

RESEARCH ARTICLE

PKC δ deficiency inhibits fetal development and is associated with heart elastic fiber hyperplasia and lung inflammation in adult PKC δ knockout mice

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Abstract

Protein kinase C-delta (PKC δ) has a caspase-3 recognition sequence in its structure, suggesting its involvement in apoptosis. In addition, PKC δ was recently reported to function as an anti-cancer factor. The generation of a PKC δ knockout mouse model indicated that PKC δ plays a role in B cell homeostasis. However, the *Pkcrd* gene, which is regulated through complex transcription, produces multiple proteins via alternative splicing. Since gene mutations can result in the loss of function of molecular species required for each tissue, in the present study, conditional PKC δ knockout mice lacking PKC δ I, II, IV, V, VI, and VII were generated to enable tissue-specific deletion of PKC δ using a suitable Cre mouse. We generated PKC δ -null mice that lacked whole-body expression of PKC δ . PKC δ +/- parental mice gave birth to only 3.4% PKC δ -/- offsprings that deviated significantly from the expected Mendelian ratio ($\chi^2(2) = 101.7, P < 0.001$). Examination of mice on embryonic day 11.5 (E11.5) showed the proportion of PKC δ -/- mice implanted in the uterus in accordance with Mendelian rules; however, approximately 70% of the fetuses did not survive at E11.5. PKC δ -/- mice that survived until adulthood showed enlarged spleens, with some having cardiac and pulmonary abnormalities. Our findings suggest that the lack of PKC δ may have harmful effects on fetal development, and heart and lung functions after birth. Furthermore,

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our study provides a reference for future studies on PKC δ deficient mice that would elucidate the effects of the multiple protein variants in mice and decipher the roles of PKC δ in various diseases.

Introduction

Protein kinase C (PKC) is a phospholipid-dependent serine/threonine kinase, first identified in 1977 by Nishizuka et al., and it plays critical roles in intracellular signal transduction [1–3]. Mammalian PKCs form a large family, categorized based on their molecular structures and activation mechanisms as conventional PKC (cPKC), requiring calcium, phosphatidylserine, and diacylglycerol for activation (α , β , and γ isoforms); novel PKC (nPKC), which do not require calcium for activation (δ , ϵ , η , and θ isoforms); and atypical PKC, which do not require calcium or diacylglycerol for activation (ζ and λ/i isoforms). PKC δ belongs to the nPKC family, and PKC δ I was the first PKC δ molecular species to be reported. PKC δ I is expressed ubiquitously in various tissues and cells, suggesting a general role rather than a tissue- or cell-specific function [4]. PKC δ is involved in many cellular processes [5,6], including cell growth [7], apoptosis [8,9], tumor inhibition [10], and cell migration [11]. cPKCs and nPKCs, including PKC δ , are activated by the oncogenic promoter phorbol 12-myristate 13-acetate and, hence, are considered as drivers of tumorigenesis [4]. However, despite over 30 years of clinical trials investigating PKC inhibitors as anti-cancer agents, PKC inhibitors have failed to show tumor-suppressive effects and, in some cases, have worsened symptoms [12]. Detailed studies have identified PKC δ as a tumor suppressor [13]. Additionally, PKC δ has a caspase-3 cleavage sequence, called the DILD motif, in its V3 domain, indicating a role in apoptosis [6,7,14–16].

Studies based on PKC δ knockout mice (PKC δ KO) have demonstrated that PKC δ is involved in the maintenance of smooth muscle homeostasis and that PKC δ KO mice develop normally and are fertile [17]. PKC δ plays a critical role in B cell homeostasis and tolerance, highlighting its potential role in the treatment of autoimmune diseases [18,19]. Although these studies reported significant findings, the PKC δ KO mice used in these studies lacked only the PKC δ I and δ II isoforms. Studies have shown that several PKC δ isoforms are generated by the alternative splicing from single PKC δ gene (*Prkcd*), and that PKC δ itself forms a family (Fig 1). To date, the expression levels of PKC δ I, II, IV, V, VI, VII, and IX have been reported in mice [20–22], those of PKC δ I and III in rats [23], and those of PKC δ I and VIII in humans [24]. In mice, caspase-3 recognition sequences are present in PKC δ I, IV, and VI. PKC δ II, V, and VII have a 78-bp insertion sequence with no frameshift in the DILD motif; therefore, PKC δ II, V, and VII are not cleaved by caspase-3 [21]. PKC δ VIII, the human homolog of the mouse PKC δ II [24], exerts anti-apoptotic effects in NT2 cells [25,26]. Moreover, insulin enhances cell growth by further promoting alternative splicing of PKC δ II in HT22 cells [27]. It has been suggested that the PKC δ variants may have distinct functions.

Exons are indicated by orange, red, light blue, and yellow boxes and numbered according to a report on the mouse genome by Suh et al. [28]. Red, exon 4a; blue, 78 bp insertion in the caspase-3 recognition site; yellow, specific exon for PKC δ VI and VII; M, the first methionine; and asterisks, the termination codon. Introns are shown with horizontal lines. PKC δ isoform-characteristic exons are shown under the schematic structures of the entire PKC δ genomic DNA.

The complex regulation of gene transcription and the production of multiple splice variants, each with their own functions, indicates the efficiency of gene usage. However, mutations

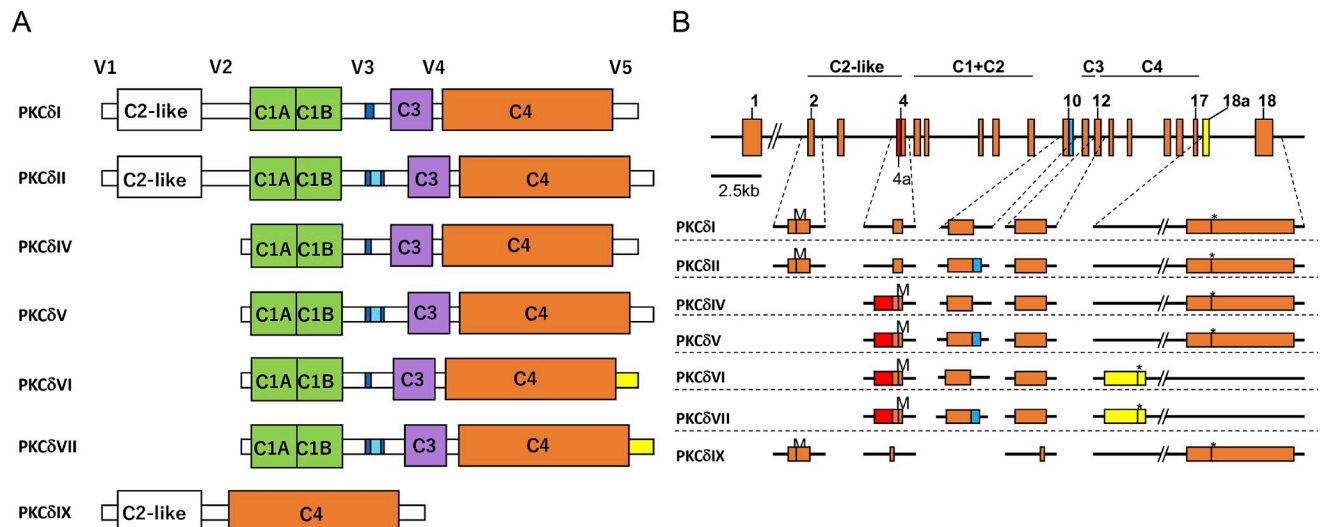


Fig 1. Schematic structures of mouse protein kinase C delta (PKC δ) isoforms and genomic DNA. (A) PKC δ I and II cDNAs consist of variable domains V1, V2, V3, V4, and V5, and conserved domains C2-like, C1a, C1b, C3, and C4. The C1 regulatory domain contains two cysteine-rich zinc finger motifs. PKC δ IV, V, VI, and VII consist of C1a, C1b, V3, C3/V4/C4, and V5 domains. Dark blue, DILD caspase-3 recognition site; light blue, 78 bp insertion in DILD caspase-3 recognition site; and yellow, specific exon for PKC δ VI and VII [20–22]. (B) Schematic structure of the PKC δ genome.

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at key points in a gene can result in the simultaneous loss of multiple proteins. In the present study, we generated mice with comprehensive deletion of the PKC δ -encoding gene, to elucidate the functions of PKC δ . Thus, in this study, a new PKC δ -deficient mouse model was generated by deleting the exon shared by PKC δ I, II, IV, V, VI, and VII. We report that this mouse model yielded different data than what has been previously described.

Materials and methods

Mouse breeding

All mice procedures were carried out in accordance with the guidelines laid down by the Institutional Animal Care and Use Committees and the Ethics Committees of Tokyo Metropolitan Institute of Medical Science (approval number 15085, 16018, 17026, 18010), Niigata University (approval number SA00542), and Showa University (approval number 50040, 51014, 52016) approved this study.

Generation of PKC δ -flox mice

Three genomic DNA fragments of the PKC δ gene (*Prkcd*): 4.6 kb of the 5' arm, 0.85 kb of the region including exon 7, and 6 kb of the 3' arm were amplified using polymerase chain reaction (PCR) from the mouse genomic bacterial artificial chromosome RP23-283B12 (Thermo Fisher Scientific, Waltham, Massachusetts, USA). The targeting vector contained the 5' arm gene fragment upstream of the first loxP sequence, pgk-1 promoter-driven neomycin phosphotransferase gene, and exon 7 (86 bp) of the PKC δ gene downstream of the first loxP sequence, with the 3' arm gene fragment downstream of the second loxP sequence. Additionally, the targeting vector contained a diphtheria toxin gene for negative selection (Fig 2A), and was electroporated into the C57BL/6N-derived embryonic stem (ES) cell line, RENKA [29], after linearization by SalI digestion. Homologous recombinants were identified by Southern blotting. After digestion with EcoRI and BamHI, the genomic DNA samples were hybridized

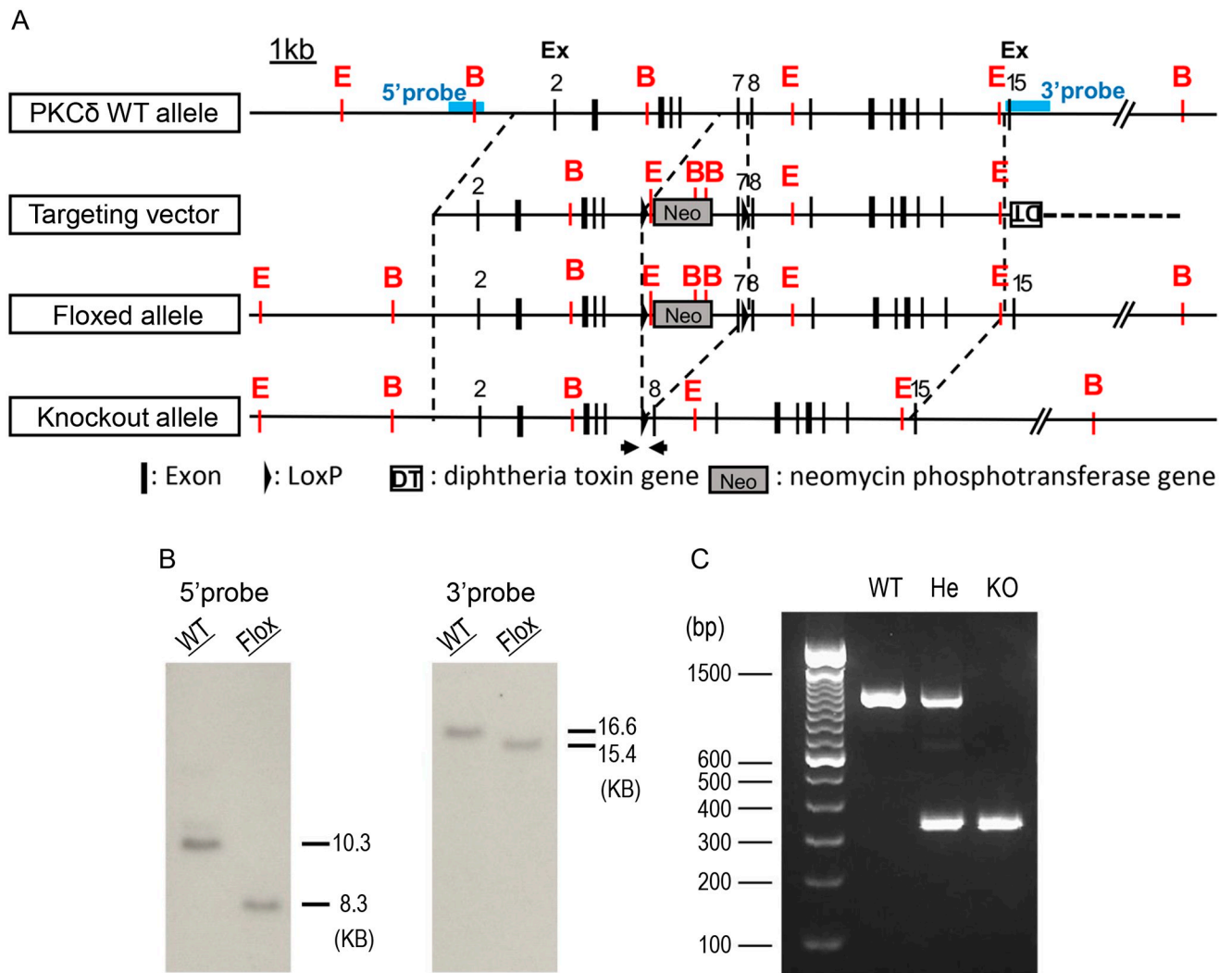


Fig 2. Generation of protein kinase C delta (PKC δ)-deficient mice. (A) Schematic representation of murine *Prkcd* genes, targeting vectors, targeted allele, and knockout allele. Numbers denote the exon numbers, and light blue boxes denote 5' - and 3' - probes for Southern blotting. Polymerase chain reaction (PCR) primers for genotyping are indicated by arrows. Red letters E and B represent EcoRI and BamHI, respectively. Dashed line indicates vector sequence. (B) Southern blotting using genomic DNA of wild type (WT) and flox+/flox+ mouse. DNA digested using EcoRI was used for 5'-probes and shows sizes of 10.3 KB for WT and 8.3 kb for flox+/flox+ mice. DNA digested using BamHI was used for 3'-probes and shows sizes of 16.6 kb for WT and 15.4 kb for flox+/flox+ mice. (C) Genotypic analysis by PCR. PCR analysis was performed using genomic DNA extracted from 4-week-old mice tails and the amniotic membranes of fetuses. WT shows bands of 1078 bp, KO of 315 bp, and He of both 1078 and 315 bp.

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with 5' - and 3' -probes, as shown in Fig 2A. DNA was separated on a 0.6% agarose gel, followed by transfer to a Biotrans Plus membrane (Pall Corp., New York, USA). The membrane was hybridized with DNA probes labeled with the PCR DIG probe synthesis kit (Roche, Basel, Switzerland). The probes were detected with an alkaline phosphatase-conjugated anti-DIG antibody and visualized with a DIG luminescent detection kit (Roche). EcoRI digested genomic DNA hybridized with a 5'-probe showed 10.3 kb for wild type (WT) and 8.2 kb for the targeted allele. BamHI digested genomic DNA hybridized with the 3'-probe showed 16.6 kb for the WT allele and 15.4 kb for the targeted allele (Fig 2B). ES cells with the correct recombination were used to produce chimeric mice.

Generation of PKC δ deficient mice

PGK2-Cre mice [Tg (Pkg2-cre) 24Shb] [30] maintained at Showa University, specifically express Cre recombinase in spermatocytes, and were used for mating with the PKC δ -flox mice to obtain PKC δ -deficient mice. From the offsprings, we selected PKC δ +/- (He, hetero-type knockout) and Cre-/- mice and mated them to obtain homozygous PKC δ -/- mice (KO, PKC δ knockout).

Genotyping PKC δ deficient mice using PCR

Genotyping was performed by PCR using genomic DNA from the tail of 4-week-old offsprings, and the amnion from the embryos using the Tks Gflex DNA polymerase (TAKARA, Kusatsu, Shiga, Japan) under the following conditions: 2 min at 94°C, 35 cycles of 20 sec at 94°C, 1 min at 68°C, followed by 2 min at 68°C. The primers used for genotyping are given in Fig 2, and their sequences are as follows: forward, 5'-GCAGGTGGTGAGTGTTCCCTT-3'; reverse, 5'-GGCATGTTCGATGTTGAAGCG-3'. The size of the PCR products was 1078 bp for WT and 315 bp for KO allele.

Histological analysis

Hearts, lungs, and spleens were removed from WT (PKC δ +/+) and KO mice at 16 and 24 weeks of age for histological analysis. After mating 11 pairs of PKC δ He mice, E11.5 embryos were collected after confirming the number of embryo sacs. Genomic DNA from the amniotic membranes from E11.5 was used for fetal genotyping. Murine tissues and E11.5 embryos were fixed using 4% paraformaldehyde solution at 4°C overnight, dehydrated, defatted through an ethanol- and xylene-series, and then paraffin-embedded. Each tissue was sliced into 5- μ m thick sections and stained with Hematoxylin and Eosin (H&E stain). The heart and lung sections were stained using Masson's trichrome (MT) and Elastica van Gieson staining (EVG) to stain collagen fibers and elastic fibers, respectively. We used Hematoxylin solution made by Muto Pure Chemicals Co. Ltd. (Tokyo) and Eosin solution by Sakura Finetek Japan Co. Ltd. (Tokyo). For both MT and EVG, we used staining solutions made by Muto Pure Chemicals Co. Ltd. (Tokyo).

Statistical analysis

The statistical difference between our results and Mendel's law was analyzed using the chi-square test. The Kaplan–Meier method was used to analyze the survival of PKC δ KO mice.

Results

Generation of PKC δ -deficient mice

In the present study, PKC δ -deficient mice were generated to analyze the function of PKC δ . PKC δ is reported to generate eight molecular species (PKC δ I, II, IV, V, VI, VII, and IX) from a single gene (*Prkcd*) in mice [20–22]. The structures of the mouse PKC δ molecular species in mice are shown in Fig 1A. Studies on PKC δ knockout mice (KO) reported that KO mice were not deficient in all PKC δ molecular species [17,18]. Therefore, to fix the lineage of PKC δ knockout mice into C57BL6/N mice, we generated conditional knockout C57BL6/N mice with deficient PKC δ I, II, IV, V, VI, and VII using the RENKA, C57BL/6N ES cell line.

The relationship between the PKC δ gene structure and the different molecular species generated by alternative splicing is shown in Fig 1B, with the genetic architecture previously described by Suh et al. [28]. In the present study, we inserted LoxP sequences in the upstream and downstream regions of exon 7 of 86 base pairs, that could reliably frameshift due to the

defect and prevent protein generation (Fig 2A). Homologous recombination was performed using the RENKA C57BL/6N ES cell line [29] to generate PKC δ conditional knockout mice (PKC δ flox/+). PKC δ flox/flox mice were further generated by self-mating PKC δ flox/+ mice, which were then enrolled (registered: C57BL/6 Prkcd<tm1Shb>). Confirmation of this genetic insertion of PKC δ flox/ flox mice was performed by genomic Southern blotting with C57BL/6N wild-type mice (Fig 2B). PKC δ flox mice were crossed with PGK2–Cre mice [Tg (Pkg2-cre) 24Shb] [30] to generate conditional knockout of PKC δ genes (PKC δ KO). Since PGK2–Cre mice express Cre recombinase in spermatocytes and spermatids in the testis, the resulting male mice have a PKC δ gene deletion in their spermatozoa. PKC δ +/- mice with Cre-/- genotypes obtained by crossing with wild-type PKC δ +/+ (WT) were enrolled as PKC δ +/-, which means PKC δ gene Heterozygous mice (He) (registered: C57BL/6-Prkcd<tm1.1Shb>).

Male and female PKC δ He mice were mated, offspring were weaned four weeks after delivery, and the genotypes were confirmed using PCR. Representative PCR data are shown in Fig 2C. The genotypes of the resulting offspring at weaning are indicated in Table 1. The proportion of genotypes in the offspring mice was expected to be WT:He:KO = 1:2:1, whereas the observed proportion was WT:He:KO = 104 (27.1%):267 (69.5%):13 (3.4%). The genotypes of all 384 offspring born from pairs of PKC δ He mice did not follow Mendelian rules (By χ^2 test, $\chi^2 = 101.7$, $p < 0.001$).

The lifespan of all PKC δ KO mice born to pairs of PKC δ He mice was as follows: 2.5 months (n = 1), 3 months (n = 2), 6 months (n = 1), 7 months (n = 1), 11 months (n = 2), 13 months (n = 1), and 15 months (n = 1). Four other knockout mice were used in the experiment. These nine PKC δ KO mice were kept simultaneously with their littermate WT and He mice, and their Kaplan-Meier survival curves are shown in Fig 3. However, since WT and He mice were euthanized at 1 year (52 weeks old), the study by Yuan R. et al. [31] was cited as supplementary data for WT. No spontaneous death was observed in WT and He mice within 1 year. In contrast, KO mice started to die when they were 10 weeks old. As a result, it was found that the 50% survival rate of the KO mice was 31.8 weeks, which was 1/4 that of WT mice.

Histological analysis

Eight mice were euthanized for histological analysis. Four PKC δ KO mice (one male and two female 24-week-old mice, one male 16-week-old mouse), and four gender- and age-matched WT mice descend from a pair of PKC δ He parents. Visual observations revealed that all PKC δ KO mice had enlarged spleens compared to those of PKC δ WT mice, and the hearts were enlarged in three of the four PKC δ KO mice (Fig 4A). Additionally, conspicuous deformation was observed in the lungs of three out of four KO mice (Fig 4A). These findings are summarized in Table 2. No visible differences were observed in the other organs. Histological analysis

Table 1. Genotypic analysis of 4-week-old offspring mice at weaning from heterozygous parents.

	WT	He	KO
Male	53	144	3
Female	51	123	10
Total	104	267	13
Rate of offspring	27.1%	69.5%	3.4%

The genotypes of all 384 offspring born from pairs of PKC δ . He mice did not follow Mendelian rules ($\chi^2(2) = 101.7$, $p < 0.001$).

Genotypes: WT, wild type mice; He, hetero-type knockout mice; KO, homozygous PKC δ -/- mice (PKC δ knockout).

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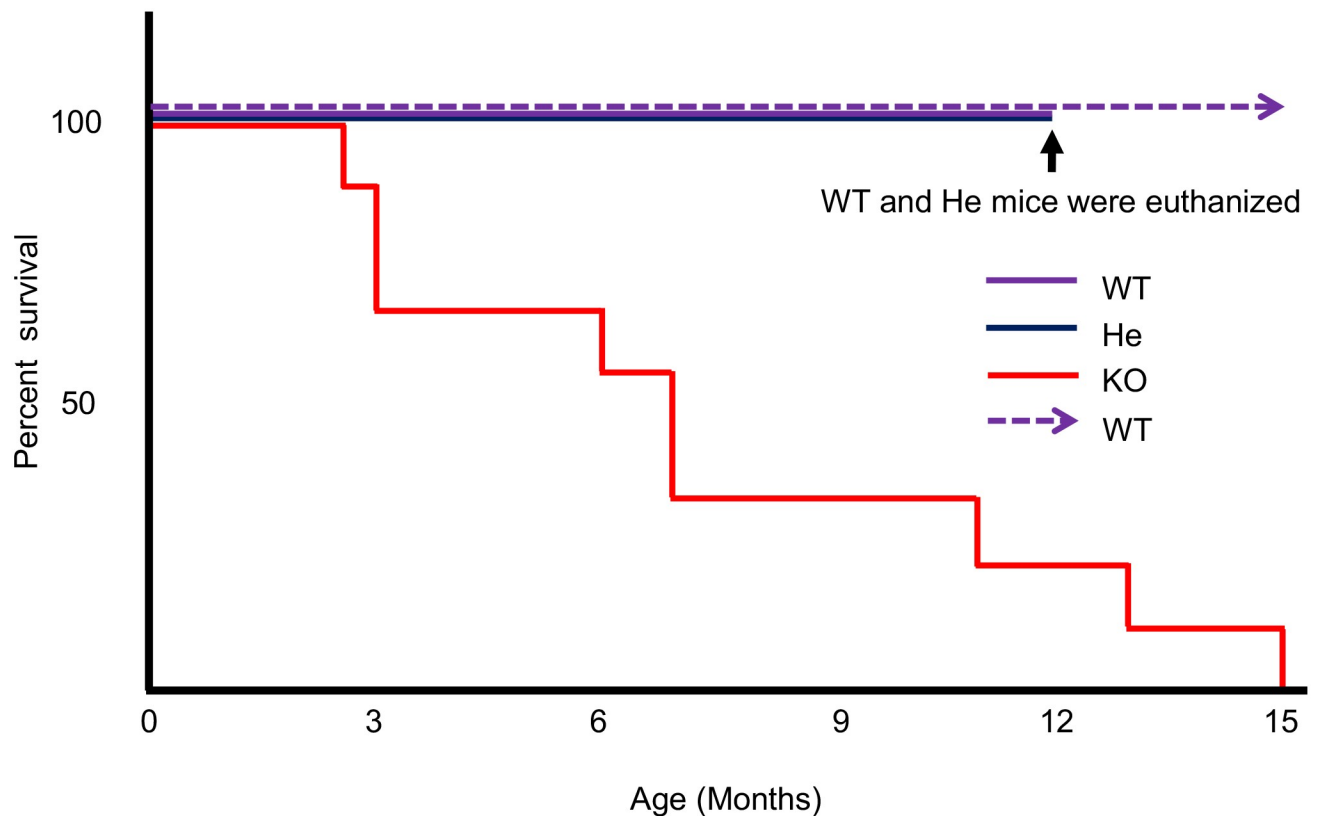


Fig 3. Survival curve of PKC δ KO mice (Kaplan-Meier plots). PKC δ He mice were spontaneously bred, the genotypes of the offsprings were determined, and their survival was monitored. KO mice (all nine KO mice) were observed until all mice died, whereas WT and He mice were euthanized at 52 weeks of age. Therefore, the study by Yuan et al. [31] is cited as supplementary data for WT and is indicated by the dashed arrow.

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was performed for each of the above organs. Calcification was observed in the heart of two out of four PKC δ KO mice, and inflammation was observed in the lungs of three out of four KO mice, indicating poorly air space, as observed by H&E staining (Fig 4B). In the two cases with a cardiac abnormality, a calcific mass above the left ventricle in one mouse (Fig 4B and 4C) and calcification was observed at the base of the cardiac mitral valve in the other mouse (Fig 4D) were observed. The calcification in both cases observed in the hearts of male mice. MT and EVG staining were performed on heart sections of mice with calcification in the heart to examine the reason for the calcification. In MT staining, collagen fibers were stained blue with aniline blue solution and the cytoplasm was stained red with Masson's solution. In EVG staining, elastic fibers were stained black with resorcinol fuchsin solution, collagen fibers were stained red with Sirius red solution, and muscle fibers were stained yellow with picric acid. The staining indicated a clear increase in elastic fibers in the hearts of PKC δ KO mice compared to the hearts of WT mice (Fig 5A). In both PKC δ KO mice hearts, EVG staining revealed that there was a hyperplasia of the elastic fibers in the endocardium. In addition, there was a slight trend in the blood vessels of the heart. Furthermore, the results of MT staining showed an increase in collagen fibers; however, the changes in elastic fibers were more intense and predominant. This may have resulted in lime-induced thrombus; however, a causal relationship could not be established. An increase and calcification of elastic fibers in the endocardium were observed in the hearts of two of the four PKC δ KO mice. However, it was difficult to determine whether this change was significant. In addition, MT and EVG staining were

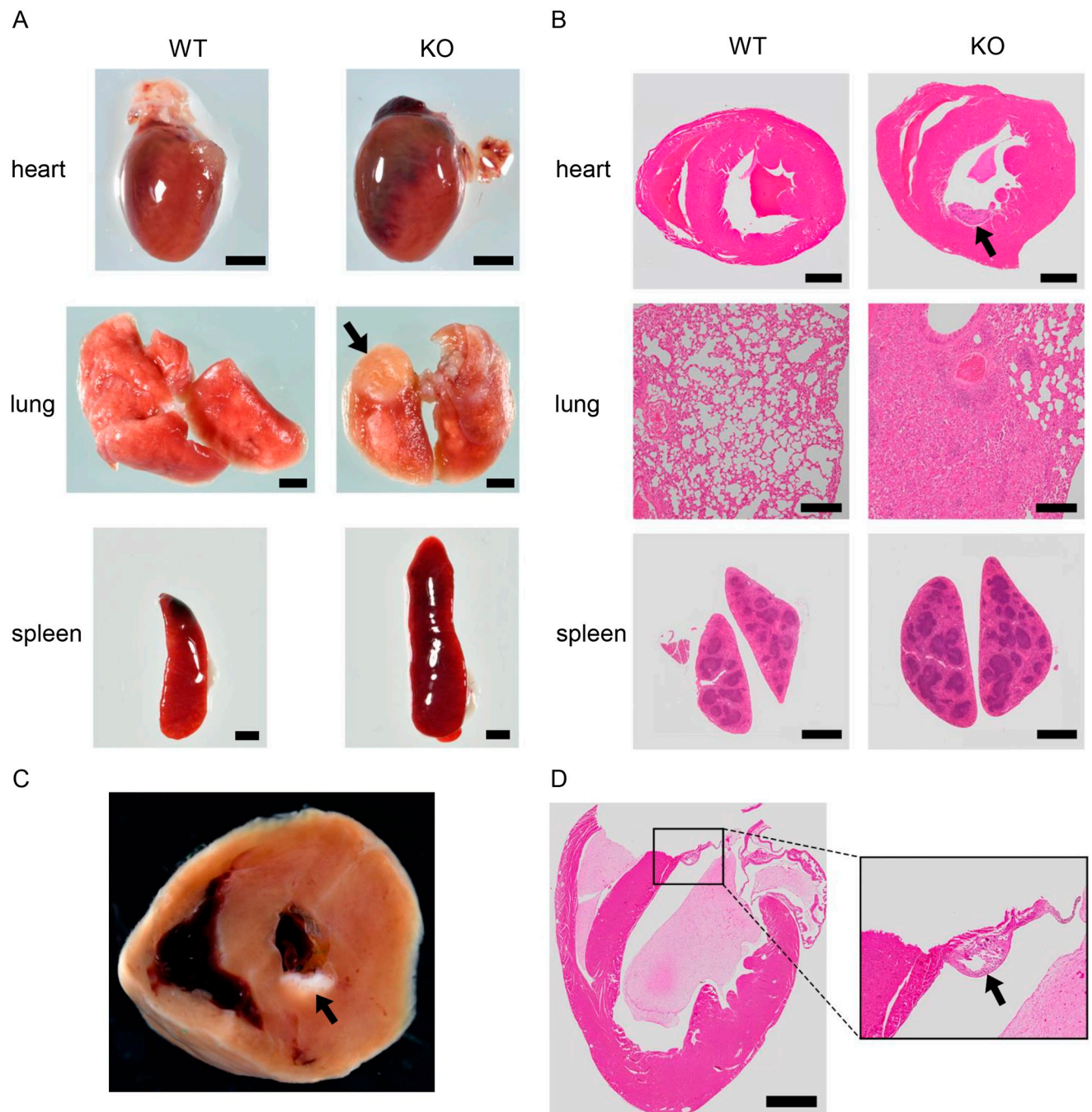


Fig 4. Histological analysis of WT and KO mice. (A) Comparison of heart and spleen sizes between WT and KO mice. Arrow indicates the site of deformity in the lung. Scale bars represent 2 mm. (B) Hematoxylin and eosin-stained images of hearts, lungs, and spleens from WT and KO mice. The heart image shows a cross-section. Scale bars represent 1 mm for the heart and spleen and 200 μ m for the lung. Arrow indicates the site of calcification. (C) Heart of a KO mouse with internal calcification. Arrow indicates the site of calcification. Inset shows calcification at the base of the mitral valve. Scale bars represent 1 mm. (D) Hematoxylin and eosin-stained images of vertically cut hearts from KO mice. Arrow indicates the site of calcification.

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Table 2. Phenotypic analysis of KO-mice.

	Age	Sex	Heart		Lung	Spleen
			Enlarged	Calcification	Inflammation	Enlarged
1	16 w	Male	+	+	+	+
2	24 w	Male	+	+	+	+
3	24 w	Female	+	-	+	+
4	24 w	Female	-	-	-	+

Age, sex, and phenotype of the mice (n = 4) subjected to histological analysis are shown.

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performed on the hearts of PKC δ KO mice without cardiac enlargement and there was no difference was observed compared to the WT mice. Examination of lung abnormalities revealed that three PKC δ KO animals showed more eosinophilic exudates and macrophages in the alveoli, and more round cell infiltrates compared with those of the PKC δ WT mice. Macrophages were also found in the bronchi. Despite strong congestion and inflammation in the lungs, the increase in fiber was inconspicuous. Lung abnormalities appeared to be a change resulting from decreased cardiac function (Fig 5B).

Analysis of 11.5-day-old embryos

To verify the reason for the low production rate of approximately 3.4% of PKC δ KO mice from crosses between PKC δ He mice, we analyzed fetuses on embryonic day 11.5 (E11.5). PKC δ He mice were self-crossed, and 11 pregnant mice were examined. The uterus was removed, and the fetus was collected after confirming the number of fetal sacs. The total number of fetal sacs was 104, and 103 fetuses were identified. One fetal sac contained no fetus, as it had already been resorbed. The fetal implantation rate of each genotype is shown in Table 3A. Genotypes were determined by PCR using genomic DNA from the amniotic membrane. The PCR patterns of WT, He, and KO mice are shown in Fig 2C. Since the PKC δ IV, V, VI, and VII isoforms were expressed in a testis-specific manner, we thought that PKC δ knockout mice might have some problems in the early stages of development such as fertilization, early cleavage, and implantation. However, our results demonstrated the presence of 8–12 fetal sacs in the uterus of 10 out of 11 mother mice. The genotype of the fetuses in the fetal sacs at E11.5 is shown in Table 3A. It suggests that all genotypes of WT, He, and KO follow the Mendelian rule ($\chi^2(2) = 2.077$, n.s.), and fertilization, early cleavage, and implantation (placenta confirmed) were not problematic. However, as shown in Table 3B, the survival rate of KO fetuses was significantly lower than that of the WT and He embryos at E11.5 ($\chi^2(2) = 30.892$, $p < 0.001$). This suggests that KO fetuses have problems developing in the mother's body, which explains the one possible reason for the low birth rate of KO mice. The implantation rate of PKC δ KO fetuses was 23.1% of the total, but the survival rate decreased in the mother's body and further decreased to 3.4% at 4 weeks of age (Table 1).

As representative E11.5 samples, fetuses of mother numbers 8 and 11 are shown in Fig 6A and 6B. Fetuses with heartbeats on E11.5 (fetus numbers 2 (WT) and 7 (KO) in Fig 6A, and 3 (WT) and 10 (KO) in Fig 6B) were fixed using 4% paraformaldehyde solution and stained with HE for histological examination (Fig 7A and 7B). The fetuses of PKC δ KO mice were slightly smaller than those of WT mice, and their Sclerotome and spinal cord appeared to be immature compared with those of the WT mice (Figs 6 and 7). Although there was no difference in the size of the heart between KO and WT mice, the ventricular wall was thin in the KO mice. These are especially mesoderm-derived organs. Additionally, there were no obvious

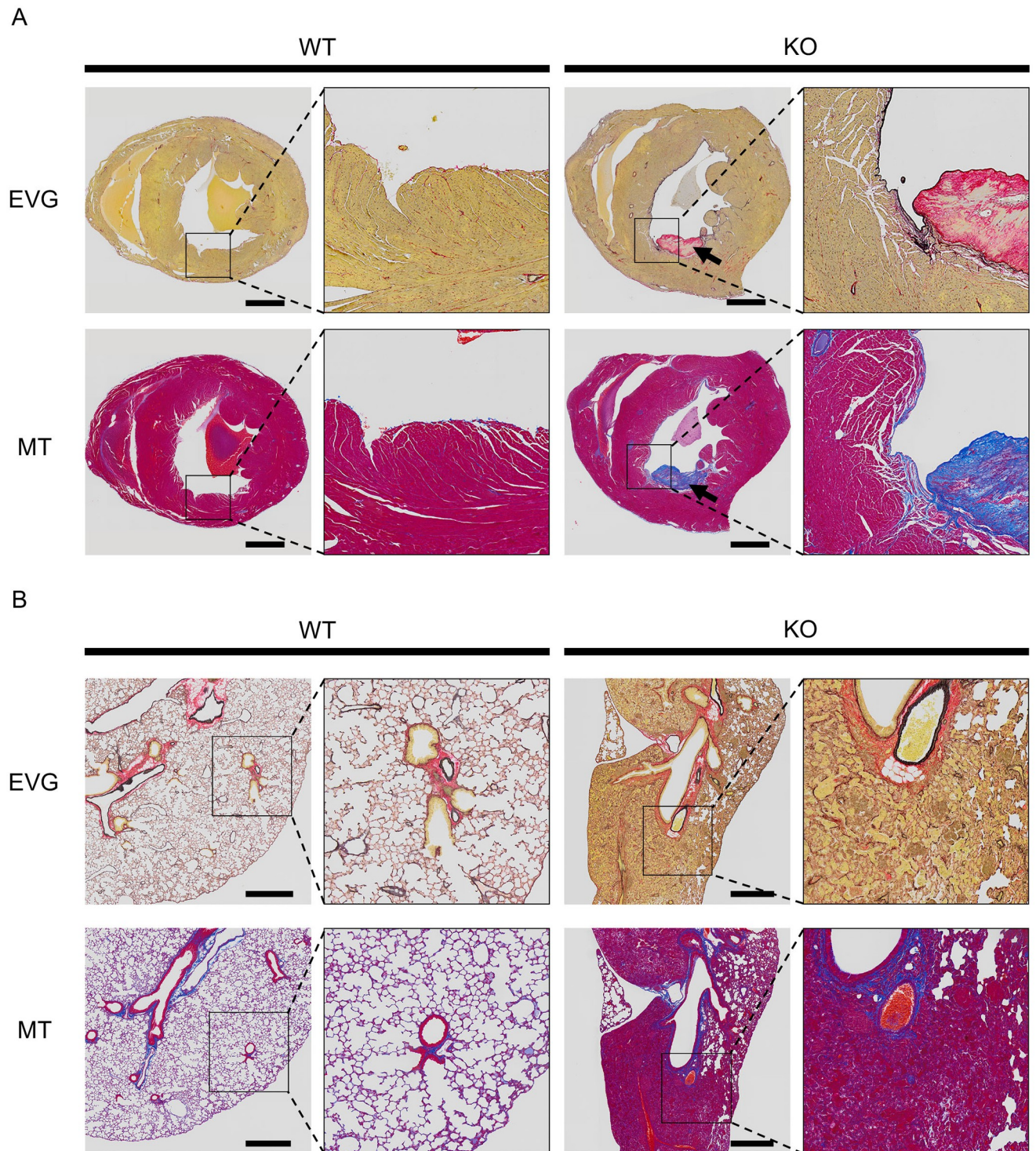


Fig 5. Images of elastic fiber staining of hearts from WT and KO mice. (A) Elastica van Gieson (EVG) and Masson's trichrome (MT) staining were performed using horizontally cut hearts from WT and KO mice. An enlarged view is shown on the right. Scale bars represent 1 mm. Arrow indicates the site of calcification. (B) EVG and MT staining images of lungs from WT and KO mice. An enlarged view is shown on the right. Scale bars represent 500 μ m.

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Table 3. Number of fetal sacs and embryos at E11.5.

(A)			
	Number of fetal sacs holding fetuses of each indicated genotype at E11.5		
	WT	He	KO
Total	22	58	24*
Pregnancy Rate	21.2% (n = 22/104)	55.8% (n = 58/104)	23.1% (n = 24/104)
			* One case was absorbed.
			$\chi^2(2) = 2.077$, n.s.
(B)			
	Number of fetuses of each indicated genotype at E11.5		
	WT	He	KO
Alive	16	52	7
Dead	6	6	17
Surviving rate of fetus	72.7% (n = 16/22)	89.7% (n = 52/58)	29.2% (n = 7/24)
			$\chi^2(2) = 30.892$, p < 0.001

After mating of PKC δ He mice, 11.5-day-old embryos from mothers (n = 11) were analyzed. (A) Genetic analysis of fetal sac was carried out. A Chi-squared test was applied to test whether the rate of pregnancy follows Mendelian rules. (B) The embryos with confirmed heartbeat were denoted as surviving embryos. Difference in the rate of survival among the genotypes was tested using a Chi-squared test.

Genotypes: WT, wild type mice; He, heterozygous knockout mice; KO, homozygous PKC δ ^{-/-} mice (PKC δ knockout).

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differences between the PKC δ He and PKC δ WT fetuses. These findings suggest that at E11.5, PKC δ KO mice have an overall poor developmental status, and their cause of death is considered to be poor overall growth rather than organ-specific abnormalities. These results indicate that the absence of the PKC δ gene is detrimental to fetal growth.

Discussion

In the present study, we designed a murine model lacking exon 7 of *Prckd*, which is utilized by most of the reported PKC δ splicing variants when generating PKC δ conditional knockouts. Therefore, LoxP sequences were inserted in the 5' upstream and 3' downstream regions of exon 7 of the *Prckd* gene encoding PKC δ , to generate PKC δ -flox mice using the C57BL/6N-derived RENKA ES cell line. By mating these mice with Cre mice according to each research objective, it is possible to produce mice with a conditional knockout of PKC δ molecules. Other studies on PKC δ knockout mice have been reported; however, they only knocked out PKC δ I and δ II, and did not achieve a complete knockout of all PKC δ family of proteins [17,18]. For instance, considering that the first exon of the PKC δ gene is a non-coding exon [28], Leitges M et al. generated PKC δ KO mice by eliminating the function of the second exon of the PKC δ gene. The resulting mice had a 129/SV background, developed normally, and were fertile. Moreover, these PKC δ KO mice had markedly increased arteriosclerotic lesions in the vein grafts compared with WT mice. Those mice with atherosclerotic lesions also had substantially more smooth muscle cells than the WT animals [17]. Additionally, Miyamoto et al. generated PKC δ knockout mice lacking exons 2 and 3. These mice had a 129/Sv background, were fertile, and survived up to 12 months despite the excessive proliferation of B cells and autoimmunity that was observed due to the lack of PKC δ [18]. Furthermore, in 2002, Mecklenbrauker et al. reported that PKC δ is an essential component of signaling pathways specific for inducing tolerance in B cells [19].

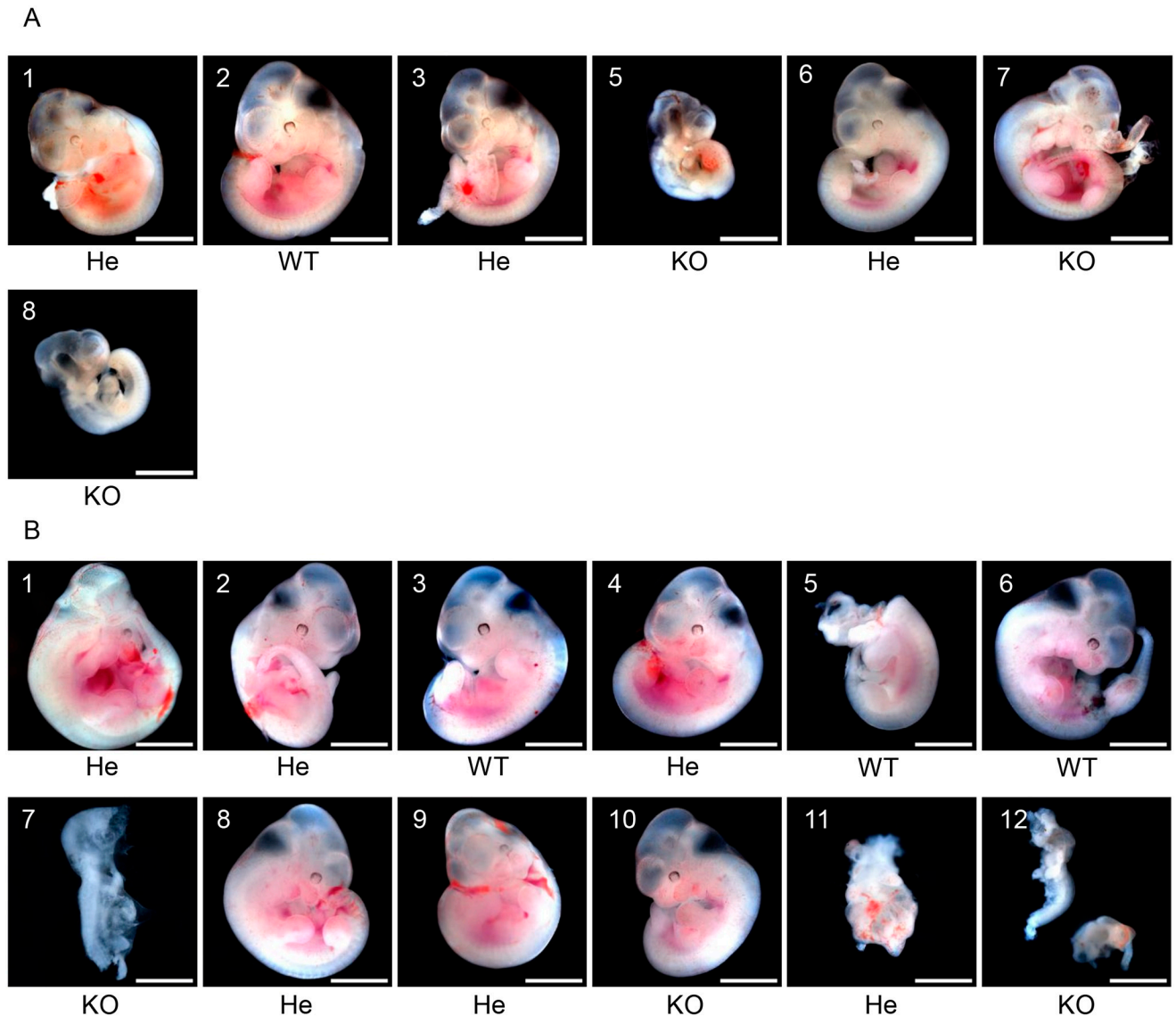


Fig 6. 11.5-day-old embryo (E11.5) generated by crossing He x He. The embryos of two mice out of 11 mother mice in Table 2 are shown in (A) and (B). The photographs show each individual and genotypes determined by their amniotic membranes. In (A), fetus number 4 was already absorbed. Scale bars represent 2 mm.

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Considering these other related studies, ours is the first to generate PKC δ KO mice with a C57BL/6N background that can reliably delete the six PKC δ species (PKC δ I, II, IV, V, VI, and VII). Many reports have described the importance of PKC δ functions, including its role in cell proliferation, cell death, and as a tumor suppressor [9–11,32–34]. In addition, the PKC δ gene produces several molecules from a single gene [20–24]. These findings suggest that a single mutation deletes more than one PKC δ molecular species, indicating the importance of this mouse model. Moreover, murine PKC δ contains a recognition sequence (DILD) for caspase-3, a member of the caspase family of cysteine proteases in its V3 domain, indicating its involvement in apoptosis [16,34–36].

Previous studies using PKC δ I- and δ II-null mice have investigated its role in apoptosis [37,38], defective osteoblast differentiation during embryonic development [39], defective

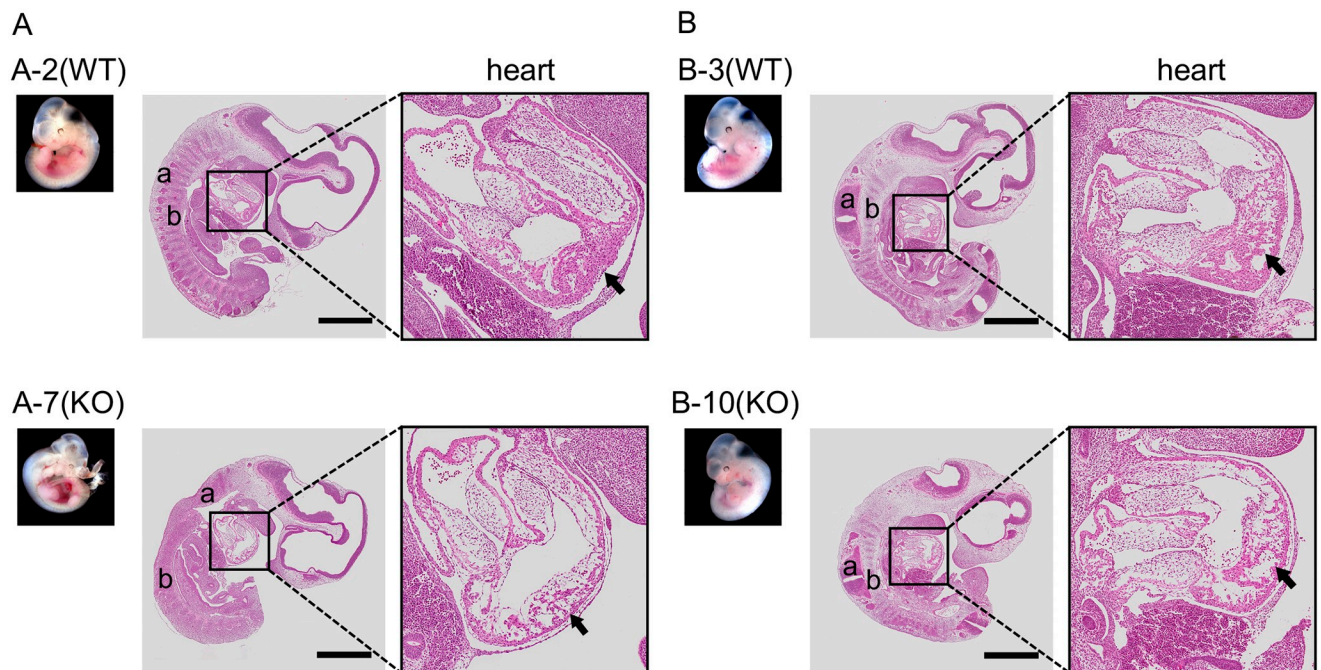


Fig 7. Histological analysis of WT and KO mice on E11.5. (A) Hematoxylin and eosin-stained image obtained by vertically cutting embryo numbers 2 (WT) and 7 (KO) in Fig 5A. (B) Hematoxylin and eosin-stained image obtained by vertically cutting embryos 3 (WT) and 10 (KO) in Fig 5B. Scale bars represent 1 mm. a, spinal cord; b, sclerotome.

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osteoclastic bone resorption [40], as well as an increased proliferation of, and autoimmunity toward, B cells [11,18]. Sites summarizing studies on PKC δ KO mice can be found online (<http://www.informatics.jax.org/marker/phenotypes/MGI:97598>).

In the present study, whole-body PKC δ deficiency resulted in the survival of only 3.4% offspring, which differs significantly from previous studies, which reported no apparent issues in PKC δ KO fertility. The discrepancy in these findings may be attributed to differences in the missing PKC δ molecular species due to either differences in the gene knockout position or the mouse strain. Here we have generated a mouse that lacks exon 7. This may have resulted in the expression of a new PKC δ molecular species, resulting in a phenotypic difference from previous PKC δ KO mice. Moreover, considering that PKC δ effectively utilizes the PKC δ gene, it may be possible to discover that another splicing variant is expressed, which functions in the biological reactions of mice having exon deficiency.

It also has been reported that the experiments with sperm from PKC δ KO male mice and oocytes from WT female mice resulted in lower fertilization rates and lower early cleavage rates [41]. However, in our study, results of *in vitro* fertilization with eggs from three WT females using one male PKC δ KO mouse spermatozoa showed no differences from WT male spermatozoa when comparing fertilization rates and developmental conditions up to two cells. Together with the results of several genotypic mating combinations (Table 2), the reproductive performance of PKC δ KO males did not appear to be affected. However, due to the small number of cases, further research is expected on the reproductive ability of PKC δ KO mice.

Examination of the number of implantations due to the mating of PKC δ He showed that the numbers of WT, He, and KO fetuses were almost consistent with Mendelian law, and the implantation of PKC δ KO fertilized eggs was not low. The main reason for the birth of only 3.4% PKC δ KO mice, may be due to the developmental failure of the fetuses as observed by the

analysis of E11.5, generated by mating PKC δ He. However, PKC δ KO did not result in complete embryonic lethality. Future studies are required to determine the reason for the low PKC δ KO birth rate, including the potential use of other PKC molecules to successfully complement its function. Moreover, this low birth rate in mice may also suggest that similar issues will arise in human fetuses with PKC δ KO, given the small number of pregnancies.

PKC δ may be associated with fetal underdevelopment during pregnancy. A possible difference in the present study from previous reports on PKC δ KO mice may be the neglect of the presence of PKC δ IV, V, VI, and VII produced by alternative splicing. PKC δ IV, V, VI, and VII are expressed in adult mice in a testis-specific manner and have not been studied in fetal development [21]. PKC δ IV and V, and PKC δ VI and VII are paired with normal and disrupted caspase-3 recognition sequences. These molecules are also deficient in the KO mice in this study. The mice generated in the present study will also be useful for further studies on PKC δ IV, V, VI, and VII species.

Notably, of the nine KO mice that exhibited spontaneous death, born from crosses between PKC δ He mice, seven died within the first year of life, while all PKC δ KO mice had a short lifespan. Analysis of three 24-week-old PKC δ KO mice and one 16-week-old PKC δ KO mouse showed enlarged hearts in three animals, with two exhibiting calcification in the left ventricle or mitral valves. Moreover, staining of these heart tissues revealed an increase in elastic fibers in the endocardium, which may have led to the calcification within the heart. In addition, inflammation was observed in the lungs of three of the four PKC δ KO mice; however, an increase in fiber was not obvious. Although no calcifications were observed in the hearts of two cases, one had an enlarged heart. It was suspected that inflammation in the lungs might have resulted from a decrease in cardiac function. This is the first study to report calcification in the heart and inflammation in the lungs of PKC δ KO mice. Moreover, as many of the PKC δ KO mice died while they were still fetuses, resulting in small litter sizes, the analysis of adult animals is not considered to be sufficient. In the future, the function of PKC δ in heart and lung lesions, as well as other organs, should be investigated in a larger population.

It has been previously reported that PKC δ is expressed in the mouse heart [42]. Recently, PKC δ and PKC ϵ double knockout mice demonstrated cardiac hypertrophy and thickening of the ventricular wall of the fetal heart [43]. However, in the present study, we observed that some mice had cardiac enlargement and calcification due to a deficiency present only in the PKC δ gene. This enlargement of the heart in PKC δ KO mice may correspond with the hypertrophy reported previously by Song et al. [43]. However, further detailed analysis is required to unambiguously ascertain the cause of the observed enlargement.

It has also been reported that PKC δ gene-deficient mice have altered immune function. PKC δ KO mice develop autoimmune diseases with aging, indicating that PKC δ plays a role in B cell tolerance of autoantigen induction [19]. Research on human lesions has advanced as well. For instance, a missense mutation (c.1528G>A) in the human PKC δ gene via a single nucleotide substitution replaces the amino acid glycine with serine (p. G510S) in children born to parents heterozygous for this missense mutation of PKC δ gene. Children homozygous for this mutation are reported to have systemic lupus erythematosus, caused by a marked decrease in PKC δ function [44]. Furthermore, patients with a homozygous missense mutation (c.1840C>T) in the human PKC δ gene, involving the substitution (p. R614W) from arginine to tryptophan, have several autoantibodies, systemic lymphadenopathy, and hepatosplenomegaly [45]. Since single mutations can cause severe symptoms, gene products must be comprehensively analyzed, particularly for genes like PKC δ , which encode multiple proteins by alternative splicing. Therefore, the mice produced in the current study have the potential to elucidate the effects of the multiple protein variants in mice, as well as humans.

PKC δ also plays a role in inflammatory diseases such as sepsis [46,47], and is reportedly associated with diabetes, with PKC δ regulating glucose production in rats [48], and increased PKC δ expression and activation in diabetic rats [49]. Moreover, continuous hyperglycemia is a common event in patients with type I and type II diabetes. It is reported that PKC δ is involved in advanced glycation end product-induced apoptosis produced by this hyperglycemic exposure [50,51]. Hence, it is expected that the mice produced in the present study will contribute to the elucidation of important functions of PKC δ and its role in various diseases.

Supporting information

S1 Raw images. Original gel images of Fig 2B-1 (DIG), Fig 2B-2 (EtBr) and Fig 2C.
(PDF)

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