

# aPKC $\lambda/\iota$ promotes growth of prostate cancer cells in an autocrine manner through transcriptional activation of interleukin-6

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**Understanding the mechanism by which hormone refractory prostate cancer (HRPC) develops remains a major issue. Alterations in HRPC include androgen receptor (AR) changes. In addition, the AR is activated by cytokines such as interleukin-6 (IL-6). Atypical protein kinase C (aPKC $\lambda/\iota$ ) has been implicated in the progression of several cancers. Herein, we provide evidence that aPKC $\lambda/\iota$  expression correlates with prostate cancer recurrence. Experiments in vitro and in vivo revealed aPKC $\lambda/\iota$  to be involved in prostate cancer cell growth through secretion of IL-6. Further, aPKC $\lambda/\iota$  activates transcription of the IL-6 gene through NF $\kappa$ B and AP-1. We conclude that aPKC $\lambda/\iota$  promotes the growth of hormone independent prostate cancer cells by stimulating IL-6 production in an autocrine manner. Our findings not only explain the link between aPKC $\lambda/\iota$  and IL-6, implicated in the progression a variety of cancers, but also establish a molecular change involved in the development of HRPC. Further, aPKC $\lambda/\iota$  expression might be a biomarker for prostate cancer progression.**

IL-6 | PSA | recurrence

Despite earlier detection and recent advances in surgery and radiation, prostate cancer is the second leading cause of cancer deaths in men in western countries (1). Hormone therapy in the form of medical or surgical castration remains the mainstay of systemic prostate cancer treatment. However, despite initial favorable responses to hormone therapy, hormone refractory tumors develop for which there is as yet no effective treatment (2). The androgen receptor (AR) and its signaling remain intact, as demonstrated by the expression of prostate specific antigen (PSA), in androgen-independent cancer cells. Alterations in these cells include AR amplification, AR point mutations, and changes in the expressions of AR co-regulatory proteins (3). In addition, AR can be activated in a ligand-independent fashion by compounds including growth factors and cytokines, such as interleukin-6 (IL-6) (4–6). Understanding the mechanism of androgen-independent prostate cancer development is essential not only for diagnosis but also more effective therapy.

Atypical protein kinase C (aPKC $\lambda/\iota$  and aPKC $\zeta$ ) is a protein kinase C isozyme distinct from other classes of this enzyme, structurally and functionally (7, 8). It plays multifunctional roles in cellular maintenance and growth of epithelial cells, for example, signal transduction and cell polarity (9–15). Studies on lung, ovary, colon, and breast cancers have demonstrated a relationship between aPKC $\lambda/\iota$  expression and cancer progression and suggest that aPKC $\lambda/\iota$  expression might predict poor survival (16–22). There are several reports showing enhanced aPKC expression in human prostate cancer tissues, but the relationship between aPKC $\lambda/\iota$  and prostate cancer progression remains unclear (23, 24). Furthermore, the mechanism by which

aPKC promotes the progression of a variety of cancers remains uncertain. Then, we focus on aPKC $\lambda/\iota$  expression and its roles in prostate cancer.

IL-6, a cytokine involved in immune and hematopoietic activities, has been implicated in the progression of a variety of human cancers (25). In prostate cancer, IL-6 has been suggested to play a role in cancer progression, especially that of hormone refractory cancer (26–30). Serum IL-6 is elevated in patients with prostate cancer (28). The IL-6 receptor is expressed in prostate cancer cell lines and IL-6 is secreted only by androgen independent prostate cancer cell lines (27, 29, 30). Although these observations suggest the importance of IL-6 in prostate cancer progression, the mechanism by which IL-6 expression is regulated in prostate cancer cells is not fully understood.

Herein, we investigated aPKC $\lambda/\iota$  expression in 29 clinical prostate cancer tissue specimens and found a correlation between aPKC $\lambda/\iota$  expression and PSA failure, a clinical hallmark of recurrence. Experiments, both in vitro and in vivo, on androgen-independent cancer cells, employing siRNA-mediated depletion of aPKC $\lambda/\iota$ , revealed this isozyme to be involved in the proliferation of prostate cancer cells. The in vitro experiments further showed aPKC $\lambda/\iota$  involvement in the secretion of IL-6 into the culture medium, suggesting cancer cell growth to occur via an autocrine mechanism. This finding demonstrates the link between aPKC $\lambda/\iota$  and IL-6, both of which have been implicated in cancer progression. We also demonstrated that aPKC $\lambda/\iota$  is involved in transcription of the IL-6 gene promoter through activation of NF $\kappa$ B and AP-1, both of which have been implicated in transcription of the IL-6 gene in prostate cancer cells. We conclude that aPKC $\lambda/\iota$  promotes prostate cancer cell growth in an autocrine manner via stimulation of IL-6 production and secretion.

## Results

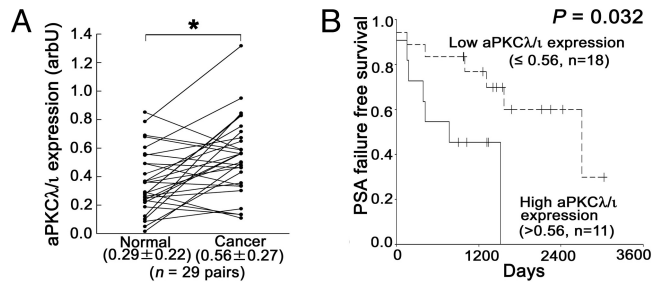
**Correlation Between aPKC $\lambda/\iota$  Expression and PSA Failure, a Clinical Marker of Prostate Cancer Recurrence.** To clarify the relationship between aPKC $\lambda/\iota$  expression and prostate cancer, we first evaluated aPKC $\lambda/\iota$  expression at the mRNA levels in 29 specimens of human prostate cancer tissues. Clinicopathological features are listed at [Table S1](#). Real-time quantitative PCR (qPCR) analyses revealed that aPKC $\lambda/\iota$  mRNA was more highly expressed in cancer tissue samples than in paired normal controls

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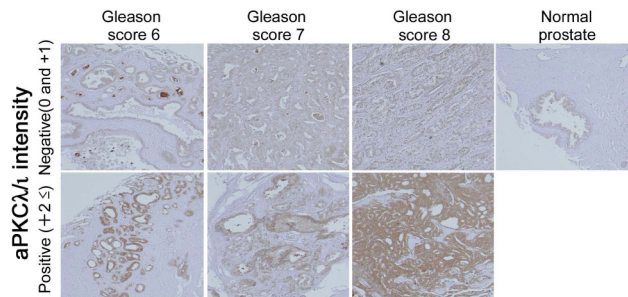
The authors declare no conflict of interest.

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**Fig. 1.** Relationships between aPKC $\lambda/i$  expression in prostate cancer tissues. (A) aPKC $\lambda/i$  expression was compared between paired prostate cancer and normal (BPH) prostate tissues obtained from same patients ( $n = 29$  pairs). \*,  $P = 0.001$  by paired  $t$  test. arbU: arbitrary units. Values indicate medians  $\pm$  SD. (B) Kaplan-Meier and log rank analysis of aPKC $\lambda/i$  and PSA failure time. aPKC $\lambda/i$  expressions in prostate cancer tissues ( $n = 29$ ) were divided into two groups according to the median value (High:  $n = 11$ ,  $>0.56$  and Low:  $n = 18$ ,  $\leq 0.56$ ), and analyzed ( $P = 0.032$  by log rank test).



**Fig. 2.** Representative examples of immunohistochemistry of aPKC $\lambda/i$  expression. Expression intensities of aPKC $\lambda/i$  were divided in two groups (positive;  $+2\leq$  and negative;  $0$  and  $+1$ ). Gleason scores are indicated in the figures and aPKC $\lambda/i$  expression in normal prostate tissue is also shown in the figure.

from same patients (Fig. 1A,  $P = 0.001$ ). There were no associations between aPKC $\lambda/i$  mRNA expression and certain clinical features (Fig. S1). On the other hand, when the samples were divided into two groups by setting a cut-off at the median aPKC $\lambda/i$  value (high:  $>0.56$ ,  $n = 11$  and low:  $\leq 0.56$ ,  $n = 18$ ), we recognized a statistically significant correlation between aPKC $\lambda/i$  mRNA expression and PSA failure (Fig. 1B,  $P = 0.032$ ). There was no correlation between other clinical features and PSA failure (Fig. S2). Serum PSA was measured every 2–3 months after radical prostatectomy. PSA failure was defined as a continuous elevation with a PSA level greater than 0.2 ng/mL. PSA failure has been suggested to be associated with cancer specific death (31). Thus, aPKC may be a prognostic biomarker for prostate cancer. In univariate and multivariate analyses, only aPKC $\lambda/i$  mRNA expression showed statistical significance (Table 1,  $P = 0.039$  in univariate and  $P = 0.033$  in multivariate analysis). Subsequent immunohistochemical analysis of aPKC $\lambda/i$  in 43 prostate specimens (cancer tissues;  $n = 40$ , and normal tissues;  $n = 3$ ) confirmed aPKC $\lambda/i$  expression at the protein level in normal and tumor tissues, with a variety of intensities (Fig. 2 and Table S2). Immunohistochemical analysis also revealed enhanced staining of aPKC $\lambda/i$  to be localized to the cytoplasm in epithelial cells of the prostate, but not in stromal cells, suggesting the importance of the specific expression of aPKC $\lambda/i$  protein in epithelial cells of the prostate.

**Suppression of aPKC $\lambda/i$  Expression Reduces Prostate Cancer Cell Growth In Vitro and In Vivo.**

The correlation of aPKC $\lambda/i$  expression in prostate cancer tissue samples with PSA failure prompted us to clarify the role of aPKC $\lambda/i$  in prostate cancer cell lines. Western blot clearly showed aPKC $\lambda/i$  expression to be higher in prostate cancer cell lines, LNCaP, PC-3 and DU145 cells, than in normal prostate cells, PrEC, as expected (Fig. 3A). To evaluate the role of aPKC $\lambda/i$  in cell growth, we introduced siRNA for aPKC $\lambda/i$  into the DU145, an androgen-independent cell line, and established a mixture of cell lines expressing siRNA for aPKC $\lambda/i$  (DU-P), as well as vector control (DU-C) cells. As shown in Fig. 3B, we confirmed the reduced expression of aPKC $\lambda/i$  in the pooled transfectant (DU-P), as compared to control cells (DU-C). We found that DU-P cells grew more slowly than control cells (Fig. 3C,  $P < 0.05$  at days 3 and 4, and  $P < 0.01$  at days 5 and 6). We next transplanted the cell lines into nude mice and monitored the tumor volume in vivo. As shown in Fig. 3D, the xenografts of aPKC $\lambda/i$ -depleted DU-P cells showed slower growth than those of control DU-C cells ( $P = 0.04$  at 5 weeks,  $P = 0.012$  at 6 weeks). The suppression of aPKC $\lambda/i$  expression in xenografts was confirmed by RT-PCR analysis (Fig. 3E,  $P = 0.001$ ). Thus, the suppression of aPKC $\lambda/i$  expression leads to the inhibition of prostate cancer growth in vitro and in vivo, clearly indicating a positive role of aPKC $\lambda/i$  in the growth of prostate cancer cells.

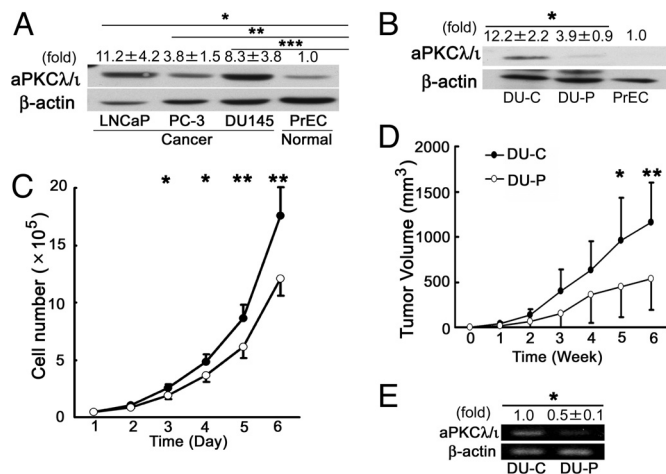
**aPKC $\lambda/i$  Mediates the Growth of Prostate Cancer Cells in an Autocrine Manner Through IL-6 Secretion.**

To explore the mechanism involved in aPKC $\lambda/i$ -dependent growth of prostate cancer cells, we

**Table 1. Relative hazard of recurrence free survival in univariate and multivariate analysis**

		Univariate			Multivariate		
		HR	95% CI	P	HR	95% CI	P
aPKC $\lambda/i$ expression	$\leq 0.56$ ( $n = 18$ )						
	$> 0.56$ ( $n = 11$ )	3.914	1.071–14.305	0.039	6.768	1.161–39.443	0.033
Stage	pT2 ( $n = 16$ )						
	pT3 ( $n = 13$ )	2.866	0.747–10.992	0.125	3.928	0.781–19.748	0.097
	Well ( $n = 5$ )						
Histology	Moderately ( $n = 13$ )	1.129	0.218–5.848	0.885	1.12	0.173–7.275	0.905
	poorly ( $n = 11$ )	0.892	0.161–4.932	0.895	0.887	0.081–9.669	0.922
	$\leq 6$ ( $n = 11$ )						
Gleason score	7 ( $n = 10$ )	1.087	0.291–4.060	0.901	1.562	0.275–8.876	0.615
	8 and 9 ( $n = 8$ )	0.842	0.184–3.866	0.825	4.346	0.346–54.581	0.255
	$\leq 68$ ( $n = 16$ )						
Age	$> 68$ ( $n = 13$ )	1.362	0.393–4.718	0.626	1.946	0.488–7.761	0.346
	$\leq 10$ ( $n = 15$ )						
PSA	$> 10$ ( $n = 14$ )	1.168	0.375–3.639	0.789	0.931	0.233–3.721	0.920

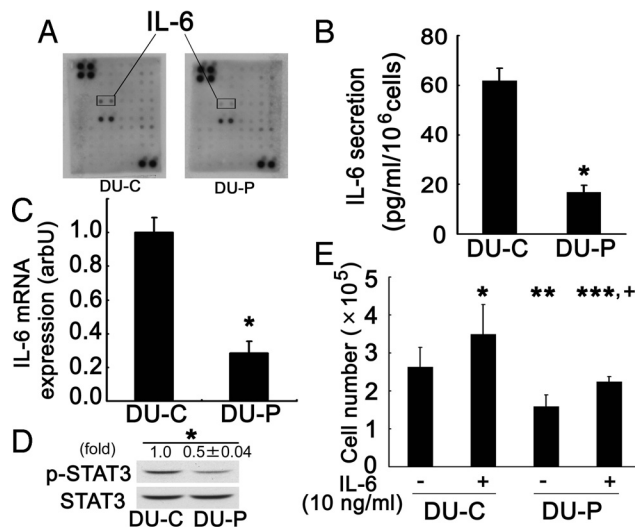
CI, confidence interval; HR, hazard ratio.



**Fig. 3.** aPKC $\lambda/i$  expression in prostate cancer cell lines and growth inhibition of aPKC $\lambda/i$  siRNA transfected DU145 cells in vitro and in vivo. (A) aPKC $\lambda/i$  expression in prostate cell lines was analyzed by Western blot.  $\beta$ -actin was used as an internal control. Values indicate means  $\pm$  SD at least three independent experiments (set as 1.0 in PrEC). \*,  $P = 0.017$ , \*\*,  $P = 0.032$  and \*\*\*,  $P = 0.031$  by unpaired  $t$  test. (B) DU145 cells transfected with siRNA for aPKC $\lambda/i$  expression vector (DU-P cells) and empty vector (DU-C cells) were confirmed by Western blot.  $\beta$ -actin was used as an internal control. Values indicate means  $\pm$  SD from at least three independent experiments (set as 1.0 in PrEC). \*,  $P = 0.004$  by unpaired  $t$  test. (C) The inhibition of cell growth of siRNA transfected cells in vitro. DU-C (filled circles) and DU-P (open circles) cells were seeded onto 12-well plates at  $4 \times 10^4$  cells and counted until day 6 ( $n = 4$  in each group). Points and bars indicate means  $\pm$  SD from at least three independent experiments. \*,  $P < 0.05$  \*\*,  $P < 0.01$  by unpaired  $t$  test. (D) The inhibition of cell growth of siRNA transfected cells in vivo. DU-C (filled circles) and DU-P (open circles) cells were implanted s.c. into the right and left flanks of male nude mice ( $n = 7$  in each group) and tumor growth was calculated at 6 weeks (The day of injection was taken as week 0). Points and bars indicate means  $\pm$  SD. \*,  $P = 0.04$ ; \*\*,  $P = 0.012$  by unpaired  $t$  test. (E) aPKC $\lambda/i$  mRNA expression extracted from xenografts. Total RNA was extracted from the same tumor specimens and aPKC $\lambda/i$  mRNA expression was examined. Values indicate means  $\pm$  SD from at least three independent experiments (set as 1.0 in DU-C derived tumors) \*,  $P = 0.001$  by unpaired  $t$  test.

focused on cytokines which have been implicated in the growth of a variety of cancer cells. Comparison of the conditioned medium between DU-C and DU-P cell lines by cytokine membrane array revealed IL-6 to be a candidate cytokine, that is, that its regulation might be under the control of aPKC $\lambda/i$  (Fig. 4A). The ELISA results confirmed that IL-6 is secreted into conditioned medium of control DU-C cells and decreased in aPKC $\lambda/i$ -depleted DU-P cells (Fig. 4B,  $P = 0.002$ ). Similarly, qPCR revealed IL-6 mRNA expression to also be suppressed in aPKC-depleted DU-P cells (Fig. 4C,  $P < 0.001$ ). These results show that aPKC $\lambda/i$  regulates IL-6 secretion in prostate cancer cells.

IL-6 is the cytokine reportedly expressed in androgen-independent prostate cancer cell lines including parental DU145 (26, 30). We next examined the effect of aPKC $\lambda/i$  depletion on phosphorylation of STAT3, one of the downstream mediators of the possible IL-6 involvement in prostate cancer cells (25, 29, 32, 33). As shown in Fig. 4D, STAT3 phosphorylation was down-regulated in DU-P cells as compared to DU-C cells, indicating that IL-6 signaling is altered in aPKC $\lambda/i$ -depleted cells. Given that aPKC $\lambda/i$  depletion results in the decreased production of IL-6, these results raise another question as to whether aPKC $\lambda/i$  depletion also affects IL-6 signaling, versus only IL-6 production. To clarify this issue, we next evaluated the response of cells to ectopic IL-6. When both cell types were treated with recombinant IL-6 (10 ng/ml) for 3 days, IL-6 increased cell growth, as we expected (Fig. 4E,  $P = 0.001$  and  $P = 0.016$ ). Moreover, DU-P

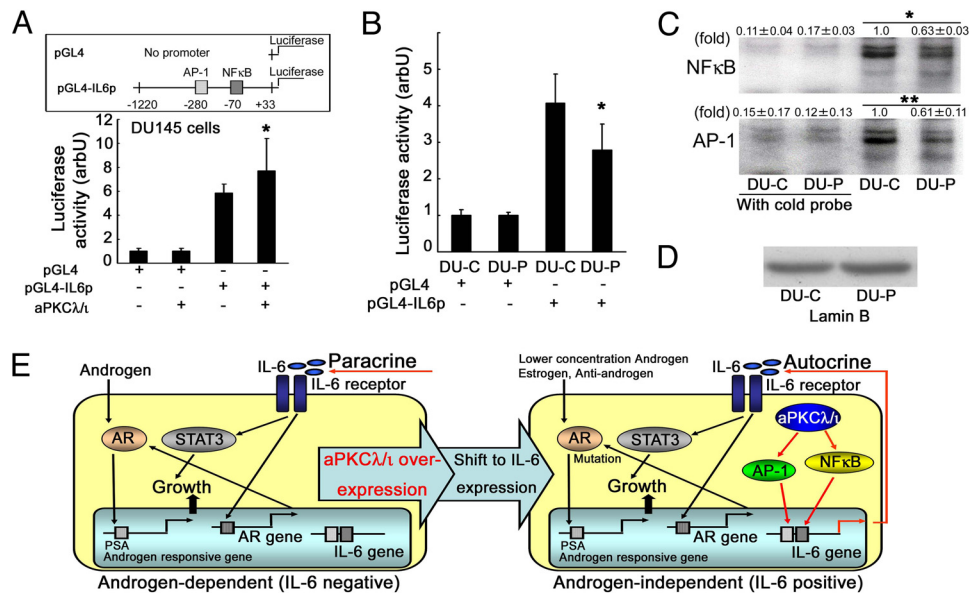


**Fig. 4.** IL-6 expression was suppressed in siRNA transfected prostate cancer cells. (A) Conditioned media from DU-C and DU-P cells were analyzed using a membrane array. Arrows show IL-6 spots. Since this representative data are screening test only, it is performed once. This data had confirmed using ELISA and qPCR. (B) Expressions of IL-6 protein in DU-C and DU-P cells were confirmed using ELISA. Conditioned media from DU-C and DU-P ( $n = 2$  in each group) were analyzed using an IL-6 ELISA kit from R&D Systems. Bars represent means  $\pm$  SD from at least three independent experiments. \*,  $P = 0.002$  by unpaired  $t$  test. (C) Expressions of IL-6 mRNA in DU-C and DU-P cells. IL-6 mRNA expression was investigated by qPCR ( $n = 3$  in each group). Bars represent means  $\pm$  SD from at least three independent experiments. \*,  $P < 0.001$  by unpaired  $t$  test. arbU: arbitrary units. (D) STAT3 phosphorylations in DU-C and DU-P cells were analyzed by Western blot. After phospho-STAT3 protein had been detected, membranes were re-probed for STAT-3, and then used as controls. Values indicate means  $\pm$  SD at least three independent experiments (set as 1.0 in DU-C). \*,  $P = 0.002$  by unpaired  $t$  test. (E) Recombinant IL-6 stimulated growth of DU-C and DU-P cells. DU-C and DU-P cells were stimulated with IL-6 (10 ng/ml). At day 3 after stimulation, cells were treated with trypsin and counted ( $n = 4$  in each group). Bars represent means  $\pm$  SD from at least three independent experiments. \*,  $P = 0.001$ , \*\*,  $P < 0.001$  and \*\*\*,  $P = 0.393$  (vs. DU-C without IL-6 stimulation) +,  $P = 0.016$  (vs. DU-P without IL-6 stimulation) by ANOVA followed by Bonferroni test.

cells treated with IL-6 showed growth similar to that of DU-C cells not treated with IL-6 (Fig. 4E,  $P = 0.393$ ). Thus, aPKC $\lambda/i$  depletion does not affect the growth response of cells to IL-6. Taking these observations together, we conclude that secretion of IL-6 enhanced by aPKC $\lambda/i$  expression plays a role in promoting growth of the prostate cancer cell line DU145. Our results suggest that this growth promotion is mediated in an autocrine manner.

**aPKC $\lambda/i$  Mediates IL-6 Gene Transcription Through NF $\kappa$ B and AP-1 in Prostate Cancer Cells.** To analyze the mechanism by which aPKC $\lambda/i$  enhances IL-6 secretion, we next examined the effect of aPKC $\lambda/i$  on transcription of the IL-6 gene by luciferase reporter assay and EMSA. The luciferase reporter gene, pGL4-IL6p, contains a 1.2-kb 5'-flanking region of genomic DNA isolated from DU145 cells. This region contains the regulatory sequences recognized by AP-1, NF $\kappa$ B, MRE and other factors (30). As shown in Fig. 5A, overexpression of aPKC $\lambda/i$  in DU145 cells enhanced activation of the IL-6 promoter (Fig. 5A,  $P = 0.016$ ). On the other hand, depletion of aPKC $\lambda/i$  (DU-P cells) resulted in decreased reporter gene expression (Fig. 5B,  $P < 0.001$ ). These results strongly support the notion that aPKC $\lambda/i$  is involved in transcription of the IL-6 gene. Transcription of this gene is regulated by two major transcription factors, AP-1 and NF $\kappa$ B, in prostate cancer cells (30). To examine the involvement of these transcription factors in aPKC $\lambda/i$ -mediated IL-6 gene





**Fig. 5.** Involvement of aPKCλ/i in activation of the IL-6 gene promoter and its transcription factors, NFκB and AP-1. (A) IL-6 promoter activation was induced by wild type aPKCλ/i in DU145 cells. DU145 cells were transfected with pGL4, pGL4-IL6p, empty vector and wild type aPKCλ/i. After a 48 h incubation, luciferase activity was analyzed in a luminometer ( $n = 4$  in each group). Each control was given a value of 1.0 and each bar indicates the means  $\pm$  SD of at least three independent experiments. \*,  $P = 0.016$  (vs. pGL4-IL6p without wild-type aPKCλ/i expression vector) by ANOVA following Bonferroni test. (B) IL-6 promoter activation was reduced in DU-P cells. DU-C and DU-P cells were transfected with pGL4 and pGL4-IL6p. After a 12-h incubation, luciferase activity was analyzed in a luminometer ( $n = 4$  in each group). Each control was given a value of 1.0 and each bar indicates the means  $\pm$  SD of at least three independent experiments. \*,  $P < 0.001$  (vs. DU-C with pGL4-IL6p transfection) by ANOVA following Bonferroni test. (C) DNA binding properties of NFκB and AP-1 were examined by EMSA. Each nuclear extract was reacted with  $^{32}$ P-labeled specific probes. To confirm specific binding of nuclear extracts and probes, reactions were also carried out using a cold probe. Values indicate means  $\pm$  SD at least three independent experiments (set as 1.0 in DU-C). \*,  $P < 0.001$  and \*\*,  $P = 0.026$  by ANOVA following Bonferroni test. (D) Nuclear extracts for EMSA were confirmed by lamin B immunoblot. (E) Autocrine mechanism of prostate cancer cell growth involving aPKCλ/i and IL-6. Overexpression of aPKCλ/i in prostate cancer cells leads to up-regulation of IL-6 transcription through NFκB and AP-1 activation. Secreted IL-6 stimulates cell growth through STAT3 activation in an autocrine manner.

transcription, we used EMSA. The results showed both NFκB and AP-1 to be reduced upon aPKCλ/i depletion (Fig. 5 C and D,  $P < 0.001$  and  $P = 0.026$ ). These results support the notion that both AP-1 and NFκB are involved in aPKCλ/i-mediated transcriptional activation of the IL-6 gene.

## Discussion

aPKCλ/i overexpression has been implicated in the progression and invasiveness of several tumor types including non-small cell lung cancer, ovarian cancer, breast cancer and glioma (16–22, 34, 35). Gene amplification of aPKCλ/i is also observed in some cases (16, 18). In this study, we obtained evidence supporting a statistically significant correlation between aPKCλ/i overexpression and a clinical marker of prostate cancer recurrence, PSA failure. A very recent study demonstrated aPKCλ/i expression is required for cell survival (36). Thus, the aPKCλ/i expression level and its activity might be prognostic factors for PSA failure.

IL-6, as noted above, has been implicated in the progression of a variety of tumors including prostate cancer (25–27, 29). Its overexpression has been detected in tissues (37, 38) and serum (28) from cancer patients. Preoperative plasma IL-6 level is related to PSA failure after radical prostatectomy (39). In vivo study indicates an important role of IL-6 in prostate cancer cells. Inhibition of IL-6 by anti-IL-6 antibody decreases the growth of PC-3 cells, which is one of an androgen-independent prostate cancer cells, in vivo (40). Therefore, IL-6 may play a role in androgen-independent growth of prostate cancer. Our findings on the molecular link between aPKCλ/i and IL-6 raise an intriguing question as to the correlation between aPKCλ/i expression and IL-6 expression in prostate cancer patients. We evaluated our tissue samples for IL-6 expression by qPCR. However, there was no statistically significant correlation be-

tween aPKCλ/i expression and IL-6 expression (Fig. S3). One explanation is that the examined tissues contained not only cancer cells but also stromal cells. In support of this possibility, most prostate cancer tissues that express high levels of aPKCλ/i protein (30/43, 70%) show a specific overexpression in epithelial cells but not in the surrounding stromal cells. However, IL-6 is reportedly expressed not only in prostate cancer epithelial cells, but also in stromal cells (37, 38). Thus, the use of laser capture microdissection (LCM) might be required to clarify this point. Unfortunately, we did not have sufficient sample quantities for LCM. We are currently conducting more extensive clinical studies aimed at clarifying the correlation between aPKCλ/i expression and IL-6 expression.

Experiments using the prostate cancer cell line DU145 revealed that aPKCλ/i is involved in prostate cancer growth both in vivo and in vitro. Furthermore, depletion of aPKCλ/i in DU145 cells suppressed NFκB and AP-1 activities, transcription and secretion of IL-6, as well as suppressing cell growth, but not IL-6 signaling. We conclude that enhanced aPKCλ/i expression in prostate cancer cells results in overproduction and secretion of a prostate growth factor, IL-6, at the transcriptional level. This forms an autocrine loop contributing to the growth of prostate cancer (Fig. 5E). The aPKC-dependent expression of IL-6 mRNA is also observed for another androgen-independent prostate cell line, PC-3, suggesting the generality of this regulation (Fig. S4). The specific overexpression of aPKCλ/i in epithelial cells but not in stromal cells of the prostate further supports such an autocrine mechanism. It is known that most of androgen-independent prostate cancer tissues overexpress or/and express mutated AR that is still activated by lower concentration androgen, estrogen and anti-androgen drug (3). Taken together, the pathway might cooperate with the deregulated AR

system to regulate proliferation of hormone-independent prostate cancer cells (Fig. 5E).

The following reports strongly support our conclusion. Combined constitutive activation of NF $\kappa$ B and AP-1 has been reported to mediate deregulated expression of IL-6 in DU145 cells (30). Ectopic expression of IL-6 in IL-6-negative LNCaP prostate cancer cells results in stimulation of the STAT3 signaling pathway as well as cell growth (29). However, the mechanism by which IL-6 expression is deregulated is not yet fully understood. As for aPKC $\lambda/\iota$ , several studies have shown that aPKC $\lambda/\iota$  activates NF $\kappa$ B and AP-1 (9, 10, 13–15, 41–44). Furthermore, aPKC activates NF $\kappa$ B in prostate cells (45). aPKC deregulates the growth of mouse prostate cancer cells (46). While the mechanism by which aPKC affects the growth of cancer cells remains obscure, the involvement of Rho B suppression in aPKC-dependent invasive properties in glioblastoma (35) and that of activation of the Rac1/Erk pathway in aPKC-dependent growth and tumorigenicity have been reported (17, 22). Our present results may explain the link between aPKC $\lambda/\iota$  and IL-6, two molecules implicated in the progression of a variety of malignancies, and establish a molecular mechanism underlying prostate cancer development and/or progression, thereby providing insights into the prognosis and treatment of prostate cancer.

How aPKC $\lambda/\iota$  expression was regulated in prostate cancer cells? Amplification of aPKC $\lambda/\iota$  gene is reported in lung and ovarian cancer (16, 18, 19) and amplification of chromosome 3q including aPKC $\lambda/\iota$  gene in prostate cancer cell lines (47). Another possibility is that aPKC $\lambda/\iota$  expression is up-regulated through the transcriptional activation of aPKC $\lambda/\iota$  promoter during hormone refractory process. Gustafson et al. reported aPKC $\lambda/\iota$  promoter analysis using luciferase (48). They show Bcr-Abl regulates aPKC $\lambda/\iota$  expression through the MEK-dependent activation of an Elk1 element within aPKC $\lambda/\iota$  promoter in leukemia cell line. We are ongoing the investigation of the mechanism of aPKC $\lambda/\iota$  overexpression involving in hormone refractory process. Further, other molecules expression, such as PAR-4, might be affected the alteration of aPKC $\lambda/\iota$  activity during hormone refractory process (49).

One of the most important clinical aspects of prostate cancer is dramatically decreased androgen-dependent cell growth, a typical indicator of prostate cancer progression. Possible mechanisms include androgen receptor overexpression and other mutations that result in hypersensitivity to androgens and/or other growth factors (3–6). IL-6 is overexpressed in hormone refractory prostate cancer patients and is one of the factors implicated in this process (26–28, 37). Importantly, IL-6 can stimulate androgen receptor activation independently of androgens with the induction of androgen receptor expression (4, 5). A recent study on LNCaP cells, which are sensitive to androgens and do not usually express IL-6, suggested IL-6 to be involved in the progression of prostate cancer cells from androgen dependence to androgen independence (50). It has been reported androgen-dependent prostate cancer cells obtained from xenografts treated with the anti-IL-6 antibody retained in androgen-dependence. In contrast, cancer cells obtained from xenografts untreated with the anti-IL-6 antibody are converted to androgen-independence in vitro and in vivo experiments, which means that IL-6 contributes to the development of androgen independence in prostate cancer (51). Taking these observations and our results together, we speculate that the molecular link between aPKC $\lambda/\iota$  and IL-6 revealed in the present study supports the notion that aPKC $\lambda/\iota$  is involved in this transition from androgen dependent to androgen independent growth of prostate cancer. As for the relationship between androgens and aPKC, there is an interesting report suggesting that aPKC $\zeta$ , another aPKC isotype, is involved in the growth of androgen dependent LNCaP cells

(52). In LNCaP cells, androgen stimulates aPKC $\zeta$  through an unknown mechanism, and AILNCaP, a LNCaP subline established after androgen depletion, showed constitutive activation of aPKC $\zeta$ . Another study on a breast cancer cell line, MCF7, showed the involvement of aPKC $\zeta$  in estradiol-dependent cell growth (53). These reports further suggest a close relationship between aPKC and hormone-dependent cell growth. Our present findings provide additional evidence clarifying this point and are anticipated to facilitate understanding the progression of hormone-related malignancies including prostate cancer. Furthermore, aPKC $\lambda/\iota$  expression might be a biomarker for prostate cancer progression.

## Materials and Methods

**Cell Lines, Patients, and Tissues Sample.** LNCaP, PC-3, and DU145 cells were obtained from the American Type Culture Collection. PrEC cells were obtained from Clonetics. All cell lines were maintained with suitable medium (F-12 supplemented with 10% FCS (FCS) for LNCaP and PC-3, MEM supplemented with 10% FCS for DU145, PrEGM for PrEC) under 5% CO<sub>2</sub>.

Paired human untreated primary prostate cancer tissues and normal (or benign prostatic hypertrophy (BPH)) ( $n = 29$ ) tissues from same patients were obtained during radical prostatectomy at Yokohama City University Hospital and its affiliates. The sampling and usage of all prostate tissues in this study were approved by the ethical committee of Yokohama City University Graduate School of Medicine and performed only after obtaining informed consent from each patient. For details, see *SI Text* and *Table S1*.

**Reagents.** Human recombinant IL-6 was purchased from R&D Systems. G418 was purchased from Invitrogen Corp. Anti-aPKC $\iota$  antibody was purchased from BD Biosciences. Anti-lamin-B antibody (c-19) was purchased from Santa Cruz Biotechnology Inc. Anti-actin antibody (AC-15) was purchased from Sigma-Aldrich. Anti-phospho-STAT3 and anti-STAT3 (#9131 and #9132, respectively) antibodies were purchased from Cell Signaling Technology Inc. Anti-rabbit and anti-mouse horseradish peroxidase conjugates were purchased from GE Healthcare U.K. Ltd.

**Generation of Stable Transfectant-Induced siRNA for aPKC $\lambda/\iota$ .** To investigate the role of aPKC $\lambda/\iota$  in prostate cancer, we generated aPKC $\lambda/\iota$  knock-down cells using siRNA for aPKC $\lambda/\iota$ . The pEB6-Super vector (54) encoding the shRNA sequence for aPKC $\lambda/\iota$  RNAi with the target sequence 5'-CAA GTG TTC TGA AGA GTT T-3' (DU-P cells) or empty vector (DU-C cells) was transfected into DU145 cells using Nucleofactor electroporation methods (Amaxa AG). Then, transfected cells were selected by G418 (800  $\mu$ g/mL) over a 3-week period. After the specific down-regulation of aPKC $\lambda/\iota$  had been confirmed by Western blot, the cells were used for further experiments.

**RNA Extraction and Real-Time Quantitative PCR (qPCR).** Total RNA from cell lines, prostate tissues and xenografts were extracted using ISOGEN (Nippon-Gene) according to the manufacturer's instructions. After cDNA had been synthesized with random hexomers and MMLV (Moloney Murine Leukemia Virus), qPCR was performed with an iCycler and SYBR Green Supermix (Bio-Rad). For details, see *SI Text*.

**Immunohistochemistry.** Immunohistochemistry was performed for aPKC $\lambda/\iota$  protein expression according to the previous report (20). For details, see *SI Text*.

**Cell Growth Analysis.** DU-C or DU-P cells ( $4 \times 10^4$ ) were incubated in 12-well plates (day 0). Incubated cells were harvested with trypsin and counted till 6 days (from day 1 to day 6) using a hemacytometer (Beckman Coulter, Inc.). For the IL-6 stimulation experiment,  $4 \times 10^4$  DU-C or DU-P cells were seeded onto 12-well plates and incubated for 24 h. The medium was then changed to phenol red-free RPMI1640 with 0.1% BSA (BSA), IL-6 (10 ng/mL) was added and incubation was continued for another 3 days. Then, cells were harvested with trypsin and counted.

**In Vivo Tumor Growth.**  $5 \times 10^6$  cells (DU-C and DU-P cells) were injected into the flank regions of athymic nude mice (4–6 weeks old,  $n = 7$ ). Tumors were measured weekly with a caliper (for comparison with the week 0 value). The tumor volume was calculated using the formula: tumor volume (mm<sup>3</sup>) = 0.5  $\times$  length  $\times$  (width)<sup>2</sup>. After 6 weeks, tumors were isolated and aPKC $\lambda/\iota$  expression was confirmed by RT-PCR.

**Cytokine Membrane Array.** Cytokines in the conditioned medium were detected using Human Cytokine Array III (Ray Biotech) according to the manufacturer's instructions. For details, see [SI Text](#).

**ELISA (ELISA) for IL-6 Secretion.** IL-6 secretion in the collected medium was measured using a human IL-6 ELISA kit according to the manufacturer's instructions (R&D Systems). For details, see [SI Text](#).

**Western Blot.** Cell lysates were prepared and subjected to Western blot. For details, see [SI Text](#).

**Luciferase Assay.** Approximately 1.2 kb of the IL-6 5'-flanking region was generated using PCR from genomic DNA extracted from DU145 cells, and cloned into the pGL4.0 [*Luc2*] vector (pGL4-IL6p) (Promega). Wild type  $\text{aPKC}\lambda_i$  was obtained as described in previous reports (9, 10).  $\text{pRL-SV40}$  was used as the internal control for the luciferase assay (Promega). After cells transfected

each plasmid vector were incubated and lysed, luciferase activity was measured using the dual-luciferase reporter assay system (Promega) and a luminometer, TD-20/20 (Turner Design). For details, see [SI Text](#).

**Electrophoretic Mobility Shift Assay (EMSA).** Nuclear proteins of DU-C or DU-P were extracted using NE-PER (Pierce Biotechnology Inc.) and subjected to EMSA using gel shift assay systems (Promega). For details, see [SI Text](#).

**Statistical Analysis.** All statistical analyses were performed using SPSS for windows (SPSS Inc.). For details, see [SI Text](#).

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