

Research article

DIFFERENCES BETWEEN GROUP X AND GROUP V SECRETORY PHOSPHOLIPASE A₂ IN LIPOLYTIC MODIFICATION OF LIPOPROTEINS

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Abstract: Secretory phospholipases A₂ (sPLA₂s) are a diverse family of low molecular mass enzymes (13-18 kDa) that hydrolyze the *sn*-2 fatty acid ester bond of glycerophospholipids to produce free fatty acids and lysophospholipids. We have previously shown that group X sPLA₂ (sPLA₂-X) had a strong hydrolyzing activity toward phosphatidylcholine in low-density lipoprotein (LDL) linked to the formation of lipid droplets in the cytoplasm of macrophages. Here, we show that group V sPLA₂ (sPLA₂-V) can also cause the lipolysis of LDL, but its action differs remarkably from that of sPLA₂-X in several respects. Although sPLA₂-V released almost the same amount of fatty acids from LDL, it released more linoleic acid and less arachidonic acid than sPLA₂-X. In addition, the requirement of Ca²⁺ for the lipolysis of LDL was about 10-fold higher for sPLA₂-V than sPLA₂-X. In fact, the release of fatty acids from human serum was hardly detectable upon incubation with sPLA₂-V in the presence of sodium citrate, which contrasted with the potent response to sPLA₂-X. Moreover,

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Abbreviations used: Ab – antibody; apoB – apolipoprotein B; BSA – bovine serum albumin; COX – cyclooxygenase; FCS – fetal calf serum; HDL – high density lipoprotein; HPLC – high-performance liquid chromatography; LDL – low density lipoprotein; lysoPC – lysophosphatidylcholine; PBS – phosphate-buffered saline; PC – phosphatidylcholine; PLA₂ – phospholipase A₂; SDS-PAGE – SDS-polyacrylamide gel electrophoresis; sPLA₂ – secretory PLA₂; sPLA₂-IB – group IB sPLA₂; sPLA₂-IIA – group IIA sPLA₂; sPLA₂-V – group V sPLA₂; sPLA₂-X – group X sPLA₂

sPLA₂-X, but not sPLA₂-V, was found to specifically interact with LDL among the serum proteins, as assessed by gel-filtration chromatography as well as sandwich enzyme-immunosorbent assay using anti-sPLA₂-X and anti-apoB antibodies. Surface plasmon resonance studies have revealed that sPLA₂-X can bind to LDL with high-affinity ($K_d = 3.1$ nM) in the presence of Ca²⁺. Selective interaction of sPLA₂-X with LDL might be involved in the efficient hydrolysis of cell surface or intracellular phospholipids during foam cell formation.

Key words: Secretory phospholipase A₂, Low-density lipoprotein, High-density lipoprotein, Phospholipids, Calcium ion

INTRODUCTION

Phospholipase A₂s (PLA₂s) are a diverse family of lipolytic enzymes that hydrolyze the *sn*-2 fatty acid ester bond of glycerophospholipids to produce free fatty acids and lysophospholipids [1, 2]. Over the past three decades, a number of PLA₂s have been identified and classified into different families based on biochemical features and primary structure [3, 4]. Among them, secretory PLA₂s (sPLA₂s) have several characteristics including a low molecular mass (13-18 kDa) and an absolute catalytic requirement for millimolar concentrations of Ca²⁺ [4, 5]. At present, nine different groups of sPLA₂s have been identified in humans (IB, IIA, IID, IIE, IIF, III, V, X and XII) [4, 6-8]. Group IB sPLA₂ (sPLA₂-IB) has been thought to act as a digestive enzyme, given its abundance in digestive organs [9]. Besides a role in lipid digestion, this sPLA₂ has been shown to play a role in cell proliferation, lipid mediator release, acute lung injury, and endotoxic shock through binding to a specific receptor known as the phospholipase A₂ receptor (PLA₂R) [10-13]. Group IIA sPLA₂ (sPLA₂-IIA) is thought to play a pivotal role in the progression of inflammatory conditions, since its local and systemic levels are elevated in numerous inflammatory diseases [14, 15]. It was previously shown that sPLA₂-IIA is expressed in the atherosclerotic arterial intima and is associated with extracellular matrix structures and lipid droplets [16-19]. Group V sPLA₂ (sPLA₂-V) was identified as a homologue of another group II family PLA₂ [20]. However, Bingham *et al.* [21] have unraveled the difference in the subcellular location of these sPLA₂s in bone-marrow-derived mast cells and demonstrated that these enzymes exert different functions and thus are not redundant. It was also reported that exogenously added sPLA₂-V was internalized and localized to perinuclear membranes and induced fatty-acid release leading to leukotriene synthesis [22]. Secretory PLA₂-X has 16 cysteine residues located at positions characteristic of the classical types of sPLA₂-IB and sPLA₂-IIA, and also has an amino acid C-terminal extension that is typical of group II sPLA₂ subtypes [23]. We have shown that sPLA₂-X can induce the release of arachidonic acid leading to cyclooxygenase (COX)-dependent prostaglandin formation, as well as marked production of lysophosphatidylcholine (lysoPC) in various cell types, including macrophages, spleen cells, and colon cancer cells [24-26]. Moreover, we

demonstrated that sPLA₂-X, as well as sPLA₂-IB, is also an endogenous ligand of PLA₂R and the soluble form of the receptor could regulate the biological functions of secretory phospholipase A₂ [27].

Although some of the biological roles of these sPLA₂s are expressed through binding to PLA₂R, most of their functions are exerted by their lipid-digesting property to produce free fatty acids and lysophospholipids. Notably, sPLA₂-V and X were shown to induce the release of fatty acids from various types of cells as expected from their higher specificity for phosphatidylcholine (PC), which is a major lipid component in the outer membrane of cells [28, 29, 30-32]. Moreover, it was reported by us and other groups that PLA₂-V and X efficiently hydrolyze PC, which is a major component in serum lipoproteins, as well as that on cell membrane [28, 33]. For example, we have shown that sPLA₂-X can induce potent lipolysis of LDL, leading to the production of large amounts of unsaturated fatty acids and lysoPC [34]. The sPLA₂-X-modified LDL was efficiently incorporated into macrophages to induce the accumulation of cellular cholesterol ester and the formation of non-membrane-bound lipid droplets in the cytoplasm [34]. Furthermore, sPLA₂-X was found to induce lipolytic modification of HDL linked to the loss of its anti-atherogenic property [35]. Moreover, we have found that sPLA₂-X is expressed markedly in foam cell lesions in the arterial intima of high fat-fed apolipoprotein E-deficient mice. Meanwhile, Gesquiere *et al.* [33] have reported that sPLA₂-V shows strong hydrolyzing activity toward PC and can induce the release of fatty acids from both LDL and HDL more potently than sPLA₂-IIA. These observations strongly suggest that these sPLA₂s play important roles in the development of atherosclerosis through their modification of lipoproteins. However, the difference in enzymatic and biological properties of these sPLA₂s in lipolysis of lipoproteins has not been well characterized.

Therefore, we compared here the potency and characteristics of human sPLA₂-V and PLA₂-X in the hydrolysis of LDL. Although both sPLA₂s efficiently hydrolyzed PC in LDL to the same extent, sPLA₂-X released more arachidonic acid and less linoleic acid than sPLA₂-V. We also found a difference of Ca²⁺ dependency on the hydrolysis of phospholipids between these two sPLA₂s. Finally, we found a specific association of sPLA₂-X, but not sPLA₂-V, with LDL in human serum by means of gel filtration and a specific sandwich enzyme-linked immunosorbent assay (ELISA) system. High-affinity interaction between sPLA₂-X and LDL was also confirmed by analysis with BIAcore instruments.

MATERIALS AND METHODS

Materials

Purified recombinant human sPLA₂-IB, sPLA₂-X and sPLA₂-X-HisTag proteins were prepared as described previously [28, 25]. The recombinant human sPLA₂-V protein was also prepared as described previously [23]. Recombinant human sPLA₂-IIA was generously provided by Dr. Ruth Kramer (Eli Lilly, USA).

Mouse anti-human sPLA₂-X was prepared by the standard method. Rabbit anti-human sPLA₂-X antibody (Ab) was prepared as described previously [28], and anti-sPLA₂-IIA Ab was purchased from Cayman Chemicals (Ann Arbor, MI). Bovine serum albumin (BSA) and other chemicals were purchased from Sigma (St. Louis, MO). The sensor chip NTA was purchased from BIAcore AB (Uppsala, Sweden).

Preparation of human LDL and HDL

Very low-density lipoprotein (VLDL; density less than 1.006 g/ml), LDL (d = 1.019-1.063 g/ml), and high-density lipoprotein (HDL; d = 1.085-1.210 g/ml) were isolated from plasma of healthy and fasting donors by sequential ultracentrifugation, as described previously [36]. LDL and HDL were used as a substrate from a plasma pool. This study was approved by our institutional review committee and the procedures followed were in accordance with our institutional guidelines. All subjects gave informed consent with this study.

Measurement of released fatty acids, PC and lysoPC in sPLA₂-treated lipoproteins

Human LDL (1 mg/ml) was pre-incubated for 10 min at 37°C, and stimulated with various concentrations of sPLA₂ enzymes in a final volume of 40 µl. The reaction was stopped by the addition of 160 µl of Dole's reagent, and the released fatty acids were extracted, labeled with 9-anthryldiazomethane (Funakoshi, Japan), and then analyzed by reverse-phase high-performance liquid chromatography (HPLC) on a LiChroCART 125-4 Superspher 100 RP-18 column (Merck), as described previously [36, 37]. In some experiments, total amounts of released fatty acids were measured using NEFA C-test Wako (Wako, Japan). For measurement of the amounts of PC and lysoPC in LDL, lipids were extracted with organic solvent as described previously [23].

Gel-filtration analysis of human serum modified with sPLA₂s by FPLC

For the modification of human serum (or plasma) with sPLA₂s, human serum was incubated with 50 nM sPLA₂-IB, IIA, V or X at 37°C in a buffer composed of 12.5 mM Tris-HCl (pH 8.0), 0.25 M NaCl and 0.0125% BSA. The reaction was stopped by addition of EDTA at a final concentration of 5 mM. The sPLA₂ modified human serum was analyzed with Superose HR 6 10/30 (Amersham-Pharmacia, Uppsala, Sweden) by fast protein liquid chromatography (FPLC). A degassed and filtered solution of 0.15 M sodium chloride, 0.3 mM disodium-EDTA and 3.1 mM sodium azide, pH 7.3, was used for pre-equilibration of the columns. Samples were chromatographed with the same solution at a flow rate of 0.5 mL/min, and fractions of 500 µl each were collected. Individual fractions were assayed for PLA₂ activity and immunoblotting as described below.

Immunoblotting analysis

Each fraction was separated with sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) using either a 4-20% gradient gel (Daiichi

Chemical Co., Ltd.) for detection of Apolipoprotein B-100 (ApoB-100) or a 15-25% gradient gel for detection of sPLA₂s. The separated proteins were electroblotted onto nitrocellulose membranes (Atto, Japan) using a semidry blotter (Atto, Japan), according to the manufacturer's instructions. The membranes were probed with the relevant antibodies and visualized using a chemiluminescent detection reagent (ECL Western blotting detection reagents, Amersham Pharmacia Biotech) according to the manufacturer's instructions and analyzed using a Fluor-S MAX MultiImager (Bio-Rad).

PLA₂ activity assay by chromogenic assay

Secretory PLA₂ activity was measured using diheptanoyl thio-PC as a substrate according to the method of Reynold *et al.* [38]. Briefly, mixed micelles consisting of 1 mM diheptanoyl thio-PC and 0.3 mM Triton X-100 were used as a substrate. The assay mixture contained 25 mM Tris-HCl buffer (pH 7.5), 0.12 mM DTNB, 10 mM CaCl₂, 0.1M KCl, and 1 mg/ml BSA. The reactions were initiated by addition of each gel filtration fraction to the assay mixture containing the substrate in a final volume of 200 μ l. The reaction was monitored at an absorbance wavelength of 405 nm with a microplate reader.

PLA₂ activity assay by the PG/Chol method

Phospholipase A₂ activity in each gel filtration fraction was measured by the method of Tojo *et al.* [39] with some modifications. Briefly, the substrate used was the mixed micelles of 1 mM 1-palmitoyl-2-oleyl-sn-glycero-3-phosphoglycerol (Avanti Polar Lipid, Inc., AL) and 2 mM sodium cholate, and each gel filtration fraction was incubated at 40°C for 30 min in 100 mM Tris-HCl buffer (pH 8.0) containing 150 mM NaCl, 10 mM CaCl₂, 1 mg/ml fatty acid-free BSA (Sigma) and substrate (final volume of 100 μ l). The reaction was almost linear during this incubation. The enzymatic reaction was stopped by the addition of 400 μ l of Dole's reagent (2-propanol : heptane : 2N H₂SO₄ = 40:10:1 (v/v/v)), and 6 nmol of margaric acid (Nu-Chek Prep, Inc., Elysian, MN) was added as an internal standard. Fatty acids were extracted by a modified Dole's extraction procedure [40]. The heptane layer containing fatty acids was dried *in vacuo* and the residue was dissolved in 50~100 μ l of 0.05% anthryldiazomethane (Funakoshi Co., Tokyo, Japan) in 10% ethyl acetate and 90% methanol followed by incubation at room temperature for 15 min. An aliquot (10~20 μ l) of the sample was injected into a reverse-phase HPLC system. A LiChroCART 75-4 Superspher 60 RP-8 column was employed (Merck, Darmstadt, Germany). Sample components were eluted isocratically at a flow rate of 1.1 ml/min with a mobile phase of acetonitrile-water (93:7, v/v) and UV absorbance was monitored at a wavelength of 254 nm. The column oven temperature was 26°C.

Detection of sPLA₂-X/LDL complexes with sandwich ELISA

ELISA plates (NUNK MAXISORP #442404) were coated with 100 μ l/well of anti-human sPLA₂-X IgG dissolved in Tris-buffered saline (TBS) (10 mM Tris-HCl, pH 7.4, 150 mM NaCl). The plates were incubated for 18 h at 4°C and then

blocked with 25% Block Ace (Dainippon Pharmaceutical) in H₂O for 2 h at room temperature. The wells were washed with TBS containing 2 mM CaCl₂ or 2 mM EDTA and incubated with samples diluted with 1% BSA/TBS containing 2 mM CaCl₂ or 2 mM EDTA. The plate was washed as above, followed by incubation with a 1:1000 dilution of peroxidase-conjugated anti-human ApoB antibody (The Binding Site). After washing, the plate was developed with 100 µl/well of tetramethylbenzidine, and the reaction was stopped by adding 100 µl/well of 1N H₂SO₄. Absorbance was read at 450 nm.

Surface plasmon resonance (SPR) analysis

The binding of sPLA₂-X to LDL and HDL was analyzed on a Biacore 3000 surface plasmon resonance system (Uppsala, Sweden) using an NTA chip. All the binding experiments were carried out at 25°C and a flow rate of 20 µl/min except nickel loading (5 µl/min). SPR buffers and solutions were as follows: eluent buffer: 10 mM HEPES, 0.15 M NaCl, 0.05 mM EDTA, 0.005% Tween-20 pH 7.4; nickel solution: 0.5 mM NiCl₂ in eluent buffer; regeneration solution: 10 mM HEPES, 0.15 M NaCl, 0.35 M EDTA, 0.005% Tween-20 pH 8.3. Secretory PLA₂-X-HisTag and lipoproteins were diluted with eluent buffer. If the analyses were performed in the presence of CaCl₂, all the buffers contained 1 mM CaCl₂ other than the nickel solution. The NTA sensor chip was loaded with 20 µl of the nickel solution to saturate the NTA surface with Ni²⁺ after washing with the regeneration solution followed by eluent buffer. Forty microliters of 200 nM sPLA₂-X-HisTag was injected to immobilize the ligand. Subsequently, 50 µl of each lipoprotein (LDL or HDL) at 0.2 µM was injected to analyze the interaction between sPLA₂-X and lipoproteins, and the sensorgram was allowed to run for up to 300 s, following which the chip was regenerated. For kinetic analysis, the analyte concentration of LDL were 10, 25, 50, 100, and 125 µg/ml. The kinetic parameters for lipoproteins were estimated by BIAevaluation 3.0 software (Biacore AB) using A + B = AB (kinetic 1:1 binding with drifting baseline) by separate fitting to fit the data.

RESULTS

Hydrolysis of the phospholipids in LDL by sPLA₂-V and sPLA₂-X

We have demonstrated that sPLA₂-X markedly induced the lipolysis of LDL [34] and another group reported that sPLA₂-V caused the lipolysis of both LDL and HDL [33]. As shown in Fig. 1A, a large amount of free fatty acid was dose-dependently released upon the incubation of LDL with both sPLA₂-V and sPLA₂-X, while little was released on the incubation of LDL with sPLA₂-IB and IIA. Total amounts of fatty acids released by sPLA₂-V were comparable to those released by sPLA₂-X at the same time point (Fig. 1B). To dissect the profiles of lipolysis by both sPLA₂-V and sPLA₂-X, we first examined the phosphatidylcholine (PC) content in LDL after treatment since both sPLA₂s have been shown to have higher specificity to hydrolyze PC, which is a major component of phospholipids in LDL [34, 33]. PC content decreased with time after treatment

with 50 nM of both sPLA₂s, to an equal extent (data not shown). Corresponding to this reduction, the production of lysoPC was detected in LDL at 3 h and increased up to 24 h after both treatments (data not shown). We next determined the composition of free fatty acids (FFA) released from LDL with the respective sPLA₂s at a concentration of 50 nM. As shown in Fig. 1C, sPLA₂-X elicited the marked release of various types of unsaturated fatty acids from human LDL in the following order: linoleic acid (C18:2) > arachidonic acid (C20:4) > oleic acid (C18:1) > docosahexaenoic acid (C22:6) > linolenic acid (C18:3), whereas little palmitic acid (C16:0), a saturated fatty acid, was observed at 3 h (Fig. 1C) though slightly more was released at 24 h (data not shown). The amount of fatty acid increased time-dependently. Secretory PLA₂-V released various types of unsaturated fatty acids from human LDL in the following order: linoleic acid (C18:2) > oleic acid (C18:1) > arachidonic acid (C20:4) = docosahexaenoic acid (C22:6) > linolenic acid (C18:3) (Fig. 1C). Again there was little release of saturated fatty acids such as palmitic acid (C16:0) from LDL after the sPLA₂-V treatment. Moreover, much less arachidonic acid than linoleic acid was released from LDL by sPLA₂-V, and the amount of arachidonic acid was only about 30% of that in

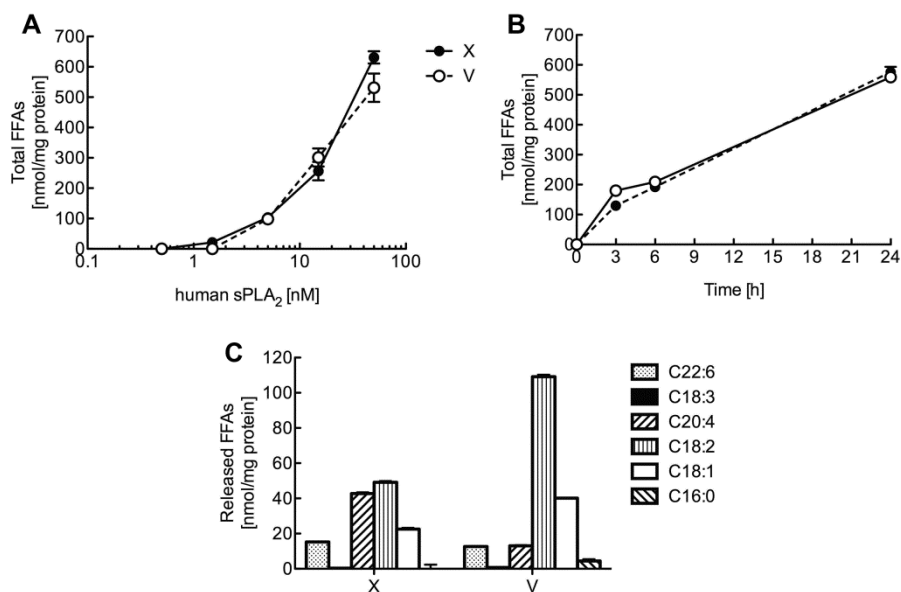


Fig. 1. Fatty acids released from LDL by human sPLA₂s. A – LDL (1 mg/ml) was incubated with the indicated concentration of purified human sPLA₂s at 37°C for 24 hours. B – LDL (1 mg/ml) was incubated with 50 nM purified human sPLA₂s at 37°C for the period indicated, C – LDL (1 mg/ml) was incubated with 50 nM purified human sPLA₂-V or X for 3 hours at 37°C. The released fatty acids were quantified as described in Materials and Methods. Each point represents the mean ± S.D. of triplicate measurements. The data are representative of three experiments.

the case of sPLA₂-X (Fig. 1C). Secretory PLA₂-V was shown to release about 7-fold more linoleic acid than arachidonic acid from LDL (Fig. 1C), which was consistent with its fatty acid specificity using a synthetic substrate [41]. In contrast, sPLA₂-X induced marked release of arachidonic acid from LDL with almost the same level of linoleic acid (Fig. 1C). No dose-dependent difference in the released fatty acid profile was observed on treatment with the two sPLA₂s (data not shown). Also, longer incubation increased the amount released, but did not change the profile (data not shown). These findings showed that although both sPLA₂s release unsaturated fatty acids from LDL effectively, there is an obvious difference in the fatty acids they prefer, especially arachidonic acid.

Release of fatty acids from whole human serum and plasma by sPLA₂-V and X

Next we investigated the effect of both sPLA₂s on whole serum or plasma prepared using sodium citrate to clarify the differences between the lipolysis by these sPLA₂s under physiological conditions. Both whole human serum and plasma were incubated with either sPLA₂-V or sPLA₂-X at 37°C for 24 h. A large amount of fatty acid was released from human serum and plasma on treatment with sPLA₂-X (Fig. 2). By contrast, much less fatty acid was released by sPLA₂-V from human plasma than from human serum (Fig. 2). Whole human serum contained a sufficient (>1 mM) concentration of calcium cations (Ca²⁺), while human plasma prepared using sodium citrate (0.38%) buffer contained much less calcium due to quenching. Then human serum containing the same concentration of sodium citrate was prepared and incubated with both sPLA₂s and the release of fatty acids was determined. The amount released by sPLA₂-X was almost the same in human plasma and human serum with or without sodium citrate. However, sodium citrate decreased the amount of fatty acid released by sPLA₂-V from human serum to near that released from plasma.

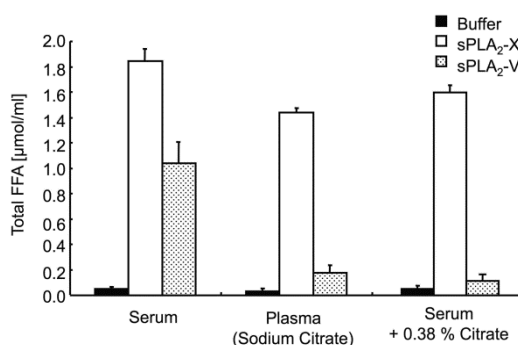


Fig. 2. The release of fatty acids from whole human serum and plasma by sPLA₂-V or sPLA₂-X. Whole human serum and plasma were incubated with 50 nM purified human sPLA₂-V or sPLA₂-X for 24 hours at 37°C. The total amount of fatty acids released was quantified as described in Materials and Methods. In the case of human serum, incubation was performed in the absence and presence of 0.38% sodium citrate. Each point represents the mean ± S.D. of triplicate measurements. The data are representative of three experiments.

Difference of calcium dependency between sPLA₂-X and sPLA₂-V

The difference in the release of fatty acids by these two sPLA₂s in Fig. 2 was mostly caused by the concentration of Ca²⁺ within reaction mixtures and implied a difference in Ca²⁺ dependency between these two sPLA₂s. Therefore, we examined the PLA₂ activity using PLPC (1-palmitoyl, 2-linoleoyl-phosphatidyl choline) liposomes and LDL in various Ca²⁺ concentrations. Under our experimental conditions, these sPLA₂s showed almost the same maximum. As shown in Fig. 3A, the activity of sPLA₂-X to lyse PLPC reached a maximum at 1 mM and was still about 20% at 0.03 mM. The activity of sPLA₂-V did not reach a plateau even at 10 mM of CaCl₂, and was slight at less than 0.1 mM. The difference in PLA₂ activity between them was more remarkable for LDL hydrolysis (Fig. 3B). After incubation at 37°C for 24 h, sPLA₂-X displayed about 40% activity even at 0.1 mM and reached maximum activity at 1 mM. Even if CaCl₂ was not contained in the reaction mixture, sPLA₂-X still exhibited residual PLA₂ activity compared to that in the presence of EDTA (data not shown). In contrast, sPLA₂-V hardly showed any PLA₂ activity at low concentrations though sPLA₂-V activity was increased to about 60% at 1 mM and sPLA₂-V did not reach a plateau even at more than 1 mM. Accordingly, sPLA₂-V requires about a 10-fold concentration of calcium cation compared to sPLA₂-X, and sPLA₂-X is capable of acting at low concentration of calcium.

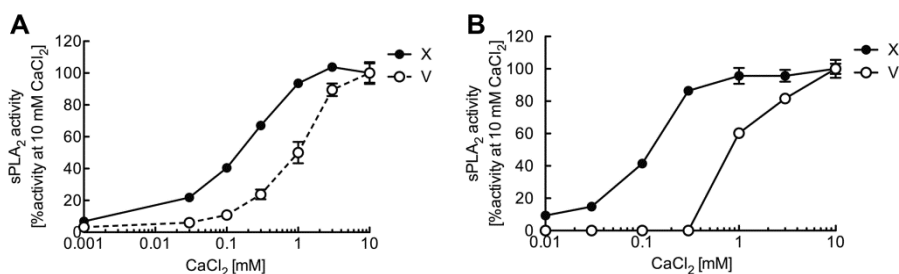


Fig. 3. The difference of calcium dependency between sPLA₂-V and X for PLA₂ activity. PLPC (A) and LDL (B) were incubated with sPLA₂-V or X (50 nM) at 40°C for 30 min (A) and 37°C for 24 hours (B) respectively under various concentrations of CaCl₂. Enzymatic activity was respectively measured as described in Materials and Methods. Each point represents the mean \pm S.D. of triplicate measurements. The data are representative of three experiments.

Selective association of sPLA₂-X with LDL in human serum

Under physiological conditions such as in vascular vessels, concentration of calcium cations in the extracellular space is considered as more than 1 mM. Thus, it is unknown which sPLA₂ hydrolyzes lipoproteins *in vivo*. It has been shown that some enzymes such as platelet activating factor acetylhydrolase bind to lipoproteins [42-44]. In our study we tried to determine the interactions between lipoproteins and sPLA₂s. To investigate whether sPLA₂s associate with lipoproteins in physiological conditions, they were incubated with human serum

at a physiological temperature. After 24 h of incubation, each reaction mixture was subjected to a gel filtration analysis. As shown in Fig. 4A, for the reaction mixture of human serum/sPLA₂-X, PLA₂ activity was concentrated between fraction 17 and 25 where authentic purified LDL also eluted. In contrast, the same fractions chromatographed from human serum incubated with either sPLA₂-IB or IIA did not show any PLA₂ activity (Fig. 4A). As shown in Fig. 4A, sPLA₂-IIA as well as sPLA₂-IB was eluted in the fraction corresponding to low molecular weight proteins including serum albumin. Activity of sPLA₂-V ranged broadly from the LDL fraction to the HDL fraction. Also, the peak activity of sPLA₂-V was not consistent with the LDL fraction. Further, the existence of sPLA₂-X in the LDL fraction was confirmed by immunoblotting using antibody

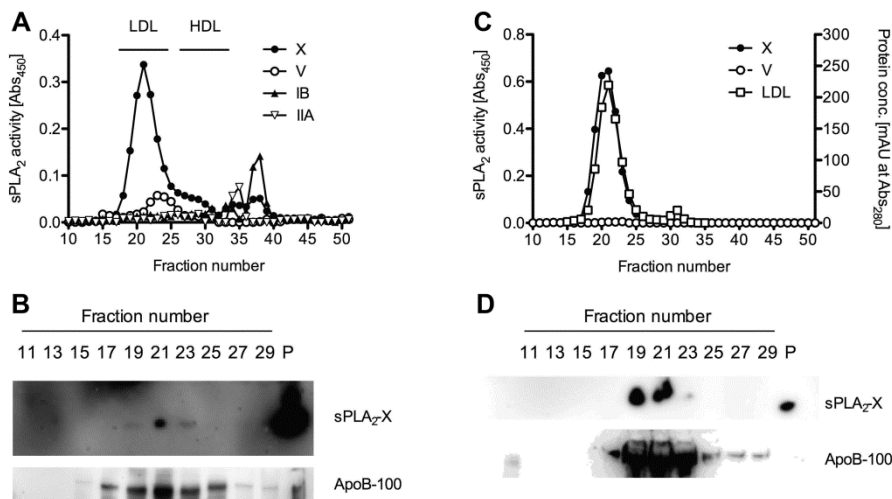


Fig. 4. Human sPLA₂-X selectively binds to LDL on incubation with human serum. Human serum (A, B) or 1 mg/ml isolated LDL (C, D) was incubated with various types of 50 nM purified human sPLA₂ at 37°C for 24 hours. After 24-h incubation, the reaction mixture was separated by gel filtration and subsequently each fraction was subjected to a PLA₂ activity assay (A, C) and western blotting (B, D) using anti-ApoB-100 antibody or anti-sPLA₂-X antibody as described in Materials and Methods. The protein concentration of each fraction was monitored by Abs₂₈₀ with Pharmacia FPLC (C). The data are representative of three experiments.

against sPLA₂-X and apoB protein, which is a major component of LDL particles (Fig. 4B). Immunoblotting also showed that no other isoforms of sPLA₂s including sPLA₂-IIA were detected in the LDL fraction (data not shown). Moreover, these results were ascertained in the case of human plasma (data not shown). Next, we examined the interaction between sPLA₂s and the isolated LDL. After 24-h incubation of LDL with either sPLA₂-X or -V each reaction mixture was subjected to gel filtration analysis. As shown in Fig. 4CD, PLA₂ activity of sPLA₂-X, not sPLA₂-V, was detected in the LDL fraction,

determined by protein concentration. Western blotting by using anti-sPLA₂-X and anti-apoB antibodies also revealed the coexistence of sPLA₂-X with LDL. Taken together, these results indicated that among these sPLA₂s, only sPLA₂-X selectively binds to LDL in human blood. To confirm that only sPLA₂-X binds specifically to LDL, purified LDL was incubated with the sPLA₂s and chromatographed and analyzed as above. PLA₂ activity was detected in the LDL fraction from the reaction mixture incubated with sPLA₂-X but not other subtypes including sPLA₂-IB, IIA, and V (data not shown). Further, it was confirmed that sPLA₂-X existed in the LDL fraction by immunoblotting (data not shown). Secretory PLA₂-X from the reaction without LDL was chromatographed around fraction 35 where low molecular weight proteins were eluted (data not shown). Therefore sPLA₂-X, but not other sPLA₂s, specifically binds to LDL retaining its enzymatic activity.

Detection of sPLA₂-X/LDL complexes with sandwich ELISA

To confirm directly that sPLA₂-X associated with LDL to form sPLA₂-X/LDL complexes, we set up a sandwich ELISA system. We immobilized anti-human sPLA₂-X antibodies on the plate to trap sPLA₂-X/LDL complexes, and detected the trapped complexes by recognition of apolipoprotein B (ApoB) on the surface of LDL particles with peroxidase-conjugated anti-ApoB antibody. In the presence of CaCl₂, we detected strong signals in LDL treated with sPLA₂-X by sandwich ELISA (Fig. 5A). In contrast, in native LDL alone or in sPLA₂-X alone, we did not detect any signals (data not shown). These results show that we could specifically detect sPLA₂-X/LDL complexes by sandwich ELISA.

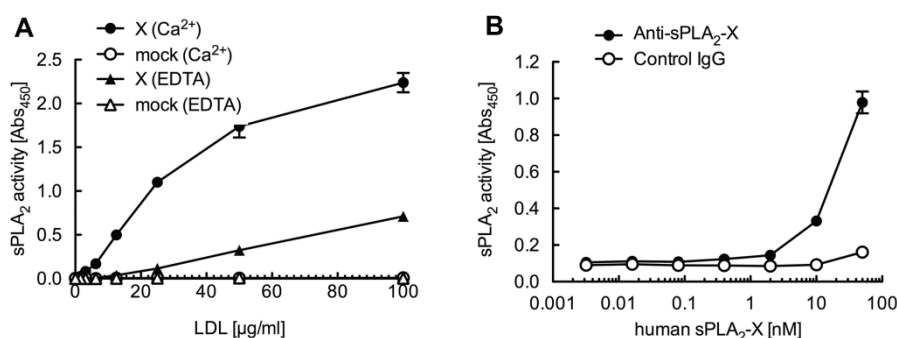


Fig. 5. Detection of the binding between sPLA₂-X and LDL by sandwich ELISA system. A – LDL (1 mg/ml) was incubated with (sPLA₂-X) or without (mock) 50 nM human sPLA₂-X in the presence of 1 mM CaCl₂ (Ca²⁺) or 5 mM EDTA (EDTA) for 24 hours at 37°C and diluted to various concentrations as described in “Experimental Procedures”. The binding between sPLA₂-X and LDL was detected by sandwich ELISA with anti-human sPLA₂-X IgG and peroxidase-conjugated-anti-human ApoB antibody. B – Human serum was incubated with various concentrations of human sPLA₂-X for 24 hours at 37°C and diluted 5 times with 1% BSA/TBS containing 2 mM CaCl₂. The binding between PLA₂-X and LDL was detected by sandwich ELISA. Each point represents the mean ± S.D. of triplicate measurements. The data are representative of three experiments.

However, in the absence of Ca^{2+} condition during both the pre-incubation and ELISA procedure, we detected weak but significant signals in LDL incubated with sPLA₂-X (Fig. 5A). This result suggests that sPLA₂-X associated weakly with LDL under Ca^{2+} -free conditions, and Ca^{2+} acted to strengthen this association. Furthermore, we treated human serum with various concentrations of sPLA₂-X, and tried to detect the binding of sPLA₂-X to LDL in serum. As shown in Fig. 5B, the binding of sPLA₂-X to LDL was found dose-dependently with an increasing amount of sPLA₂-X, and the binding of sPLA₂-X to LDL could be detected even at 2 nM sPLA₂-X. This result shows that the binding of sPLA₂-X to LDL is found in human serum, and that when endogenous sPLA₂-X exists in the bloodstream, sPLA₂-X could exist in the LDL-bound state.

Kinetic analysis of the interaction between sPLA₂-X and LDL

To investigate the fashion in which sPLA₂-X and LDL associate, a SPR analysis was performed. A Recombinant sPLA₂-X-His Tag protein was immobilized to the nickel ion-activated NTA sensor chip, and LDL at various concentrations was applied as described in the materials and methods. In the absence of CaCl_2 , the sensorgram of LDL was increased but rapidly decreased to baseline soon after the injection of LDL (Fig. 6A). This suggested that LDL could be associated with but easily dissociated from immobilized sPLA₂-X-HisTag protein. In contrast, in the presence of 1 mM CaCl_2 , the sensorgram of LDL increased but decreased very slowly after the injection of LDL (Fig. 6B). The sensorgrams of HDL using as a negative control of the binding to sPLA₂-X were hardly increased and decreased to baseline and were not changed regardless of either the absence or presence of CaCl_2 (data not shown). Further, the association and dissociation constants were calculated by the BIAevaluation 3.5 program (Table 1). The apparent

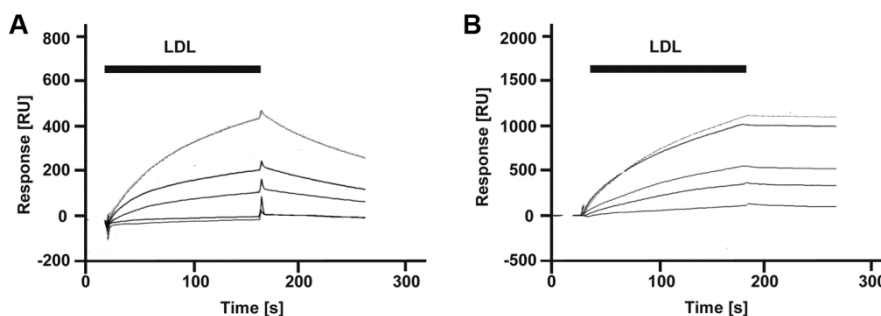


Fig. 6. Study of interaction between human sPLA₂-X and LDL by surface plasmon resonance (SPR) analysis. Recombinant human sPLA₂-X-HisTag at 200 nM was immobilized to a nickel-activated NTA sensor chip, and then 10, 25, 50, 100 and 125 µg/ml of LDL was respectively injected in the absence (A) and presence (B) of 1 mM CaCl_2 without control. Nonspecific interaction was excluded by subtracting the sensorgram of the control cell (without immobilized sPLA₂-X). Thick lines show the injection periods of LDL. The data are representative of three independent experiments.

dissociation constant of LDL for sPLA₂-X was about 1.10×10^{-7} M in the absence of CaCl₂, but became two- orders of magnitude lower in the presence of CaCl₂ ($K_d = 3.10 \times 10^{-9}$ M), following tight binding. The association constant of LDL was not significantly changed in the the presence of CaCl₂. These data indicated that Ca²⁺ plays an important role in the binding of LDL to sPLA₂-X.

Table 1 . Kinetic binding constant of interaction between sPLA₂-X and lipoprotein.

Lipoprotein	CaCl ₂ (mM)	k _a (1/Ms)	k _d (1/s)	K _d (M)
LDL	0	$8.25 \pm 2.41 \times 10^4$	$8.86 \ 86 \pm 5.98 \times 10^{-3}$	$1.10 \pm 0.62 \times 10^{-7}$
	1	$6.54 \pm 3.44 \times 10^4$	$0.24 \pm 0.23 \times 10^{-3}$	$0.031 \pm 0.017 \times 10^{-7}$

The data shows the average \pm SD of calculated value from each analyte concentration.

DISCUSSION

The present study demonstrated differences between sPLA₂-V and sPLA₂-X in substrate specificity in the lipolysis of lipoprotein, Ca²⁺ dependency on PLA₂ activity, and specificity to bind LDL. We have previously shown that sPLA₂-X can induce the release of arachidonic acid leading to cyclooxygenase (COX)-dependent prostaglandin formation, as well as marked production of lysophosphatidylcholine (lyso-PC) in various cell types, including macrophages, spleen cells, and colon cancer cells [24-26]. Also, we have recently shown that sPLA₂-X was expressed in various pathogenic states including atherogenic plaques in some animal models of atherosclerosis [34]. In this study, both sPLA₂-X and sPLA₂-V induced lipolytic modifications of LDL to the same extent. However, sPLA₂-X released arachidonic acid from LDL more effectively than did sPLA₂-V (Fig. 1). This result indicated that the substrate specificity of both sPLA₂s was preserved even on LDL, as found previously with synthetic phospholipids as substrates [45]. Eicosanoid production following the arachidonic acid release plays pivotal roles in various pathological conditions such as inflammation and atherosclerosis [28, 46, 47]. Therefore, sPLA₂-X might play a more important role than sPLA₂-V by inducing the production of proinflammatory lipid mediators, such as prostaglandins (PGs) and leukotrienes (LTs). When whole human serum and plasma were treated with sPLA₂-V and sPLA₂-X, the amount of fatty acids released by sPLA₂-X from whole human plasma was 10-fold higher than that by sPLA₂-V. Generally, human plasma is prepared with sodium citrate or EDTA, which is a chelating reagent of divalent cations, especially Ca²⁺, and therefore the concentration of Ca²⁺ in the reaction mixture using plasma is very low. In accordance with these results, activation of sPLA₂-V required a higher concentration of Ca²⁺ than sPLA₂-X for the lipolysis of synthetic substrates and LDL. Secretory PLA₂-X can be activated at 0.03 mM up to about 60%, while sPLA₂-V remained inactivated at similar concentrations. In previous reports [34, 33], the concentration of Ca²⁺ in the reaction mixture was mostly more than 1 mM, which is possibly why no distinct difference in

Ca²⁺ dependency between these sPLA₂s was observed. This clarified difference in Ca²⁺ dependency for enzymatic activity between sPLA₂-V and sPLA₂-X could be important for considering their biological roles *in vivo*. In this study, Ca²⁺ played an important role in the interaction between sPLA₂-X and LDL *in vitro*. We showed that Ca²⁺ facilitated the binding of sPLA₂-X to the LDL. Surface plasmon resonance (SPR) analysis revealed that sPLA₂-X immediately associated with LDL and gradually dissociated in the absence of CaCl₂ (Fig. 6). The K_d value was 1.1x10⁻⁷ M. In the presence of CaCl₂, the apparent association rate of sPLA₂-X for LDL was almost the same as that in the absence of CaCl₂, while the apparent dissociation rate remarkably decreased. Thus, the binding of sPLA₂-X to LDL became more rigid in the presence than the absence of CaCl₂, and the K_d value was about 3.1x10⁻⁹ (an increase of more than 30-fold). Generally, sPLA₂s are considered to change their conformation as calcium cations bind to their Ca²⁺ loop. Therefore calcium-dependent conformational change of sPLA₂-X may lead to tight binding to LDL.

It was previously reported that some kinds of phospholipase bind to serum proteins [30, 31]. For instance, sPLA₂-IIA bound to human factor Xa and that interaction was largely dependent on the basic properties of sPLA₂-IIA [30]. Another study has shown that human sPLA₂-IIA was selectively associated with HDL by a gel-filtration analysis of serum in transgenic mice overexpressing human sPLA₂-IIA and apoA-I [48]. From the results in Fig. 4, however, sPLA₂-IIA was detected not in the HDL fraction but in the lower molecular weight fraction. To identify the difference in eluted fractions among sPLA₂s, we measured PLA₂ activity for each fraction from each enzyme reaction mixture by another method (PG/Chol method). Then sPLA₂s without sPLA₂-IIA were found to have the same profiles of PLA₂ activity in Fig. 4A, but sPLA₂-IIA was mainly detected in the HDL fraction (data not shown). Thus sPLA₂-IIA probably associated with HDL in our experiments. Our results show that sPLA₂-IB and IIA were not bound to LDL in human whole serum (Fig. 4) and isolated LDL (data not shown). As for sPLA₂-V, broad PLA₂ activity was found from the LDL fraction to the HDL fraction (Fig. 4A). Although sPLA₂-V was not detected by western blotting using commercially available anti-human sPLA₂-V antibody, we found that sPLA₂-V did not associate with the purified LDL by measuring PLA₂ activity (Fig. 4CD). These findings indicated that only sPLA₂-X was selectively associated with LDL in our experimental conditions (Fig. 4CD). The specificity with which sPLA₂-X bound to LDL was very different from that of sPLA₂-V. Meanwhile, it has been shown that LDL binds to various proteins other than membrane receptors [49-53]. Among them, lipoprotein lipase (LPL) is well characterized [49, 54]. Each lipoprotein is composed of specific apoproteins and lipid molecules such as phospholipids and cholesterol and triglyceride. Boren *et al.* [54] concluded that the LPL-apoB interaction is dependent on lipids but not on apoB because chemical modification of apoB did not abolish the interaction, and partial delipidation of LDL markedly decreased the binding to LDL. In contrast, Goldberg *et al.* proposed that protein-protein

interaction between LPL and apoB is more important than the interaction between LPL and LDL lipids [55-59] using mutant apoB proteins. Thus, whether apoB or lipids contribute to the interaction with LPL has been controversial. Likewise, whether the interaction between sPLA₂-X and LDL is primarily mediated by the apoprotein (perhaps apoB-100) or by lipids is an important subject. We did not detect specific binding between sPLA₂-X and HDL which did not contain Apo B-100 [60]. These results suggest the importance of Apo B-100 in the association of these molecules. Whether ApoB-100 or phospholipid interacts with sPLA₂-X is unclear in the present study. The mature sPLA₂-X protein is acidic (pI 5.3) in contrast to the basic properties of group IIA and V sPLA₂s. If the interaction is ascribed to this acidic characteristic, basic residues including lysine or arginine within ApoB-100 might be essential. In the future, what is bound to sPLA₂-X should be clarified by means of LDL including mutational recombinant ApoB-100 or other methods. The biological roles of the association between sPLA₂-X and LDL remain unclear. One hypothesis is that LDL acts as a carrier protein for binding to sPLA₂-X, and such concentrated sPLA₂-X rapidly hydrolyzes LDL. Further, we have shown that LDL modified by sPLA₂-X was efficiently incorporated into macrophages to induce the accumulation of cellular cholesterol ester and formation of non-membrane-bound lipid droplets in the cytoplasm [34]. Therefore, sPLA₂-X itself was also possibly incorporated due to the high affinity of sPLA₂-X for LDL. Because sPLA₂-X showed substantial activity even in the absence of Ca²⁺ (Fig. 3), it could be active after incorporation into cells, where the concentration of Ca²⁺ is much lower than in the extracellular space. Thus, sPLA₂-X in the modified LDL may also elicit the release of arachidonic acid from the cell surface and/or cytoplasm in macrophages.

In conclusion, we have demonstrated here marked differences between sPLA₂-V and sPLA₂-X compared to other sPLA₂s: sPLA₂-X introduces large amounts of arachidonic acid following eicosanoid biosynthesis, is activated less dependently on the calcium cation concentration, and specifically and tightly binds to LDL. These results indicate sPLA₂-X to play a pivotal role in the modification of lipoproteins related to atherosclerosis.

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