipids rapidly enter lipid rafts in the cell membrane promoting raft fission [45] and promote net cholesterol efflux [46], which might explain the increase in cholesterol efflux properties of sPLA₂-HDL. In line with that notion, sPLA₂-HDL preparations, but not control HDL, markedly disrupted neutrophil lipid rafts. Interestingly, the increased ability of sPLA₂-HDL to mobilize cell cholesterol seems to be independent of ABCA1 and SR-BI, proteins known to mediate cholesterol efflux (Supplementary figure X). Moreover, the improved cholesterol efflux ability of sPLA₂-HDL strikingly differs between cell types, given that sPLA₂-HDL mediated cholesterol efflux of from J774 macrophages was not different from native HDL (Supplementary figure XI). The reason for that surprising observation is not clear, but different lipid raft content and/or lipid raft composition of freshly isolated neutrophils compared with a macrophage cell line might be an explanation.

Of particular interest is the recent observation that a large fraction of phosphatidylcholine moiety of administered reconstituted HDL is rapidly hydrolyzed by sPLA₂ isoforms in blood and causes a remarkable increase of HDL-associated lysophospholipid content reaching levels of 300 μmol/L 4 hours post administration [47;48], which raises the possibility that formed lysophospholipids contributed to the potent anti-inflammatory activity of administered reconstituted HDL.

Previous studies supported an important role for a subset of sPLA₂ isoforms (sPLA₂-IIA, -III, -V, and -X) from initiation and progression to cardiovascular complications in the pathophysiology of atherosclerosis [49]. Hydrolysis of phosphatidylcholine in low-density lipoprotein by sPLA₂ isoforms has been implicated in the promotion of atherosclerosis [50;51]. These observations have stimulated interest in the potential value of sPLA₂ inhibition as a cardioprotective strategy. Initial studies demonstrated that varespladib, a nonspecific pan-sPLA₂ inhibitor, reduced activity of sPLA₂-IIA by more than 90%, in addition to lowering low-density-lipoprotein cholesterol and C-reactive protein in patients with stable coronary disease and acute coronary syndrome. Surprisingly, varespladib-induced sPLA₂ inhibition did not reduce cardiovascular ischemic complications and resulted in an excess rate of myocardial infarction and stroke in patients with acute coronary syndromes [8]. Moreover, inhibition of sPLA₂ showed no overall survival benefit in septic patients [9]. Interestingly, also a large phase III study testing an inhibitor of lipoprotein-associated phospholipase A₂ has failed to lower the risk of cardiovascular events in coronary heart disease patients [52], providing further evidence that the physiologic and pathophysiologic roles of phospholipases are not well understood.

A potential limitation of the present study is that we only assessed the impact of sPLA₂ on HDL functionality, while *in vivo* HDL particles are subjected to an array of compositional changes and factors acting in concert. Hydrolysis of HDL phospholipids by sPLA₂ alters the structure and composition of HDL particles with the metabolic consequences of increased catabolism. Experimental studies have shown that the addition of LPS to plasma resulted in a marked decrease in cholesterol ester transfer protein activity, leading to changes in HDL levels considered as an adaptive response to preserve or increase HDL [53]. But also other factors alter HDL composition and function under inflammatory conditions. The concentrations of the acute phase protein SAA markedly increase during the acute-phase reaction. SAA associates with HDL, altering structure and function of HDL [54]. Interestingly, overexpression of SAA in the absence of generalized inflammation does not result in changes of HDL cholesterol or apoA-I levels *in vivo* [55] whereas overexpression of sPLA₂ results in alterations of HDL metabolism *even* in the absence of generalized inflammation [56]. Although increased sPLA₂ and decreased CETP activities might be beneficial for host defense in the setting of acute infection, replacement of HDL-apolipoproteins by SAA might contribute to increased incidence of cardiovascular disease [54].

In conclusion, our data provide novel evidence that the increased cholesterol mobilizing activity and suppression of rise in intracellular Ca²⁺ levels of sPLA₂-HDL might effectively counteract agonist-induced activation of neutrophils. Our results raise the possibility that

sPLA₂-induced modification of HDL composition and function modulates neutrophil trafficking and effector responses during inflammation.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgements

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Abbreviations

C5a	complement component 5a
FFA	free fatty acid
fMLP	N-formyl-methionyl-l-leucyl-l-phenylalanine
HDL	high-density lipoprotein
ICAM-1	intracellular adhesion molecule-1
IL-8	interleukin 8
NET	neutrophil extracellular traps
LPC	lysophosphatidylcholine
PC	phosphatidylcholine
PMNL	polymorphonuclear leukocytes
sPLA₂	secretory phospholipase A ₂
sPLA₂-HDL	secretory phospholipase A ₂ treated HDL

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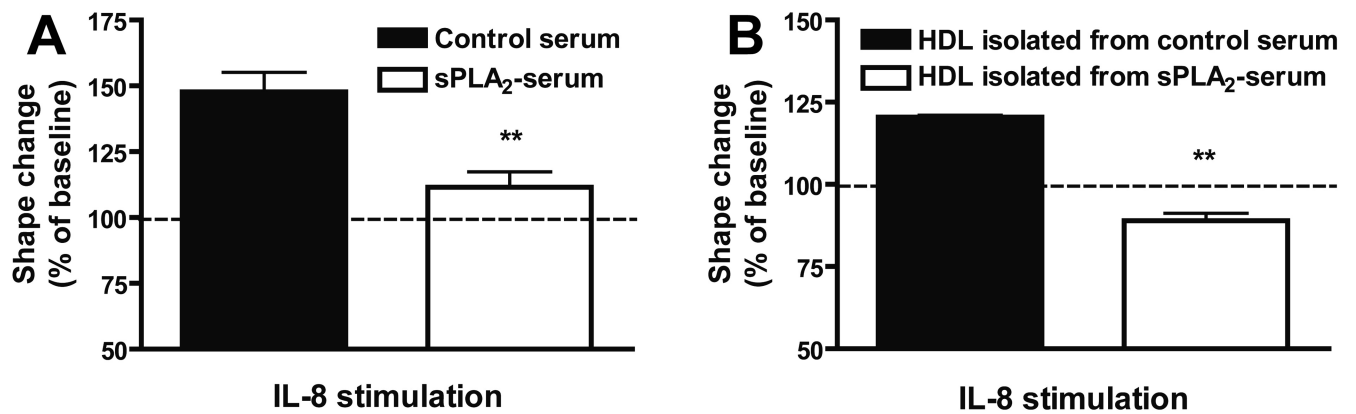


Figure 1. sPLA₂-treated serum inhibits neutrophil activation

Freshly isolated human serum was incubated in the absence or presence of sPLA₂ type III for 24 hours at 37°C. (A) Treated and untreated serum (20 % v/v) was subsequently added followed by stimulation with IL-8 (3 nmol/L) for 4 min. (B) HDL was isolated from sPLA₂ treated serum and control serum and subsequently added to human neutrophils (125 µg HDL-protein/ml, 15 min at 37°C) followed by stimulation with IL-8 (3 nmol/L) for 4 min. Neutrophil shape change was assessed by flow cytometry. Values are expressed as % of baseline. The results are shown as mean ± SD of three separate experiments using PMNL from different healthy donors. Statistical analysis was performed using Student's t-test. *p < 0.05, **p < 0.01 versus control.

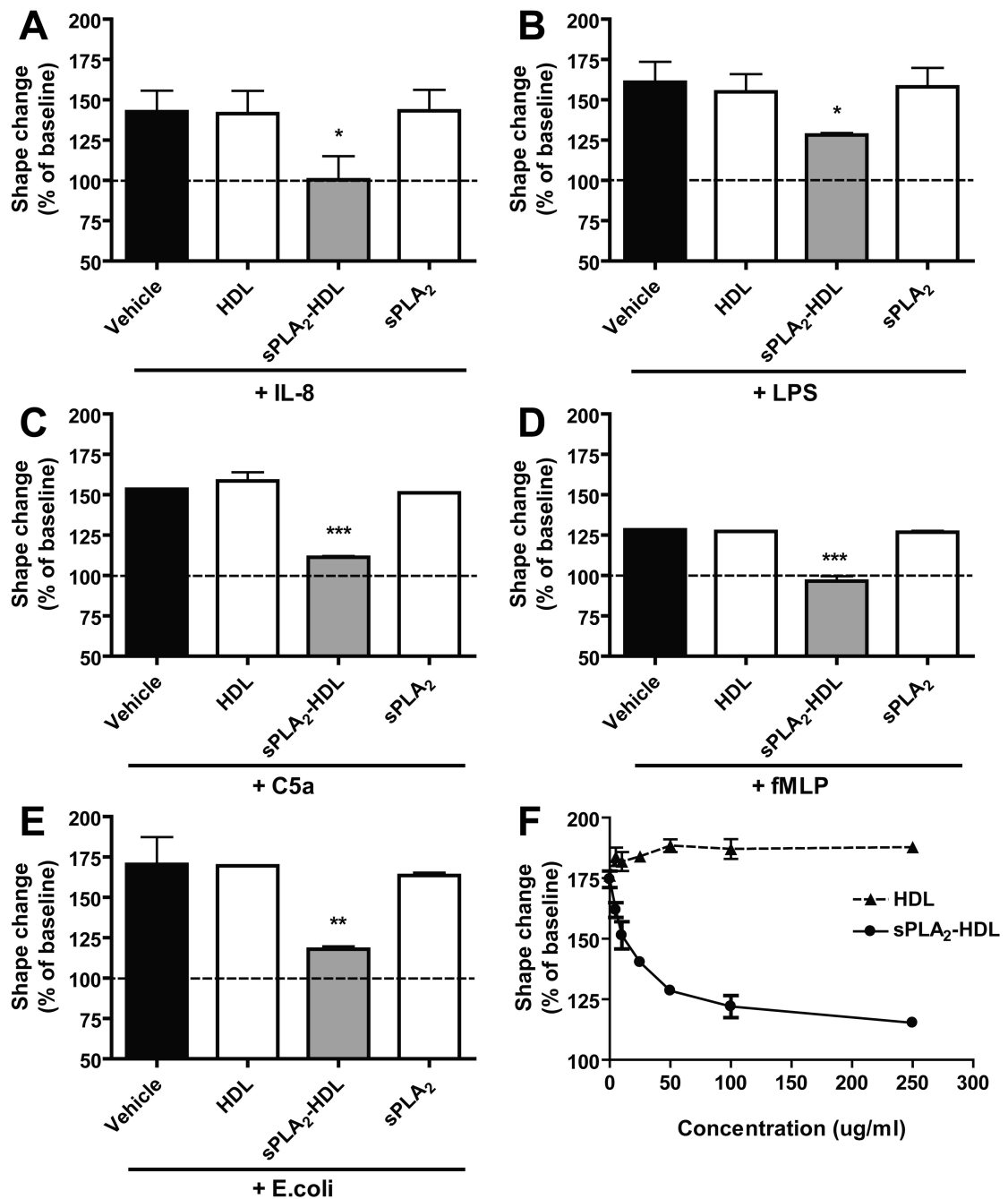


Figure 2. sPLA₂-treated HDL inhibits IL-8, LPS, C5a, fMLP and E. coli-induced neutrophil activation

Isolated neutrophils were incubated with vehicle, native HDL (100 μ g protein/ml), sPLA₂-treated HDL (100 μ g protein/ml) or sPLA₂ alone for 15 min at 37°C, and stimulated with (A) IL-8 (3 nmol/L, 4 min, 37°C) (B) 1 ng/ml LPS in the presence of 2 % serum (90 min, 37°C) (C) C5a (30 nmol/L, 4 min, 37°C) (D) fMLP (5 nmol/L, 4 min, 37°C) or (E) E. coli bacteria (1 h, 37°C, the ratio of neutrophils and bacteria was 1:5). Neutrophil shape change was measured by flow cytometry. Values are expressed as % of baseline. Results shown are mean \pm SD. The data represent the results from three to five experiments done using PMNL

from different healthy donors. Statistical significance was assessed by one-way ANOVA with Tukey multiple-comparison post hoc test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ versus vehicle. (F) Concentration-response curve with sPLA₂-treated HDL. Neutrophils were incubated with increasing concentrations of native HDL or sPLA₂-treated HDL (0 up to 250 $\mu\text{g/ml}$) for 15 min at 37°C and stimulated with 1 ng/ml LPS (90 min, 37°C). Neutrophil shape change was measured by flow cytometry and results are shown as mean \pm SD (n=3).

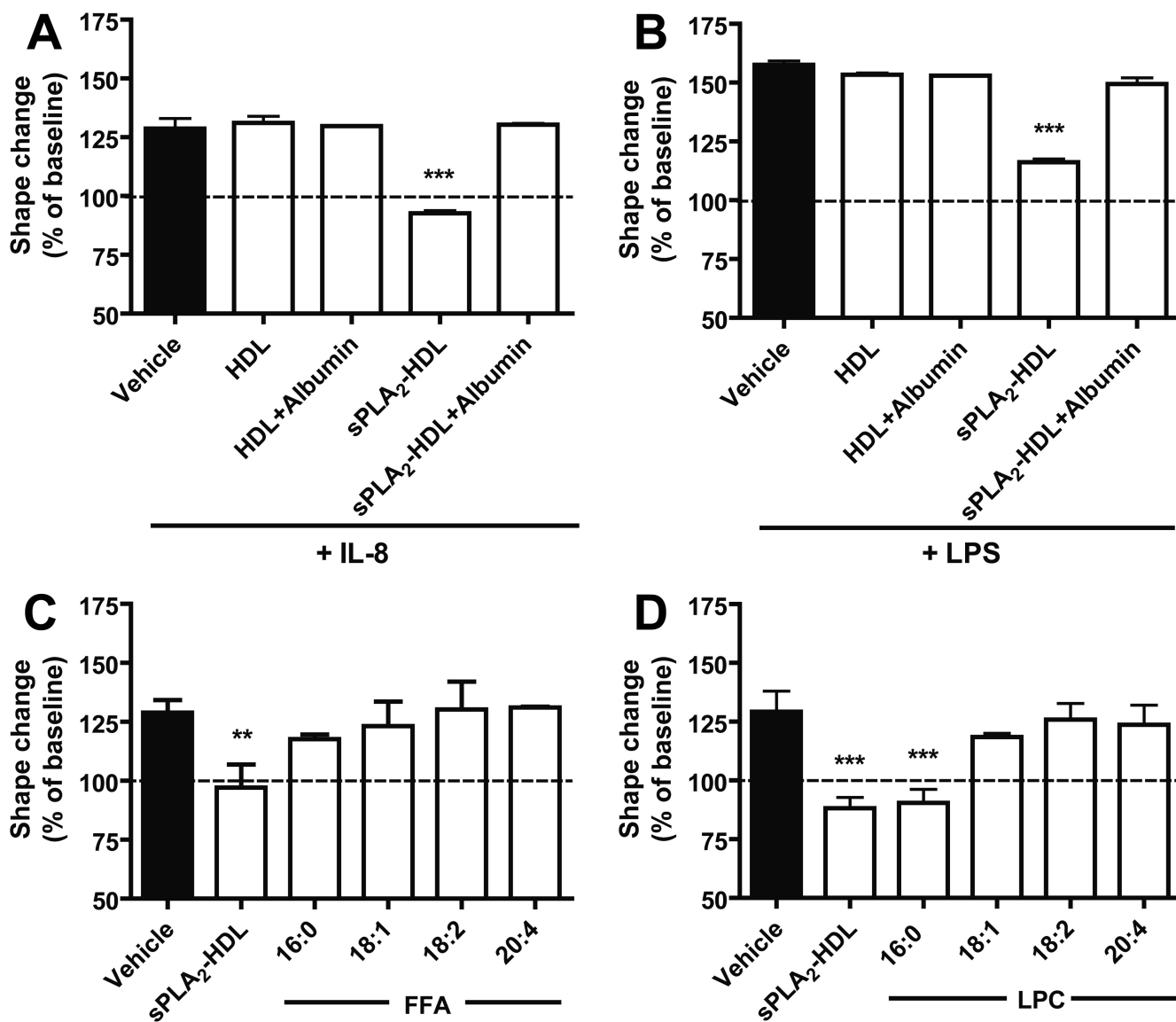


Figure 3. Presence of LPC on sPLA₂-treated HDL is necessary for the inhibition of neutrophil shape change

Native or sPLA₂-treated HDL were incubated in the presence or absence of 50 mg/ml albumin (1 h, 37°C) and HDL was re-isolated by density gradient ultracentrifugation to remove FFA and lysophospholipids as described in Materials and Methods. 100 µg/ml HDL protein was used in the shape change assay with (A) IL-8 (3 nmol/L, 4 min) or (B) 1 ng/ml LPS (90 min, 37°C). **Enrichment of HDL with FFA and LPC.** Neutrophils were preincubated (15 min, 37°C) with sPLA₂-HDL (100 µg/ml protein) and (C) HDL enriched with FFA (16:0, 18:1, 18:2, 20:4) or (D) HDL enriched with LPC (16:0, 18:1, 18:2, 20:4). Neutrophils were then stimulated with IL-8 (3 nmol/L, 4 min, 37°C). Neutrophil shape change was measured by flow cytometry. Values are expressed as % of baseline. Results shown are mean ± SD of three separate experiments using cells from different healthy donors. Statistical significance was assessed by one-way ANOVA with Tukey multiple-comparison post hoc test. *p < 0.05, **p < 0.01, ***p < 0.001 *versus* vehicle.

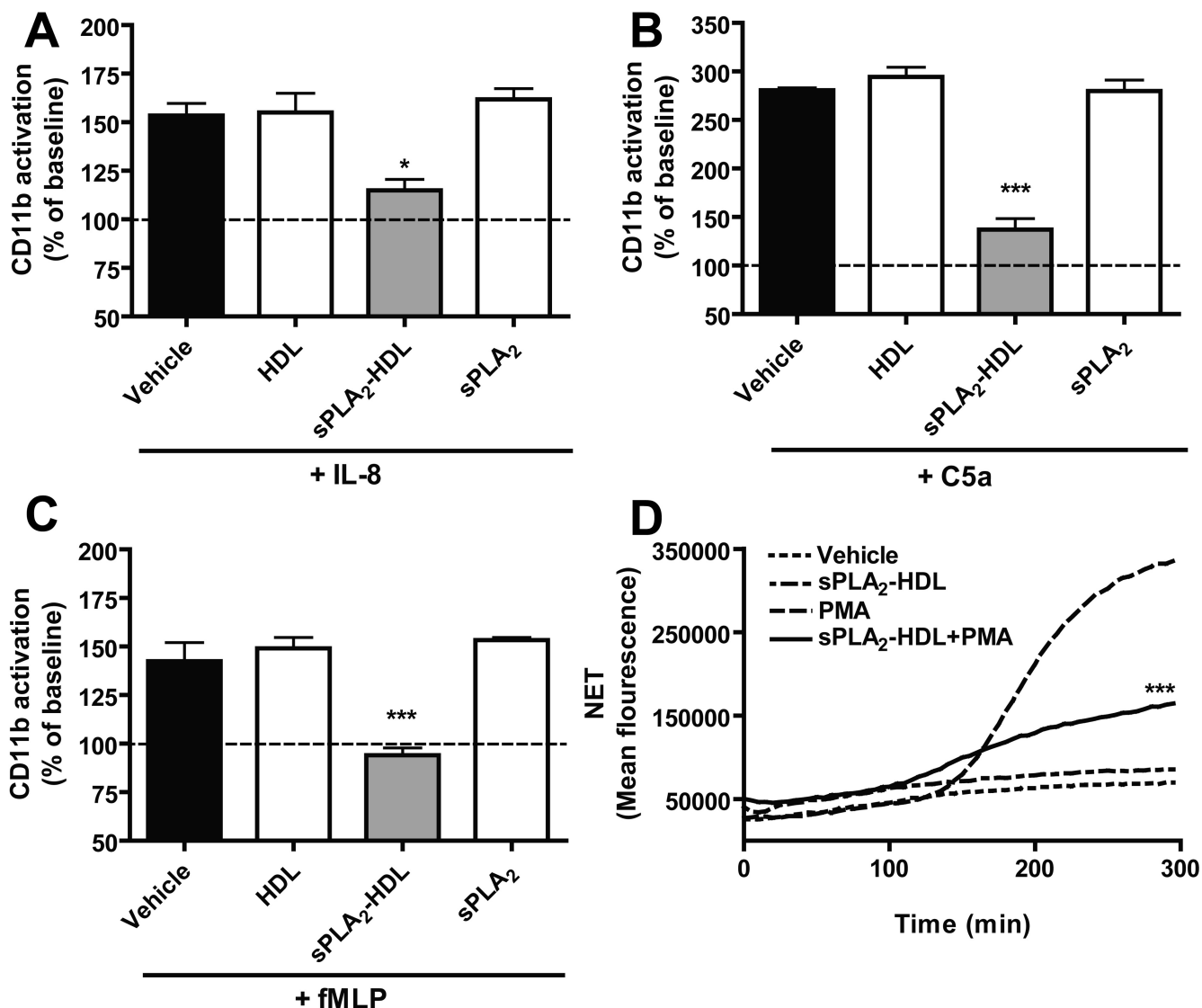


Figure 4. sPLA₂-treated HDL inhibits CD11b activation in neutrophils

Isolated neutrophils were preincubated with vehicle, HDL (100 µg protein/ml), sPLA₂-HDL (100 µg protein/ml) or sPLA₂ alone for 15 min at 37°C in the presence of antibody to the active epitope of CD11b and stimulated with (A) IL-8 (3 nmol/L), (B) fMLP (10 nmol/L) or (C) C5a (30 nmol/L) for 4 min at 37°C. CD11b activation was measured by flow cytometry. Vehicle treated (unstimulated) control was set as baseline and values are expressed as % over baseline. The results are shown as mean ± SD of 4 separate experiments using cells from different healthy donors. Statistical significance was assessed by one-way ANOVA with Tukey multiple-comparison post hoc test. *p < 0.05, **p < 0.01, ***p < 0.001 *versus* vehicle. (D) Isolated neutrophils were treated with vehicle or sPLA₂-HDL (100 µg protein/ml) in the presence or absence of 10 nmol/L PMA and NET formation was measured as an increase in fluorescence of the SYTOX Green dye every 10 min for 5 hours at 37°C. Results show a representative experiment of three independent experiments. Values are

expressed as mean fluorescence. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ versus PMA at 5 hours time point.

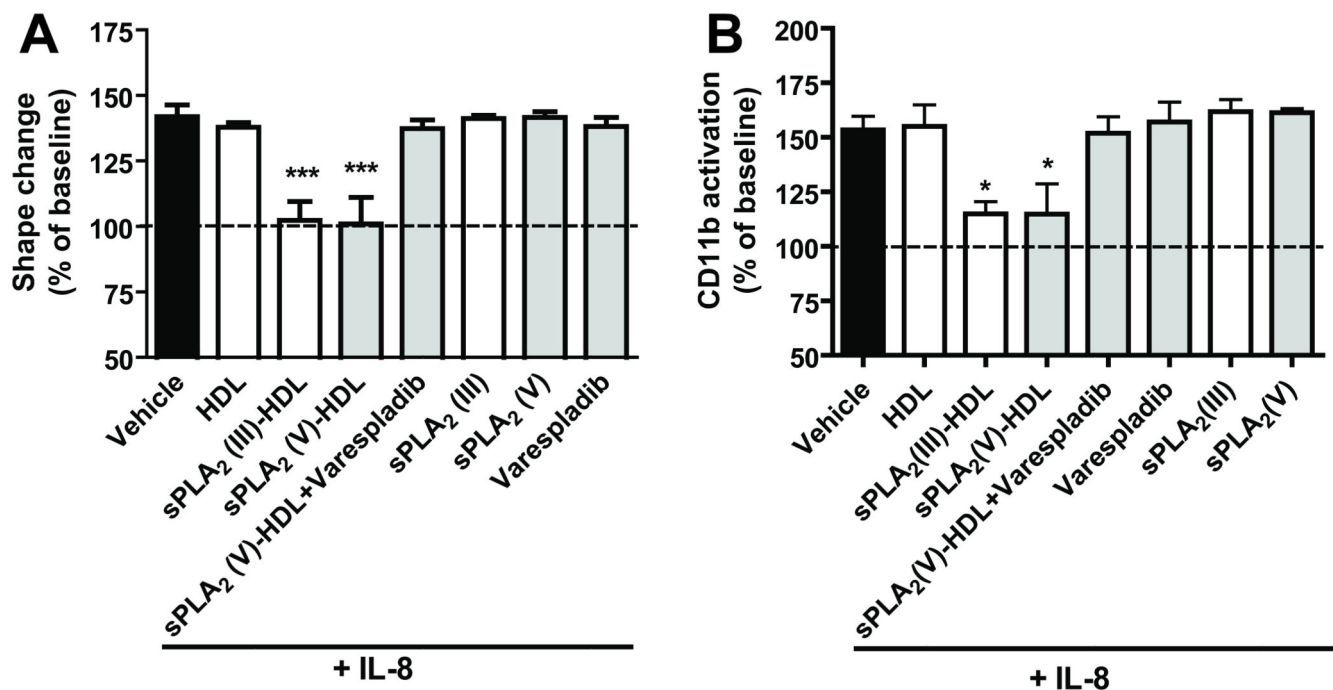


Figure 5. The sPLA₂ inhibitor varespladib inhibits the effects of sPLA₂-treated HDL on neutrophils

HDL was incubated with 200 ng/ml sPLA₂ type III or 400 ng/ml sPLA₂ type V in the presence or absence of varespladib (1 μ mol/L), overnight, at 37°C. These samples were used for neutrophil treatment. Isolated neutrophils were preincubated (15 min at 37°C) with vehicle (assay buffer), HDL (100 μ g protein/ml), sPLA₂ (III)-HDL (100 μ g protein/ml), sPLA₂ (V)-HDL, sPLA₂ (V)-HDL + varespladib, varespladib, sPLA₂ (III) or sPLA₂ (V) alone. Neutrophils were then stimulated with IL-8 (3 nmol/L, 4 min, 37°C) to assess shape change (A) or CD11b activation (B) by flow cytometry. Values are expressed as % of baseline. Results represent means \pm SD (n=3). Statistical significance was assessed by one-way ANOVA with Tukey multiple-comparison post hoc test. *p < 0.05, **p < 0.01, ***p < 0.001 *versus* vehicle.

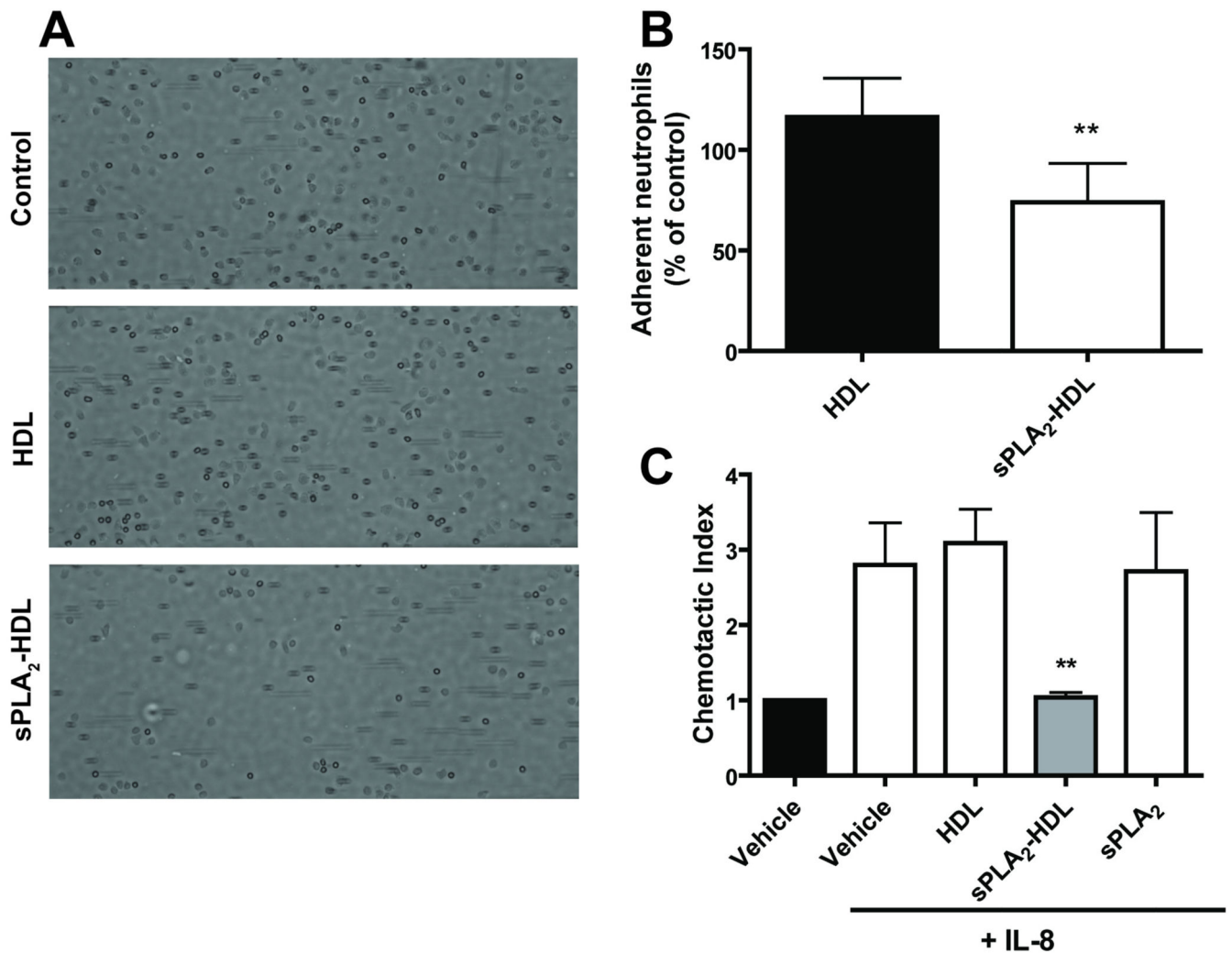


Figure 6. sPLA₂-HDL inhibits neutrophil adhesion under flow conditions and neutrophil migration

Isolated human neutrophils were treated with vehicle (assay buffer), HDL (100 μ g protein/ml) or sPLA₂-HDL (100 μ g protein/ml) for 15 minutes at 37°C before perfusion over ICAM-1 coated microchannels. (A) Representative images were taken 5 minutes after the start of the perfusion with neutrophils. (B) Images were quantified by computerized image analysis. Values are expressed as % of vehicle treated control. Data are shown as mean \pm SD of 6 experiments with neutrophils from different donors. Statistical analysis was performed using Student's t-test. * $p < 0.05$, ** $p < 0.01$ versus HDL. (C) Neutrophils were preincubated with vehicle, HDL (100 μ g protein/ml), sPLA₂-HDL (100 μ g protein/ml) or sPLA₂ for 15 minutes at 37°C and seeded into the upper wells. Cells were allowed to migrate towards IL-8 (10 nmol/L) for 1 hour at 37°C. Cells that had migrated to the lower chamber were enumerated by flow cytometry. The results are shown as mean \pm SD of 3 separate experiments using PMNL from different healthy donors. Statistical significance was assessed by one-way ANOVA with Tukey multiple-comparison post hoc test. * $p < 0.05$, ** $p < 0.01$ versus vehicle.

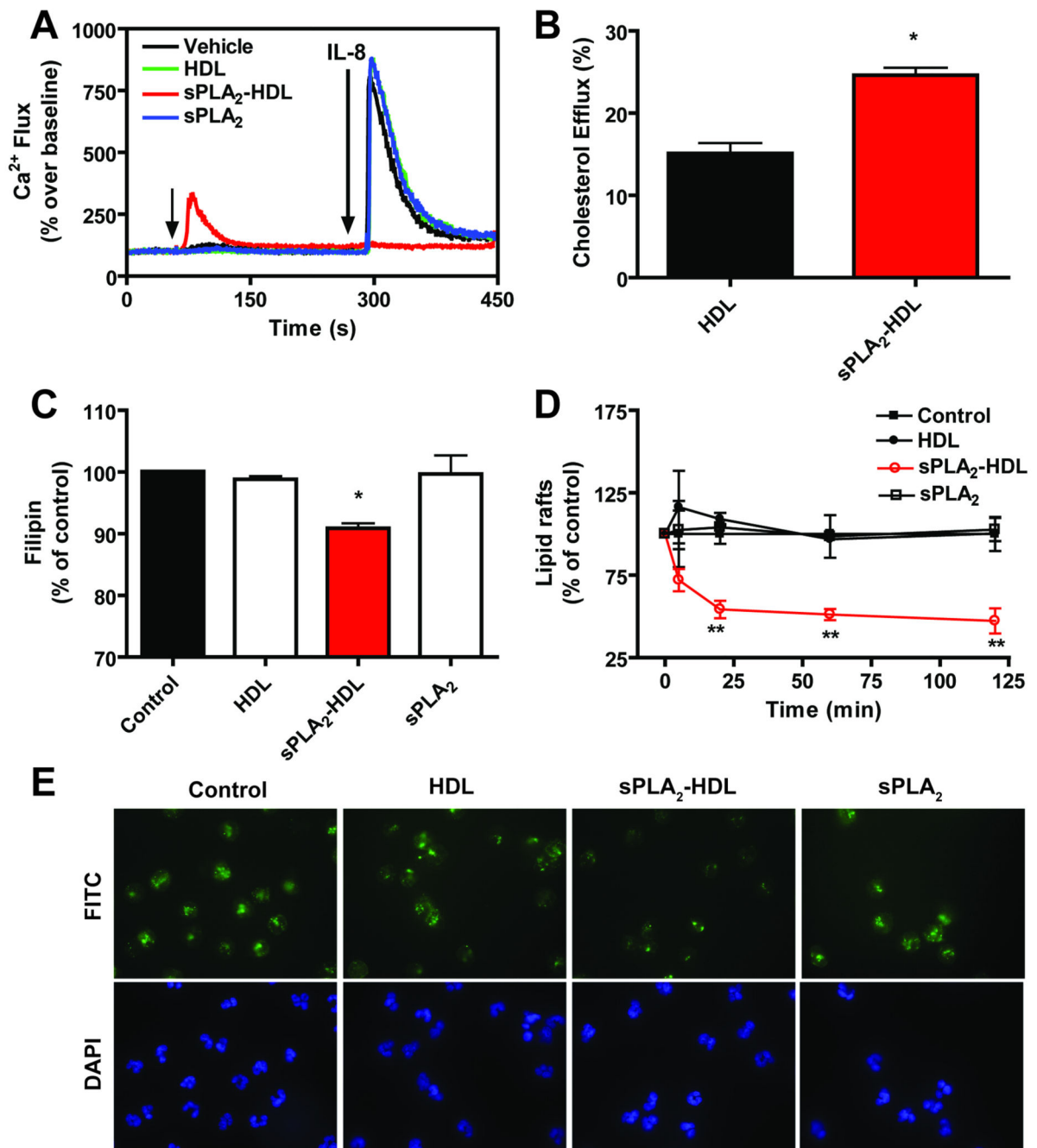


Figure 7. sPLA₂-HDL abolishes agonist-induced Ca²⁺ flux and efficiently removes unesterified cholesterol from neutrophils

(A) Baseline Ca²⁺ levels were measured for 1 min and then neutrophils were treated with vehicle, HDL (100 µg protein/ml), sPLA₂-HDL (100 µg protein/ml) or sPLA₂ for 4 min and Ca²⁺ flux was induced with IL-8 (10 nmol/L). Ca²⁺ flux was detected at room temperature by flow-cytometry as an increase in fluorescence intensity of the Ca²⁺-sensitive dye Fluo-3-AM in the FL-1 channel. Results show a representative of 4 independent experiments done with cells from different donors. Values are expressed as % over baseline. (B) Isolated neutrophils were preloaded with [³H]-cholesterol for 3 hours and incubated with HDL (100

μg protein/ml) or sPLA₂-HDL (100 μg protein/ml protein) for 60 minutes, and percentage of cholesterol efflux was assessed. Cholesterol efflux is expressed as the radioactivity in the medium relative to total radioactivity in medium and cells. The results are shown as mean \pm SD of three separate experiments using PMNL from different healthy donors. Statistical analysis was performed using Student's t-test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ versus HDL. (C) Neutrophils were incubated with vehicle, HDL (100 μg protein/ml), sPLA₂-HDL (100 μg protein/ml), sPLA₂ alone for 30 min at 37°C and stained with Filipin. The amount of unesterified cholesterol was assessed by flow-cytometry. Vehicle treated control was set at 100 % and values are expressed as % of control. Results are shown as mean \pm SD (n=3). Statistical significance was assessed by one-way ANOVA with Tukey multiple-comparison post hoc test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ versus control. (D) Neutrophils were incubated with vehicle, HDL (100 μg protein/ml), sPLA₂-HDL (100 μg protein/ml) or sPLA₂ alone for 5 up to 120 min at 37°C. Lipid raft abundance was determined by assessing FITC-cholera toxin B stained cholesterol-containing microdomains by flow cytometry. Control was set at 100 % and values are expressed as % of vehicle-treated control. Results are shown as mean \pm SD (n=3). Statistical significance was assessed by one-way ANOVA with Tukey multiple-comparison post hoc test. * $p < 0.05$, ** $p < 0.01$ versus control. (E) Fluorescence microscopy analysis of lipid rafts (green staining). DAPI (blue staining) was used to visualize the nuclei. Original magnification X60 was used. Representative images from 3 independent experiments with neutrophils from different donors are shown.