Group X Phospholipase A2 Stimulates the Proliferation of Colon Cancer Cells by Producing Various Lipid Mediators


Institut de Pharmacologie Moléculaire et Cellulaire, Université de Nice Sophia Antipolis et Centre National de la Recherche Scientifique, Valbonne, France (F.S., I.J., E.B., C.P., C.M.M., G.L.); Departments of Chemistry and Biochemistry, University of Washington, Seattle, Washington (J.G.B., M.H.G.); and Department of Pathology, University of Turku, Turku, Finland (K.A.T., V.J.O.L., T.J.N.)

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ABSTRACT

Among mammalian secreted phospholipases A2 (sPLA2s), the group X enzyme has the most potent hydrolyzing capacity toward phosphatidylcholine, the major phospholipid of cell membrane and lipoproteins. This enzyme has recently been implicated in chronic inflammatory diseases such as atherosclerosis and asthma and may also play a role in colon tumorigenesis. We show here that group X sPLA2 [mouse (m)GX] is one of the most highly expressed PLAs in the mouse colon and that recombinant mouse and human enzymes stimulate proliferation and mitogen-activated protein kinase activation of various colon cell lines, including Colon-26 cancer cells. Among various recombinant sPLA2s, mGX is the most potent enzyme to stimulate cell proliferation. Based on the use of sPLA2 inhibitors, catalytic site mutants, and small interfering RNA silencing of cytosolic PLA2α and M-type sPLA2 receptor, we demonstrate that mGX promotes cell proliferation independently of the receptor and via its intrinsic catalytic activity and production of free arachidonic acid and lysophospholipids, which are mitogenic by themselves. mGX can also elicit the production of large amounts of prostaglandin E2 and other eicosanoids from Colon-26 cells, but these lipid mediators do not play a role in mGX-induced cell proliferation because inhibitors of cyclooxygenases and lipoxygenases do not prevent sPLA2 mitogenic effects. Together, our results indicate that group X sPLA2 may play an important role in colon tumorigenesis by promoting cancer cell proliferation and releasing various lipid mediators involved in other key events in cancer progression.

Phospholipases A2 (PLAs) catalyze the hydrolysis of the sn-2 ester bond of glycerophospholipids to generate free fatty acids and lysophospholipids (Schaloske and Dennis, 2006; Lambeau and Gelb, 2008). Over the past few years, it has been realized that PLAs constitute a superfamily of enzymes comprising several intracellular enzymes and secreted PLAs (sPLAs).

The group IVA cytosolic PLA2 (cPLA2α) is the best known intracellular PLA2, and it clearly plays an important, yet not exclusive role in the release of arachidonic acid (AA) and subsequent production of eicosanoids in various biological settings (Kita et al., 2006). In contrast, the biological functions of the different sPLAs are slowly being unraveled. sPLAs have been implicated in lipid digestion and obesity; activation of immune cells; asthma; atherosclerosis; acute respiratory distress syndrome; and host defense against bac-

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ABBREVIATIONS: PLA2, phospholipase A2; sPLA2, secreted phospholipase A2; cPLA2, cytosolic phospholipase A2; Apc, adenomatous polyposis coli; AA, arachidonic acid; m, mouse; MAP, mitogen-activated protein; DMEM, Dulbecco’s modified Eagle’s medium; FCS, fetal calf serum; PGE2, prostaglandin E2; FAF BSA, fatty acid-free bovine serum albumin; ITS, insulin, transferrin, selenious acid; MAPK, mitogen-activated protein; MK886, 3-[1-(p-chlorophenyl)-5-isopropyl-3-tert-butylthio-1H-indol-2-yl]-2,2-dimethylpropanoic acid; ATX, autotaxin; ECF, enhanced chemiluminescence; siRNA, small interfering RNA; h, human; RT-qPCR, reverse transcription quantitative polymerase chain reaction; DIG, digoxigenin; SSC, standard saline citrate; PBS, phosphate-buffered saline; TBS, Tris-buffered saline; GFP, green fluorescent protein; OS2, Oxyuranus scutellatus scutellatus sPLA2-2; LPA, lysophosphatidic acid; LPC, lysophosphatidylcholine; ANOVA, analysis of variance; LPE, lysophosphatidylethanolamine; COX, cyclooxygenase; LOX, lipoxygenase; Ss2826, [4-(tetradecanoylamo)benzyl]phosphonic acid; PA, phosphatidic acid; LY329722, 3-(3-aminooxalyl-1-benzyl-2-ethyl-6-methyl-1H-indol-4-yl)propionic acid.
teria, viruses, and parasites (Touqui and Wu, 2003; Triggiani et al., 2006; Lambeau and Gelb, 2008; Nevalainen et al., 2008). Besides their catalytic activity, sPLA₂s are also able to bind to specific soluble and membrane-binding proteins, including the M-type receptor (Rouault et al., 2007).

Several PLA₂s have also been implicated in various cancers. Disruption of the cPLA₂α gene decreases initiation and growth of intestinal tumors in Apc-mutated mice (Takaku et al., 2000) but increases the number of tumors in the carcino- 
gen azoxymethane-induced mouse model of colon cancer (Islesy et al., 2005). The gene coding for mouse group IIA (mGIIA) sPLA₂ was identified as a gene modifier that reduces the number of intestinal polyps in Apc(+/Min) mice, but the mechanism of action is still unclear (Fijneman and Cormier, 2008). Distinct roles have also been proposed for human group IIA sPLA₂ in various cancers (Sved et al., 2004, Cummings, 2007). Recent data have shown differential expression of sPLA₂s IGD, III, and IV but not X in human colon cancer (Murakami et al., 2005; Mounier et al., 2008). In Apc-/- and azoxymethane mouse models of colon cancer (Murakami et al., 2005; Mounier et al., 2008). In Apc-/- and azoxymethane mouse models of colon cancer, the expression of group X sPLA₂ may be increased in small intestinal tumors but not in colon tumors (Takaku et al., 2000; Islesy et al., 2005). In vitro, group CB, IA, and III sPLA₂s have been reported to stimulate cell proliferation and activation of MAP kinases in various cancer cells (Kinoshiba et al., 1997; Sved et al., 2004; Murakami et al., 2005). Finally, subcutaneous injection into nude mice of colon tumor cells overexpressing mGIIA sPLA₂ resulted in a 2.5-fold increase in tumor size (Belinsky et al., 2007).

Over the past few years, the group X sPLA₂ has emerged as the most potent sPLA₂ capable of hydrolyzing phosphatidyl-
choline and acting extracellularly on cellular membranes and noncellular phospholipid substrates such as lipopolysaccharides (Lambeau and Gelb, 2008). Consequently, group X sPLA₂ has been proposed to play a role in atherosclerosis (Lambeau and Gelb, 2008), asthma (Henderson et al., 2007), and colon cancer (Morioka et al., 2000).

The above-mentioned findings plus the fact that group X sPLA₂ is expressed at very high levels in human (Cuppillard et al., 1997; Morioka et al., 2000; Mounier et al., 2008) and mouse colon (Eerola et al., 2006; this study) prompted us to analyze the proliferative effect of group X sPLA₂ on various colon cancer cells, including the adenocarcinoma-derived mouse Colon-26 cancer cells. Among different mouse sPLA₂s, group X sPLA₂ was the most potent enzyme to stimulate cell proliferation. Using a combination of tools and methods, we found that its proliferative effect does not depend on binding to the M-type receptor or activation of cPLA₂α but rather on its intrinsic catalytic activity and ability to release free fatty acids and lysophospholipids, which most likely act in concert to stimulate cell proliferation.

Materials and Methods

Materials. The mouse adenocarcinoma cell line Colon-26 was obtained from Cell Lines Service (Heidelberg, Germany). AJ02-nm0 cells (Belinsky et al., 2007), YAMC cells (young adult mouse colon) (Whitehead et al., 1993), and Apc(+/Min) cells (Forest et al., 2003) were generous gifts from Drs. D. W. Rosenberg (University of Connecticut Health Center, Farmington, CT), R. H. Whitehead (Melbourne, Australia), and F. Pierre (Ecole Nationale Vétérinaire de Toulouse, Toulouse, France) and J. Menanteau (Ecole Nationale Vétérinaire de Nantes, Nantes, France), respectively. RPMI 1640 medium and Dulbecco’s modified Eagle’s medium (DMEM) were from Invitrogen (Cergy Pontoise, France). Fetal calf serum (FCS) was from Dominique Dutcher (Brumath, France). Oleic acid, linoleic acid, linolenic acid, oleoyl-L-α-lysophosphatidic acid sodium salt, 1-oleyl-sn-glycero-3-phosphocholine (PGE₂, AA, aspirin, ibuprofen, indomethacin, and FAF BSA were from Sigma-Aldrich (L’île d’Abeau Chesnes, France). [methyl-3H]Thymidine, [3H]oleic acid, and [3H]AA were from PerkinElmer Life and Analytical Sciences (Courtaboeuf, France). Recombinant murine interferon-γ was from PeproTech EC (London, UK). Insulin, transferrin, selenious acid (ITS) was from BD Biosciences (Le Pont-De-Claix, France). Phospho-
specific p42/p44 MAPK antibodies (clone E-4), p42/p44 antibodies, and cPLA₂α antibodies were from Santa Cruz Biotechnology, Inc. (Tebu-Bio SA, Le Perray en Yvelines, France). MK886, baicalein, and ATX antibody were from Cayman Chemicals (Interchim, Montluçon, France). Alkaline phosphatase-conjugated secondary antibodies were from Jackson ImmunoResearch Laboratories Inc. (Immuno-
tech, Marseille, France), and enhanced chemiluminescence (ECF) substrate was from GE Healthcare (Orcay, France). PGE₂ enzyme-linked immunosorbent assay kit was from Assay Designs (Eurome-
dex, Strasbourg, France). Prevalidated siRNAs were from QIAGEN (Courtaboeuf, France). Hoechst was from Invitrogen (Cergy Pontoise, France). LY294392 and Me-indoxam sPLA₂ inhibitors were prepared as described previously (Singer et al., 2002; Smart et al., 2006). Lynsing Matrix D beads were from Qiogene (Illkirch, France). Nucleosipin RNA extraction kit was from Macherey Nagel (Hoerdt, France). All recombinant mouse and human sPLA₂s used in this study and catalytically inactive hGX and mGX H48Q sPLA₂ mutants were prepared as described previously (Singer et al., 2002; Rouault et al., 2006, 2007).

sPLA₂ Expression in Mouse Intestinal Tract Analyzed by RT-qPCR. Tissue samples from duodenum, jejunum, ileum, and proximal and distal colon from two C57BL/6J female or three BALB/c female mice were snap-frozen in liquid nitrogen. Total RNA was isolated using the Nucleosipin RNA extraction kit with DNase I treatment and Lysing Matrix D beads to homogenize the samples. Reverse-transcription and reverse transcription quantitative poly-
mers chain reaction (RT-qPCR) were performed as described pre-
viously (Eerola et al., 2006). The abundance of each sPLA₂ mRNA target was calculated relative to the expression of glyceraldehyde-3-
phosphate dehydrogenase mRNA, which was used as a reference 
gene. The tissue distribution of sPLA₂s and cPLA₂α was further normalized to the lowest cycle threshold value accurately measured. A relative abundance of 1 (arbitrary unit = 1) was assigned to the expression level of the pancreatic group IB sPLA₂ mRNA in the colon, which was detected with cycle threshold values of 33 or higher. Standard deviation calculation was made according to User Bulletin 2 from Applied Biosystems, Foster City, CA.

Expression of mGX sPLA₂ by In Situ Hybridization. Samples for in situ hybridization were obtained from small and large intesti-
tines of C57BL/6J mice, fixed in 4% phosphate-buffered formalin, and embedded in paraffin. A 0.45-kilobase mGX sPLA₂ cDNA insert was cloned into the pReCMV vector (Invitrogen) and used as a template to prepare digoxigenin (DIG)-labeled antisense (test) and sense (control) RNA probes by in vitro transcription using a com-
mmercial kit (DIG RNA labeling kit; Roche Diagnostics, Meylan, France) according to the manufacturer’s instructions. Nonradioactive in situ hybridization has been described in detail previously (Haapamäki et al., 1999). In brief, 5-μm-thick paraffin sections were placed on silanated slides, dewaxed in xylene, and rehydrated in a graded alcohol series. Pretreatment was performed sequentially with 200 

m HCl for 20 min, 20 μg/ml proteinase K at 37°C for 15 min, 0.2% 
glycine in phosphate-buffered saline, pH 7.4, two times for 3 min
each, freshly prepared 0.4% acetic anhydride in 100 mM triethanol-
amine two times for 5 min each, and finally with ficin (Digest-All kit; Zymed Laboratories, South San Francisco, CA) at 37°C for 20 min. The sections were then blocked in hybridization solution without probe. The sections were washed in 2× standard saline citrate (SSC) for 5 min, in 0.2× SSC, 30% formamide at 37°C three times for 5 min each, and 2× SSC two times for 5 min each. The sections were blocked with 3% bovine serum albumin in Tris-buffered saline, and the hybridized probe was detected with alkaline phosphatase-labeled anti-DIG Fab-fragments (1/2000; Roche Diagnostics) using a substrate solution containing 0.18 mg/ml 5-bromo-4-chloro-3-indolyl-phosphate, 0.34 mg/ml 4-nitro blue tetra-
zolium phosphate (Roche Diagnostics), and 0.24 mg/ml levamisole (Vector, San Francisco, CA) in 100 mM Tris, pH 9.5, 100 mM NaCl, and 50 mM MgCl₂. The color reaction was allowed to develop at 4°C overnight and at 25°C for 6 h. The tissue sections were briefly counterstained with hematoxylin, with washed, and mounted in aqueous mounting medium.

Cell Culture. Colon-26 cells were grown at 37°C in DMEM/10% FCS, antibiotics, and 2 mM glutamine. AJ02-nm0 cells were grown in RPMI 1640 medium/10% FCS with antibiotics, glutamine, and supplementation with insulin (6.25 ng/ml), transferrin (6.25 μg/ml), and selenious acid (6.25 ng/ml; ITS). YAMC and Apc(/H9251)/Min) cells were found to contain two mutated pla2g2a alleles by polymerase chain reaction on genomic DNA (see Supplemental Data). YAMC cells were found to be heterogeneous for the pla2g2a gene, Colon-26 cells were found to be wild type, and AJ02-nm0 and Apc(+/Min) cells were found to contain two mutated pla2g2a alleles (Supplemental Fig. 1S).

[3H]Thymidine Incorporation and Cell Growth Assays. Colon-26 (5000 cells/well), YAMC (10,000 cells/well), AJ02-nm0 (5000 cells/well), and Apc(+/Min) (4000 cells/well) cells were plated into 48-well culture plates (Falcon; BD Biosciences Discovery Labware, Bedford, MA) that were precoated (Colon-26 and YAMC) or not (AJ02-nm0 and Apc(+/Min)) with rat tail collagen to ensure tight adhesion of cells. Fifty percent (Colon-26) or 80% [YAMC, AJ02-nm0, and Apc(+/Min)] confluent cells were made quiescent by incubating them for 24 h in serum-free medium with 0.02% FAF BSA at 37°C. It is important that we observed that the percentage of cell conflu-
ence can dramatically affect the ability of the sPLA₂ to trigger cell proliferation. Quiescent cells were treated with medium containing 0.02% FAF BSA and the effectors and were further incubated for 24 h (Colon-26) or 48 h [YAMC, AJ02-nm0, and Apc(+/Min)] at 37°C (Colon-26 and AJ02-nm0) or 33°C [YAMC and Apc(+/Min)]. During the last 4 h, [methyl-³H]Thymidine was added to wells for a final concentration of 1 μCi/ml, 1 μM unlabeled thymidine. In some experiments, cell supernatant of Colon-26 cells was collected for PGE₂ assays. The cells were then washed twice with ice-cold phosphate-buffered saline, incubated for 30 min in 5% ice-cold trichloroacetic acid, solubilized in 0.2 N NaOH, and analyzed for [³H]thymidine incorporation into DNA. For cell growth assays, cells were seeded as described above, starved at low density, and then treated with sPLA₂. Cells were dissociated and counted after trypan blue stain-
ing. Assays were done in triplicate.

Immunoblot Analysis. Colon-26 cells grown and made quiescent as described above were treated with effectors for various times, washed with PBS, lysed in hot Laemmli buffer, and boiled for 5 min at 95°C. Sonicated lysates were subjected to 10% SDS-polyacry-
mine gel electrophoresis and transferred to Immobilon-FL polyvi-
nylidene fluoride membrane (Millipore, Saint-Quentin-en-Yvelines, France). Membranes were blocked for 2 h at room temperature in NETQ buffer (150 mM NaCl, 5 mM EDTA, 50 mM Tris, pH 7.4, 0.05% Triton X-100, and 0.25% gelatin) and probed for 1 h with phospho-specific p42/44 MAPK antibodies or cPLA₂α antibodies (1/2000). Blots were washed five times with PBS/0.1% Tween 20, incubated with alkaline phosphatase-conjugated secondary antibod-
ies diluted in PBS/Tween (1/10,000), visualized by incubation for 5 min with ECF substrate, and scanned with a Pro-Xpress imager (excitation, 480 nm; emission, 590 nm). For Western blot of the mouse M-type receptor, cells were lysed in ice-cold 20 mM Tris, pH 7.4, 2 mM EDTA, and 1 mM phenylmethylsulfonyl fluoride and sonicated. Unboiled proteins (60 μg) prepared in nonreducing Lae-
mmi buffer were separated on a 7% SDS-polyacrylamide gel ele-
trohoresis gel and transferred to an Immobilon-FL membrane. The membrane was blocked with 5% blocking agent (GE Healthcare) dissolved in TBS/Tween (25 mM Tris, pH 7.8, 150 mM NaCl, and 0.15% Tween 20) for 1 h and then incubated with the anti-mouse M-type receptor antibody (1/5000 in TBS/Tween; Rouault et al., 2007) overnight at 4°C. The membrane was washed six times for 5 min each and incubated with secondary alkaline phosphatase goat anti-rabbit antibody (1/10,000 in TBS/Tween). The immunocomplex was visualized with ECF substrate as described above.

siRNA Transfection. siRNAs were designed using the program HP GenomeWide siRNA (HiPerformance Design Algorithm; QIAGEN). Best siRNAs respectively targeting the mouse M-type receptor and mouse cPLA₂α were selected from two QIAGEN-de-
signed siRNAs after validation by binding assays and immunocyto-
chemistry for the M-type receptor and Western blot analysis for cPLA₂α. For siRNA transfection, duplex siRNAs were first heated for 1 min at 90°C and cooled at 37°C for 30 min to allow for annealing. Cells were seeded on day 0 and transfected on day 1. Duplex siRNAs were preincubated with HiPerFect reagent (QIAGEN) and added dropwise to each well to a final concentration of 5 nM. On day 2, cells were made quiescent for 24 h before stimulation with sPLA₂ and additional Western blotting, binding assays, proliferation experi-
ments, or immunocytochemistry. Transfection efficiency was evalu-
ated by transfection with an Alexa546-coupled siRNA and fluores-
cence visualization with a microscope. The effect of the best-selected siRNAs for M-type receptor [forward, r(GGU ACA CUC GAU ACA UUA AddTdT; reverse, r(UUAUGUUACUGUGUACUUdTdT) and cPLA₂α [forward, r(GGA GAU UAA UGA GCU A)AddTdT; reverse, r(UAG CUC UUC AUU AAA CUC C UdTdT)] were compared with the effect of the nonrelevant GFP-directed siRNA (forward, GCA AGC UGA CCC UGA AGU UCA U; reverse, GAA CUG CAC CUG GCC G).

Binding Assays and Immunocytochemistry. The expression of the mouse M-type receptor in various cell lines was determined by binding experiments using iodinated OS₁ as described previously (Rouault et al., 2007). Cells were seeded in six-well plates, and 90% confluent cells were incubated for 1 h in 1 binding buffer (140 mM NaCl, 1 mM CaCl₂, 20 mM Tris, pH 7.4, and 0.1% BSA) at room temperature, with 200,000 cpm iodinated OS₁ in the absence (total binding) or presence (nonspecific binding) of 100 nM unlabeled OS₁. Cells were washed, lysed with 0.2 M NaOH, and analyzed for bound iodinated OS₁. The expression of the mouse M-type receptor after siRNA transfection was evaluated by immunocytochemistry as follows. Colon-26 cells were seeded in 24-well culture plates (4000 cells/well) on coverslips coated with rat tail collagen. Twenty-four hours after transfection with siRNA, the medium was changed, and cells were incubated in FCS-free medium for 24 h to mimic condi-
tions of proliferation assays. Cells were then fixed with 3.7% para-
formaldehyde for 25 min at room temperature and washed twice
with PBS. Free aldehyde groups were quenched with 50 mM NH4Cl in PBS, and cells were washed once with PBS and permeabilized with 0.5% saponin (two times for 5 min each). Nonspecific sites were blocked with 10% horse serum/0.05% saponin. Anti-mouse M-type receptor rabbit serum (Rouault et al., 2007) was diluted 1/1000 in 5% PBS/0.05% saponin. After three washes with PBS/0.05% saponin, fluorescein isothiocyanate-linked secondary antibodies were diluted 1/600 in 5% PBS/0.05% saponin and incubated for 30 min at room temperature. After three washes with PBS/0.05% saponin, nuclear sections were stained with Hoechst 33342 (5 μg/ml). Cells were washed once in PBS/0.05% saponin, twice in PBS, and once in H2O, and then they were mounted with Dako mounting medium (Dako, Trappes, France) and visualized under a microscope.

AA Release. AA release was performed under the same conditions as those used for proliferation assays. During the starvation time, cells were labeled with [3H]AA (0.2 μCi/well) in FCS-free DMEM/0.02% FAF BSA for 24 h at 37°C. Cells were washed twice with FCS-free medium/0.1% FAF BSA and incubated for the indicated periods in 200 μl of serum-free medium with 200 mM sPLA2. Cell-free supernatants and cell monolayers lysed in 400 μl of 0.2 N NaOH were submitted to scintillation counting. [3H]AA release is expressed as the percentage of cpm in cell supernatant versus total cpm (supernatant and cell lysate).

Eicosanoid Assays. Colon-26 cells (10^6 cells) were grown in six-well culture plates for 2 days, starved for 24 h, and stimulated with 24 h with 200 mM mgX sPLA2 in 1 ml of serum-free medium containing 0.02% FAF BSA. Total eicosanoid production in combined supernatant and cell pellet was assayed by high-performance liquid chromatography/mass spectrometry according to the procedure described previously (Kim et al., 2005; Henderson et al., 2007). For specific PGE2 assay, cell supernatants from proliferation assays were centrifuged, appropriately diluted (1/10–1/500), and PGE2 levels were determined using the PGE2 enzyme immunoassay kit (Assay Designs) according to the manufacturer’s instructions.

Lysophospholipid Analysis. Colon-26 cells (2 x 10^6 cells) were seeded in 10-cm Petri dishes and grown for 2 days. Then, cells were starved for 24 h in FCS-free medium/0.02% FAF BSA and stimulated with effectors in 4 ml of FCS-free medium/0.02% FAF BSA for 6 and 24 h. After incubation, the cell supernatants were collected and lyophilized. The cell monolayer (approximately 10^7 cells) was scraped in PBS, centrifuged, and stored as a dried pellet at −70°C. The following procedure was adapted from previously published methods of lysophosphatidic acid (LPA) analysis (Baker et al., 2001). Lyophilized supernatants were resuspended in 0.5 ml of Milli-Q water (Millipore), and cell pellets were thawed on ice in 0.5 ml of Milli-Q water. To each sample, 1.5 ml of 30 mM citric acid and 40 mM Na2HPO4, pH 4.0, was added and vortexed for 30 s. Samples were then spiked with 500 pmol of 17:0 LPA and 1 nmol of d31-16:0 LPC and extracted using 4 ml of 1-butanol followed by re-extraction with 2 ml of 1-butanol. Organic phases were dried in vacuo overnight. The resulting residues were reconstituted in 1.5 ml of CHCl3/MEOH (2:1), dried again, and finally reconstituted in 100 μl of CHCl3/MEOH (2:1). Liquid chromatography/tandem mass spectrometry analysis was performed using a Micromass Quattro Micro tandem quadrupole mass spectrometer (Waters, Milford, MA) coupled with a 2795 Chromatography System (Waters). Twenty microliters of concentrated lipid extract was injected onto a Luna silica column (5-μm particle size, 4.6 x 250 mm; Phenomenex, Torrance, CA) equilibrated with 65% solvent A (30:40, n-hexane/isopropanol) and 35% solvent B (30:60:15, n-hexane/isopropanol/H2O) at 30°C with a flow rate of 1 ml/min. After injection, 35% solvent B was maintained for 6 min. A gradient of 35 to 100% solvent B was run over 24 min. The gradient was held at 100% solvent B for an additional 20 min. Mass transitions, cone voltages, and collision energies are available on request. Transitions were monitored in the negative mode for the first 30 min, with 40-ms dwell times, and in the positive mode for the final 20 min, with 100-ms dwell times. In both modes, 10 ms interchannel and 100-ms interscan delays were used.

Statistics. Data are expressed as mean ± S.D. Statistical analyses were performed with Prism (GraphPad Software Inc., San Diego, CA) using Student’s t test and one-way analysis of variance (ANOVA), with Bonferroni adjustment for multiple comparisons. P values <0.05 were considered as statistically significant.

Results

Group X sPLA2 Is Expressed at High Levels in Mouse Colon. We have recently found that mgX sPLA2 is expressed at high levels in the small intestine and colon at both mRNA and protein levels (Eerola et al., 2006). To further analyze the expression of mgX sPLA2 and other sPLA2 genes in the different sections of the intestine of C57BL/6J mice, we have measured by RT-qPCR the relative abundance of mRNAs coding for the full set of mouse sPLA2s (except IIC), for cPLA2α (Fig. 1A) and for the M-type receptor (data not shown). Most sPLA2s were expressed in the different gut sections, but marked differences were found. Among catalytically active sPLA2-s, mgX sPLA2 is by far the most highly expressed enzyme in the intestine of C57BL/6J mice, with highest expression in the distal colon. For example, its level of expression in the colon is approximately 10-fold higher than that of mgV sPLA2. It is interesting that mgX sPLA2 mRNA levels vary along the intestine with up to 120-fold increase in expression from duodenum to distal colon. Second, the levels of mRNA for the catalytically inactive group XIIB sPLA2-like protein are similar to those of mgX sPLA2, but the expression pattern is the inverse. Third, group IID, IIF, III, and XIIA sPLA2-s are expressed in the intestine at significant but low levels, and group IB and IIE sPLA2-s are barely detectable. Expression of group XIIB sPLA2 is rather constitutive all along the intestine. Conversely, that of group IIF, III, and V sPLA2-s increases, whereas that of group IID sPLA2 decreases when moving from the proximal to the distal part of the gut. Fourth, the expression of cPLA2α is 5-fold lower than that of mgX sPLA2, but follows a similar gradient of expression along the intestine. Finally, the expression of the M-type receptor was low along the gut sections (data not shown), but, interestingly, it follows the same gradient of expression of mgX sPLA2.

It is important that C57BL/6J mice have a disrupted mGIIA sPLA2 gene (MacPhee et al., 1995), which probably explains the low and aberrant amount of mRNA observed in the small intestine (Fig. 1A). When RT-qPCR assays for the different sPLA2-s were done as described above on intestinal sections from BALB/c mice expressing a functional gene, the expression of mGIIA sPLA2 in jejunum and ileum was found to be much higher than that of all other sPLA2-s, including mgX sPLA2 (Fig. 1B). However, the mRNA levels of mGIIA were fairly low in proximal and distal colon, where mgX sPLA2 remained the most abundantly expressed sPLA2 (Fig. 1B).

To determine more precisely the cellular site of mgX sPLA2 expression in small and large intestine, we performed in situ hybridization. Figure 1, C and D, shows strong labeling of columnar epithelial cells of mucosal villi, Paneth cells in the crypts of Lieberkühn, and ganglion cells of the myenteric plexus between smooth muscle cell fibers. Figure 1E illustrates the absence of labeling from a section of small intestine hybridized with sense probe. Figure 1F shows labeling of epithelial cells in the mucosal glands of the large intestine adjacent to empty-looking goblet cells.
We also analyzed the expression level of group X sPLA₂ in the intestine sections using specific sets of PLA₂ primers. To facilitate the analysis, we used RT-qPCR in both normal colon and colon adenocarcinoma (Cupillard et al., 1997; Morioka et al., 2000; Takaku et al., 2000; Österström et al., 2002; Ilalley et al., 2005; Mounier et al., 2008), we compared the expression between the different sPLA₂s, the data were normalized to glyceraldehyde-3-phosphate dehydrogenase mRNA, which was used as a reference gene, and then expressed relative to the lowest expression level that can be accurately measured in our RT-qPCR assay conditions [i.e., the expression of pancreatic group IB sPLA₂ in the colon (relative abundance of 1 (arbitrary unit = 1))]. Note that two different ordinate axes have been used in B. mGIIE sPLA₂ could not be detected in all intestine sections from BALB/c mice (C–F). In situ hybridization of mGX sPLA₂ in small intestine (ileum) showing labeling of columnar epithelial cells in mucosal villi (V), Paneth cells (P), and ganglion cells (G) of the myenteric plexus. Hybridized with antisense probe. D, absence of reaction product from ileal tissue when hybridized with sense probe. E, in situ hybridization of mGX sPLA₂ in large intestine (occum) showing labeling in epithelial cells. Hybridized with antisense probe.

Expression of the different mouse sPLA₂s in the intestine of Expression of group X sPLA₂ in the RNA level by RT-qPCR and in situ hybridization of mGX sPLA₂, A and B, RT-qPCR in C57BL/6J (A) and BALB/c (B) mouse intestine sections using specific sets of PLA₂ primers. To facilitate the comparison of expression between the different sPLA₂s, the data were first normalized to glyceraldehyde-3-phosphate dehydrogenase mRNA, which was used as a reference gene, and then expressed relative to the lowest expression level that can be accurately measured in our RT-qPCR assay conditions [i.e., the expression of pancreatic group IB sPLA₂ in the colon (relative abundance of 1 (arbitrary unit = 1))]. Note that two different ordinate axes have been used in B. mGIIE sPLA₂ could not be detected in all intestine sections from BALB/c mice (C–F). In situ hybridization of mGX sPLA₂ in small intestine (ileum) showing labeling of columnar epithelial cells in mucosal villi (V), Paneth cells (P), and ganglion cells (G) of the myenteric plexus. Hybridized with antisense probe. D, absence of reaction product from ileal tissue when hybridized with sense probe. E, in situ hybridization of mGX sPLA₂ in large intestine (occum) showing labeling in epithelial cells. Hybridized with antisense probe.

Group X sPLA₂ Stimulates the Proliferation of Mouse Colon Cells. Because group X sPLA₂ is expressed in both normal colon and colon adenocarcinoma (Cupillard et al., 1997; Morioka et al., 2000; Takaku et al., 2000; Österström et al., 2002; Ilalley et al., 2005; Mounier et al., 2008), we analyzed the effect of exogenous recombinant group X sPLA₂ on the proliferation of four different mouse colon cell lines (YAMC, Apc⁺/Min, AJ02-nm0, and Colon-26). Both mGX and hGX sPLA₂ were able to stimulate [³H]thymidine incorporation in the four tested cell lines (Fig. 2). At 200 nM, group X sPLA₂ increased [³H]thymidine incorporation by up to 3-fold, depending on the cell line and conditions of stimulation. This proliferative effect is important because it corresponds to approximately 50% of the maximal effect observed with fetal calf serum. We decided to use the Colon-26 cell line for the experiments described below because these cells are adenocarcinoma-derived and were easier to grow and transfect than the other cell lines. We also found that these cells do not express mGX sPLA₂ at the mRNA level by RT-qPCR and at the protein level by both a highly specific and sensitive time-resolved fluorimunoassay (Eerola et al., 2006) and Escherichia coli sPLA₂ enzymatic assays (data not shown). Compared with various mouse sPLA₂s, we found that group X sPLA₂ has the strongest mitogenic capacity on Colon-26 cells (Fig. 3A) and dose-dependently induced cell proliferation (Fig. 3B). mGV and mGIIE sPLA₂ also significantly induced cell proliferation, whereas the other sPLA₂s were relatively poor inducers. To further demonstrate that group X sPLA₂ can trigger cell proliferation, we measured the growth of Colon-26 cells treated with mGX and hGX sPLA₂s by direct cell counting. As indicated in Fig. 3C, cells proliferated more rapidly over time in the presence of group X sPLA₂. In accordance with their proliferative effects, both mGX and hGX sPLA₂s were capable of inducing a time-dependent phosphorylation of p42/44 MAPK, with a sustained induction at 4 h (Fig. 3D).

Group X sPLA₂ Stimulates the Proliferation of Colon-26 Cells via Its Catalytic Activity and Independently of Binding to the M-Type Receptor. We next analyzed the molecular mechanisms involved in the proliferative effect of group X sPLA₂. To evaluate the role of sPLA₂ enzymatic activity, we first used the specific sPLA₂ inhibitors Me-indoxam (Singer et al., 2002) and LY329722 (compound B in Smart et al., 2006), which inhibit the catalytic activity of mGX sPLA₂, with IC₅₀ values of 500 and 75 nM, respectively. Both inhibitors clearly suppressed the proliferation and MAP kinase phosphorylation induced by mGX sPLA₂ (Fig. 4A and B). We then tested the mitogenic activity of catalytically inactive mutants of mGX and hGX sPLA₂s as well as of OS₂, a snake venom sPLA₂ with high catalytic activity on phosphatidylycholine (Rouault et al., 2006). The H48Q mutants of mGX and hGX sPLA₂s have less than 0.1% of wild-type catalytic activity (data not shown), and the D49K mutant of OS₂ is fully inactive (Rouault et al., 2006). Results shown in Fig. 4C further demonstrate that the catalytic activity of group X sPLA₂ is required for its mitogenic effect. Finally, we evaluated the role of cPLA₂α and found that the mitogenic effect of group X sPLA₂ on Colon-26 cells does not depend on cPLA₂α activation because siRNA silencing of cPLA₂α efficiently suppressed protein expression but has little effect on the fold increase factor of [³H]thymidine incorporation induced by hGX sPLA₂ (Fig. 4D). However, our results also suggest that cPLA₂α is involved in the proliferation of Colon-26 in the absence of exogenously added sPLA₂ because siRNA silencing slightly decreased the basal level of incorporation of [³H]thymidine (Fig. 4D).

Because the M-type sPLA₂ receptor was proposed previ-
ously to play a role in cell proliferation (Kinoshita et al., 1997) and because both mGX and hGX sPLA2s bind to the mouse M-type receptor (Rouault et al., 2007), we sought to determine whether this receptor is expressed in the above-mentioned different colon cells and plays a role in the proliferative effect of group X sPLA2. We first screened the four colonic cell lines for the expression of the receptor using iodinated OS1, the snake venom sPLA2 that binds to the M-type receptor with very high affinity and specificity (Rouault et al., 2007). Apc(+/Min) cells were found to express high levels of the receptor, whereas Colon-26 and YAMC cells express low but clearly detectable levels (Fig. 5A). Conversely, AJ02-nm0 cells do not express the receptor. Scatchard plot analysis indicated that Colon-26 cells contain a single population of binding sites for iodinated OS1, with a $K_d$ value of 90 pM and a maximal binding capacity ($B_{max}$) of 0.016 pmol/mg total protein (data not shown). These binding data were confirmed by Western blot analysis (Fig. 5B) and RT-qPCR analysis (data not shown). The fact that group X sPLA2 stimulates the proliferation of AJ02-nm0 cells (Fig. 5A), which do not express the M-type receptor (Fig. 5A), represents a first indication that the M-type receptor is not required for the proliferative effect of group X sPLA2. This view was further supported by siRNA experiments targeting the M-type receptor expressed in Colon-26 cells. As shown in Fig. 5C, we observed by both binding assays and immunocytochemistry that the expression of the M-type receptor was dramatically reduced at 48 and 72 h after siRNA transfection, at the time window where the Colon-26 cells were stimulated with exogenous hGX and mGX sPLA2s for cell proliferation (Fig. 5D). The specificity of siRNA silencing of the receptor was validated using a nonrelevant GFP-siRNA that did not suppress M-type receptor expression (Fig. 5C). The capacity of group X sPLA2 to increase Colon-26 proliferation was unaffected by knocking-down the M-type receptor, indicating that receptor binding is not required for the proliferative effect of group X sPLA2 (Fig. 5D).

**Group X sPLA2 Produces Various Lipid Mediators from Colon-26 Cells.** Based on the above-mentioned findings, we sought to determine whether group X sPLA2 can release various lipid mediators, including free AA, eicosanoids, and lysophospholipids from Colon-26 cells. As found previously in other cells (Singer et al., 2002), exogenously added mGX, but not mGIB, mGIIA, mGV, and mGXIIA sPLA2s, can release significant amounts of free AA from Colon-26 cells radiolabeled with $[^3]$H]AA (Fig. 6A). Similar results were obtained when cells were radiolabeled with $[^3]$H]oleate (data not shown). We also monitored the ability of group X sPLA2 to produce various eicosanoids derived from free AA by combined liquid chromatography/mass spectrometry after stimulation of Colon-26 cells by the sPLA2 for 6 h (data not shown) and 24 h (Fig. 6B). Group X sPLA2 increases the production of several eicosanoids, with PGE2 being by far the most prominent eicosanoid product, with up to 74 ng/10^6 cells. PGD2, PGE2, and 5-, 12-, and 15-hydroxyeicosatetraenoic acids, and to a lesser extent leukotriene C4 and leukotriene E4, are also produced by group X sPLA2 at 6 h (data not shown) and 24 h.

**Fig. 2.** Effect of group X sPLA2 on $[^3]$H-thymidine incorporation in mouse colon cell lines. After starvation for 24 h at 37°C in serum-free and additive-free medium, all cell lines were incubated in the presence of mGX and hGX sPLA2s (200 nM) or FCS under the following conditions: Colon-26 were grown at 37°C for 24 h in DMEM and 0.02% FAF BSA (A); AJ02-nm0 cells were grown at 37°C for 48 h in RPMI 1640 medium and 0.02% FAF BSA (B); Apc(+/Min) cells were grown at 33°C in DMEM, 0.02% FAF BSA, and 10 ng/ml recombinant EGF (C); and YAMC cells were incubated at 33°C for 48 h in serum-free RPMI 1640 medium and 0.02% FAF BSA (D). $[^3]$H-thymidine was added during the last 4 h of sPLA2 stimulation, and cells were processed as described under Materials and Methods. The increase in $[^3]$H-thymidine incorporation induced by sPLA2s was compared with untreated cells (−) and FCS-treated cells. In all panels, values are representative of at least two experiments performed in triplicates. Significant differences between untreated and sPLA2-treated cells were found (**, $P < 0.001$; Student’s t test).
In parallel experiments, we monitored the ability of group X sPLA2 to release various lysophospholipids, namely, LPA, LPC, LPE, lysophosphatidylglycerol, lysophosphatidylinositol, and lysophosphatidylserine. We found that group X sPLA2 can release a variety of lysophospholipids with different fatty acids at the sn-1 position from Colon-26 cells at 6 h (Fig. 7) and 24 h (data not shown). The lysophospholipids produced in highest quantities were LPC and LPE, a result that is in accordance with the relative abundance of these lipids in cell membranes and the ability of group X sPLA2 to efficiently hydrolyze zwitterionic phospholipids (Singer et al., 2002; Mitsuishi et al., 2007). For example, treatment with group X sPLA2 for 6 h led to 25- and 36-fold increases in the release of total LPC and LPE compared with untreated cells (Fig. 7A). It is important that we also found that group X sPLA2 increases by 3.7-fold the overall production of LPA. The detailed data for production of the different acyl chain species for LPC and LPA are shown in Fig. 7B. They indicate, for example, 41- and 26-fold increases for the two main products oleoyl-LPC and palmitoyl-LPC.

Effects of Various Lipid Mediators Released by Group X sPLA2 on the Proliferation of Colon-26 Cells. The above-mentioned results led us to analyze the proliferative effect of each type of lipid mediators that are generated by the action of group X sPLA2. Because LPC can be further converted into LPA by autotaxin (Ferry et al., 2008), we also tested the effect of LPA on cell proliferation. As shown in Fig. 8, free AA, LPC, and LPA can all stimulate the incorporation of [3H]thymidine in Colon-26 in a dose-dependent manner. Various other fatty acids were also able to stimulate cell proliferation, although with a lower efficacy than free AA (data not shown). For example, oleic (50 μM), linoleic (50 μM), and α-linolenic (20 μM) acids could stimulate the incorporation of [3H]thymidine by factors of 1.4, 1.6, and 2.1, respectively. Furthermore, both AA and LPA were able to phosphorylate p42/44 MAPK (Fig. 8). To test the hypothesis that PGE2 production could explain at least in part the mitogenic effects of group X sPLA2, we treated Colon-26 cells with four nonselective or COX-2-selective inhibitors (aspirin, indomethacin, ibuprofen, and rofecoxib). We found that all four inhibitors were able to dramatically reduce PGE2 production (Fig. 9A) but were unable to suppress the cell proliferation induced by mGX sPLA2 (Fig. 9B). It is interesting that the COX inhibitors were also ineffective at suppressing the proliferative effect of free AA (Fig. 9C). Furthermore, we...
found that exogenously added PGE₂ (1–1000 nM) did not induce cell proliferation (data not shown). Finally, incubation of Colon-26 cells with MK886 or baicalein, that inhibit 5- and 12-LOXs, respectively, did not alter the effect of mGX sPLA₂ on cell proliferation (Fig. 9D). Together, these results indicate that although group X sPLA₂ can release large amounts of PGE₂ (and small amounts of other eicosanoids) from Colon-26 cells (Fig. 6), this PGE₂ is at best a minor effector of the sPLA₂ proliferative effect. Rather, the sPLA₂ proliferative effect seems to be due to the combined production and direct action of free AA and lysophospholipids, including LPC and LPA.

### Discussion

This work shows for the first time that group X sPLA₂ can stimulate the proliferation of colon cancer cells and activate the phosphorylation of p42/44 MAPK via its potent catalytic activity and ability to produce various lipid mediators. We found that the proliferative effect of group X sPLA₂ was much higher than that of other mouse sPLA₂s. This effect was as strong as or higher than that of sPLA₂-IB, -IIA, or -IIIB acting on different cell types (Kinoshita et al., 1997; Sved et al., 2004; Murakami et al., 2005; Belinsky et al., 2007). mGV and mGIIA sPLA₂s, but not other enzymes, were also able to promote cell proliferation. Except for mGIIB, the ability of the different sPLA₂s to stimulate proliferation seems to be linked to their direct capacity to hydrolyze phosphatidylcholine and release AA and LPC from cells (Singer et al., 2002; Masuda et al., 2005). As found previously in LNCaP prostatic cancer cells (Sved et al., 2004), it is possible that the proliferative effects mediated by mGIIB sPLA₂ is due to activation of cPLA₂, but we did not further investigate this point.

The proliferative effect of various sPLA₂s has been associated with binding to the M-type receptor (Kinoshita et al., 1997) or enzymatic activity of sPLA₂ and subsequent activation of cPLA₂ (Sved et al., 2004; Murakami et al., 2005). We found here that the proliferative effect of group X sPLA₂ fully depends on its intrinsic catalytic activity and not on binding to the M-type receptor or cPLA₂ activation. First, two specific sPLA₂ inhibitors suppressed the proliferative effect of group X sPLA₂. Second, the H48Q mutants of mGV and hGV sPLA₂s and the D49K mutant of OS₂ (Rouault et al., 2006), which have less than 0.1% of wild-type enzymatic activity but still bind with high affinity to the M-type receptor (data not shown; Rouault et al., 2006), had no or very modest effects on cell proliferation. Third, mGV and hGV sPLA₂s can trigger

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**Fig. 4.** The proliferative effect of group X sPLA₂ on Colon-26 cells is dependent on sPLA₂ catalytic activity. A and B, effect of sPLA₂ inhibitors on the proliferative effect of mGX sPLA₂ and MAP kinase activation. Me-indoxam (10 μM) and LY329722 (5 μM) were preincubated with mGX sPLA₂ (200 nM) in DMEM and 0.02% FAF BSA for 20 min before incubation with Colon-26 cells. Under untreated conditions, incorporation of [3H]thymidine was 54,669 dpm; mGX sPLA₂ alone but not mGX sPLA₂ preincubated with Me-indoxam or LY329722 significantly stimulated [3H]thymidine incorporation versus untreated cells (⁎⁎, P < 0.001; one-way ANOVA, with Bonferroni adjustment). Although significant (P < 0.01), the effect of Me-indoxam was probably due to a toxic effect that was not observed with LY329722 (P > 0.05). C, effect of catalytically inactive mGX H48Q, hGX H48Q, and OS₂ D49K sPLA₂s (200 nM) on the proliferation of Colon-26 cells (⁎⁎, P < 0.001; one-way ANOVA, with Bonferroni adjustment). Under untreated conditions, the incorporation of [3H]thymidine was 74,679 dpm. D, effect of cPLA₂ siRNA silencing on the proliferative effect of hGX sPLA₂. Two different commercially available siRNAs (QIAGEN) targeting mouse cPLA₂ were tested for silencing by Western blotting (inset), and the effect of the best cPLA₂ siRNA (n°1) versus an irrelevant GFP-directed siRNA was tested for incorporation of [3H]thymidine triggered by hGX sPLA₂. Colon-26 cells were transfected with siRNA on day 1, starved the next day for 24 h, and then assayed for cPLA₂ expression or incorporation of [3H]thymidine triggered by hGX sPLA₂ (200 nM). Under untreated conditions, the incorporation of [3H]thymidine was 45,825 dpm. The differences in incorporation of [3H]thymidine in the absence and presence of sPLA₂ or between untreated cells transfected with the two siRNA are statistically significant (⁎⁎, P < 0.01; Student’s t test). In all panels, data are representative of two experiments.
proliferation on cells expressing or not the M-type receptor. Fourth, siRNA silencing of the M-type receptor in Colon-26 cells did not affect group X sPLA₂-induced cell proliferation. Finally, because of the possible role of cPLA₂ (Sved et al., 2004), we analyzed its role by siRNA silencing and found that group X sPLA₂ stimulates cell proliferation independently of cPLA₂α.

In line with the role of group X sPLA₂ enzymatic activity in proliferation of Colon-26 cells, we found that group X sPLA₂ can release relatively large quantities of free AA, LPC, and LPA, which are all able to stimulate cell proliferation and activation of MAP kinase phosphorylation. AA is the precursor of numerous eicosanoids, including PGE₂ and leukotriene D₄, which contribute to colon cancer cell proliferation (Wang and Dubois, 2006). Group X sPLA₂ increases the production of various prostaglandins, leukotrienes, and hydroxyeicosa- tetraenoic acids, but these mediators do not explain the sPLA₂ mitogenic effect (Fig. 9). It has been shown that Colon-26 cells express the four PGE₂ receptors (EP1–4) and that PGE₂ is mitogenic on these cells (Pozzi et al., 2004). That exogenous PGE₂ or sPLA₂-produced PGE₂ did not trigger the proliferation of our Colon-26 cells may be linked to the high basal concentrations of PGE₂ (approximately 30 nM) that already saturate EP receptors. Treatment of Colon-26 cells with 5- and 12-LOX inhibitors also does not prevent the mitogenic effect of sPLA₂ or free AA, indicating that the proliferative effect of the sPLA₂ is in part mediated by free AA without conversion into prostaglandins or leukotrienes.

Based on lysophospholipid analyses of cells treated with hGX sPLA₂ (Fig. 7), we could estimate that the sPLA₂ can release LPA and LPC at nanomolar and low micromolar concentrations, respectively. Although these concentrations are lower than those of exogenously added LPA and LPC (Fig. 8), lysophospholipids produced endogenously may be more effective because they are released locally at the cell surface. The proliferative effect of LPA is probably explained by its binding to receptors LPAR1, LPAR2, and LPAR4, which are expressed in Colon-26 cells (Supplemental Fig. 3S). Whether the proliferative effects of exogenous LPC is due to conversion into LPA by ATX and subsequent binding to LPA receptors or to a direct action via other G protein-coupled receptors and/or transactivation of tyrosine kinase receptors (Ikeno et al., 2005; Fujita et al., 2006) remains an open question. We analyzed Colon-26 cells for expression of ATX at the mRNA and protein levels (Supplemental Fig. 3S). Although detectable amounts of ATX mRNA were observed by RT-qPCR analysis, no ATX protein could be detected by Western blot analysis using two distinct antibodies and a radioactive lysophospholipase D enzymatic assay (Ferry et al., 2008). We also analyzed the effect of the recently described ATX inhibitor S32826 (Ferry et al., 2008) on the production of LPA triggered by hGX sPLA₂ in conditions identical to those used on incorporation of [³H]thymidine triggered by hGX sPLA₂. Colon-26 cells were transfected with siRNAs on day 1, starved on day 2 for 24 h, and then stimulated on day 3 for 24 h in the absence (−) or presence of mgX and hGX sPLA₂ (200 nM). [³H]Thymidine was added during the last 4 h of sPLA₂ stimulation, and cells were processed as described under Materials and Methods. Under untreated conditions, the incorporation of [³H]thymidine was 42,395 dpm. Incorporation of [³H]thymidine in sPLA₂-treated cells over untreated is statistically significant (***, P < 0.05; Student’s t test), but no significant difference was found between the two siRNAs. Data are representative of two experiments.
The ATX inhibitor S32826 had no effect on LPA production at 1 and 10 μM (Supplemental Fig. 3S), further indicating that Colon-26 cells do not express ATX. It thus seems unlikely that the LPA produced by group X sPLA₂ comes from released LPC being converted into LPA by ATX. Although we cannot rule out the presence of another lysophospholipase D-like activity in Colon-26 cells, our finding raises the possibility that group X sPLA₂ directly hydrolyzes cellular PA to generate LPA by acting either at the plasma membrane or after shuttling into intracellular compartments enriched into PA. It is noteworthy that the total amount of LPA released by group X sPLA₂ was less than 3% of the LPC produced (Fig. 7A), suggesting that the sPLA₂ may have

Fig. 6. Release of AA by various mouse sPLA₂s and production of eicosanoids triggered by mGX sPLA₂ from Colon-26 cells. A, release of AA by various mouse sPLA₂s was measured on quiescent Colon-26 cells prelabelled with [3H]AA for 24 h as described under Materials and Methods. Mouse recombinant sPLA₂ (200 nM) were incubated with cells for 6 h in DMEM with 0.02% FAP BSA. [3H]AA release is expressed as the percentage of radioactivity present in cell medium relative to the total radioactivity incorporated into cells. sPLA₂-X-treated cells release significantly more [3H]AA (**, P < 0.001; one-way ANOVA, with Bonferroni adjustment). A representative experiment of at least two experiments is shown. B, eicosanoid production triggered by mGX sPLA₂. Colon-26 cells (10⁶ cells) were starved for 24 h and treated with mGX sPLA₂ (200 nM) for 24 h in DMEM and 0.02% BSA. The cell medium and the cell monolayer were collected and analyzed for eicosanoid production by liquid chromatography/mass spectrometry as described under Materials and Methods. A representative experiment of two experiments is shown.

Fig. 7. Release of lysophospholipids by group X sPLA₂ from Colon-26 cells. A, total amount of LPX acyl chain species released by hGX sPLA₂. B, release of the different LPC and LPA acyl chain species by hGX sPLA₂. Colon-26 cells were starved for 24 h and treated with hGX sPLA₂ (200 nM) for 6 h in DMEM and 0.02% BSA. The cell medium and the cell monolayer were collected, extracted with organic solvent, combined, and analyzed for lysophospholipid production by liquid chromatography/mass spectrometry as described under Materials and Methods. A representative experiment of two experiments is shown.
access to little amounts of PA substrate. We have shown that group X sPLA₂ can efficiently hydrolyze PA in mixed phospholipid vesicles (Singer et al., 2002), but there has been so far no report on the capacity of this enzyme to release LPA from cells. Only a few studies have suggested that group IIAsPLA₂ may be involved in LPA release (Fourcade et al., 1995; Snitko et al., 1997). Together, the above-mentioned data indicate that the proliferative effects of group X sPLA₂ are likely to be dependent on the combined production of both free fatty acids, including AA, and lysophospholipids, including LPC and LPA.

We also found by RT-qPCR that group X sPLA₂ is expressed at very high levels in the small intestine and more particularly in the colon of C57BL/6J and BALB/c mice. The cellular sites of mGX sPLA₂ expression include columnar epithelial cells, Paneth cells, and ganglion cells. It should be noted that the very low level of expression measured for mGIAsPLA₂ mRNA in C57BL/6J mice is due to the natural disruption of the pla2g2a gene. In the small intestine of BALB/c mice harboring a functional gene, the expression level of mGIAsPLA₂ was much higher than those of group X sPLA₂ and other sPLA₂s (Fig. 1B). However, the expression of mGIAsPLA₂ dramatically decreased in the colon, whereas that of mGX sPLA₂ increased, making mGX sPLA₂ among the most highly expressed sPLA₂ gene in the proximal and more particularly in the distal colon of both C57BL/6J and BALB/c mice (Fig. 1, A and B). In good accordance, group IIA and X sPLA₂s are also the most highly expressed sPLA₂s in human colon (Mounier et al., 2008). These observations raise the question as to whether the two enzymes play redundant or divergent functions within the small intestine and colon. Based on their unique molecular and functional features, it is likely that the two sPLA₂s have distinct roles. First, group IIA sPLA₂ is a very basic protein, whereas group X is the most acidic sPLA₂ (Lambeau and Gelb, 2008). Second, group IIA sPLA₂ binds tightly to anionic phospholipid interfaces but not zwitterionic interfaces, whereas group X sPLA₂ shows similar binding (Singer et al., 2002). Third, their expression in normal intestine is different in both mouse and human species (see above). Fourth, there is a strong up-regulation of group IIA sPLA₂ at either mRNA or protein levels in inflammatory bowel diseases and probably in mouse and human colon tumors (Isley et al., 2005; Cummings, 2007). Alternatively, there is no conclusive evidence for up-regulation of group X sPLA₂ in the small intestine and colon of ApcΔ716 (Takaku et al., 2000) and ApcΔ14 mice (Supplemental Fig. 2S), two models of human familial adenomatous polyposis. No up-regulation was observed in the colon tumors of mice treated with the carcinogen azoxymethane (Isley et al., 2005) and in human colorectal adenocarcinomas (Osterstroem et al., 2002; Mounier et al., 2008). This is probably reminiscent of the fact that the catalytic activity of group X sPLA₂, but not group IIA, may be regulated at the post-translational level by maturation of its N-terminal propeptide by a still poorly defined proteolytic mechanism (Cupillard et al., 1997; Masuda et al., 2005). Finally, group IIA sPLA₂ probably plays an important antibacterial role in the intestine and is accordingly highly expressed in Paneth cells of the small intestine and epithelial cells of colonic mucosa (Nevalainen et al., 2008). Its antitumoral role in colon cancer has been proposed to be linked to this antibacterial activity, but the mechanism is still enigmatic (Fijneman and Cormier, 2008). Conversely, group X sPLA₂ has been proposed to play a central role in AA release and PGE₂ production in the colon but not in the small intestine where cPLA₂α would play the major role (Morioka et al., 2000; Takaku et al., 2000). This hypothesis fits well with our RT-qPCR data showing a 5-fold

Fig. 8. Effect of AA, LPA, and LPC on Colon-26 proliferation. Effect of free AA (A), LPA (B), and LPC (C) on [³H]thymidine incorporation and p42/44 MAPK phosphorylation. Quiescent Colon-26 cells were stimulated with various concentrations of the different lipid mediators, and [³H]thymidine incorporation was evaluated as described under Materials and Methods. Under untreated conditions, the incorporation of [³H]thymidine was 42,493 dpm. The effect of AA and LPA on p42/44 phosphorylation was measured by incubating Colon-26 cells for the indicated periods at 37°C with 20 μM AA and 5 μM LPA as described in Fig. 3 legend. A representative experiment of at least two separate experiments is shown (∗, P < 0.05; **, P < 0.001 over untreated cells; one-way ANOVA, with Bonferroni adjustment).
higher expression of group X sPLA2 over cPLA2α in the colon but not in the small intestine (Fig. 1). Whether autoxidation and/or LPA receptors, which are overexpressed in cancer (Parrill and Baker, 2008), contribute to the effects of group X sPLA2 is unknown. That group X sPLA2 can produce various lipid mediators suggests that this enzyme may also regulate other key events in tumorigenesis (Wang and Dubois, 2006). Finally, the presence of mGX sPLA2 mRNA in ganglion cells may suggest additional functions in the enteric nervous system, including neurotogenesis (Masuda et al., 2005), peristaltic reflex, or nociception.

In conclusion, we have shown that group X sPLA2 can stimulate the in vitro proliferation of colon cancer cells via its enzymatic activity and production of free AA and lysophospholipids. The high expression of group X sPLA2 in normal colon and tumors in both mouse and human species and its ability to produce various lipid mediators suggests that this enzyme plays a similar role in vivo.

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Address correspondence to: Dr. Gérard Lambeau, Institut de Pharmacologie Moléculaire et Cellulaire, Université de Nice Sophia Antipolis et Centre National de la Recherche Scientifique, 680 route des Lucioles, Sophia Antipolis, 06560 Valbonne, France. E-mail: lambeau@ipmc.cnrs.fr.
Group X Phospholipase A₂ Stimulates the Proliferation of Colon Cancer Cells
by Producing Various Lipid Mediators


Institut de Pharmacologie Moléculaire et Cellulaire, Université de Nice Sophia Antipolis et Centre National de la Recherche Scientifique, Valbonne 06560, France (F.S., I.J., E.B., C. P., C.M.M., G.L.)
Departments of Chemistry and Biochemistry, University of Washington, Seattle, Washington 98195, USA (J.G.B., M.H.G.)
Department of Pathology, University of Turku, Kiinamyllynkatu10, FIN-20520, Turku, Finland (K.A.T., V.J.O.L., T.J.N.)
Supplemental methods

Genotyping of *pla2g2a* gene in mouse colon cell lines. Mouse colon cell lines (2 x 10^6 cells) and 3-mm tail tissue from C57BL/6J and Balb/C mice were lysed in 150 µl of tissue lysis buffer comprising 100 mM Tris-HCl (pH 8.0), 200 mM NaCl, 5 mM EDTA, 0.2% SDS, and 0.2 mg/ml proteinase K at 55°C overnight. DNA was purified from the clarified lysates by precipitation with 1 volume of isopropanol. Approximately 0.1 µg of genomic DNA was subjected to PCR amplification with the Uptitherm Taq DNA polymerase (Uptima, Interchim) and a set of primers of mGIIA-Com-F (5’- CAGAGCTGACAGCATGAAGGTCCTC-3’) and mGIIA-WT-R (5’- TCTGTGGCATCCTTGGGGGAT-3’) for the mGIIA WT allele or a set of mGIIA-KO-F (5’- GCCCTGGGTTGGCAAGGGATT-3’) and mGIIA-Com-R (5’- AGGCCGCTTGTAGCAACAGTTCAGAG-3’) for the mGIIA KO allele. The PCR conditions were 95°C for 2 min followed by 35 cycles of 95°C for 30 s, 64°C (for WT) or 70°C (for KO) for 30 s and 72°C for 60 s on a thermal cycler (Hybaid). The PCR products were analyzed by 1.2% agarose gel electrophoresis with ethidium bromide staining. mGIIA-Com-F and mGIIA-WT-R give the wild-type band of 374 bp, and mGIIA-KO-F and mGIIA-Com-R give the mutated band of 310 bp.

Expression of sPLA_2s in Apc^{Δ14} mice and azoxymethane-treated SWR/J and A/J mice. Apc^{Δ14} mice were obtained from Dr. Christine Perret (Paris, France). Total RNA was isolated from tumors and adjacent tissues from the colon and ileum of 3.5 months old mice. RNA extraction was performed as described in the main methods section. Total RNA from the distal colon and tumors of SWR/J and A/J mice injected with azoxymethane were prepared as described (Papanikolaou et al., 2000). RT-qPCR analysis was performed as described in the main methods section. Primers for COX2 and PLA_2R are available on request. The mRNA fold variation between normal matched tissues and tumor samples was calculated as previously described (Mounier et al., 2008).

Expression of autotaxin and LPA receptors in Colon-26 cells by RT-qPCR and western-blot analysis. Total RNA from Colon-26 cells and RT-qPCR analysis was performed as described in the main methods section. qPCR primer sequences for autotaxin (mATX-F, 5’-GACCCCTAAAGCCATTATTGCTAA-3’; mATX-R, 5’-GGGAAGGTGCTGTTTCATGT-3’) were provided by Dr. Saulnier-Blache (Toulouse, France). qPCR primers for LPA1 to LPA4 were from Masiello et al. (Masiello et al., 2006). Conditioned medium from Colon-26 cells grown in 10-cm Petri dishes was prepared by incubating subconfluent cells at 37°C for 72 h in DMEM serum-free medium. Cell supernatant was collected, centrifuged to eliminate cellular debris, and concentrated about 40-fold using an Amicon YM-10 centricron device (Millipore) and stored at -20°C until analysis. Western blot analysis with the rabbit polyclonal autotaxin antibody (1/10,000) from Cayman Chemicals was performed as recommended by the manufacturer. Western blot analysis with the chicken polyclonal antibody (1/20,000) was performed as described (Giganti et al., 2008).
Legend for Supplemental Figures

Fig. 1S. Genotyping of mouse colon cell lines. Genomic DNA was extracted from YAMC, Colon-26, Apc\textsuperscript{Min} and AJ02- nm0 mouse colon cell lines and from tails of C57BL/6J and Balb/C mice and subjected to PCR analysis as indicated in supplemental methods. The PCR fragments were analyzed by agarose gel electrophoresis with ethidium bromide staining.

Fig. 2S. Expression of group X sPLA\textsubscript{2} and others in mouse models of colon cancer. RNA was isolated from the colon of azoxymethane-treated SWR/J and A/J mice or the colon and ileum of Apc\textsuperscript{A14} mice. The expression of sPLA\textsubscript{2}s, cPLA\textsubscript{2}α, PLA\textsubscript{2}R and COX2 was analyzed by RT-qPCR as described in methods.

Fig. 3S. Expression of autotaxin and LPA receptors in Colon-26 cells and effect of S32826 on LPA production triggered by hGX sPLA\textsubscript{2}. A, RT-qPCR analysis for COX2, autotaxin (ATX) and LPA receptors. The relative gene expression was normalized to the expression (Cp values were 21 to 22) of topoisomerase I which was used as a housekeeping gene. B, Western blot analysis of ATX in Colon-26 cell medium. Conditioned media from Colon-26 cells was prepared as described in methods. 50 ng of recombinant ATX (lane 1) and 80 µg of total protein from conditioned medium (lane 2) were loaded under reducing conditions on a 8% SDS-PAGE gel. The presence of ATX was analyzed by western blot using a commercially available rabbit polyclonal ATX-peptide antibody (Cayman Chemicals) or a polyclonal chicken antiserum raised against the recombinant full-length ß isoform of ATX (Giganti et al., 2008). C, Effect of the ATX inhibitor S32826 on LPA release by hGX sPLA\textsubscript{2} from Colon-26 cells. Cells were treated in conditions identical to those indicated in Fig. 7 in the main text. The ATX inhibitor was preincubated with cells for 15 min at the indicated concentrations before sPLA\textsubscript{2} addition.

References


Fig. 1S
(supplemental data)

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<th>PCR for WT allele</th>
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<td>1: YAMC</td>
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Fig. 2S
(supplemental data)