

Identification of a 42-kDa Group IV cPLA₂-activating protein, cPLAP γ , as a GTP-binding protein in the bovine brain

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Jae Hyung Bach, Kwang Mook Jung,
Jae Sin Choi, Sung Yun Jung, Mi
Reyoung Chin, Kyong Hoon Ahn,
Seok Kyun Kim and Dae Kyong Kim*

Department of Environmental & Health Chemistry, College of
Pharmacy, Chung-Ang University, 221 Huksuk-dong, Dongjak-ku,
Seoul 156-756, South Korea

*Dae Kyong Kim, Department of Environmental & Health
Chemistry, College of Pharmacy, Chung-Ang University, 221
Huksuk-Dong, Dongjak-Ku, Seoul 156-756, South Korea.
Tel: +82-2-820-5610, Fax: +82-2-825-7920,
email: proteinlab@hanmail.net

Brain tissue contains multiple forms of Phospholipase A₂ (PLA₂) whose activities are involved in intracellular and intercellular signalling related to normal functions such as long-term potentiation, neurotransmitter release, cell growth and differentiation. Among them, we focused on regulatory mechanism of cPLA₂ α (Group IVA cytosolic PLA₂) in brain tissue. In the present study, we report the identification of a cPLA₂-activating protein (cPLAP) in the bovine brain. cPLAP activity appeared as two major peaks with molecular masses of 200 and 42 kDa in a Superose 12 gel filtration FPLC column. The 42-kDa form of cPLAP, designated cPLAP γ , was further purified using a Mono S FPLC column to near homogeneity and characterized to as a GTP-binding protein (G protein). Metabolic labelling and immunoprecipitation studies revealed that cPLAP γ associates with cPLA₂ *in vitro* and co-immunoprecipitates with [³⁵S]-cPLA₂. Notably, cPLAP γ rendered cPLA₂ fully activated at submicromolar concentrations of Ca²⁺. These results suggest that cPLAP γ may act as a G protein, activating cPLA₂ prior to reaching full intracellular Ca²⁺ concentrations.

Keywords: calcium/cPLA₂-activating protein/Group IV cytosolic phospholipase A₂/GTP-binding protein/purification.

Abbreviations: 2-[1-¹⁴C]AA-GPC, 1-stearoyl-2-[1-¹⁴C]arachidonoyl-*sn*-glycero-3-phosphocholine; 2-[1-¹⁴C]AA-GPE, 1-acyl-2-[1-¹⁴C]arachidonoyl-*sn*-glycero-3-phosphoethanolamine; AA, arachidonic acid; AppNHp, adenylylimidotriphosphate; CaLB domain, Ca²⁺-dependent lipid-binding domain; cPLA₂, cytosolic phospholipase A₂; cPLAP, cPLA₂-activating protein; GDP, guanosine diphosphate; G protein, GTP-binding regulatory protein; GTP, guanosine triphosphate; MAP kinase, mitogen-activated protein kinase; PC, phosphatidylcholine; PE, phosphatidylethanolamine; PI-PLC,

phosphatidylinositol-specific phospholipase C; PLA₂, phospholipase A₂ (EC 3.1.1.4); PLAP, PLA₂-activating protein.

Phospholipase A₂ (PLA₂; EC 3.1.1.4) are ubiquitous enzymes that specifically hydrolyze the acyl ester bond at the *sn*-2 position of membrane phospholipids to produce a free acid and lysophospholipid (1).

Brain tissue contains multiple forms of PLA₂, whose activities are involved in intracellular and intercellular signalling related to normal functions such as long-term potentiation, neurotransmitter release, cell growth, and differentiation (2). Increased levels or activity of PLA₂ and excessive production of arachidonic acid (AA) and lysophospholipids may lead to disease states and neuronal injury or neurodegenerative diseases (3), including Alzheimer's disease (4), multiple sclerosis (5), olivopontocerebellar atrophy (6), as well as psychiatric disorders such as schizophrenia (7).

Recent advances in molecular biology of PLA₂ have led to the identification of more than 20 isoforms with PLA₂ activity. The PLA₂ can be divided into several groups depending upon enzymatic properties, subcellular localization, their structure and cellular function (2). These are secretory PLA₂ (sPLA₂), cytosolic PLA₂ (cPLA₂) and calcium-independent PLA₂ (iPLA₂). Among them, the most thoroughly investigated group of PLA₂ is cPLA₂ (Group IV PLA₂) in brain tissue. Our laboratory has focused on the study of the group IVA cPLA₂ (Group IVA PLA₂ or cPLA₂ α). Although brain tissue contains cPLA₂ activity, it has never been purified to homogeneity and characterized from brain (2). Thus, not only its regulatory mechanism of AA release in the brain but also the existence of cPLA₂ as an active enzyme in the brain remains largely uncertain.

Although to date extensive studies have focused on the regulatory mechanism of cPLA₂, the precise mechanism remains not fully understood. At present, three major mechanisms have been elucidated. First, Ca²⁺ is an important regulator of cPLA₂ activity. This enzyme has a Ca²⁺-dependent lipid-binding (CaLB) domain, which is responsible for translocation of cPLA₂ from the cytosol to the membrane compartment in the presence of 10⁻⁶ M Ca²⁺ (8, 9). That is, cPLA₂ requires Ca²⁺ only for translocation and association with its substrate membrane via the CaLB domain. Second, phosphorylation of cPLA₂ is also postulated as a

possible mechanism of activation. Diverse agonists that provoke the release of AA also activate a variety of kinases (10–12). Third, ceramide can interact directly with cPLA₂ via the CaLB domain and thereby serves as a membrane-docking device that facilitates cPLA₂ action in inflammatory diseases (13). Moreover, ceramide-1-phosphate not only increases the membrane affinity of cPLA₂ but also may act as an allosteric activator of the membrane-associated enzyme (14). However, detailed mechanisms of its regulation remain to be elucidated.

In a signal transduction concept first proposed by Rodbell *et al.* (15), GTP-binding regulatory protein (G protein) was proposed as a transducer or a molecular switch. Many phospholipases have been found to interact with and be directly regulated by specific G proteins, *e.g.* phospholipase C (PLC) by heterotrimeric G proteins (16, 17) and phospholipase D by ADP-ribosylation factor (18). In the brain, PLA₂ activity and AA release are linked to the receptors for dopamine, glutamate, serotonin, P₂-purinergics, cytokines and growth factors through various coupling mechanisms (19–23). In the case of PLA₂, there have been many trials to identify the corresponding G protein that regulates PLA₂s, especially high-molecular weight cPLA₂ α , which is hormonally regulated (24). Using the non-hydrolysable guanine nucleotide analogues GTP γ S/GDP β S and G protein-sensitive bacterial toxins such as cholera toxin and pertussis toxin, there are an increasing number of reports suggesting that AA release in various cells and tissues is regulated by G proteins (25). cPLA₂ is regulated by Ras-related C3 botulinum toxin substrate 1 (Rac1), a member of the Rho family of G proteins, which acts as a molecular switch by cycling between active GTP bound and inactive GDP-bound forms (26). Rac1 binds to effectors or molecules in its GTP bound form, and influences cPLA₂ activation and subsequent release of AA (27–29). However, at present, there is no evidence for direct interaction of cPLA₂ with a G protein either *in vivo* or *in vitro*.

On the other hand, 28-kDa PLA₂-activating protein (PLAP) has been identified and characterized (30). The PLAP stimulated sPLA₂ but not cPLA₂. The purified protein selectively stimulated sPLA₂ when phosphatidylcholine (PC) was used as a substrate but had no effect on activity with phosphatidylethanolamine (PE) as a substrate. Interestingly, a homology search identified that PLAP has a domain exhibiting significant homology with β -transducin, a member of the G β superfamily. However, the importance of this domain for PLAP activity remains unknown, as PLAP activity has not been shown to be affected by GTP γ S or GDP β S.

In the present study, we identified a cPLA₂-activating protein (cPLAP) in the bovine brain that specifically enhanced the *in vitro* activity of porcine spleen cPLA₂. The cPLAP activity appeared as two major peaks with molecular masses of 200 and 42 kDa on a Superose 12-gel filtration FPLC column. We also demonstrate here that a polyclonal antibody generated against the cPLA₂ α protein immunoprecipitated 42 kDa cPLAP (cPLAP γ) from bovine brain as well

as human promyelocytic leukaemia HL60 and human leukaemic monocyte lymphoma U937 cell lines in metabolic labelling study. These results strongly suggest that cPLAP γ may be a G protein which fully activates cPLA₂ at sub-micromolar concentrations of Ca²⁺.

Materials and Methods

Materials

1-Stearoyl-2-[1-¹⁴C]arachidonoyl-*sn*-glycero-3-phosphocholine (2-[1-¹⁴C]AA-GPC) (55 mCi/mol), 1-acyl-2-[1-¹⁴C]arachidonoyl-*sn*-glycero-3-phosphoethanolamine (2-[1-¹⁴C]AA-GPE) (55.6 mCi/mol) and [α -³²P]GTP (1 mCi/ml, 3000 Ci/mmol) were purchased from Amersham Life Science, Ltd (Buckinghamshire, England). DEAE-cellulose gel (DE52) was purchased from Whatman Co (Maidstone, England) and Phenyl-Sepharose CL-4B gel, Resource S column and PD-10 were purchased from Pharmacia LKB Biotechnology AB (Uppsala, Sweden). A Phenyl-5PW column was purchased from Tosoh (Tokyo, Japan). A Centricron YM10 centrifuge was purchased from Millipore (Bedford, MA, USA). Porcine pancreatic PLA₂ was purchased from Sigma Chemical Co. (St Louis, MO, USA). cPLA₂ α was purified from porcine spleen as described previously (31). Rat platelet group II PLA₂ was partially purified from rat platelets as described previously (32).

Anti-cPLA₂ antibody was prepared as described previously (31). GTP γ S and GDP β S were purchased from BIOMOL Research Laboratories, Inc (Plymouth, PA, USA) and guanosine 5'-triphosphate (GTP), ATP and adenylylimidotriphosphate (AppNHp) from Sigma Chemical Co. All other chemicals were of the highest purity or molecular biology grade available from commercial sources.

Assay for cPLA₂, iPLA₂ and cPLAP activities

To assay cPLA₂ activity, the substrate, 2-[1-¹⁴C]AA-GPC or 2-[1-¹⁴C]AA-GPE, was dried under nitrogen and resuspended in ethanol. The standard incubation system (100 μ l) for assay of cPLA₂ activity contained the cPLA₂ preparation, 75 mM Tris-HCl at pH 7.5, 5 mM CaCl₂ and 4.5 nmol of the substrate (~55,000 cpm). iPLA₂ activity assay was performed in the standard reaction buffer (100 μ l) contained iPLA₂ preparations, 200 mM Tris-HCl, pH 7.5, 5 mM EDTA and 4.5 nmol of the substrate. The reactions were carried out at 37°C for 5 min and stopped by adding 560 μ l of modified Dole's reagent (*n*-heptane/isopropyl alcohol/1 N-H₂SO₄; 400/390/10, by volume) (33) and 110 μ l of water, vortex mixed and centrifuged. Then, 150 μ l of the upper phase was transferred to a new microtube, to which 800 μ l of *n*-heptane and silica gel (~5 mg) was added. The sample was vortex mixed and centrifuged again and an aliquot (800 μ l) of the supernatant was removed and added to 2.5 ml of β -scintillation solution (INSTA GEL-XF) and counted for radioactivity with a Packard Tri-Carb liquid scintillation counter (Packard Instrument Co, Meriden, CT, USA). In order to characterize GTP effect, cPLA₂ activity was assayed in the presence of 2.5 mM of MgCl₂. The cPLA₂ activity itself was not affected by Mg²⁺. To measure cPLAP activity, an aliquot of the sample preparation was added to the assay system for cPLA₂ activity and expressed as an *n*-fold increase in activity.

Detection and purification of cPLAP γ and iPLA₂

Bovine brain was freshly obtained from a local slaughterhouse and frozen at -70°C. A 600 g portion of brain was thawed in five volumes of homogenizing buffer V (50 mM Tris-HCl at pH 7.5, 50 mM KCl, 3 mM MgCl₂, 1 mM EDTA and 10 mM 2-mercaptoethanol) with a Polytron PT-MR6000 (Kinematica AG, Rittau, Switzerland). The homogenates were centrifuged at 10,000g for 60 min. The resulting supernatants were adjusted to pH 5.0 with acetic acid and centrifuged again at 10,000g for 15 min. After centrifugation, precipitates were resuspended in buffer V using Polytron at 4°C. The resuspension of pH 5.0-precipitate was loaded onto a DEAE-cellulose anion exchange column (DE52, 2.5 mm \times 12.5 cm, bed volume 200 ml; Whatman Ltd) designed for use with a P-3 peristaltic pump (Pharmacia LKB Biotechnology AB) with buffer A (50 mM Tris-HCl at pH 7.5, 1 mM EDTA and 10 mM 2-mercaptoethanol) at a flow rate of ~25 ml/min, and the column

was washed with buffer A. The washed column was eluted at a flow rate of 25 ml/min with a stepwise gradient of 0–1.0 M NaCl. The protein fractions eluted with 1.0 M NaCl were fractionated at 30 ml per fraction. A 20- μ l aliquot of each fraction was assayed for iPLA₂ and cPLA₂-activating activities. The active fractions with iPLA₂ activity were pooled.

The active fractions from the DEAE-cellulose column were pooled and adjusted to 0.5 M (NH₄)₂SO₄ by adding 4.0 M (NH₄)₂SO₄ solution and applied to a Phenyl-Sepharose CL-4B column (30 mm \times 25 cm, bed volume 30 ml; Pharmacia LKB Biotechnology) pre-equilibrated with buffer A containing 0.5 M (NH₄)₂SO₄. The column was eluted at a flow rate of 10 ml/min with sequential stepwise elution of buffer A containing 0.25 M (NH₄)₂SO₄ and buffer B (25 mM glycine–NaOH at pH 9.0, 1 mM EDTA, and 10 mM 2-mercaptoethanol). A 20- μ l aliquot of each 10 ml fraction was assayed for iPLA₂ and cPLA₂-activating activities at 37°C for 5 min. The active pool of the Phenyl-Sepharose CL-4B column was adjusted to 0.75 M (NH₄)₂SO₄ and applied to Phenyl-5PW HPLC column, which was pre-equilibrated with buffer A containing 0.75 M (NH₄)₂SO₄. The column was eluted at a flow rate of 5 ml/min with a linear gradient with water. The active fractions with cPLA₂-activating activity were pooled, dialyzed with buffer S (25 mM imidazole–HCl at pH 6.6, 3 mM MgCl₂, 1 mM EDTA and 10 mM 2-mercaptoethanol) for 18 h. The dialyzed sample was applied to a Resource S FPLC column (1 ml; Pharmacia LKB Biotechnology AB) pre-equilibrated with buffer S. The column was eluted at a flow rate of 1 ml/min with a linear gradient of 0.0–1.0 M NaCl. A 20- μ l aliquot of each 1 ml fraction was assayed for cPLA₂-activating activity. The active fractions (~3.0 ml) were concentrated to 400 μ l using a Centricon YM10 (Millipore). The active concentrate was applied to a Superose 12 gel filtration FPLC column (Pharmacia LKB Biotechnology AB) pre-equilibrated with buffer G (50 mM Tris–HCl at pH 7.5, 1 mM EDTA, 10 mM 2-mercaptoethanol and 0.1 M NaCl). The column was eluted at a flow rate of 0.5 ml/min. A 50- μ l aliquot of each 0.5 ml fraction was assayed for cPLA₂-activating activity at 37°C for 5 min. The 42-kDa peak fractions were pooled and dialyzed with buffer S (25 mM imidazole/HCl at pH 6.6, 3 mM MgCl₂, 1 mM EDTA and 10 mM 2-mercaptoethanol) for 18 h. The dialyzed sample was applied to a Mono S FPLC column (1 ml/bed volume; Pharmacia LKB Biotechnology AB) pre-equilibrated with buffer S. The column was eluted at a flow rate of 1 ml/min with a linear gradient of 0.0–1.0 M NaCl. A 20- μ l aliquot of each 1.0 ml fraction was assayed for cPLA₂-activating activity.

Photoaffinity labelling of cPLAP γ with [α -³²P]GTP

To determine whether GTP binds to cPLAP γ by photoreaction, photoaffinity labelling of cPLAP γ with [α -³²P]GTP was performed as previously described (34, 35) with a minor modification. Briefly, the samples were mixed with 5–10 mCi of [α -³²P]GTP, 0.1 mM AppNHp and 2 mM MgCl₂ and pre-incubated for 5 min at 4°C. The samples were then placed in an ice bath and irradiated with UV light (254 nm) for 5–10 min using a UV cross-linker (SPECTROLINKER XL-1000, Spectronics Co, Manukau City, New Zealand). After irradiation, the samples were mixed with Laemmli's stopping solution (36) and allowed to stand at room temperature for 1 h. The samples were subjected to SDS–PAGE using 8–12% gels, and the gels were dried and exposed overnight on Kodak X-OMAT XAR-5 film using DuPont image intensifying screens.

Co-immunoprecipitation of cPLAP γ with cPLA₂ in HL60 and U937 cells metabolically labelled with L-[³⁵S]methionine

The cells cultured in DMEM (10% FCS) were collected by centrifugation at 400g for 5 min. The resulting pellets were resuspended in DMEM (10% FCS) without L-methionine supplemented with L-glutamine (2%) as a wash buffer and centrifuged at 400g at room temperature for 5 min. After removing the wash buffer as completely as possible, the cells were resuspended in methionine-free medium (1 \times 10⁷ cells/ml). After transferring to a small tissue culture plate (16 mm, 24 wells/plate), the cells were incubated for 30 min at 37°C to exhaust the remaining methionine.

Then 100 μ Ci [³⁵S]methionine was added to the plates and the cells were allowed to incorporate the radiolabel into total protein for 3 h. The cells were treated with 1 μ M A23187 for 30 min, transferred to a test tube and washed twice with PBS. The resulting pellets were

resuspended in 1 ml of a RIPA buffer (150 mM NaCl, 1% NP-40, 0.5% sodium deoxycholate, 0.1% SDS, 50 mM Tris–HCl at pH 7.5 and 1 mM EDTA) per 1 \times 10⁷ cells for lysis. The cells were kept on ice for 30 min with occasional mixing and spun down for 10 min at 10,000g at 4°C. The resulting supernatants were centrifuged at 100,000g for 30 min and the supernatants used as the soluble cell-free fractions.

To pre-clear proteins in the soluble fractions that bind non-specifically to immune complexes or the solid phase, 1 ml fractions were incubated with 50 μ l of normal serum on ice for 1 h. Then 500 μ l of 10% washed suspension of fixed *Staphylococcus aureus* Cowan I (SAC) was added to the lysates and further incubated for 30 min on ice. The resulting supernatants were incubated with 20 μ l of anti-cPLA₂ antibody bound to Protein A-Sepharose beads for 12 h. Immune complexes bound to the beads were pelleted at 1,000g at 4°C for 3 min and washed five times with 1 ml of the lysis buffer. The cPLA₂ and associated proteins were eluted by incubating for 5 min at 100°C with 70 μ l of 2 \times Laemmli's buffer and subjected to 10% SDS–PAGE. After washing twice for 30 min with DMSO, the gel was reacted with 22% PPO in DMSO for 1 h and extensively washed for 30 min with water in a circulating chamber. Then, the gel was dried at 80°C for 40 min and exposed to film overnight at –70°C for fluorography.

Results

Detection and partial purification of cPLAP γ

Recently, we identified and purified a novel form of iPLA₂ in the pH 5.0-precipitate of the brain homogenate (Eui Man Jeong *et al.*, manuscript in preparation). The brain homogenates were resuspended with buffer V and applied to a DEAE-cellulose anion exchange column, where the iPLA₂ activity was eluted with a stepwise gradient of 1.0 M NaCl with an 82% yield and ~10.5-fold purification (Fig. 1A). The active fractions of iPLA₂ were adjusted to 0.5 M (NH₄)₂SO₄ and then applied to a Phenyl-Sepharose CL-4B hydrophobic column. The bound proteins were eluted with stepwise gradients of buffer A containing 0.25 M (NH₄)₂SO₄ and buffer B containing no salt, respectively. The iPLA₂ activity was eluted in the glycine-eluted fractions with a yield of <20%, but not in 0.25 M (NH₄)₂SO₄ elution (Fig. 1B). This low yield prompted us to examine an existence of an activator in other fractions. First, we observed a small increase by ~1.2–1.5-fold in iPLA₂ activity in the 0.25 M (NH₄)₂SO₄-elution fractions, but this result could not explain the low yield in this step (data not shown). Since we cannot exclude the possible existence of cPLA₂ in the glycine-eluted fraction, we added a highly purified cPLA₂ from porcine spleen to the 0.25 M (NH₄)₂SO₄-elution fractions. Interestingly, as shown in Figure 1B, a significant cPLA₂-activating activity was observed in the 0.25 M (NH₄)₂SO₄-elution fractions. One unit of cPLA₂-activating activity was defined as the ability that doubles 52 nmol/min of the total activity of the purified spleen cPLA₂ for 5 min/mg of each protein in the samples. For purification, the fractions showing cPLA₂-activating activity were re-adjusted to 0.75 M (NH₄)₂SO₄ and then applied to a Phenyl-5PW HPLC column (Fig. 1C). The active fractions were pooled and dialyzed for 18 h. Then, the dialyzed sample was applied to a Resource S FPLC column (Fig. 1D). The active fractions had 55% less protein and a total activity recovery of ~60% compared with the loaded sample. Finally, a 3.0-ml pool of the active fractions from the Resource S FPLC

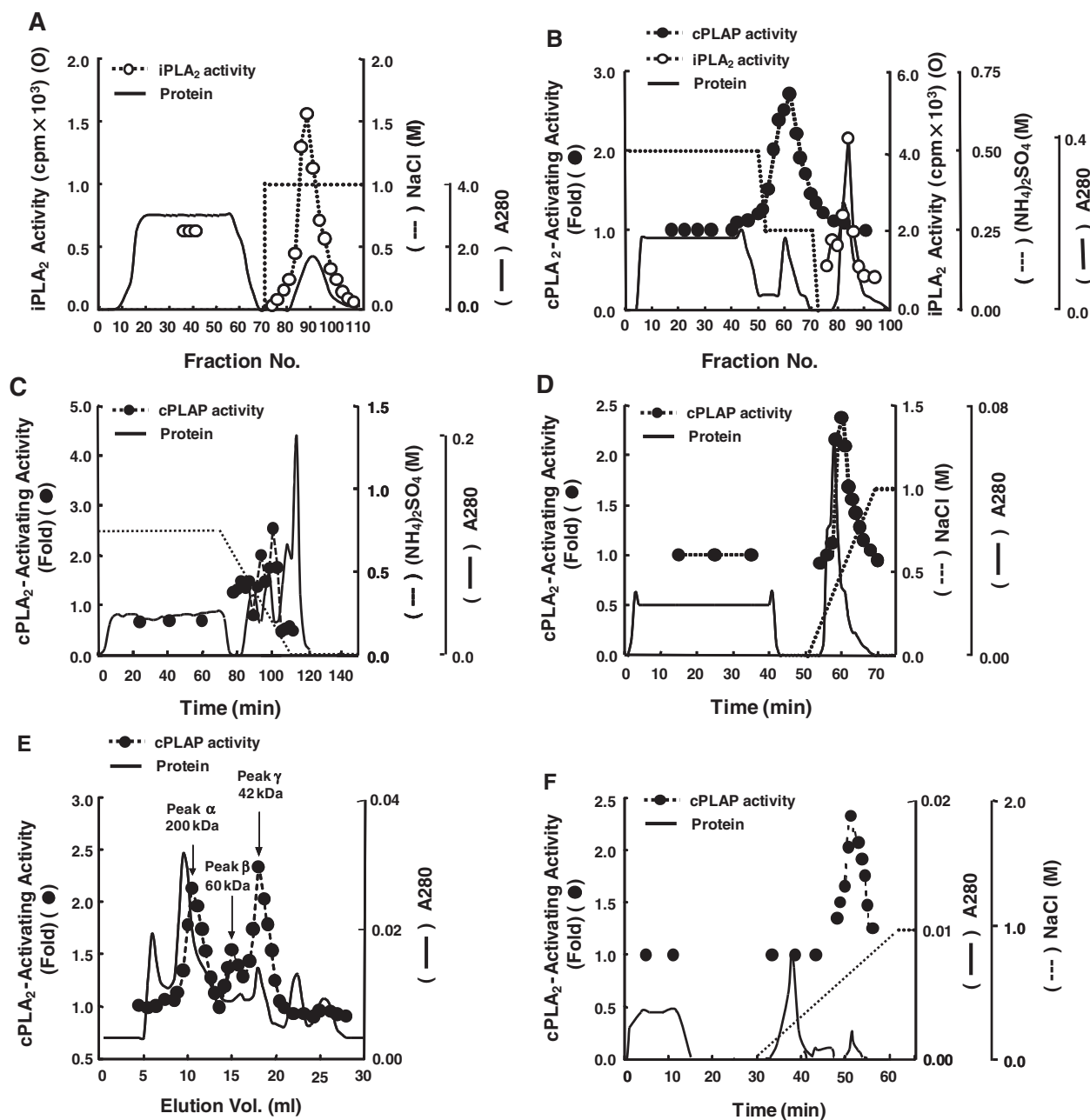


Fig. 1 Detection and purification of cPLAP. (A) Detection of iPLA₂ in bovine brain samples. The pH 5.0-precipitate of bovine brain homogenate was applied to a DE52 column pre-equilibrated with buffer A. The column was eluted and each fraction was assayed for iPLA₂ activity as described in 'Materials and Methods' section. The active fractions (numbers 83–96) were pooled for the next step. (B) Detection of cPLAP by Phenyl-Sepharose CL-4B hydrophobic column chromatography. The pool of DE52 was applied to a Phenyl-Sepharose CL-4B column pre-equilibrated with buffer A containing 0.5M (NH₄)₂SO₄. The column was eluted with a sequential stepwise elution and each fraction was assayed for cPLAP activity as described in 'Materials and Methods' section. The iPLA₂ activity was eluted in the elution step without salt. (C) Phenyl-5PW HPLC column profile of cPLAP. The pool of the active cPLAP fractions of the Phenyl-Sepharose CL-4B column was applied to a Phenyl-5PW HPLC column pre-equilibrated with buffer A. The column was eluted and each fraction was assayed for cPLAP activity as described in 'Materials and Methods' section. (D) Resource S FPLC profile of the active pool from the Phenyl-5PW column. The active pool was dialyzed and applied to a Resource S FPLC column pre-equilibrated with buffer S. The column was eluted and assayed for cPLAP activity as described in 'Materials and Methods' section. Fraction numbers 59–61 were pooled for the next step. (E) Superose 12-gel filtration FPLC profile of cPLAP. The active pool of Resource S was applied to a Superose 12 FPLC column pre-equilibrated with buffer T. The column was eluted and assayed for cPLAP activity as described in 'Materials and Methods' section. (F) Mono S cation exchange FPLC profile of cPLAP. The active pool of Superose 12 FPLC column was applied to a Mono S FPLC column, which was eluted and assayed for cPLAP activity as described in 'Materials and Methods' section.

column was concentrated to ~400 μ l and applied to a Superose 12 gel filtration FPLC column. The cPLA₂-activating activity was separated into two major peaks of apparent molecular masses of 200 and 42 kDa,

termed peak α and γ , respectively (Fig. 1E). The 42-kDa protein of peak γ was further purified using a Mono S cation exchange FPLC column (Fig. 1F).

Effects of cPLAP γ on various types of PLA₂

A 20- μ l aliquot of the active pool obtained from the Mono S FPLC column was assayed for cPLAP activity. To examine the substrate specificity of the acyl chains at the *sn*-2 position, PLA₂-activating activity was assayed for 2-PA-PC, 2-LA-PC and 2-AA-PC. There was no preferential activity of cPLA₂ toward AA at the *sn*-2 position (Figure 2A). The effects of cPLAP γ on various types of PLA₂ were then examined. cPLAP γ enhanced group IVA PLA₂ activity by \sim 2-fold, but did not affect pancreatic group I, non-pancreatic group II or partially purified iPLA₂ from the bovine brain. The results indicate that cPLAP γ activated significantly cPLA₂ by up to \sim 2.5-fold (Figure 2B). Next, we also examined the effects of GTP γ S on cPLAP γ activity. The addition of GTP γ S further increased the activating effect of cPLAP γ on cPLA₂ α (Figure 2A and B). The data suggest that cPLAP γ may be a GTP-binding protein.

To determine the nature of the cPLA₂-activating activity, the effects of boiling and digestion with trypsin on cPLA₂-activating activity were assessed (Figure 2C). A 20- μ l aliquot of the active fractions from Mono S was pre-incubated for 15 and 30 min at 75°C, followed by 0.5 μ g/ μ l of trypsin and then the trypsin inhibitor 0.25 μ g/ μ l aprotinin, or vice versa, before addition of the cPLA₂ enzyme source. The cPLA₂-activating activity was sensitive to boiling and digestion with trypsin, thus the cPLA₂-activating activity is attributable to a protein in the preparation, and is named cPLA₂-activating protein accordingly.

cPLAP γ has characteristics of G protein

To characterize the cPLAP γ , the effects of guanine nucleotides on the activity of cPLAP were investigated. Because most of the GTP proteins require an appropriate concentration of Mg²⁺ for nucleotide binding

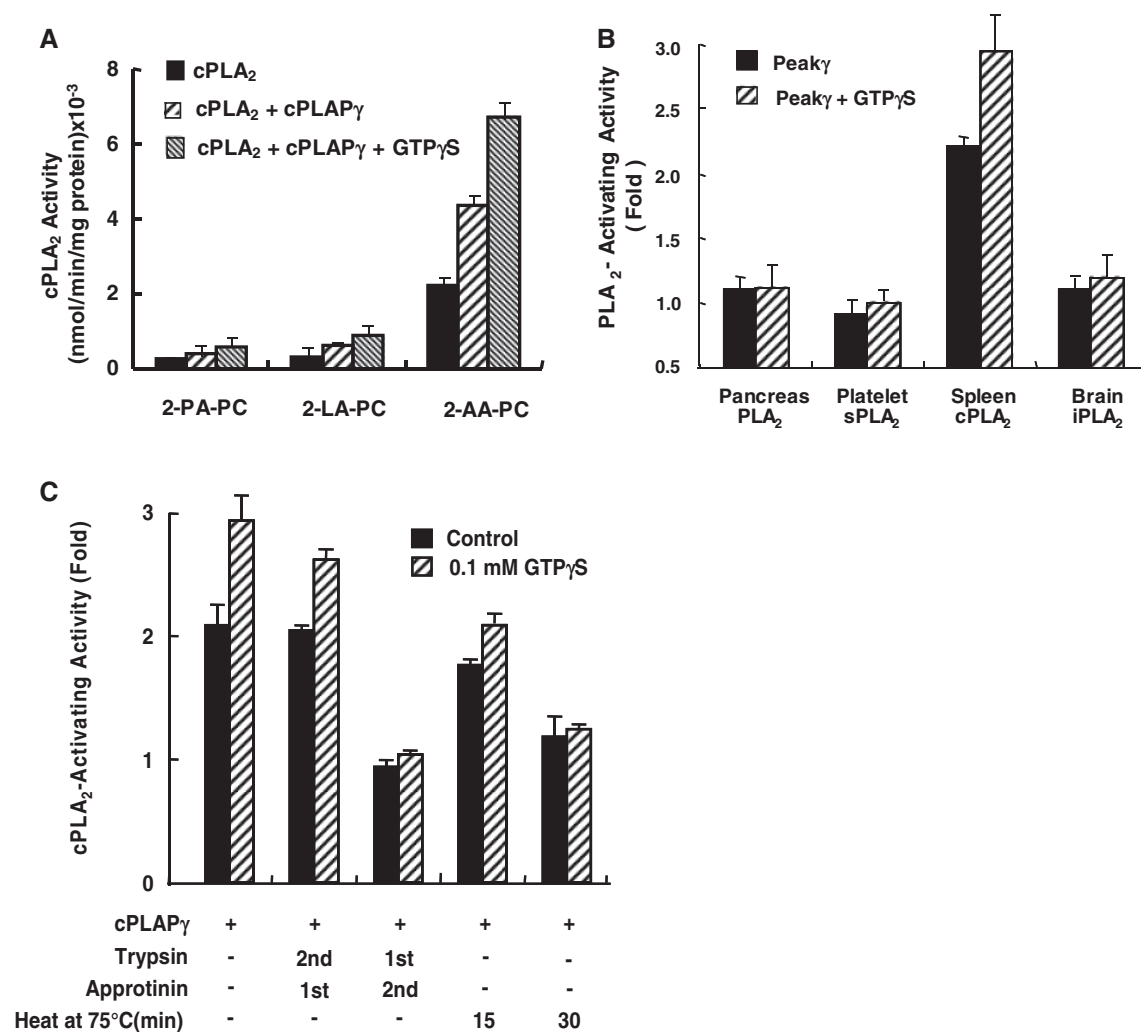


Fig. 2 Selectivity of cPLAP γ for subtypes of PLA₂ and fatty acids at the *sn*-2 position. A 20- μ l aliquot of the active pool obtained from the Mono S FPLC column was assayed for cPLAP activity. (A) cPLAP γ activity was assayed using PCs containing various fatty acids at the *sn*-2 position as described in 'Materials and Methods' section. (B) The cPLAP γ activity was assayed using various types of PLA₂ as described in 'Materials and Methods' section. Pancreas PLA₂ (Group I PLA₂), platelet sPLA₂ (Group II), spleen cPLA₂ (Group IV), and brain iPLA₂ (Ca²⁺-independent PLA₂) had 3,500, 3,200, 4,280 and 4,100 cpm under the assay conditions, respectively. (C) Effects of trypsin and heat on cPLAP γ activity. cPLAP γ activity was measured by incubating with the combination of trypsin and its inhibitor aprotinin. Activity was also assayed by boiling at 75°C for 15 and 30 min. Values are the average of duplicate assays with a variance of <10%. The result is representative of three independent experiments for each condition.

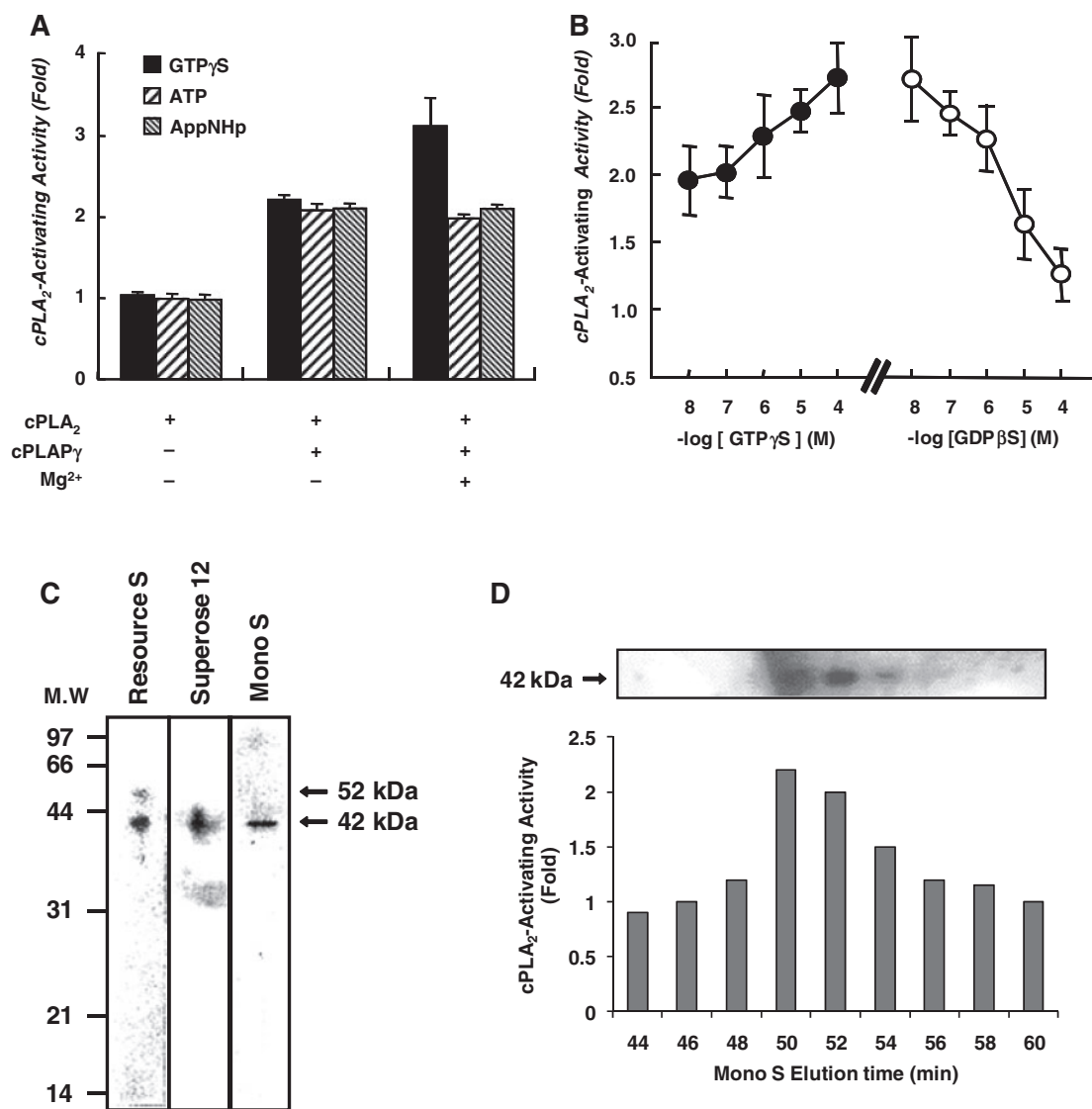


Fig. 3 Identification of cPLAP γ as a G protein. (A) Effects of Mg²⁺ ion and nucleotide analogues on cPLAP γ activity. cPLAP γ activity was assayed with 0.1 mM GTP γ S, ATP and AppNHp under standard assay conditions with or without 2.5 mM MgCl₂. After incubation for 5 min, cPLAP γ activity was measured as described in ‘Materials and Methods’ section. (B) cPLAP γ activity was increased by GTP γ S and decreased by GDP β S. A 50- μ l aliquot of the active fractions from the Superose 12 FPLC column was assayed with increasing concentrations of GTP γ S (dark circle) followed by additional treatment with increasing concentrations of GDP β S (open circle) as described in ‘Materials and Methods’ section. (C) Photoaffinity labelling of cPLAP γ with [α -³²P]GTP. A 30- μ l aliquot of the active fractions from each column was incubated with 5 mCi of [α -³²P]GTP and 0.5 mM AppNHp in the presence of 2 mM MgCl₂ and labelled with UV light (254 nm). The proteins were analyzed with SDS/PAGE (12% gel) followed by autoradiography as described in ‘Materials and Methods’ section. (D) Photoaffinity labelling of cPLAP γ with [α -³²P]GTP. A 30- μ l aliquot of the fractions from Mono S column was incubated with 5 mCi of [α -³²P]GTP and 0.5 mM AppNHp in the presence of 2 mM MgCl₂ and labelled with UV light (254 nm). The proteins were analyzed with SDS/PAGE (12% gel) followed by autoradiography as described in ‘Materials and Methods’ section.

and configuration, 2.5 mM of MgCl₂ was added to the standard assay system. The cPLA₂ activity itself was not affected by Mg²⁺ (data not shown). As shown in Figure 3A, cPLAP activity enhanced by GTP γ S required the presence of Mg²⁺. ATP and AppNHp, other purine nucleotides whose structures are similar to the guanine nucleotide, had no effect (Figure 3A). GTP γ S dose-dependently enhanced cPLAP γ activity up to ~1.5- to 2.0-fold, whereas GDP β S, a non-hydrolyzable GDP analogue, dose-dependently decreased the enhanced cPLAP γ activity (Figure 3B). To further confirm the possibility that cPLAP γ may be a G protein, a photoaffinity labelling experiment with [α -³²P]GTP was performed (Figure 3C). When the

active fractions from Resource S and Superose 12 columns were labelled with [α -³²P]GTP, proteins with molecular masses of 42 and 52 kDa were cross-linked with UV and the protein band at 42 kDa was seen at the same position as the major band from the Mono S column of the final step (Figure 3C). Furthermore, cPLAP γ was observed as a 42 kDa band, paralleled with the activity from the Mono S column (Figure 3D). These results strongly suggest that cPLAP γ may be a GTP-binding protein. The cPLAP activity appearing with ~200 kDa molecular mass in the Superose 12 gel filtration FPLC was not affected by guanine nucleotides nor was labelled by UV cross-linking (data not shown).

Identification of the 42 kDa protein which is co-immunoprecipitated with cPLA₂

cPLA₂ is found normally in a various cells including HL60 and U937 cells. Proteins from HL60 and U937 cells were radiolabelled by incubating the cells in [³⁵S]methionine-containing medium. The supernatants obtained from the [³⁵S]methionine-labelled cells were used as the soluble cell-free fractions for immunoprecipitation. As shown in Figure 4, Protein A-Sepharose beads bound to anti-cPLA₂ antibody precipitated a 42-kDa protein from both HL60 and U937 cells, but those bound to normal serum did not.

On the other hand, it is important whether interaction of cPLA₂ and cPLAP γ require an increase in intracellular Ca²⁺ concentration. Our results showed that proteins from A23187-untreated cell had no different effect on the interaction of cPLAP γ toward cPLA₂ (data not shown). Although intracellular Ca²⁺ is not required to interaction between cPLAP γ and cPLA₂, there is a possibility that cPLAP γ could make cPLA₂ more active through increase of Ca²⁺sensitivity. Next, we investigated the effects of Ca²⁺ on the increase of cPLA₂ activity by cPLAP γ .

Ca²⁺-dependency of the cPLA₂-activating activity of cPLAP γ

cPLA₂ is known to have full activity at a micromolar concentration of Ca²⁺. Interestingly, the present data indicate that cPLAP γ fully activates cPLA₂ even at sub-micromolar concentrations of Ca²⁺ (Figure 5). The effect of GTP γ S on the activity of cPLAP γ may be due to its ability to enhance cPLAP γ activity rather than to increase its sensitivity to submicromolar Ca²⁺ (Figure 5). The full activity of cPLAP γ in 10⁻⁷ M of

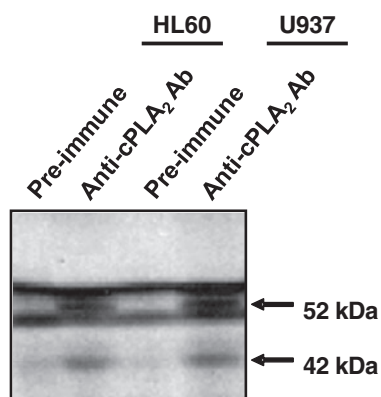


Fig. 4 Co-immunoprecipitation of cPLAP γ with cPLA₂ in HL60 and U937 cells metabolically labelled with L-[³⁵S]methionine. HL60 and U937 cells were labelled with [³⁵S]-methionine and treated with 1 μ M A23187 as described in 'Materials and Methods' section. The cells were resuspended in a RIPA buffer containing 1% NP-40, 0.5% sodium deoxycholate, 0.1% SDS and 1 mM EDTA for lysis and spun down to obtain the resulting supernatants, as described in 'Materials and Methods' section. The proteins of the soluble fractions were pre-cleared and incubated with anti-cPLA₂ antibody bound to Protein A-Sepharose beads. The cPLA₂ and associated proteins in the immune complexes were eluted and subjected to 10% SDS-PAGE as described in 'Materials and Methods' section. The gel was then dried at 80°C for 40 min and exposed to film overnight at -70°C for fluorography.

Ca²⁺ was also inhibited by GDP β S in a dose-dependent manner (data not shown).

Discussion

AA is known to serve as a chemical mediator as well as a second messenger and is generated in membrane lipids by activation of cPLA₂ (1, 2). The cPLA₂ enzyme is strictly regulated and abnormal signalling is believed to cause serious pathological disasters. The regulatory mechanisms of cPLA₂ have been extensively studied to understand the nature of its pathogenesis and to develop new therapy targets. However, the mechanism by which AA is released at an early stage of stimulation remains largely unknown.

G proteins have been proposed as a transducer and molecular switch for cPLA₂. Many phospholipases have been shown to be regulated by directly interacting with specific G proteins (16–18). There have been many attempts to discover a corresponding G protein that regulates cPLA₂ (24). Several lines of evidence are accumulating that AA release in various cells and tissues is regulated by G proteins using guanine nucleotide analogues and bacterial toxins such as cholera and pertussis toxin (25). It has also been suggested that a Rac1 G protein may serve as a mediator of cPLA₂ activation (27, 28). However, at present, there is no direct evidence for the existence of a G protein coupled with cPLA₂ either *in vivo* or *in vitro*.

The present study reports a G protein that may interact with cPLA₂ to mediate its activation based on the following results: (i) cPLAP γ selectively increased cPLA₂ activity among various mammalian types of PLA₂; (ii) cPLAP γ activity was enhanced by GTP γ S and decreased by GDP β S in a dose-dependent manner, and this effect required Mg²⁺; (iii) ATP and other purine nucleotides such as AppNHp had no effect under the same conditions; (iv) cPLAP γ was

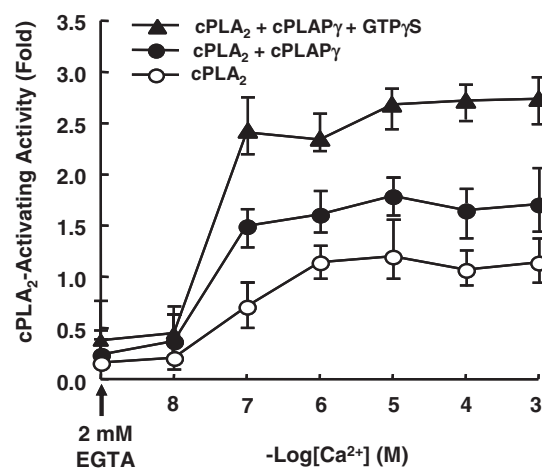


Fig. 5 Effects of cPLAP γ on Ca²⁺-dependency of the purified cPLA₂. cPLAP γ activity was assayed in 75 mM Tris-HCl buffer (pH 7.5) containing Ca²⁺ ions at the indicated concentrations. The absolute concentrations of free Ca²⁺ were calculated using an equation based on the stability constant of the EGTA/CaCl₂ system (5 mM EGTA and 5 mM CaCl₂). The cPLA₂ enzyme was purified from porcine spleen. The data represent three independent experiments, showing the mean \pm SEM of three measurements for each sample.

labelled with [α - 32 P]GTP through photoaffinity; and (v) metabolic labelling with [35 S]methionine and co-immunoprecipitation by anti-cPLA₂ antibody revealed that cPLAP γ binds cPLA₂.

Two major peaks associated with cPLAP activity were detected, peaks α and γ , from the Superose 12-gel filtration FPLC column (Figure 1E). Peak α of \sim 200 kDa was not responsive to GTP γ S or GDP β S (data not shown). Peak γ , termed cPLAP γ , was further purified with a Mono S cation exchange FPLC column to near homogeneity (Figure 1F). Based on SDS-PAGE analysis, photoaffinity labelling of cPLAP γ , and the migration profile in gel filtration chromatography, the molecular weight of cPLAP γ was estimated 42 kDa. Because the molecular mass of the α -subunit of the heterotrimeric G protein is 42 kDa and AIF4 $^-$ is an activator of the α -subunit, the effect of AIF4 $^-$ on the cPLAP γ activity was examined. Unfortunately, the effect could not be evaluated because AIF4 $^-$ significantly inhibited cPLA₂ activity in a dose-dependent manner (data not shown). Thus the cross-reactivity with antibodies against previously known six types of G proteins, Gs, Gi, Gq, Go, Gi₂ and Gi₃ proteins, was also examined through the immunoblot analysis on fraction of final purification step, but the 42 kDa cPLAP γ did not react with any of them (data not shown).

Further study should also be carried out to examine whether the 42-kDa band as shown by fluorography in Figure 4 is cPLAP γ . Metabolic labelling analysis suggested that cPLAP γ may associate with cPLA₂. However, neither cPLA₂ activity nor cPLAP γ activity was detected in the immunoprecipitates obtained from the metabolic labelling cells (data not shown), suggesting that the RIPA buffer may denature the proteins, with consequent loss of their activities. The authors' previous study showed that considerable cPLA₂ activity was detected in immunoprecipitates obtained using an anti-cPLA₂ antibody (31).

In the rat brain, cPLA₂ was ubiquitously expressed throughout the brain (37). Although cPLA₂ activity comprises a small amount of the total PLA₂ activity of the brain tissue, its specific activity is high. Such a small amount of cPLA₂ activity may have important biological and pathological functions (38). cPLA₂ is functionally linked with both COX-1 and COX-2 during immediate and delayed eicosanoid production (39). Coordinated up-regulation of cPLA₂ and COX-2 activities is also thought to contribute to brain damage following cerebral ischaemia and neuroinflammation (40). On the other hand, it was demonstrated that PLA₂ activating proteins occur in Aplysia neurons and rat cerebral cortex (41). In the context of our results and previous reports, we hypothesize that cPLAP γ is important in cPLA₂ regulation and may be involved in inflammatory response induced by G protein coupled receptors. G protein coupled receptors in the brain may activate cPLAP γ , in turn, trigger increase of cPLA₂ activity that results in neuronal apoptosis or pathological changes. Although this is an appealing possibility, we do not currently have data to support this. Further study should also be carried

out to identify and elucidate its primary structure and more precise mechanism for the cPLA₂ activation.

On the other hand, cPLA₂ activity is known to be regulated by the intracellular concentration of Ca²⁺ (9, 42). Activity was not detected at concentrations of 1 to 5×10^{-7} M Ca²⁺ observed in the resting cells, but cPLA₂ was fully activated at a concentration of $\sim 10^{-6}$ M Ca²⁺ enhanced by stimulation (9, 42). Interestingly, cPLAP γ fully activated cPLA₂ at sub-micromolar concentration of Ca²⁺ (Figure 5), suggesting that cPLA₂ may become sensitive to lower concentrations of Ca²⁺ through cPLAP γ activity. This finding strongly supports the possibility that cPLA₂ may be activated irrespective of the PI-PLC pathway, which is known to be activated through G protein-coupled receptor activation to produce IP₃ and DAG (43). It has been unclear whether cPLA₂ activation is dependent on the activation of PI-PLC, which could trigger a sharp increase in intracellular Ca²⁺ via the role of IP₃ in Ca²⁺ mobilization (44). Many lines of evidence indicate that the release of AA from membrane lipids could occur without the generation of IP₃ (45), suggesting activation of PLA₂ prior to activation of PI-PLC. However, no evidence has been provided to explain this phenomenon. The present results may shed light on this longstanding and controversial finding that cPLA₂ can be activated without the involvement of PI-PLC (46, 47).

At present, the biochemical mechanism by which cPLAP γ enhances cPLA₂ activity and allows the enzyme to fully activate at sub-micromolar concentrations of Ca²⁺ remains unknown. It has been reported that cPLA₂ contains a pleckstrin homology (PH) domain and interacts with multi-modular membrane-binding proteins that play key roles in signal transduction (8). The PH domain is known as a site of action for certain G proteins, and thus may be a site of action for cPLAP γ .

In summary, cPLAP γ was identified as a 42-kDa G protein that specifically enhances the *in vitro* activity of cPLA₂. cPLAP γ associates with cPLA₂ and allows the enzyme to fully activate at sub-micromolar concentrations of Ca²⁺. In addition, cPLAP γ is suggested to play a role in the release of AA by activating cPLA₂ at an early stage prior to activation of PI-PLC, which is known to provide the micromolar concentrations of Ca²⁺ required for full activation of cPLA₂. To verify this, the precise nature of cPLAP γ should be dissected through sequencing of its primary structure and a study of the interaction between cPLAP γ and cPLA₂.

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Conflict of interest

None declared.

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