

Localization of various secretory phospholipase A₂ enzymes in male reproductive organs

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Abstract

Current evidence suggests the presence of transcripts for several secretory phospholipase A₂ (sPLA₂) enzymes in male genital organs. In this study, we examined by immunohistochemistry the localization of group IIA, IIC, IID, IIE, IIF, V and X sPLA₂s in male genital organs. In sPLA₂-IIA-deficient C57BL/6 mouse testis, sPLA₂-IIC, -IID, -IIE, -IIF, -V and -X were diversely expressed in spermatogenic cells within the seminiferous tubules. Immunoblotting revealed the presence of these sPLA₂s in mouse spermatozoa. In addition, sPLA₂-IIF, -V and -X were localized in the interstitial Leydig cells. The same set of sPLA₂s was detected in a mouse cultured Leydig cell line, and adenovirus-mediated transfer of these sPLA₂s into Leydig cells resulted in increased prostaglandin production. sPLA₂-IIC, -IID, -IIE, -IIF, -V and -X were also detected diversely in the epithelium of the epididymis, vas deferens, seminal vesicles, and prostate. In a sPLA₂-IIA-positive FVB strain, weak expression of sPLA₂-IIA was detected in Leydig cells. Notable differences in the sPLA₂ expression profiles were found in the seminal vesicles and prostate between mice and humans. Taken together, individual sPLA₂s exhibit distinct or partially overlapping localizations in male reproductive organs, suggesting both specific and redundant functions.

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1. Introduction

Secretory phospholipase A₂ (sPLA₂) is a group of enzymes with conserved catalytic site and Ca²⁺-binding

loop, and catalyzes the release of fatty acids and lysophospholipids from the *sn*-2 position of glycerophospholipids in the presence of Ca²⁺. To date, 10 sPLA₂ enzymes (sPLA₂-IB, -IIA, -IIC, -IID, -IIE, -IIF, -III, -V, -X, and -XIIA) have been identified in mammals [1]. sPLA₂s belonging to the I/II/V/X branch may have arisen from recent and successive gene duplication events, whereas group III and XII sPLA₂s are distantly related. Since the genes for sPLA₂-IIA, -IIC, -IID, -IIE, -IIF and -V are clustered on the human chromosome 1p34–36, these enzymes have often been referred to as the group II subfamily sPLA₂s. The sPLA₂ enzymes have been implicated in diverse biological events, such as arachidonic acid

Abbreviations: PLA₂, phospholipase A₂; sPLA₂, secretory PLA₂; cPLA₂, cytosolic PLA₂; COX, cyclooxygenase; PG, prostaglandin; mPGES, microsomal PGE₂ synthase; RT-PCR, reverse transcriptase-polymerase chain reaction; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis; IL, interleukin; PBS, phosphate-buffered saline; LPC, lysophosphatidylcholine; HSPG, heparan sulfate proteoglycan

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(AA) metabolism [2–12], digestion of dietary phospholipids [13], atherosclerosis [14–16], anti-bacterial defense [17–20], and oncogenesis [21]. However, the precise physiological functions of individual sPLA₂s are still obscure.

During the last two decades, the presence of sPLA₂s in male genital organs and seminal plasma of several animal species has been demonstrated in several studies [22–33]. Initial biochemical analyses showed that a PLA₂ activity exists in spermatozoa and that PLA₂ may be an important factor for the acrosome reaction and fertilization [22–24]. An sPLA₂ purified from human seminal plasma was ascribed to sPLA₂-IIA on the basis of its N-terminal sequence [25]. Immunohistochemical analyses using antibodies raised against sPLA₂-IB or -IIA demonstrated positive staining in prostate [26, 27], cauda epididymal spermatozoa [28] and seminal vesicles [27], according to animal species. However, recent evidence for the presence of transcripts for several sPLA₂s (IIA, IIC, IID, IIE, IIF, V and X) in human and mouse male genital organs [29–31] has raised a question as to the cross-reactivity of the previously used anti-sPLA₂ antibodies among sPLA₂ isozymes. Although Langlais et al. [32] purified a 17-kDa PLA₂ from human spermatozoa, its N-terminal amino acid sequence does not fit that of any of the mammalian sPLA₂s so far known. Expression of sPLA₂-IIC, a rodent-specific enzyme, in mouse spermatogenic cells within the testicular seminiferous tubules has been demonstrated by *in situ* hybridization [33].

In this study, we evaluated the localization of various sPLA₂s (IIA, IIC, IID, IIE, IIF, V and X) in mouse male reproductive organs. We show that individual sPLA₂s display unique cellular distributions in the testis, epididymis, vas deferens, seminal vesicles and prostate. There are some species-associated differences in the expression profiles of sPLA₂s, particularly in the prostate. On the basis of the results, possible functions of each sPLA₂ in the male genital organs are discussed.

2. Materials and methods

2.1. Materials

Inbred C57BL/6 and FVB strains of mice were purchased from Shizuoka Laboratory Animal Center (Shizuoka, Japan). Procedures for animal handling and preparation of mouse and human tissue sections were approved by the ethical committee of our Faculty. Mouse Leydig cell line I-10 were obtained from Human Science Research Resources Bank (Osaka, Japan) and were cultured in RPMI 1640 medium (Nissui Pharmaceutical Co.) containing 10% (v/v) fetal calf serum (FCS; Bioserum). Specific antibodies for various mouse and human sPLA₂s were described previously [34,35]. Goat antibodies against human cyclooxygenase (COX)-1 and human COX-2, and a rabbit antibody against human group IVA cytosolic PLA₂ (cPLA₂α) were purchased from Santa Cruz. Antibodies against cytosolic PGE synthase

(cPGES) and microsomal PGE synthases (mPGES-1 and -2) were described previously [36–38]. Preparation of cDNAs for sPLA₂s and cPLA₂α was described previously [2–5]. The enzyme immunoassay kit for PGE₂ was purchased from Cayman Chemicals. Human interleukin (IL)-1β was purchased from Genzyme. Primers for reverse transcription-polymerase chain reaction (RT-PCR) were purchased from Greiner Japan. Other reagents of analytical grade were from Wako Chemicals.

2.2. RT-PCR

Synthesis of cDNA was performed using 0.5 μg of total RNA from cells and tissues and AMV reverse transcriptase, according to the manufacturer's instructions supplied with the RNA PCR kit (Takara Biomedicals). Subsequent amplification of the cDNA fragments was performed using 0.5 μl of the reverse-transcribed mixture as a template with specific primers for each sPLA₂. For amplification of sPLA₂-IIA, -IIC, -IID, -IIE, -IIF, -V, and -X, a set of 23-bp oligonucleotide primers corresponding to the 5' - and 3' - nucleotide sequences of their open reading frames were used as primers [29–31]. The PCR conditions for sPLA₂-IIA, -IIC, -IID, -IIE, -V, and -X were 94 °C for 30 s and then 30 cycles of amplification at 94 °C for 5 s and 68 °C for 4 min, using the Advantage cDNA polymerase mix (Clontech). The PCR conditions for sPLA₂-IIF were 94 °C for 30 s and then 30 cycles of amplification at 94 °C for 30 s, 58 °C for 30 s, and 72 °C for 30s, with *ExTaq* polymerase (Takara Biomedicals). The PCR products were analyzed by 1% agarose gel electrophoresis with ethidium bromide. The gels were further subjected to Southern blot hybridization with sPLA₂ cDNAs as probes, as required for the experiments.

2.3. Northern blotting

Equal amounts (10 μg) of total RNA obtained from cells by TRIzol reagent (Invitrogen) were applied to separate lanes of 1.2% (w/v) formaldehyde-agarose gels, electrophoresed, and transferred to Immobilon-N membranes (Millipore). The resulting blots were then probed with the respective cDNA probes that had been labeled with [³²P]dCTP (Amersham Bioscience) by random priming (Takara Biomedicals). All hybridizations were carried out as described previously [2–5].

2.4. SDS-PAGE/immunoblotting

Lysates from 2×10⁵ cultured cells or 50-μg protein equivalents of tissue or sperm homogenates in phosphate-buffered saline (PBS) were subjected to SDS-PAGE using 7.5% (for cPLA₂α and COXs), 12.5% (for PGESs) and 15% (for sPLA₂s) gels under reducing (for cPLA₂α, COXs and PGESs) and non-reducing (for sPLA₂s) conditions. The separated proteins were electroblotted onto nitrocellulose membranes (Schleicher & Schuell) using a semi-dry blotter

(MilliBlot-SDE system; Millipore). After blocking with 3% (w/v) skim milk in TBS containing 0.05% (v/v) Tween-20 (TBS–Tween), the membranes were probed with the respective antibodies for 2 h, followed by incubation with horseradish peroxidase-conjugated anti-goat or -rabbit immunoglobulin for 2 h, and were visualized using the ECL Western blot system (NEN™ Life Science Products) as described [2–5].

2.5. Immunohistochemistry

Immunostaining of mouse genital organs was performed as described previously [39]. After deep anesthesia with ether, mice were perfused via the left ventricle of the heart with PBS containing 4% (w/v) sucrose and 4% (w/v) paraformaldehyde and then with Bouin's solution. Genital organs were removed and soaked overnight at 4 °C in Bouin's fixative. After being embedded in paraffin, the tissues were sectioned and mounted on slides. The sections were then deparaffinized in xylene and rehydrated in ethanol with increasing concentrations of water. Endogenous peroxidase activity was quenched with 0.3% (v/v) hydrogen peroxide in PBS for 30 min at room temperature. The tissues were then treated for 1 h at room temperature with 10% (v/v) sheep serum in PBS containing 0.1% (v/v) Triton X-100 and 0.1% (w/v) sodium azide for blocking. Then the tissue sections were sequentially treated with antisera for individual mouse sPLA₂s (1:200 dilution) for 1 day at 4 °C and with biotinylated anti-rabbit antibody (1:500 dilution) for 2 h at room temperature. Primary and secondary antibodies were diluted in PBS containing 1% (v/v) sheep serum (to avoid nonspecific binding) and 0.1% Triton X-100. A Vectastain Elite avidin-biotin-peroxidase kit (Vector Laboratories, Inc.) with diaminobenzidine substrate was used according to the manufacturer's protocol. The tissues were counterstained with hematoxylin. Finally, sections were dehydrated, rinsed in xylene, and mounted with Mount-Quick (Daido Sangyo, Tokyo, Japan).

Immunohistochemical staining of human tissue sections was performed as described previously [37,38]. Briefly, the tissue sections were incubated with Target Retrieval Solution (DAKO) as required, incubated for 10 min with 3% (v/v) H₂O₂, washed three times with PBS for 5 min each, incubated with 5% (v/v) skim milk for 30 min, washed three times with PBS–Tween for 5 min each, and incubated for 2 h with anti-human sPLA₂ antibodies (1:100 dilution in PBS). Then the sections were treated with the CSA system staining kit (DAKO) with diaminobenzidine substrate.

2.6. Expression of PLA₂s by the adenovirus system

Adenoviruses bearing individual PLA₂ cDNAs were prepared with the ViraPower Adenovirus Expression System (Invitrogen) according to the manufacturer's instruction. Briefly, the full-length cDNAs for sPLA₂s and cPLA₂α were subcloned into the pENTER/D-TOPO vector by using

the pENTER Directional TOPO Cloning kit (Invitrogen). After purification of the plasmids from the transformed Top10 competent cells (Invitrogen), the cDNA inserts were transferred to the pAd/CMV/V5-DEST vector (Invitrogen) by means of the Gateway system using LR clonase (Invitrogen). The plasmids were purified and digested with *PacI* (New England BioLabs). The linearized plasmids (1–2 μg) were then mixed with 4 μl of Lipofectamine 2000 (Invitrogen) in 200 μl of Opti-MEM medium (Invitrogen) and transfected into subconfluent 293A cells (Invitrogen) in 1 ml of Opti-MEM in six-well plates (Iwaki Glass). Then the 293A cells were cultured for 1–2 weeks in RPMI 1640 medium containing 10% FCS, with replacement of the medium every 2 days. When most cells became detached from the plates, the cells and culture medium were harvested together, freeze-thawed twice, and centrifuged to obtain the adenovirus-enriched supernatants. Then aliquots of the supernatants were added to fresh 293A cells and cultured for 2–3 days to amplify adenoviruses. After two to four cycles of amplification, the resulting adenovirus-containing media were used as virus stocks. Viral titers were determined by the plaque-forming assay with 293A cells. As a control, the pAd/CMV/V5-GW/*lacZ* vector (Invitrogen) was digested with *PacI* and transfected into 293A cells to produce *LacZ*-bearing adenovirus.

Aliquots of the adenovirus-containing medium were added to I-10 cells grown in 24-well plates (Iwaki Glass) and cultured for up to 36 h in the presence of 1 ng/ml IL-1β at 37 °C in a CO₂ incubator. Then the supernatants were subjected to the sPLA₂ enzyme assay and PGE₂ enzyme immunoassay, and the cells were taken for Northern blotting to assess the expression of PLA₂s.

2.7. Measurement of sPLA₂ activity

sPLA₂ activity was assayed by measuring the amounts of radiolabeled linoleic acid released from the substrate 1-palmitoyl-2-[¹⁴C]linoleoyl-phosphatidylethanolamine (Amersham Bioscience). The substrate in ethanol was dried under a stream of N₂ and was dispersed in water by sonication. Each reaction mixture (total volume 250 μl) consisted of appropriate amounts of the required sample, 100 mM Tris–HCl (pH 7.4), 4 mM CaCl₂ and 10 μM substrate. After incubation for 20 min at 37 °C, [¹⁴C]linoleic acid was extracted by Dole's method, and the radioactivity was quantified by liquid scintillation counting, as described previously [2].

3. Results

3.1. Detection of sPLA₂s in mouse genital organs by RT-PCR and immunoblotting

Expression of transcripts for various sPLA₂s in genital organs of male C57BL/6 mice (testis, epididymis, vas

deferens, seminal vesicles and prostate) was examined by RT-PCR (Fig. 1A). sPLA₂-IIA was detected in none of these organs (data not shown), in agreement with the natural disruption of its gene in this mouse strain [21]. Intense expression of the transcript for sPLA₂-IIC, a rodent-specific enzyme [29], was detected in the testis, epididymis and seminal vesicles, whereas it was expressed only weakly in the vas deferens and prostate. sPLA₂-IID was expressed almost equally in the testis, epididymis, vas deferens and seminal vesicles, and weakly in the prostate. Although the expression of sPLA₂-IIE was lower than that of other sPLA₂s, it could be detected throughout the genital organs, with expression in the vas deferens being highest. Expression of sPLA₂-IIF was highest in the testis, followed by the epididymis and vas deferens, whereas its expression in the seminal vesicles and prostate was weak, although significant. Expression of sPLA₂-V was high in the testis and epididymis, and modest in the vas deferens, seminal vesicles and prostate. Thus, sPLA₂-IIF and -V were preferentially distributed in the upper genital organs. Expression of sPLA₂-X was intense in the testis and seminal vesicles, faint in the epididymis and vas deferens, and rather weak in the prostate. In the testis, vas deferens and seminal vesicles, the sPLA₂-X transcript appeared as a doublet; sequencing of each DNA fragment revealed that the upper band corresponded to an alternative splicing variant that has a 99-bp insertion between exons 3 and 4 [40].

We next investigated the expression of sPLA₂ proteins in the testis, epididymis, and mature spermatozoa of C57BL/6

mice by immunoblotting with antibodies raised against individual sPLA₂s. Specificity of these antibodies was confirmed by immunoblotting with authentic recombinant sPLA₂s as well as with cell lysates transfected with individual sPLA₂s [34,35]. As shown in Fig. 1B, sPLA₂-IIC, -IID, -V and -X, but not sPLA₂-IIE (probably because its expression level is too low to be detected by immunoblotting) or sPLA₂-IIA, were each detected in the testis and epididymis at the expected molecular sizes (14–18 kDa) (Fig. 1B). These four sPLA₂s were also detected in spermatozoa (Fig. 1C). Of note, sPLA₂-IIF was detected as a major 25-kDa band (larger than the expected 18-kDa size) and a minor 45-kDa band in the testis and epididymis (Fig. 1B). This result suggests that a portion of this enzyme may exist as a homodimer (as has been proposed [30]) or that this enzyme may undergo posttranslational modification (such as glycosylation) in these tissues. In contrast, only the smaller form of sPLA₂-IIF was detected in spermatozoa (Fig. 1C).

3.2. Immunohistochemistry of sPLA₂s in the testis

To determine the cellular origins of sPLA₂s in these tissues, we performed immunohistochemistry with antibodies specific for individual enzymes. sPLA₂-IIA (used as a negative control) provided no detectable staining signals in C57BL/6 mouse testis (Fig. 2A). sPLA₂-IIC was stained in the whole area of spermatogenic cells within the seminiferous tubules, with spermatogonia providing the

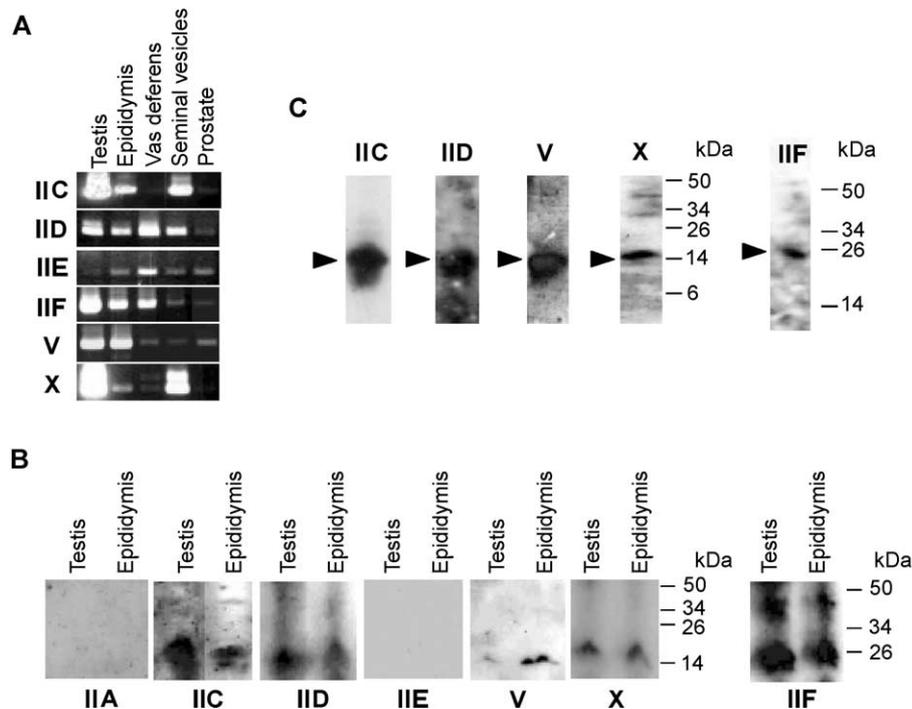


Fig. 1. RT-PCR and immunoblot analyses of sPLA₂ expression in male genital organs and spermatozoa of mice. (A) Total RNAs extracted from individual tissues were subjected to RT-PCR using primers specific for each sPLA₂. Amplified bands were visualized in agarose gels with ethidium bromide. (B and C) Immunoblotting of mouse testis and epididymis (B) as well as spermatozoa (C) with anti-sPLA₂ antibodies. Samples (50 µg protein equivalents) were applied to 15% SDS-PAGE gels and then immunoblotted with the respective antibodies.

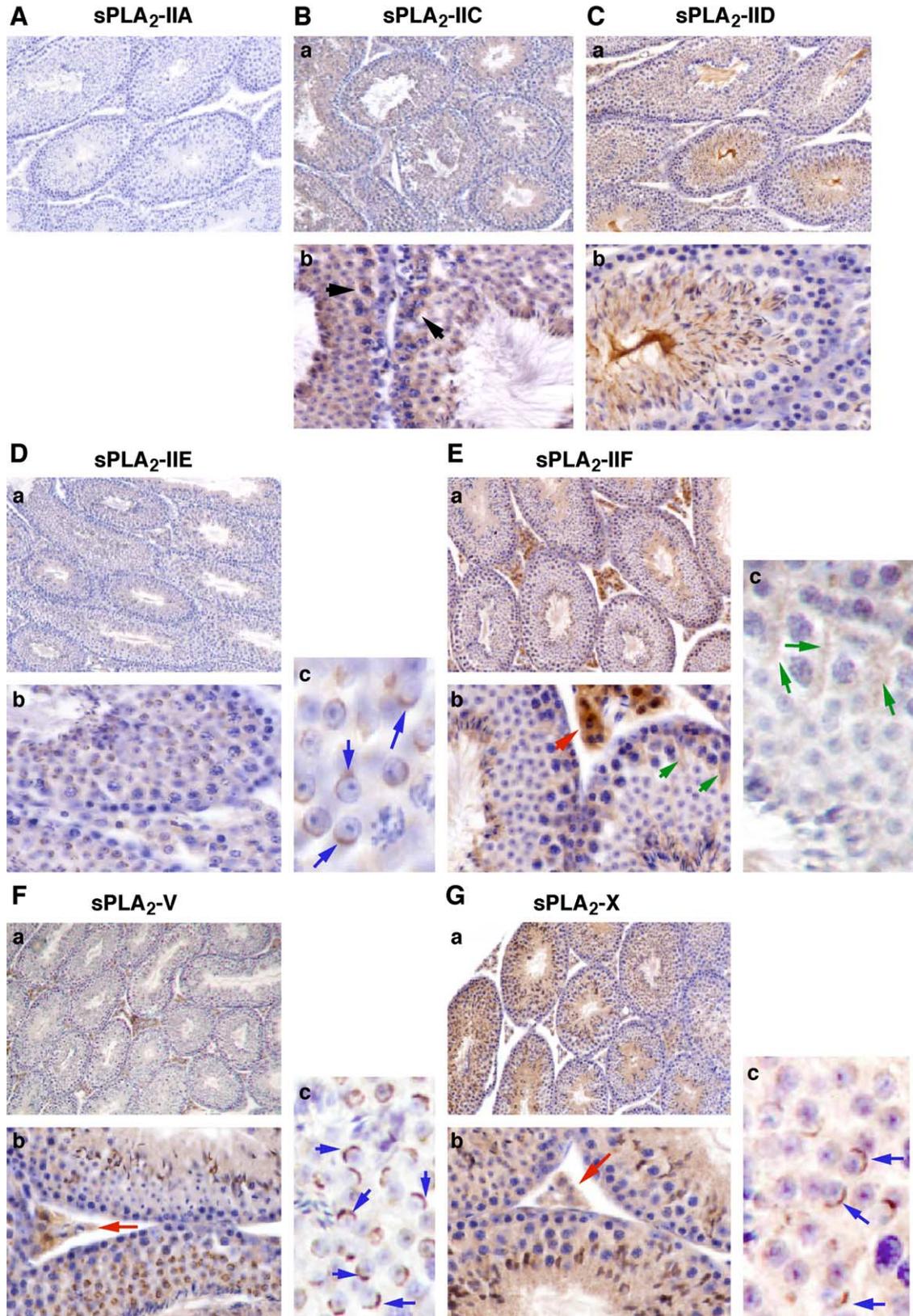


Fig. 2. Immunohistochemical localization of sPLA₂s in mouse testis. (A) sPLA₂-IIA, (B) sPLA₂-IIC, (C) sPLA₂-IID, (D) sPLA₂-IIE, (E) sPLA₂-IIF, (F) sPLA₂-V, and (G) sPLA₂-X. sPLA₂-IIC was stained in spermatogenic cells, including spermatogonia (dark arrowheads). sPLA₂-IID was detected in the central part of the seminiferous tubules, which are enriched in late spermatids and spermatozoa. Scattered signals for sPLA₂-IIE, -V and -X were detected in the acrosomes of spermatocytes (blue arrows). Leydig cells (red arrows) were positively stained for sPLA₂-IIF and -V as well as weakly for sPLA₂-X. sPLA₂-IIF was also detected in Sertoli cells (green arrows). sPLA₂-IIA was undetectable. Panels a, $\times 40$; panels b, $\times 100$; and panels c, $\times 400$.

most intense staining (Fig. 2B), consistent with a previous *in situ* hybridization study [33]. sPLA₂-IID gave string-like and elongated staining signals in the central part of the seminiferous tubules (Fig. 2C), suggesting its presence in elongating late spermatids and spermatozoa. Although staining of sPLA₂-IIE was weak, scattered signals with crescent and elongated shapes were detectable in spermatogenic cells (Fig. 2D). These signals represent a typical feature of location in the acrosomes [41]. sPLA₂-IIF (Fig. 2E) and -V (Fig. 2F) were clearly detected in Leydig cells of the interstitium. Late spermatids or spermatozoa near the central part of seminiferous tubules, as well as Sertoli cells, were also positive for sPLA₂-IIF (Fig. 2E). Crescent and elongated signals for sPLA₂-V, which were similar to but more intense than those for sPLA₂-IIE, were seen in the seminiferous tubules, suggesting its localization in the acrosomes of spermatocytes and/or round and elongating spermatids (Fig. 2F). sPLA₂-X was uniformly stained in spermatogenic cells (Fig. 2G). Again, crescent and elongated signals for sPLA₂-X were obvious in the area enriched in spermatocytes and/or round and elongating spermatids (Fig. 2G), as seen with sPLA₂-V (Fig. 2F). In addition, Leydig cells were weakly stained with sPLA₂-X (Fig. 2G).

3.3. Studies using cultured Leydig cells

Leydig cells constitute the principal endocrine component of the testis, occupying most of the extravascular space of the interstitium. The primary role of Leydig cells is to synthesize and secrete androgenic steroids in response to gonadotropic hormones. Previous studies have shown that Leydig cells produce PGE₂, which can modify their own functions and alter seminiferous tubule contractility in autocrine and paracrine fashions [42–46]. In order to assess the potential contribution of sPLA₂s to PGE₂ production in Leydig cells, we took advantage of the mouse Leydig cell line I-10. As assessed by RT-PCR, sPLA₂-IIF was most intensely detected in I-10 cells (Fig. 3A). Weak expression of sPLA₂-V and -X was also evident in these cells, whereas other sPLA₂s (IIA, IID, and IIE) were below the detection limit (Fig. 3A). Thus, detection of the transcripts for sPLA₂-IIF, -V and -X in I-10 cells was consistent with the immunohistochemical analysis of mouse testis (Fig. 2). To the best of our knowledge, this is the first demonstration of a cell line that expresses sPLA₂-IIF endogenously. As assessed by immunoblotting, I-10 cells expressed cPLA₂, COX-1 and -2, mPGES-1 and -2, and cPGES constitutively (data not shown), confirming that all components for PGE₂ biosynthesis are present in these cells.

To achieve efficient and prompt transfection of sPLA₂s into cultured cells, we constructed adenovirus harboring each sPLA₂ cDNA. Then I-10 cells were infected with adenoviruses for individual sPLA₂s for 24 h, and PGE₂ released into the supernatants was quantified. Appropriate expression of each enzyme was verified by Northern

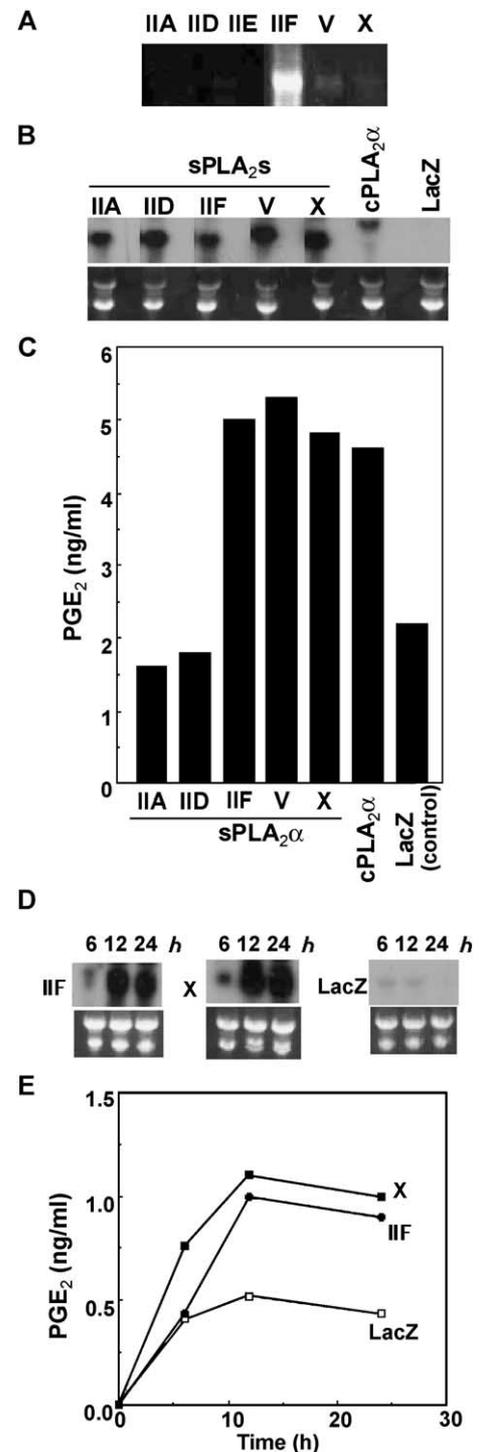


Fig. 3. Expression and function of sPLA₂s in the mouse Leydig cell line I-10. (A) Expression of endogenous sPLA₂s in I-10 cells, as assessed by RT-PCR. (B and C) PGE₂ production by I-10 cells overexpressing various sPLA₂s and cPLA₂α. Cells were infected with adenoviruses for sPLA₂s and cPLA₂α in the presence of IL-1β for 24 h. Then, the cells were subjected to Northern blotting to assess the expression of sPLA₂s and cPLA₂α (B), and the supernatants were taken for the PGE₂ assay (C). Cells infected with adenovirus harboring the *LacZ* gene were used as a control. (D and E) Time course of sPLA₂ expression and PGE₂ production by I-10 cells after infection with adenoviruses (MOI=5) for sPLA₂s and *LacZ*. Faint bands detected in *LacZ*-adenovirus-infected cells represent endogenous sPLA₂-IIF. Representative results of three to five independent experiments are shown.

blotting (Fig. 3B). The levels of individual sPLA₂s released into the supernatants of adenovirus-infected cells ranged from 10 to 20 ng/ml, as estimated by their enzymatic activities in comparison with the activities of their respec-

tive authentic recombinant sPLA₂s (data not shown). As shown in Fig. 3C, there were marked increases in PGE₂ production by I-10 cells transfected with sPLA₂-IIF, -V and -X, a group of enzymes intrinsically expressed in mouse

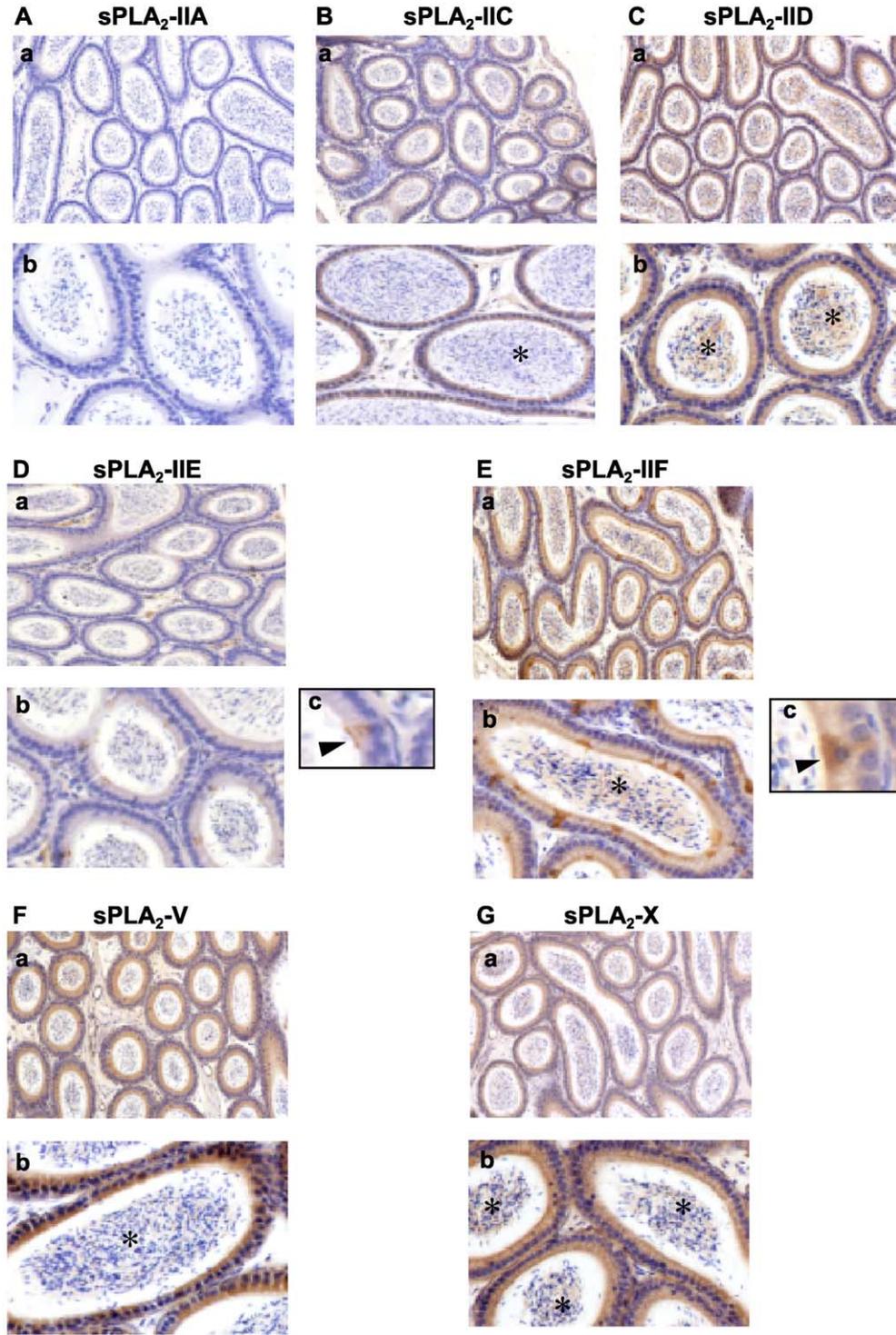


Fig. 4. Immunohistochemical localization of sPLA₂s in mouse epididymis. (A) sPLA₂-IIA, (B) sPLA₂-IIC, (C) sPLA₂-IID, (D) sPLA₂-IIE, (E) sPLA₂-IIF, (F) sPLA₂-V, and (G) sPLA₂-X. Staining of the caput portion of the epididymis is shown. Panels a, $\times 40$; panels b, $\times 100$; and panels c, $\times 400$. sPLA₂-IIC, -IID, -IIE, -IIF, -V and -X, but not sPLA₂-IIA, were located in the tubular epithelial cells of the epididymis. sPLA₂-IIE and -IIF were particularly evident in the apical epithelial cells (arrowheads). Spermatozoa (asterisks) were positive for sPLA₂-IIC, -IID, -IIF, -V and -X.

Leydig cells (see above), as well as those transfected with cPLA₂α, which was used as a positive control for increased AA metabolism [47]. sPLA₂-IIA and -IID, which were immunohistochemically undetectable in Leydig cells (Fig. 2A and C), failed to increase PGE₂ generation appreciably (Fig. 3C). Time course experiments showed that the increased production of PGE₂ after transfection with sPLA₂-IIF and -X occurred kinetically in parallel with their inducible expression (Fig. 3D and E). PGE₂ production reached a plateau peak at 12 h (Fig. 3E), at which time the expression of sPLA₂-IIF and -X also peaked (Fig. 3D).

3.4. Immunohistochemistry of sPLA₂s in other genital organs

Spermatozoa leave the testis through the efferent ductules and enter the initial segment of the epididymis. Immunostaining

of different parts of the epididymis revealed that sPLA₂-IIC (Fig. 4B), -IID (Fig. 4C), -IIF (Fig. 4E), -V (Fig. 4F) and -X (Fig. 4G) were located in the tubular epithelial cells of the epididymis. This staining pattern was similar between the caput (Fig. 4) and other parts (the initial segment, corpus and cauda) (data not shown) of the epididymis. sPLA₂-IIF often provided a very strong signal in the apical epithelial cells in the caput and cauda (Fig. 4E, panels b and c). These apical epithelial cells were also positive for sPLA₂-IIE (Fig. 4D, panels b and c). Spermatozoa within the tubules were noticeably stained with sPLA₂-IIC, -IID, -IIF, -V and -X (Fig. 4, panels b).

When small round slices of the vas deferens between the epididymis and seminal vesicles were immunostained, sPLA₂-IIC (Fig. 5B), -IID (Fig. 5C), -IIF (Fig. 5E), -V (Fig. 5F) and -X (Fig. 5G) were detected in the luminal epithelium and the cilia. More intense signals for sPLA₂-IID

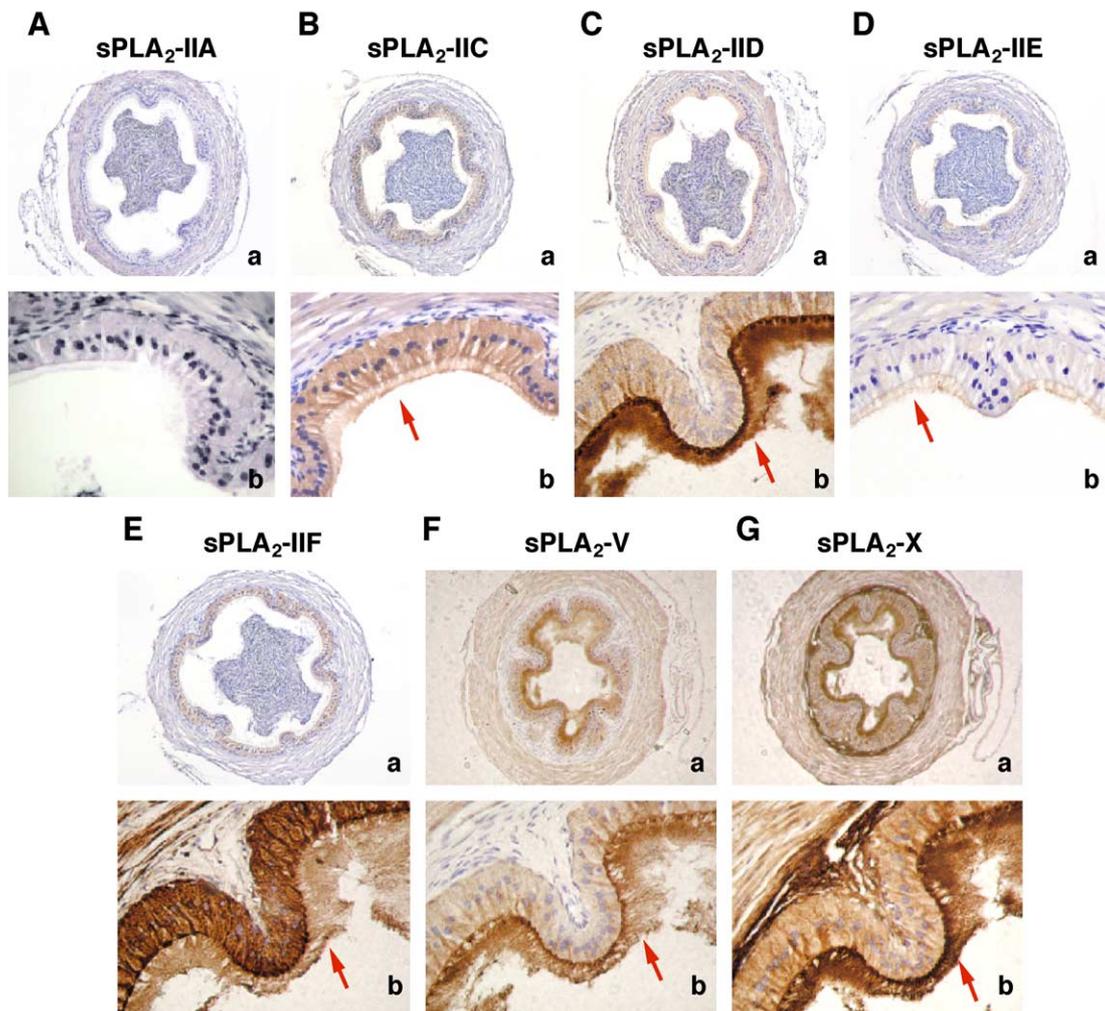


Fig. 5. Immunohistochemical localization of sPLA₂s in mouse vas deferens. (A) sPLA₂-IIA, (B) sPLA₂-IIC, (C) sPLA₂-IID, (D) sPLA₂-IIE, (E) sPLA₂-IIF, (F) sPLA₂-V, and (G) sPLA₂-X. Arrows indicate the cilia on the luminal surface of the duct epithelium. sPLA₂-IIC, -IID, -IIF, -V and -X were detected in epithelial cells, and more intense staining for sPLA₂-IID and -V was evident in the luminal cilia (arrows). The smooth muscle cell layer was also weakly positive for sPLA₂-X. Staining of sPLA₂-IIE in the apical surface of the epithelium was faint, whereas staining for sPLA₂-IIA was totally negative. Panels a, ×40; and panels b, ×100.

and -V were recognized in the cilia than in the epithelial cell layer; the reverse was true for sPLA₂-IIF. sPLA₂-X was intensely stained in both the apical and basolateral surfaces of epithelial cells, and the smooth muscle cell layer was also weakly sPLA₂-X-positive (Fig. 5G). sPLA₂-IIE was detected faintly in the apical surface of the epithelium (Fig. 5D).

In the seminal vesicular gland, sPLA₂-IIC (Fig. 6B), -IID (Fig. 6C), -IIF (Fig. 6E), -V (Fig. 6F), and -X (Fig. 6G) were expressed in epithelial cells. Expression of sPLA₂-IIE in epithelial cells was weak but significant (Fig. 6D). In the prostate, staining for sPLA₂-IIF was most intense in epithelial cells (Fig. 6E), where sPLA₂-IIC (Fig. 6B), -IID (Fig. 6C) and -IIE (Fig. 6D) were also positive.

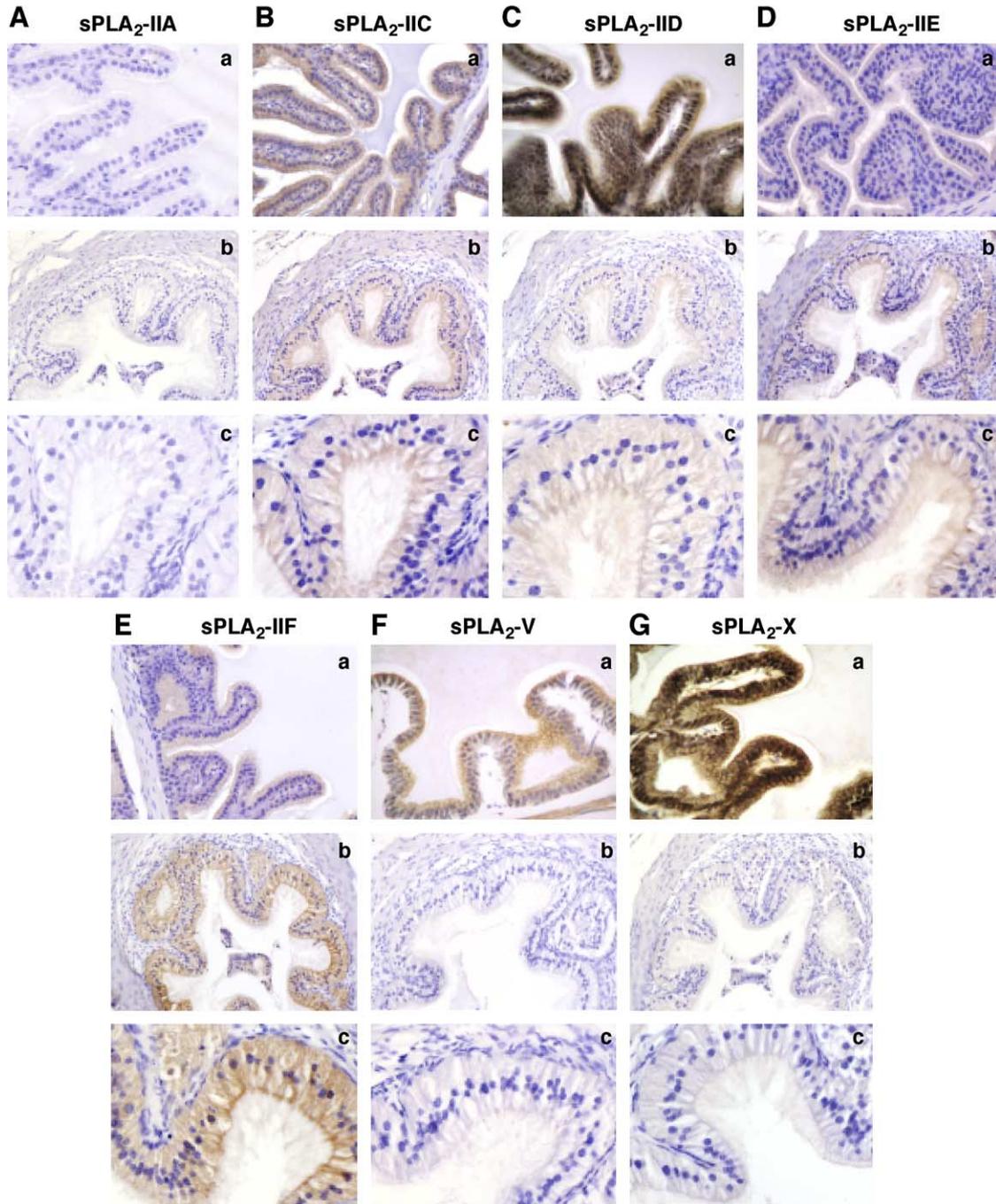


Fig. 6. Immunohistochemical localization of sPLA₂s in mouse seminal vesicles and prostate. (A) sPLA₂-IIA, (B) sPLA₂-IIC, (C) sPLA₂-IID, (D) sPLA₂-IIE, (E) sPLA₂-IIF, (F) sPLA₂-V, and (G) sPLA₂-X. Panels a indicate seminal vesicles and panels b and c show prostate. In the seminal vesicles, sPLA₂-IIC, -IID, -IIF, -V, and -X were expressed in epithelial cells. Expression of sPLA₂-IIE in epithelial cells was weak but significant, whereas sPLA₂-IIA was not detected. In the prostate, sPLA₂-IIC, -IID, -IIE and -IIF were detected in epithelial cells, whereas sPLA₂-IIA, -V and -X were undetectable. Panels a and b, $\times 40$; and panels c, $\times 100$.

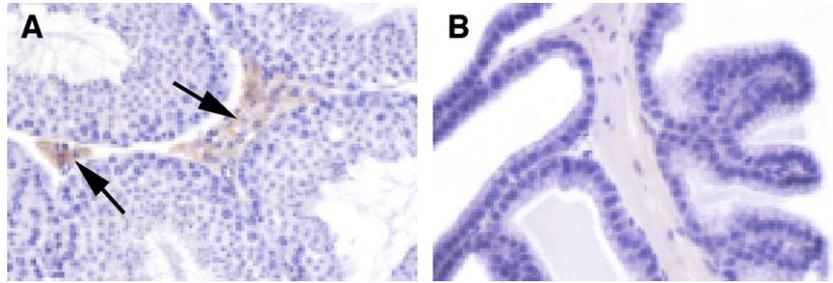


Fig. 7. Immunohistochemical localization of sPLA₂-IIA in male genital organs of FVB mice. (A) sPLA₂-IIA expression was detected in Leydig cells (arrows) in the testis. (B) sPLA₂-IIA was undetectable in seminal vesicles. Panels a, $\times 40$; and panels b, $\times 100$.

Although transcripts for sPLA₂-V and -X were faintly detected in the prostate by RT-PCR (Fig. 1), their immunoreactivity was barely detectable (Fig. 6, F & G).

As expected, sPLA₂-IIA was undetectable throughout the epididymis (Fig. 4A), vas deferens (Fig. 5A), seminal vesicles and prostate (Fig. 6A).

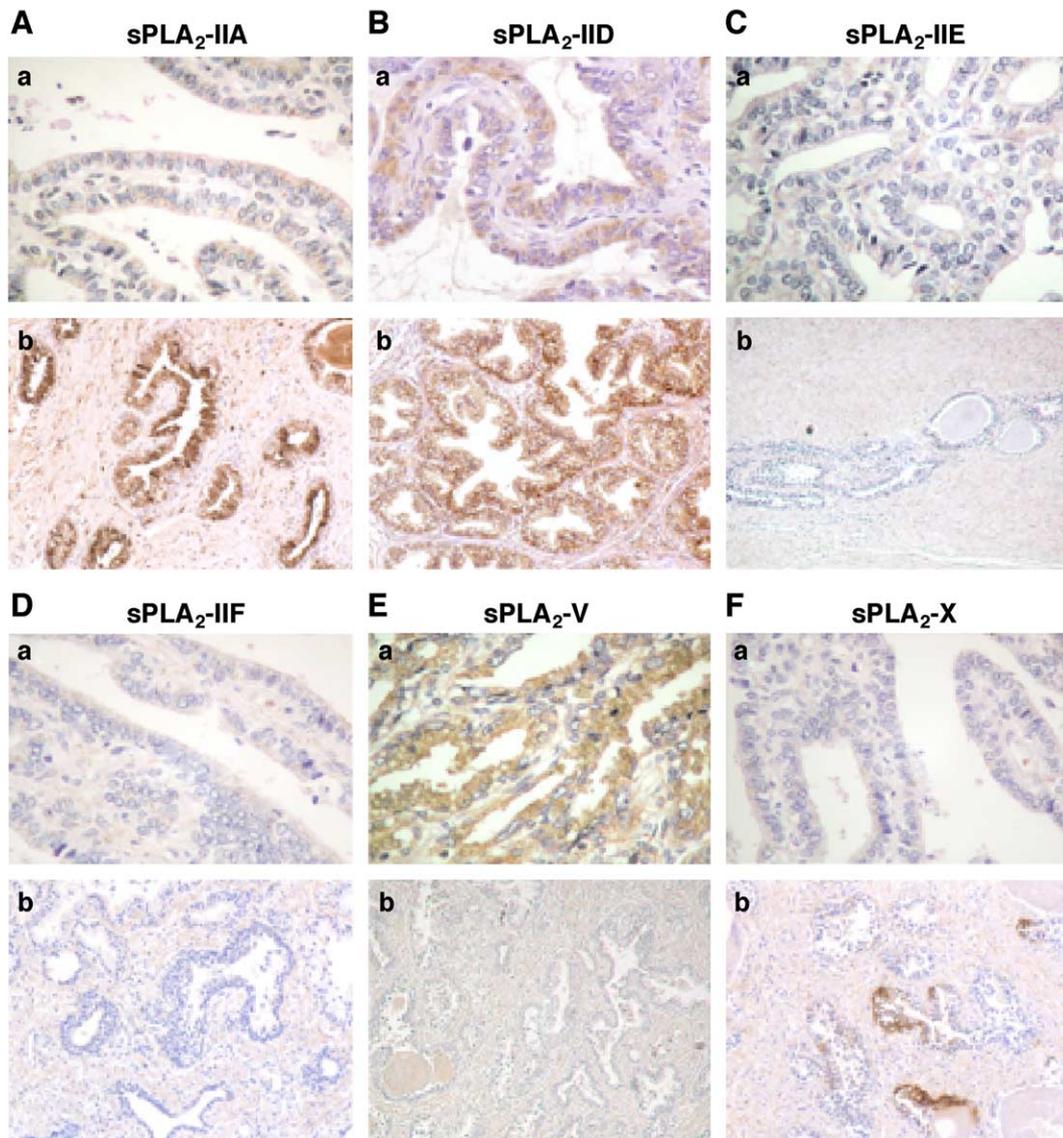


Fig. 8. Immunohistochemical localization of sPLA₂s in human seminal vesicles and prostate. (A) sPLA₂-IIA, (B) sPLA₂-IID, (C) sPLA₂-IIE, (D) sPLA₂-IIF, (E) sPLA₂-V, and (F) sPLA₂-X. Panels a and b indicate seminal vesicles and prostate, respectively ($\times 40$ magnification). In the seminal vesicles, sPLA₂-IID and -V yielded positive staining in epithelial cells. sPLA₂-IIA was also weakly positive, whereas sPLA₂-IIE, -IIF and -X were poorly expressed, in epithelial cells. In the prostate, sPLA₂-IIA and -IID, but not sPLA₂-IIE, -IIF and -V, were expressed in epithelial cells. sPLA₂-X was partly positive in these cells.

3.5. Expression of sPLA₂-IIA in male genital organs in a mouse strain with normal sPLA₂-IIA gene

Immunohistochemistry of sPLA₂-IIA in male genital organs of FVB mouse strain, in which the sPLA₂-IIA gene is intact, revealed that significant sPLA₂-IIA immunoreactivity was localized in Leydig cells in the testis (Fig. 7A). However, its expression was pretty low in the lower genital organs, as exemplified in the seminal vesicles (Fig. 7B). These results were confirmed by RT-PCR analysis of sPLA₂-IIA, where it was detected only in the testis (data not shown). The expression patterns of other sPLA₂s in these organs of FVB strain were essentially similar to those of C57BL/6 strain (data not shown).

3.6. Localization of sPLA₂s in human male reproductive organs

We next examined the locations of sPLA₂s in human male genital organs as well. Immunohistochemistry with antibodies specific for individual human sPLA₂s revealed that the expression profiles of each enzyme in the testis and epididymis of humans were similar to those of mice (data not shown)¹, except sPLA₂-IIC, which is not expressed in humans [29]. However, notable species-associated differences in the sPLA₂ expression profile were found in the seminal vesicular glands and prostate. In human seminal vesicles, sPLA₂-IID (Fig. 8B) and -V (Fig. 8E) displayed intense staining in the epithelium. Staining for sPLA₂-IIA was also weakly positive in epithelial cells (Fig. 8A), whereas sPLA₂-IIE (Fig. 8C), -IIF (Fig. 8D) and -X (Fig. 8E) were poorly expressed. Thus, in contrast to mice (Fig. 6), sPLA₂-IIA is present and sPLA₂-IIF and -X are not expressed in human seminal vesicles. In human prostate, intense immunoreactivity for sPLA₂-IIA was detected in epithelial cells and concretions of the gland (Fig. 8A). The prostate epithelium was also positive for sPLA₂-IID (Fig. 8B) and partially for sPLA₂-X (Fig. 8F), whereas sPLA₂-IIE (Fig. 8C), -IIF (Fig. 8D) and -V (Fig. 8E) were undetectable. Thus, the expression of sPLA₂s (IIA, IID and X) in human prostate is markedly distinct from that (IIC, IID, IIE, and IIF) in mouse prostate.

4. Discussion

In the present study, we performed immunohistochemistry to determine the cellular origins of several sPLA₂s (IIA, IIC, IID, IIE, IIF, V and X) in mouse and human male genital organs. We provide evidence that individual sPLA₂s exhibit unique cellular localizations in these tissues and that some enzymes are localized in the same cell types. Since the antibodies against mouse and human sPLA₂s used in this

study are highly specific (even not cross-reacting with orthologs from other animal species) [34,35] and since the staining patterns of individual orthologs in testis and epididymis are similar between mouse and human, it is likely that the observed immunoreactivities truly reflect the locations of individual sPLA₂s. Previous immunohistochemical analyses using antibodies raised against sPLA₂-IIA or -IB have shown that the immunoreactivities are located in various sites of the male genital organs, including hamster epididymal spermatozoa [28], bovine seminal vesicles and connective tissues [27], and human prostate and testicular round spermatids [26,27]. Beyond the species-associated differences in sPLA₂ expression profiles, our present results suggest that some of the previous immunohistochemical studies may have looked at multiple sPLA₂s in these organs. The localization of each sPLA₂ in mouse male reproductive organs, as clarified in the present study, is summarized in Fig. 9.

4.1. Testis

The PLA₂ reaction has been implicated in the acrosome reaction of spermatozoa and in plasma membrane fusion between sperm and oocytes [48,49]. The presence of particular sPLA₂(s) on the acrosomal surface and in the middlepiece mitochondria of hamster and human spermatozoa has been proposed [27,28]. The acrosome reaction of hamster spermatozoa and subsequent fertilization in vitro were blocked by an antibody against pancreatic sPLA₂-IB, and this effect was reversed by lysophosphatidylcholine (LPC), a major PLA₂ reaction product [50]. Accumulation of LPC occurred following in vitro incubation of spermatozoa, and LPC stimulated the fertilizing ability of spermatozoa and induced the changes in the zona pellucida and the oolemma that allowed sperm-egg fusion [51–54]. Moreover, the induction of sperm capacitation was delayed in mice lacking the receptor for platelet-activating factor, a lipid mediator that is synthesized from the *sn*-1 alkyl form of LPC [55].

We now show that several sPLA₂s are expressed in spermatogenic cells of mouse (and human) seminiferous tubules. In mouse testis, sPLA₂-IIC, -IID, -IIE, -IIF, -V and -X are expressed diversely in spermatogenic cells including spermatogonium, spermatocytes, spermatids and maturing spermatozoa (Fig. 2). It appears that sPLA₂-V, -X and -IIE are enriched in spermatocytes and/or round and elongating spermatids, whereas sPLA₂-IID and -IIF are enriched in the midpiece or tail portions of maturing spermatozoa. These sPLA₂s also provide positive staining in spermatozoa within the epididymis ducts (Fig. 4). Moreover, sPLA₂-IIC, -IID, -IIF, -V and -X are detected in mature spermatozoa by immunoblotting (Fig. 1A). Thus, it is tempting to speculate that these germ cell-associated enzymes may correspond to the sperm-associated sPLA₂ that had previously been reported [26,27]. It has been suggested that germ cell-associated sPLA₂s may play a role in the acrosome reaction

¹ Masuda, S., Murakami, M., Ishikawa, A., Ishii, T., and Kudo, I.; unpublished observation.

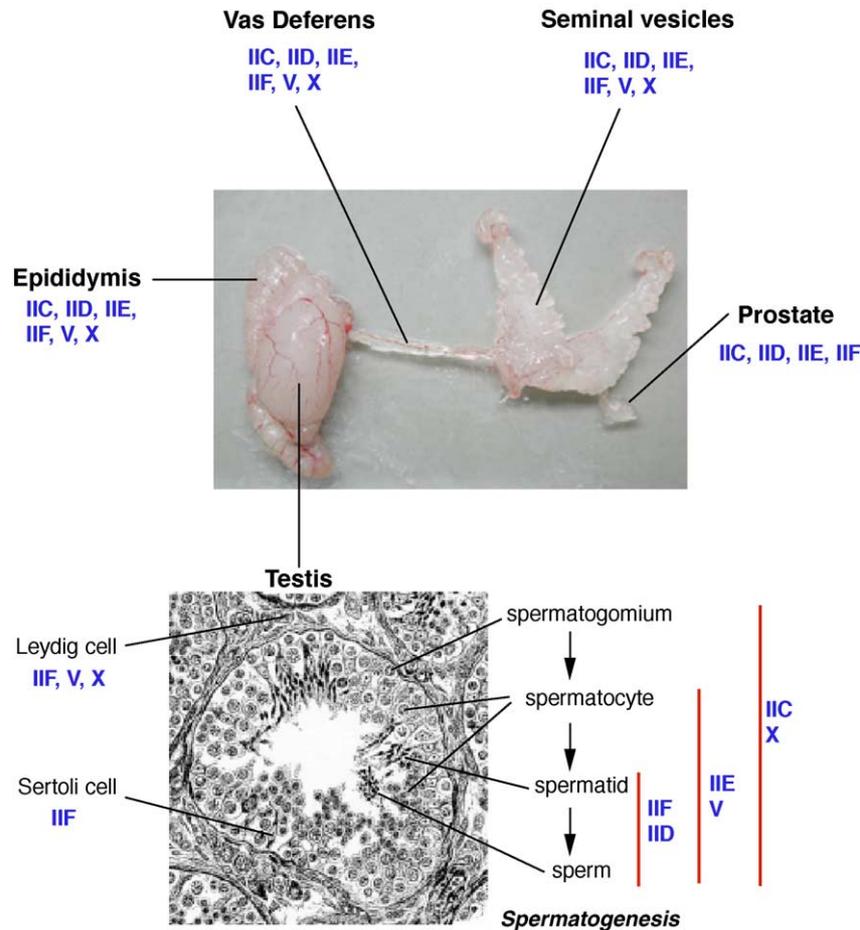


Fig. 9. Schematic diagram of the localization of sPLA₂s in mouse male genital organs. A photograph of the C57BL/6 mouse male reproductive organs, an illustration of the seminiferous tubules in the testis, and the localization of individual sPLA₂s are shown. In the testis, sPLA₂-IIC and -X are expressed virtually in the whole spermatogenic cells, sPLA₂-V and weakly sPLA₂-IIE in spermatocytes and/or spermatids, and sPLA₂-IID and -IIF in the late spermatids and/or spermatozoa. sPLA₂-IIF is the major enzyme expressed in Leydig cells, in which sPLA₂-V and -X (as well as sPLA₂-IIA in FVB mice) are also detected. sPLA₂-IIF is also expressed in Sertoli cells. In the epididymis, vas deferens, and seminal vesicles, sPLA₂-IIC, -IID, -IIE, -IIF, -V and -X are expressed in the epithelium. sPLA₂-IIC, -IID, -IIE and -IIF are detected in the prostate epithelium.

and subsequent fertilization through producing LPC [48–54]. Indeed, staining signals for sPLA₂-IIE, -V and -X in testicular spermatogenic cells (Fig. 2) could be attributed to the acrosome vesicles [41], although more detailed compartmentalization will need to be addressed by immunoelectron microscopy. The possible role of sPLA₂s in the acrosome reaction, an exocytosis event by which the acrosomal content is released, is reminiscent of the ability of sPLA₂s to promote exocytosis in mast cells and neuroendocrine cells [56,57].

We found that sPLA₂-IIF, -V and -X are expressed in mouse Leydig cells (Figs. 2 and 3), suggesting a unique role for these enzymes in this steroid hormone-producing cell in the testis. Several reports have suggested the importance of AA and its metabolites for the function of Leydig cells. For instance, stimulation of Leydig cells with gonadotropin resulted in the release of AA, which was in turn required for optimal steroidogenesis [42–44]. In contrast, another study reported that PGE₂ suppressed the gonadotropin-induced formation of testosterone in Leydig cells [45]. PLA₂/COX-2

pathway products, such as PGE₂ and PGF_{2α}, modulated interleukin (IL)-1 and IL-6 production in Leydig progenitors [49]. PGE₂ stimulated seminiferous tubule contractility through direct action on peritubular myoid cells [58]. Furthermore, immunoreactivities for the PGE₂-biosynthetic enzymes downstream of PLA₂s, such as COX-1 and mPGES-1, were detected in mouse Leydig cells [59]. The ability of sPLA₂-IIF, -V, and -X to increase cellular AA release in several cell types [3–12] supports the idea that these enzymes may be involved in augmented AA release, and thereby hormonal production, in Leydig cells.

Indeed, irrespective of the limitation of the overexpression strategy, PGE₂ production by a mouse Leydig cell line is markedly augmented by sPLA₂-IIF, -V, and -X (Fig. 3). Previous studies have shown that sPLA₂-V and -X [4,5], and probably -IIF [11], can elicit cellular AA release via acting on the outer leaflet of the plasma membrane (*the external plasma membrane pathway*). In contrast, sPLA₂-IIA and -IID act on cells through association with heparan sulfate proteoglycans (HSPGs), where these HSPG-binding sPLA₂s

bind cell surface HSPG, internalized, and then function (*HSPG-shuttling pathway*), in a cell type-specific manner [3,5]. Thus, the ability of sPLA₂-IIF, -V and -X, but not -IIA and -IID, to augment PGE₂ production in Leydig cells may indicate that the external plasma membrane pathway, but not the HSPG-shuttling pathway, is operative in these cells. Alternatively, Leydig cells might have specific machinery to be susceptible to sPLA₂-IIF, -V and -X (e.g., specific adaptors or receptors for sPLA₂s might be expressed), a possibility that needs to be addressed. Even though sPLA₂-IIA is weakly detected by immunohistochemistry in Leydig cells in FVB strain (Fig. 7), its contribution to PGE₂ generation is unclear in this context. Importantly, the main sPLA₂ expressed in Leydig cells is sPLA₂-IIF, an enzyme that is poorly expressed in other tissues [60]. Thus, it would be important to clarify the dynamics, functions, and regulatory roles of this particular enzyme in this particular cell type.

4.2. Epididymis and vas deferens

It is well established that the epididymis and vas deferens produce a large amount of prostanoids. PGE₂ is the most abundant prostanoid and exhibits potent smooth muscle relaxant activity. PGE₂ has been thought to play an important role in the transport of sperm through the epididymis and its subsequent expulsion at coitus from the vas deferens [61,62]. In rodents, PGE₂ is constitutively released into the semen from the vas deferens rather than from the prostate or seminal vesicles [63]. Expression of COX-1, COX-2 and mPGES-1 is observed in the epithelium of mouse epididymis and vas deferens [59]. Furthermore, the concentration of PGF_{2α}, another seminal prostanoid, is also higher in the vas deferens than in other parts of the male genital organs [64].

We show that sPLA₂-IIC, -IID, -IIE, -IIF, -V and -X are expressed in the epithelium of mouse epididymis (Fig. 4) and vas deferens (Fig. 5), which coincides with the localization of COXs and mPGES-1 [59]. In particular, the intense staining for sPLA₂-IIF, and to a lesser extent sPLA₂-IIE, in the apical epithelial cells of mouse epididymis coincides with the location of mPGES-1 [59]. These apical cells are considered to be responsible for absorption of excess secretion products [41]. These sPLA₂s may contribute to increased production of PGE₂ and other prostanoids by epithelial cells of the male reproductive tract. The luminal cilia of the tracts are noticeably stained with antibodies to sPLA₂-IID, -IIF, -V and -X (Fig. 5). This unique distribution may reflect that these sPLA₂s are secreted apically from the tubular epithelial cells into the lumen, or that the sPLA₂s in luminal fluids are absorbed to the luminal cilia. The latter possibility is supported by the observation that sPLA₂-V and -X proteins are clearly detected in the vas deferens by immunohistochemistry (Fig. 5), whereas their transcripts are detected only weakly by RT-PCR (Fig. 1A). A likely candidate for the sPLA₂-

binding component in the cilia is the highly sulfated glycosaminoglycan chains of proteoglycans, at least for several cationic heparanoid-binding sPLA₂s such as sPLA₂-IID and -V [2–5]. However, this scenario cannot be applicable to anionic heparin-nonbinding sPLA₂s such as sPLA₂-IIF and -X, and therefore needs further investigation.

4.3. Seminal vesicles

PGE₂ released from the seminal vesicles into the semen has been suggested to regulate sperm fertility [65], and COXs and mPGES-1 are expressed in epithelial cells of the seminal vesicles [59,66]. We show that sPLA₂-IIC, -IID, -IIE, -IIF, -V and -X are expressed in the seminal vesicular epithelium of mice (Fig. 6). sPLA₂-IID and -V are also expressed in human seminal vesicular gland, whereas sPLA₂-IIF and -X are detected only poorly (Fig. 8), thus revealing species difference. It is possible that these sPLA₂s contribute to the amplification of lipid mediator production in the seminal vesicles. Although weak expression of sPLA₂-IIA is seen in human seminal vesicular epithelium (Fig. 8), these cells do not appear to represent a major source of the sPLA₂-IIA present in the seminal plasma [25], since this enzyme is far more intensely detected in the prostate gland, as discussed below.

4.4. Prostate

Several previous immunohistochemical studies reported that sPLA₂-IIA is expressed in epithelial cells of human and bovine prostates, particularly in the posterior lobe and paraurethral glands [26,27]. Consistent with these reports, we now show that sPLA₂-IIA is strongly expressed in epithelial cells of human prostate gland (Fig. 8), thus providing additional support for the notion that the major source of the sPLA₂-IIA secreted in human seminal plasma is the prostate epithelium. It is notable that the sPLA₂ isozymes expressed in human and mouse prostates are rather distinct. Thus, in addition to sPLA₂-IIA, sPLA₂-IID and partially sPLA₂-X are expressed in human prostate epithelium (Fig. 8), whereas the sPLA₂s detected in mouse prostate are sPLA₂-IIC, -IID, -IIE and -IIF (Fig. 6). However, as the expression of these sPLA₂ mRNAs in mouse prostate is relatively low (Fig. 1A), the possibility that these sPLA₂ proteins produced from the upper genital organs are secondary associated with the prostate epithelium should be also considered. Nonetheless, the epithelial location of these sPLA₂s is rather distinct from the distribution of COX enzymes in human prostate [66], arguing against the potential contribution of these sPLA₂s to prostanoid production in this organ. Given that several sPLA₂s exhibit potent bactericidal activity [17–20], it is more likely that these sPLA₂s, particularly the sPLA₂-IIA abundantly secreted from the human prostate gland, may be a part of the anti-microbial arsenal to protect this organ and spermatozoa against microbial invasion. Similarly, various

sPLA₂s secreted from the epithelium of the upper genital organs (epididymis, vas deferens and seminal vesicles) into the lumen may also contribute to anti-microbial defense throughout the genital ducts.

4.5. Conclusion

Our present study has demonstrated, for the first time, the detailed localization of individual sPLA₂s in mouse male genital organs. Distinct sets of sPLA₂s are distributed in spermatogenic and Leydig cells in the testis and epithelial cells throughout the ducts. In general, more than one sPLA₂ is expressed in a single cell type, and these multiple enzymes may play redundant or distinct roles. As discussed above, possible functions of sPLA₂s in male reproductive organs include (i) production of lipid mediators, (ii) regulation of spermatogenesis, (iii) regulation of sperm functions, including the acrosome reaction and fertilization, and (iv) defense against microorganisms invading these tissues. The ability of particular sets of sPLA₂s (IIF, V, and X, which are intrinsically expressed in Leydig cells) to increase PGE₂ production in a mouse Leydig cell line may be a reflection of their redundant function. Nevertheless, to understand the precise roles of individual sPLA₂s in these tissues, a wide array of functional studies, including gene targeting, will be necessary. However, future studies with sPLA₂ knockout mice should be interpreted with caution, because some functions of a particular sPLA₂ may be compensated for by other sPLA₂s and because there are several differences in the expression patterns of sPLA₂s between mice and humans, as exemplified in seminal vesicles and prostate.

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