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PLOD3 suppression exerts an anti-tumor effect on human lung cancer cells by modulating the PKC-delta signaling pathway

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Abstract

Current lung cancer treatments are far from satisfactory; thus, finding novel treatment targets is crucial. We recently identified procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (*PLOD3*), which is involved in fibrosis and tissue remodeling as a radioresistance-related protein in lung cancer cells; however, its mechanism is unclear. In this study, we designed human *PLOD3*-specific short interfering (si)RNAs and tested their effects on tumor growth inhibition in vitro and in vivo. *PLOD3* knockdown overcame chemoresistance and decreased radioresistance by inducing caspase-3-dependent apoptosis in lung cancer cells. Furthermore, *PLOD3* interacted with PKC δ to activate caspase-2, 4-dependent apoptosis through ER-stress-induced IRE1 α activation and the downstream unfolded-protein response pathway. In a mouse xenograft model, *PLOD3* knockdown promoted radiation-induced tumor growth inhibition, without side effects. Moreover, lung cancer patients with high *PLOD3* expression showed poorer prognosis than those with low *PLOD3* expression upon radiotherapy, suggesting that *PLOD3* promotes tumor growth. Therefore, *PLOD3* siRNA suppresses radioresistance and chemoresistance by inducing apoptosis and renders *PLOD3* as a candidate lung cancer biomarker. *PLOD3* gene therapy might enhance the efficacy of radiotherapy or chemotherapy in lung cancer patients.

Introduction

Lung cancer is the main cause of cancer-related morbidity, and non-small-cell lung cancer accounts for 80–85% of all lung cancer cases¹. However, among these patients, only 10% achieve a complete response, and the total 5-year survival rate has remained dismal at 15%² because radiation resistance severely affects the efficacy of radiotherapy^{3,4}. Thus, we highlight the need for a greater

understanding of the cellular and molecular targets that drive tumorigenesis to achieve better treatment efficacies.

Recently, we found four proteins, including procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (*PLOD3*), which had not been previously reported to be related to radioresistance or chemoresistance⁵. *PLOD* proteins, are involved in fibrotic processes and tissue remodeling^{6,7}. Three highly homologous *PLOD* isoforms have been characterized to date, including *PLOD2*, and *PLOD3*⁸. *PLOD3* is localized on chromosome 7q36⁹, and *PLOD3* activity is critical for the biosynthesis of type IV and VI collagens¹⁰. Mutations in human *PLOD3* result in congenital disorders that influence the connective tissues of various organs¹¹, suggesting that *PLOD3* is crucial for normal collagen function. Collagen also is involved in

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tumor progression by modulating cancer cell migration, invasion¹², proliferation¹³, survival¹⁴, and metastasis¹⁵. Based on these facts, we focused on cancer cell survival with respect to PLOD3 function.

Two independent studies have reported *PLOD3* mRNA overexpression in glioma and hepatocellular carcinoma tissues^{16–18}. *PLOD3* overexpression was correlated with higher circulating protein levels in some patients¹⁹. However, the molecular mechanisms underlying the role of PLOD3 in lung cancer cell death have not been fully elucidated, and there are no data regarding the possible role of PLOD3 in lung cancer cell apoptosis. Further, the oncogenic function and prognostic value of this protein as a therapeutic and diagnostic target for lung cancer have not been revealed.

We previously found that the mechanistic target of PLOD3-induced cell death is the endoplasmic reticulum (ER)-associated stress-induced apoptosis pathway^{20,21}, which, under physiological conditions, is activated by the accumulation of misfolded proteins in the ER to maintain cell survival²². Specifically, ER stress leads to the activation of three major unfolded protein response sensors, including pancreatic eIF2- α kinase (PERK), high inositol-requiring 1 (IRE1- α), and ATF6. First, PERK phosphorylates the eukaryotic translation initiation factor-2a, resulting in both an initial decrease in general translation initiation and the selective translation of the transcription factor ATF6. Second, ATF6 induces growth arrest and DNA damage-inducible proteins (GADD153/CHOP), leading to cell-cycle arrest, hence preventing the damage to the cell^{23,24}. IRE1- α mediates the splicing of X-box-binding protein 1, which increases the transcription of ER-resident chaperones, folding enzymes, and components of the protein degradation machinery. Third, ATF6, after activating cleavage, results in both the induction of CHOP and the upregulation of protein folding and degradation²⁴. Prolonged, unresolvable ER stress overrides the salvage mechanisms of the initial unfolded protein response and eventually leads to apoptosis involving CHOP signaling, JNK activation, bcl-2 phosphorylation and depletion, and caspase cleavage (e.g., caspase-4).

Protein kinase C (PKC) isozymes comprise a family of at least 10 related serine-threonine kinases that play critical roles in the regulation of several cellular processes, including proliferation, cell-cycle regulation, differentiation, malignant transformation, and apoptosis²⁵. Based on their structures and cofactor requirements, PKC isoforms are divided into classic PKC (α , β 1, β 2, and γ), novel (δ , ϵ , η , and θ), and atypical (ζ and λ/i) groups²⁵. Members of this family are either pro-apoptotic or anti-apoptotic, depending on the isoform and cellular context. For example, PKC α and PKC ϵ inhibit apoptosis by phosphorylating or increasing the expression of the anti-apoptotic protein Bcl-2, whereas the caspase-3-dependent

and caspase-2-dependent activation of PKC δ promotes apoptosis via tyrosine phosphorylation, association with specific apoptotic proteins, and translocation of activated PKC δ to the mitochondria²⁶.

Here, to develop an anti-tumor reagent, we designed human *PLOD3*-specific short interfering (si)RNAs to knockdown the endogenous PLOD3 overexpression in lung cancer and then investigate cell proliferation and cell death. We also assessed the possible involvement of PKC δ and PLOD3 in the ER-stress-induced cell death pathway in cell culture and in an in vivo model.

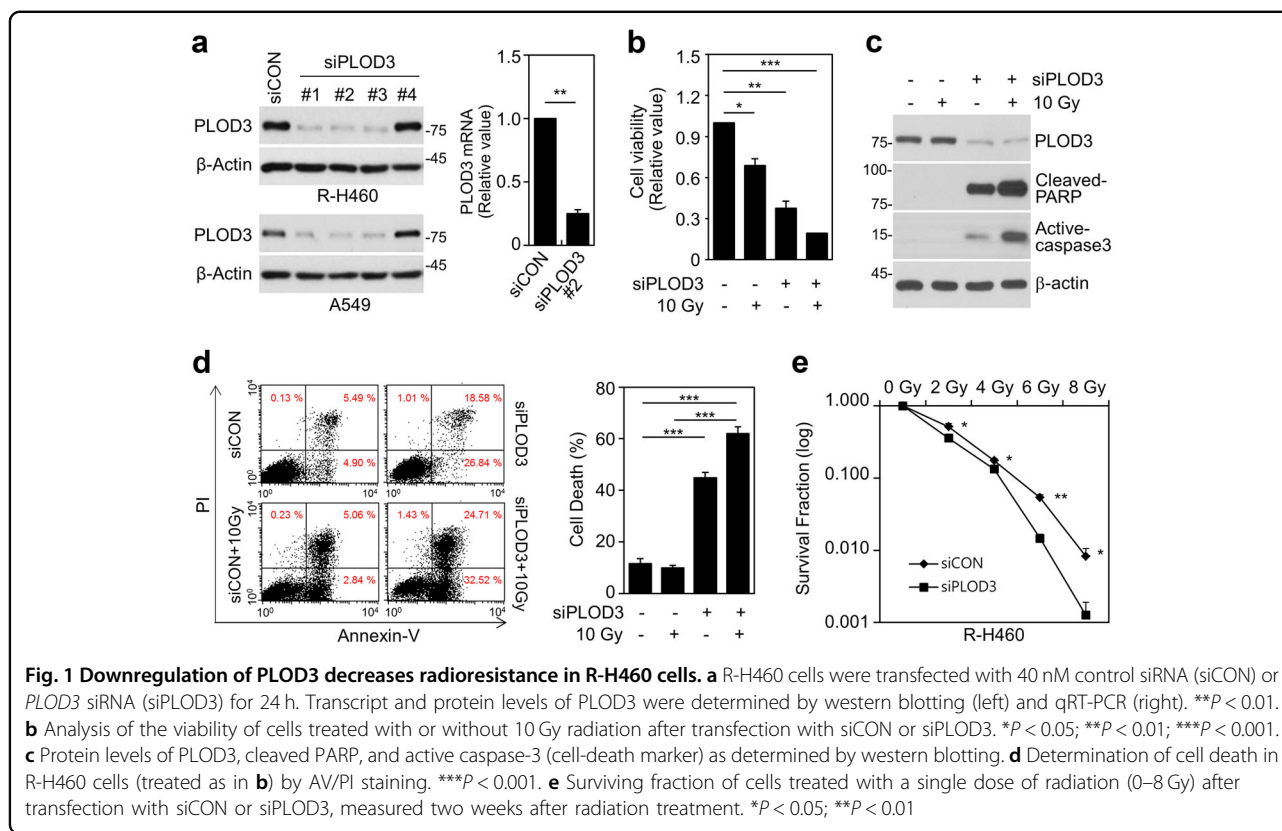
Results

PLOD3 downregulation decreases radioresistance in R-H460 cells

To determine whether PLOD3 is involved in regulating radioresistance, we tested the effects of PLOD3 depletion on cell viability and death (Fig. 1a). After siRNA transfection for 48 h, cell viability was significantly decreased in cells transfected with *PLOD3* siRNA compared to that in control siRNA-transfected cells, and this decrease was enhanced in irradiated cells (Fig. 1b, Supplementary Figure 1a). We next confirmed that PLOD3 knockdown alone or in combination with radiation led to upregulation of cleaved PARP and active-caspase-3 levels (Fig. 1c, Supplementary Figure 1b). FACS analysis showed that PLOD3 knockdown in A549 cells increased cell death (~45%) by more than 3.8-fold compared to that in control cells (~12%), and 10-Gy radiation further increased cell death slightly (Supplementary Figure 1c, d). However, in R-H469 and A549 cells, the combination of PLOD3 knockdown and 10-Gy radiation significantly increased cell death compared to that in control cells (Fig. 1d, Supplementary Figure 1c, d). Additionally, the number of colonies formed decreased with increasing radiation doses in *PLOD3* siRNA-transfected R-H460 and A549 cells, indicating a dose-dependent relationship (Fig. 1e, Supplementary Fig 1e). Therefore, our results suggest that PLOD3 depletion induces cell death to regulate radiosensitivity.

PLOD3 downregulation promotes radiosensitivity and inhibits tumorigenesis in vivo

After confirming PLOD3 transfection efficiency in vivo, we analyzed the synergistic effect of radiation and *PLOD3* siRNA on R-H460 xenograft tumors. As indicated in Fig. 2a and Supplementary Fig. 2a, tumor volumes were markedly decreased by more than ~36% and 47%, compared to those in the control siRNA group, in the *PLOD3* siRNA-treated group and radiation-treated group, respectively. The most pronounced tumor growth-inhibitory effect (60%) was observed in the *PLOD3* siRNA plus irradiation group (Fig. 2b). Further, PLOD3 protein expression was diminished in *PLOD3*



siRNA-injected mice compared to that in control siRNA-treated mice (Fig. 2b). Tumor weight measurements were in agreement with tumor volume results (Fig. 2c). Moreover, the *PLOD3* siRNA plus radiation group showed further decreased *PLOD3* protein levels compared to those in mice treated with either alone. There were possibly no visible signs of toxicity due to *PLOD3* siRNA and radiation in mice, as shown by the lack of differences in body, liver, and lung weights (Fig. 2d).

Patients with high *PLOD3*-expressing lung cancer who were treated with radiotherapy had poorer prognosis than patients with low *PLOD3*-expressing lung cancer who were treated with radiotherapy in the Kaplan–Meier Plotter database (Fig. 2e). Collectively, these findings suggest that *PLOD3* is correlated with tumor growth.

PLOD3 knockdown induces caspase-dependent apoptosis in R-H460 cells

To investigate the mechanisms associated with *PLOD3* knockdown-induced cell death, we assessed apoptosis and necrosis in R-H460 cells by flow cytometry (Fig. 3a). At 36 h, *PLOD3* siRNA significantly increased apoptosis (AV-positive) by more than 2-fold (approximately 20%) compared to control siRNA. Further, *PLOD3* siRNA-mediated apoptosis was accelerated as incubation time increased, but necrosis (AV-negative/PI-positive) was not triggered (Fig. 3b). Caspase-3 cleavage was observed at the

same time point as increased apoptosis by immunoblotting analysis. Moreover, *PLOD3* siRNA-induced apoptosis was caspase-dependent because Z-VAD-FMK, a pan-caspase inhibitor, significantly lowered the percentage of AV-positive cells and blocked the cleavage of caspase-3 (Fig. 3c). In A549 cells, FACS and western blot analysis revealed that Z-VAD-FMK treatment decreased *PLOD3* depletion-induced apoptotic cell death (Supplementary Figure 3a, b). To investigate whether cleaved caspase-3 modulates caspase activation in *PLOD3*-depleted cells, we measured the activity of caspase family members in R-H460 cells (Fig. 3d). *PLOD3* siRNA significantly increased caspase-2, 3, 4, and 6 activity by ~1.9-, 2.1-, 1.8-, and 1.7-fold, respectively, compared to control siRNA. Accordingly, to investigate whether caspase inhibitors could block *PLOD3* knockdown-induced apoptosis (Fig. 3e), R-H460 cells were treated with these compounds for 1 h and then transfected with *PLOD3* siRNA. Based on AV/PI double staining, Z-VAD-FMK, Z-VDVAD-FMK, Z-DEVD-FMK, and Z-YVAD-FMK treatment reduced apoptosis by approximately 18%, 7%, 6%, and 10%, respectively, in *PLOD3*-knockdown cells. Immunoblotting results confirmed that active caspase-3 was remarkably decreased by Z-VAD-FMK treatment, without rescuing *PLOD3* protein levels, but was only slightly reduced by Z-VDVAD-FMK, Z-DEVD-FMK, and Z-YVAD-FMK. To identify additional possible

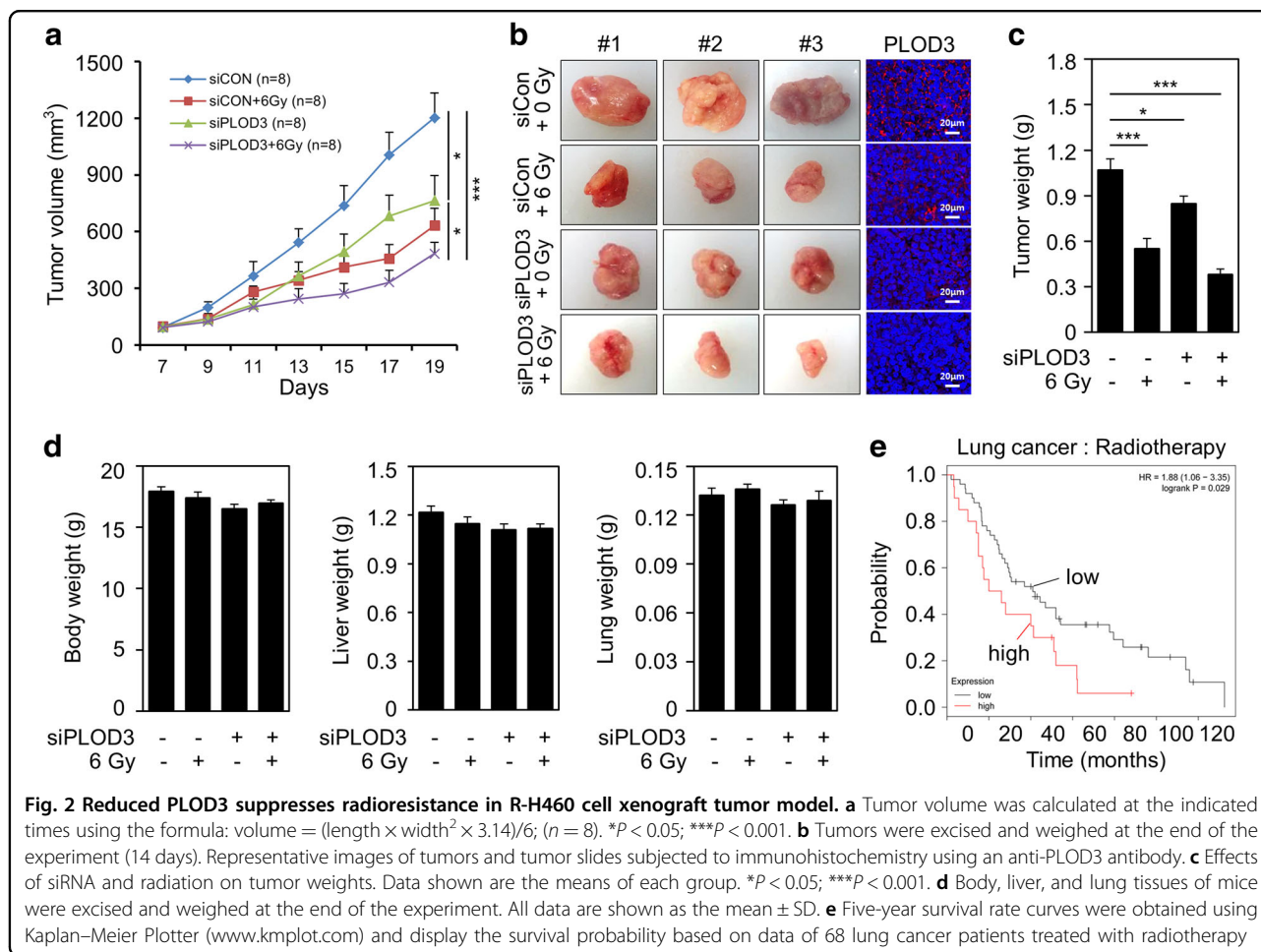


Fig. 2 Reduced PLOD3 suppresses radioresistance in R-H460 cell xenograft tumor model. **a** Tumor volume was calculated at the indicated times using the formula: volume = (length \times width² \times 3.14)/6; ($n = 8$). $*P < 0.05$; $***P < 0.001$. **b** Tumors were excised and weighed at the end of the experiment (14 days). Representative images of tumors and tumor slides subjected to immunohistochemistry using an anti-PLOD3 antibody. **c** Effects of siRNA and radiation on tumor weights. Data shown are the means of each group. $*P < 0.05$; $***P < 0.001$. **d** Body, liver, and lung tissues of mice were excised and weighed at the end of the experiment. All data are shown as the mean \pm SD. **e** Five-year survival rate curves were obtained using Kaplan–Meier Plotter (www.kmplot.com) and display the survival probability based on data of 68 lung cancer patients treated with radiotherapy

mechanisms of cellular death, we investigated autophagy and proteasome activity, which are among other mechanisms associated with anti-cancer activity. We first treated cells with *PLOD3* siRNA to examine chymotrypsin-, caspase- and trypsin-like activities in R-H460 cells. Only minor changes were observed in the levels of the three proteasomal activities after *PLOD3* siRNA treatment (Supplementary Figure 3c). After treatment with 3-MA, which is widely used to inhibit autophagy, R-H460 cells were not significantly rescued from cell death induced by *PLOD3* siRNA, and morphological changes throughout the cytoplasm and in the cell membrane were not observed (Supplementary Fig. 3d). Thus, *PLOD3* ablation induces cell death that depends on apoptosis, but not autophagy and proteasome activity.

PLOD3 depletion induces DNA damage and ER stress-dependent caspase activation in R-H460 cells

Caspase-2 has been shown to mediate DNA damage-induced apoptosis, reactive oxygen species (ROS) accumulation, and ER stress²⁷. Caspase-4 is also activated by ER stress²⁸. First, to investigate the ER localization of *PLOD3* in H460 and R-H460 cells, following treatments,

these organelles were visualized using Grp94 as an ER stress sensor protein, and nuclei were stained with DAPI. *PLOD3* was distributed homogeneously in the cytoplasm in co-localization with Grp94 (Fig. 4a). Immunoblotting revealed that *PLOD3* depletion resulted in elevated cleaved ATF6 and phosphorylation of PERK and IRE1 α as ER stress sensor proteins and also upregulated the DNA damage marker, γ H2AX (Fig. 4b). qPCR confirmed increased splicing of *XBP-1*, a downstream target of IRE1 α (Fig. 4c). Together with ER stress and the activation of the unfolded protein response system, IRE1 α is reportedly involved in caspase activation. Thus, we examined whether IRE1 α inhibition blocked caspase activation and cell death in *PLOD3*-depleted cells. The IRE1 α inhibitor APY29 reduced *PLOD3* knockdown-induced phosphorylation of IRE1 α . APY29 also reduced cleaved caspase-2 and procaspase-4 and the proportion of apoptotic cells (Fig. 4d). R-H460 cells were treated with combinations of *PLOD3*, *IRE1 α* , and control siRNA for 48 h and apoptosis was analyzed by AV/PI staining (Fig. 4e). *PLOD3* siRNA led to an increase in the apoptotic cell population (~32%); however, combined transfection with *IRE1 α* siRNA significantly reduced this by more than

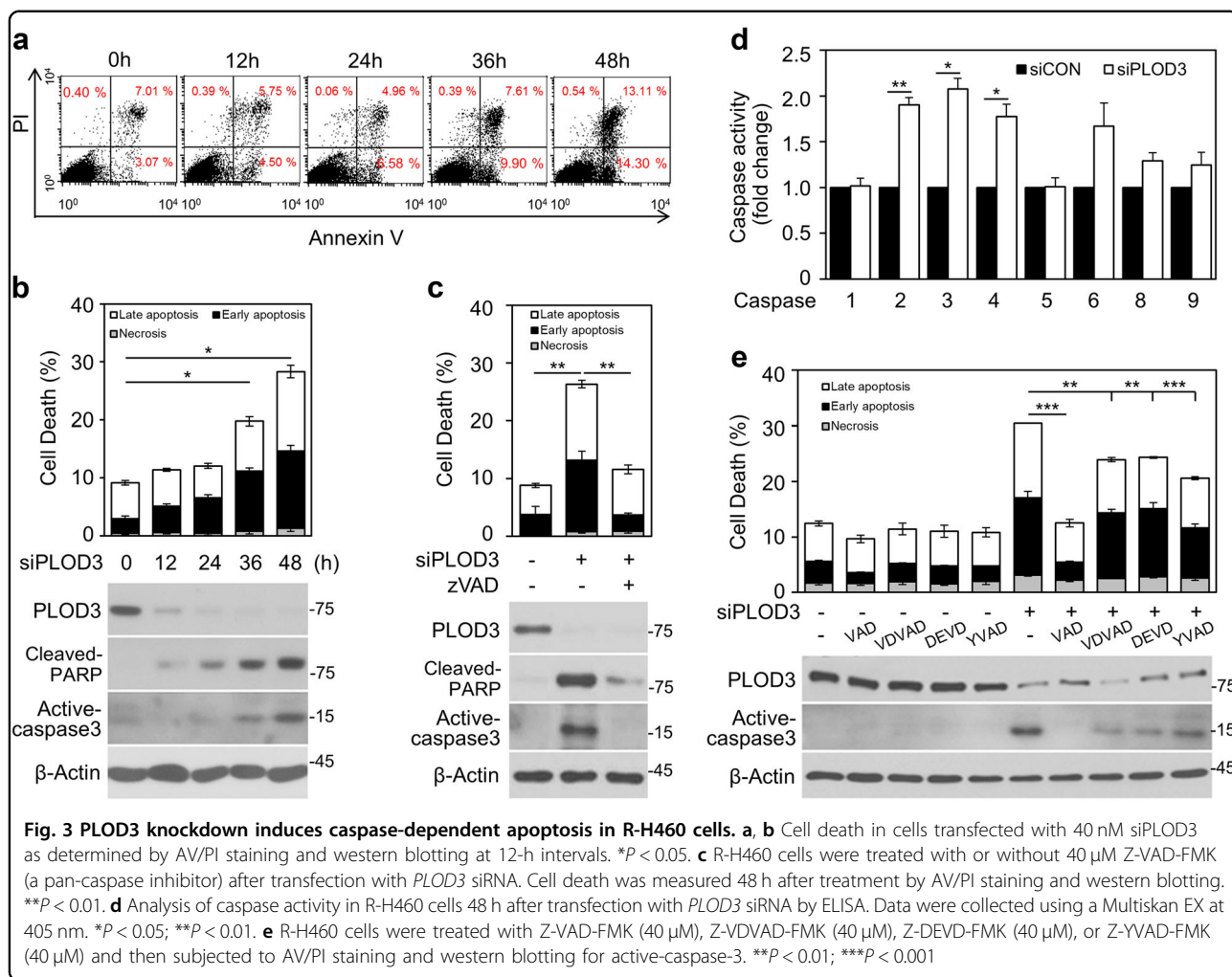


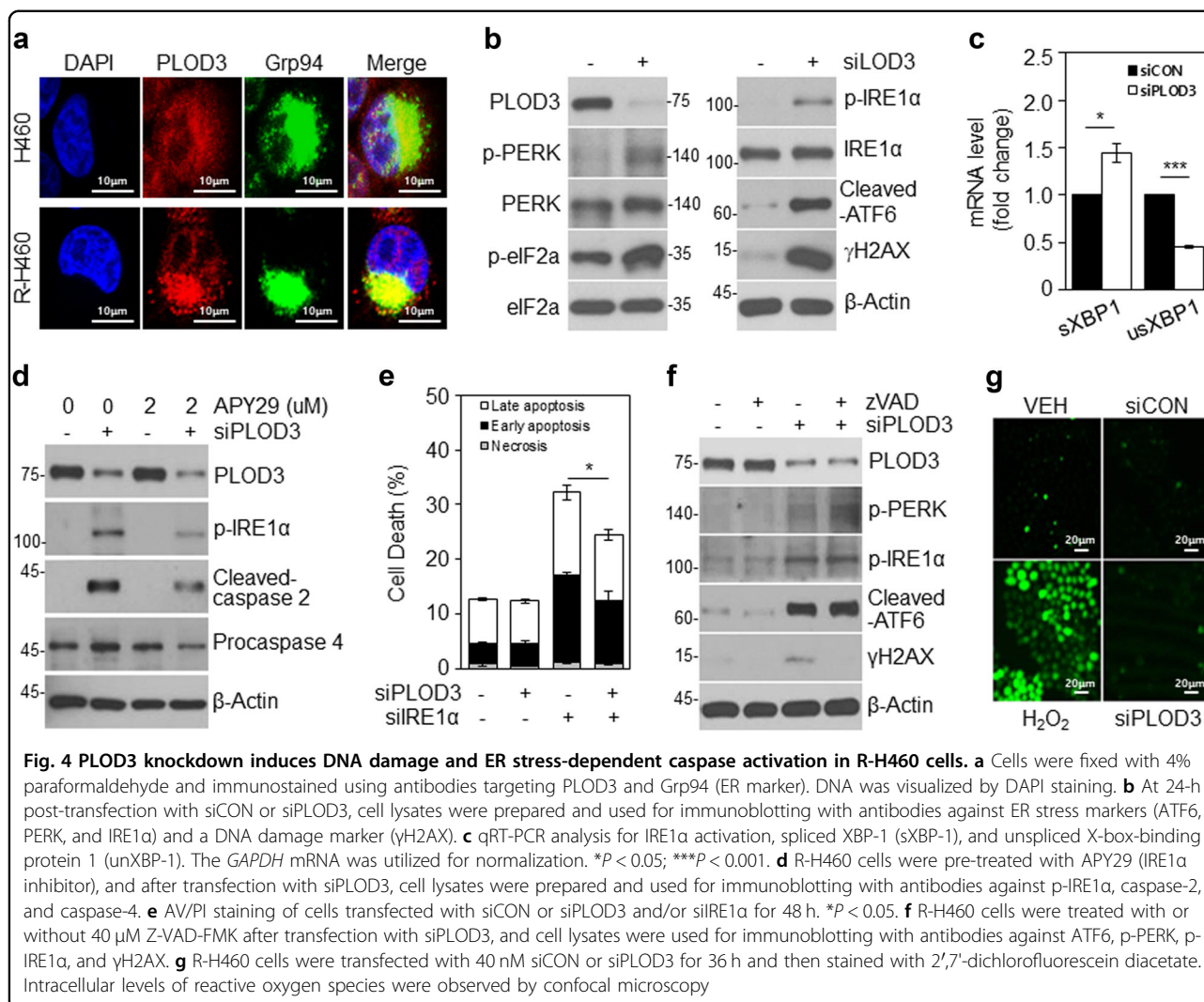
Fig. 3 PLOD3 knockdown induces caspase-dependent apoptosis in R-H460 cells. **a, b** Cell death in cells transfected with 40 nM siPLOD3 as determined by AV/PI staining and western blotting at 12-h intervals. $*P < 0.05$. **c** R-H460 cells were treated with or without 40 μ M Z-VAD-FMK (a pan-caspase inhibitor) after transfection with PLOD3 siRNA. Cell death was measured 48 h after treatment by AV/PI staining and western blotting. $**P < 0.01$. **d** Analysis of caspase activity in R-H460 cells 48 h after transfection with PLOD3 siRNA by ELISA. Data were collected using a Multiskan EX at 405 nm. $*P < 0.05$; $**P < 0.01$. **e** R-H460 cells were treated with Z-VAD-FMK (40 μ M), Z-VDVAD-FMK (40 μ M), Z-DEVVD-FMK (40 μ M), or Z-YVAD-FMK (40 μ M) and then subjected to AV/PI staining and western blotting for active-caspase-3. $**P < 0.01$; $***P < 0.001$

8% compared to PLOD3 siRNA alone. Notably, we found that caspase inhibition in PLOD3-knockdown R-H460 cells did not regulate ER stress (Fig. 4f). Furthermore, caspase inhibition using Z-VAD-FMK completely inhibited γ H2AX accumulation after PLOD3 knockdown (Fig. 4f). These results suggested that DNA damage is induced by caspase activity after PLOD3 depletion. However, PLOD3 knockdown-induced cell death was not found to depend on ROS levels (Fig. 4g). These results suggest that caspase-2 and 4 depend on IRE1 α activation in PLOD3 siRNA-treated cells.

PLOD3 regulates PKC-dependent apoptosis by activating PKC α and PKC δ via direct interactions

It is well-known that PKC α or PKC δ is involved in the ER stress response, leading to apoptosis^{29,30}. Moreover, PKC δ reportedly associates with and phosphorylates caspase-3 to promote apoptotic activity, after which activated caspase-3 cleaves PKC δ ³¹. Thus, to investigate the relationship between PLOD3 and PKC, we compared the subcellular localization of these markers in

R-H460 cells (Fig. 5a). The data showed that PLOD3 co-localized with PKC α and PKC δ in the nucleus and cytosol. Interestingly, PKC δ localized to the ER, similar to PLOD3. Next, as PLOD3 co-localized with PKCs, we evaluated the associated interactions in R-H460 cells. We immunoprecipitated exogenous PLOD3 from HA-PLOD3-, PKC α -, and GFP-PKC δ -transfected R-H460 cell lysates using a HA antibody and detected the immune complexes with HA, PKC α , and GFP antibodies (Fig. 5b, c). We observed an interaction between PLOD3 and PKCs in R-H460 cells. A proximity ligation assay confirmed that endogenous PLOD3 interacts with PKCs in vivo (Fig. 5d). A proximity ligation assay probe was used as a negative control. To examine whether PKC activity changes in the absence or presence of PLOD3, we transfected R-H460 cells with PLOD3 siRNA. PKC α and PKC δ phosphorylation was remarkably increased by PLOD3 knockdown in R-H460 cells (Fig. 5e). These results suggest that PLOD3 regulates the activity of PKC α and PKC δ , leading to PKC-dependent apoptosis. Furthermore, we examined the localization of



cleaved PKC δ , which is important for PKC δ -dependent apoptosis. Cleaved PKC δ was significantly induced in PLOD3-knockdown cells and was localized to the nucleus (Fig. 5f). Thus, we hypothesized that PKCs are related to PLOD3 depletion mediated-apoptosis in R-H460 cells. Accordingly, we investigated whether PKC δ siRNA blocks PLOD3 knockdown-induced subcellular events. R-H460 cells were treated with combinations of PLOD3, PKC δ , and control siRNA for 48 h and apoptosis was analyzed by AV/PI staining (Fig. 5g). PLOD3 siRNA led to an increase in the apoptotic cell population (~39%); however, combined transfection with PKC δ siRNA significantly suppressed this increase by more than 14%. Immunoblotting results confirmed that cleaved PARP and active caspase-3 were remarkably decreased, without rescuing PLOD3 protein levels (Fig. 5g). Based on the results of interaction with PLOD3 and PKC α and increased PKC α phosphorylation by

PLOD3 depletion, we examined whether PKC α siRNA blocks PLOD3 knockdown-induced apoptosis. To this end, R-H460 cells were treated with a combination of PLOD3, PKC α , and control siRNA for 48 h. Apoptosis was then analyzed by AV/PI staining and immunoblotting (Fig. 5h). PLOD3 siRNA increased the apoptotic population (~41%), but combined transfection with PKC α siRNA suppressed this increase by more than 13%. Immunoblotting data confirmed that cleaved PARP and active caspase-3 were remarkably decreased, without rescuing PLOD3 protein level. Further, IRE1 α and eIF2 α phosphorylation, as assessed by immunological analysis, was similar to that of cell death markers (Fig. 5i). Therefore, PKC δ siRNA inhibits the PLOD3-induced ER stress response. Next, we examined whether combined PLOD3 and PKC δ siRNAs could affect caspase activity (Fig. 5j). PLOD3 siRNA significantly increased caspase-2 activity by approximately 1.6-fold

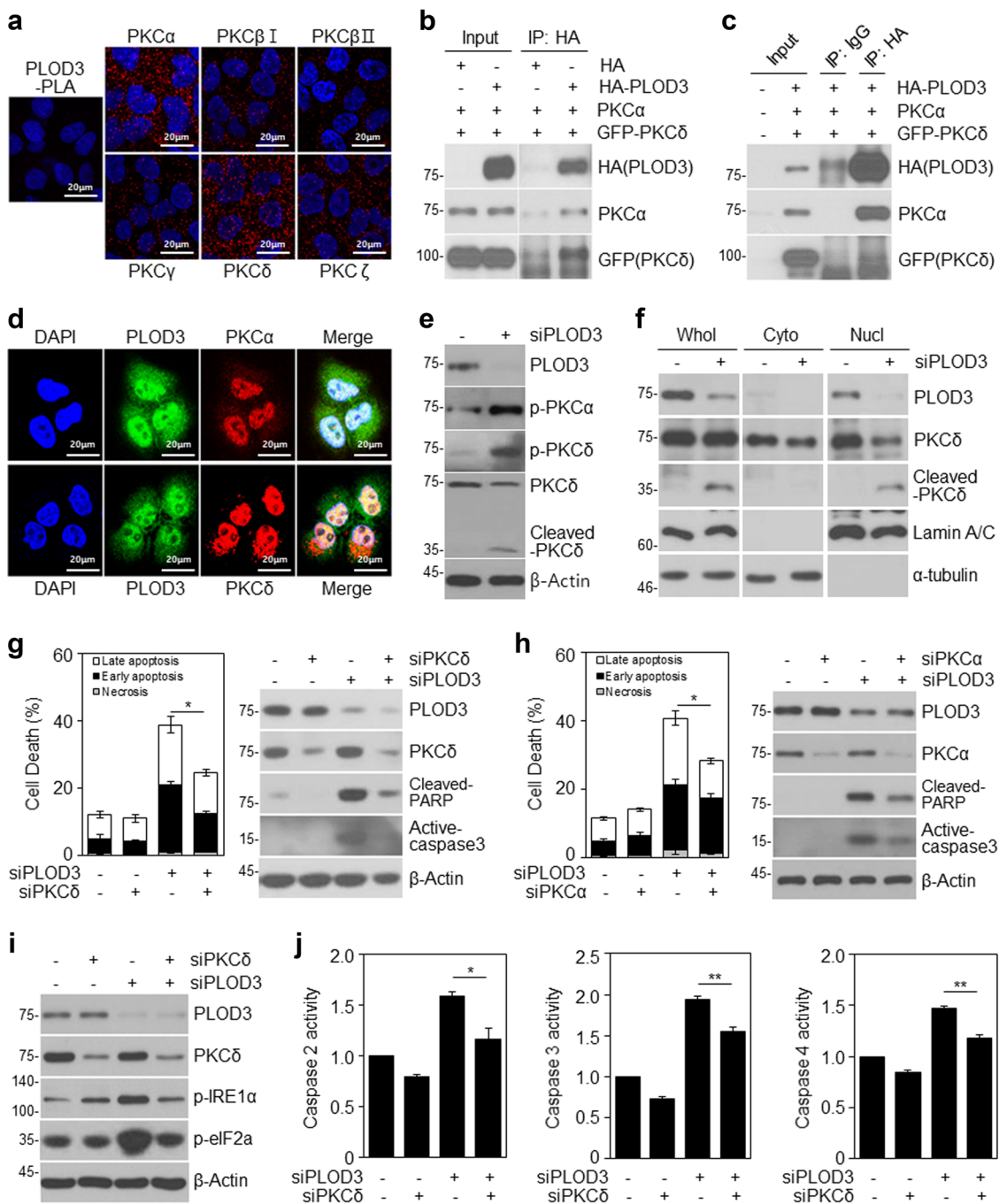


Fig. 5 (See legend on next page.)

compared to control siRNA, but combined transfection with *PKCδ* siRNA further enhanced this increase by ~1.2-fold. Further, *PLOD3* siRNA significantly increased caspase-3 activity by ~2-fold compared to that with control siRNA, but combined transfection with *PKCδ*

siRNA enhanced this increase by ~1.5-fold. Moreover, *PLOD3* siRNA significantly increased caspase-2 activity by ~1.5-fold compared to control siRNA, but combined transfection with *PKCδ* siRNA enhanced this increase by ~1.2-fold. Taken together, these results suggest that

(see figure on previous page)

Fig. 5 PLOD3 knockdown induces activation of PKCs and PLOD3 directly interacts with PKCs. **a** R-H460 cells were fixed and incubated with a mouse anti-PLOD3 antibody together with rabbit antibodies against PKCs, followed by in situ proximity ligation assay analysis. Representative confocal images of cells with proximity ligation assay-positive signals (red dots). **b, c** R-H460 cells were transfected with HA-PLOD3, PKC α , or GFP-PKC δ . Cell lysates were immunoprecipitated with normal IgG (a negative control) or HA antibody, and then immunocomplexes were resolved by SDS-PAGE and immunoblotted with antibodies against HA, PKC α , and GFP. **d** Immunostaining analysis of the localization of PKCs in R-H460 cells. Cells were fixed with 4% paraformaldehyde and immunostained for PLOD3 (green), PKC α (red), and PKC δ (red). DNA was visualized by DAPI staining. **e** PLOD3 or control siRNA was transfected into R-H460 cells; 24-h post-transfection, cell lysates were prepared and used in immunoblotting with antibodies targeting PLOD3, p-PKC α , PKC δ , p-PKC δ , and β -actin. **f** Subcellular fraction analysis of R-H460 cells following siCON or siPLOD3 treatment. Proteins in each fraction were resolved by SDS-PAGE and immunoblotted with antibodies against PLOD3, PKC δ , lamin A/C, and α -tubulin. **g** R-H460 cells were transfected with 40 nM siCON or siPLOD3 and/or siPKC δ for 48 h. Cell death in R-H460 cells was determined by AV/PI staining (left). Protein levels of the indicated proteins were determined by western blotting (right). * $P < 0.05$. **h** R-H460 cells were transfected with 40 nM siCON or siPLOD3 and/or siPKC α for 48 h. Cell death in R-H460 cells was determined by AV/PI staining (left). Protein levels of the indicated proteins were determined by western blotting (right). * $P < 0.05$. **i** Protein levels of PLOD3, PKC δ , p-eIF2 α , and p-IRE1 α were determined by western blotting. **j** Caspase-2, caspase-3, and caspase-4 activities in R-H460 cells (treated as in **b**) were determined by caspase activity assay. Data were collected using the Multiskan EX at 405 nm. * $P < 0.05$; ** $P < 0.01$

PKC δ , downstream of PLOD3, plays an important role in the PLOD3-mediated cell death mechanism.

Loss of PLOD3 overcomes chemoresistance in vitro

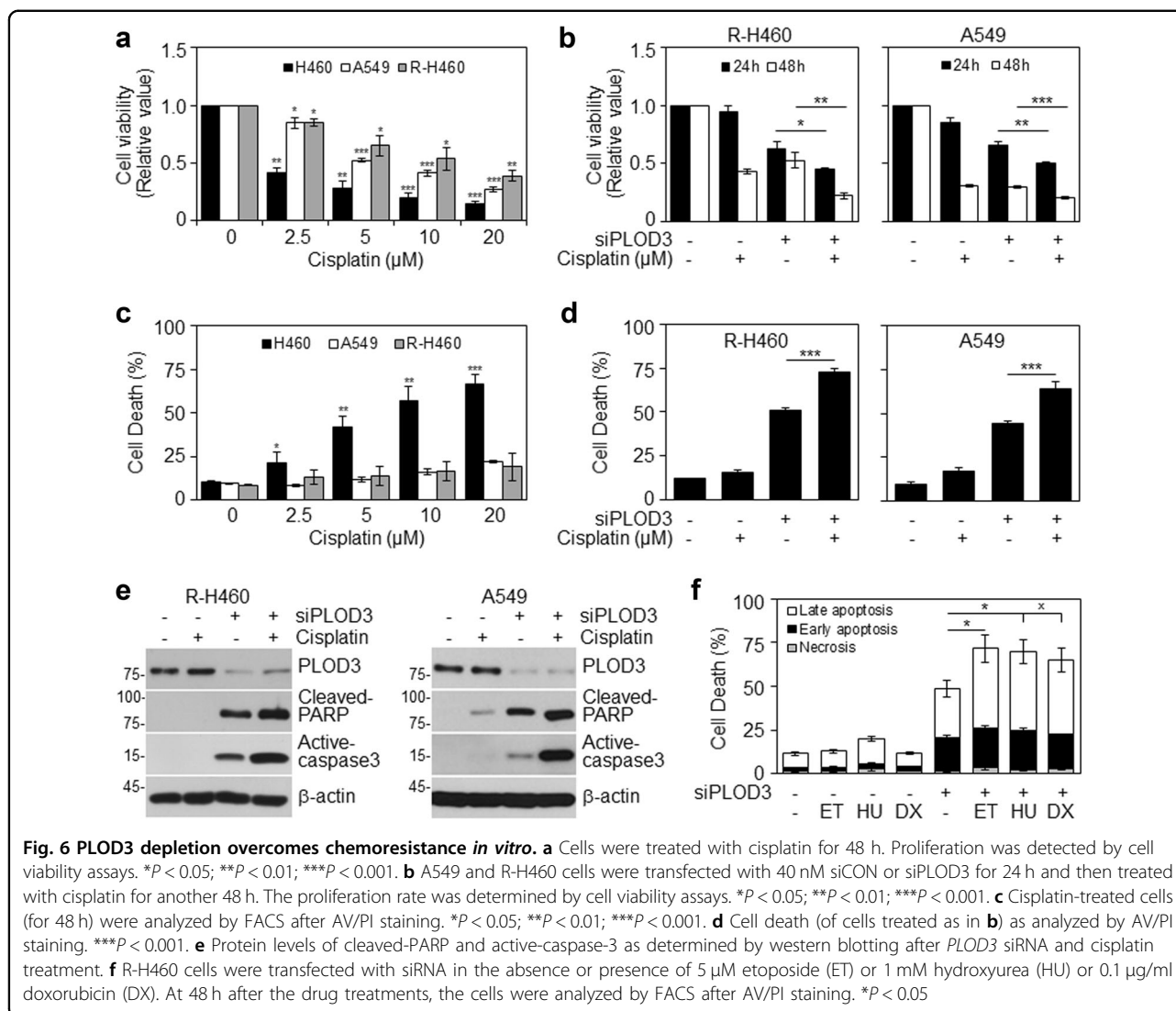
Next, we examined whether PLOD3 status influences the chemosensitivity of cancer cells. Interestingly, radio-resistant R-H460 cells were more resistant to cisplatin than their parental H460 cells, and resistance levels were similar to those of A549 cells (Fig. 6a, c). Next, we investigated the combined effect of cisplatin and PLOD3 siRNA on the viability and cell death of R-H460 and A549 cells. The combinatorial treatment attenuated cell viability and increased cell death compared to cisplatin alone (Fig. 6b, d), suggesting that PLOD3 status might influence the chemosensitivity of cancer cells to cisplatin. Next, to investigate the effect of PLOD3 knockdown on apoptosis, we measured the expression of cleaved PARP and active caspase-3 by western blot analyzes (Fig. 6e). Compared to cisplatin alone, the combination of PLOD3 siRNA and cisplatin resulted in significant increases in the levels of these two proteins in R-H460 and A549 cells. Whereas transfection of R-H460 cells with PLOD3 siRNA alone induced approximately 47% cell death compared to 10% in controls, the co-stimulation of PLOD3 siRNA-transfected cells with etoposide, hydroxyurea, or doxorubicin induced approximately 72%, 70%, and 65% cell death, respectively (Fig. 6f). Taken together, these results suggest that PLOD3 status is strongly associated with cancer cell chemosensitivity.

Discussion

PLOD3, which we identified as a radioresistance-related protein, has already been highlighted for its potential role in hepatocellular carcinoma, glioma, skin cancer, and prostate cancer^{16–19}. However, for lung cancer, the specific mechanism of PLOD3 regulation in cell death has not been elucidated.

In the current study, PLOD3 siRNA inhibited the proliferation of lung cancer cells and induced cell death, an effect that was augmented by radiation treatment, in cells and an in vivo model. We also found that (a) PLOD3 is differentially expressed during tumor progression, (b) PLOD3 expression promotes apoptosis in lung cancer by activating caspase-2,4-dependent signaling through ER stress-induced IRE1 α activation and caspase-3-dependent apoptosis, and (c) PLOD3 siRNA, combined with chemotherapeutic drugs, can increase cancer cell death, suggesting that PLOD3 is a potential therapeutic target in cancer. Thus, the present study provided the first evidence that PLOD3 inhibition can induce apoptosis in lung cancer cells, in addition to helping overcome radioresistance or chemoresistance.

Currently, PLOD2 is reportedly induced by hypoxia-inducible factor-1 α under hypoxic conditions, which in turn enhances hypoxia-induced epithelial–mesenchymal transition in glioma³² and breast cancer cells³³. Hypoxia-inducible factor-1 α also regulates PLOD1 transcription in breast cancer; however, PLOD2 activity is more critical for HIF-1-induced cancer progression³³. PLOD2 is also directly regulated by miR-26a-5p and miR-26b-5p, and PLOD2 expression is reportedly a possible prognostic marker for patients with bladder cancer³⁴ and renal cell carcinoma³⁵. Moreover, the E2Fs³⁶ and FOXA1³⁷ transcription factors have been demonstrated as regulators of PLOD2 during cancer progression. However, the regulation of PLOD1 and PLOD3 expression is not well understood. Thus, the upstream regulation of PLOD3 that could lead to the induction of cell death in our system remains to be elucidated. One report showed that miR-663a reduces PLOD3 biosynthesis by targeting the 3'-untranslated region of PLOD3 mRNA, suggesting the interrelationship of this microRNA in regulating collagen IV secretion under physiological conditions and in response to ER stress. Consistent herewith, we focused on



ER stress as an apoptosis-inducing factor as collagen is a major secretory protein in a variety of tissues, and defective secretion could result in physiological ER stress, fibrosis, and a number of disease states. As the pathway identified herein might represent a major physiological mechanism in lung cancer, it will be crucial to establish the role of the PKC δ -PLOD3-collagen axis in different tissues in future. Moreover, as PLOD3 is also present in the extracellular space, as well as in the serum and on the cellular surface¹⁹, our results suggest that PKC δ as a downstream effector of PLOD3 might also modulate other important extracellular PLOD3 functions³⁸.

PLOD3 mutations are reportedly associated with connective tissue disorders¹¹. *PLOD3* knockout in embryos and cells was reported to be associated with reduced glycosylated hydroxylysines on type IV and VI collagen and abnormal distribution³⁹. PLOD3 is overexpressed in

hepatocellular carcinoma⁴⁰ and is a potential diagnostic marker for early-stage disease⁴⁰. Further, PLOD3 knock-down controls liver tumor incidence and growth rates in a spontaneous mouse model of hepatocellular carcinoma⁴⁰. Our study showed consistent results for lung cancer and suggests that secreted PLOD3 could serve as a potential inducer of lung cancer metastasis or as a prognostic marker. We are currently expanding this mechanistic investigation in relation to metastasis using lung cancer cells and patient samples.

Interestingly, as we observed nuclear localization of cleaved PKC δ following PLOD3 knockdown, we hypothesized that *PLOD3* siRNA, via the production of ROS, might induce post-translational modifications in PKC δ , which allow its translocation to the nucleus. However, these events did not depend on ROS levels. Moreover, the hinge domain of PKC δ is the site for caspase-3 cleavage,

which happens in the nucleus and leads to the release of the δ -catalytic fragment (δ CF), corresponding to the kinase domain. Thus, in some systems, apoptotic cell death is associated with caspase 3-dependent PKC δ cleavage; the formation of the catalytically active fragment of PKC δ in our cellular model following *PLOD3* siRNA treatment should be explored further in future studies.

Recently, diverse molecular biomarkers of drug resistance have been identified. For example, the protein disulfide isomerases PDIA4 and PDIA6 regulate resistance to cisplatin-induced apoptosis in lung adenocarcinoma⁴¹. The copper transporter CTR1 is involved in the uptake of cisplatin and has been developed as a therapeutic target⁴². Thus, medical protocols should be planned according to the character of each patient and it needs to find more efficient biomarkers for identifying the molecular mechanisms of cisplatin resistance. Our results suggest that chemotherapy could be more effective in combination with RNAi-mediated knockdown of *PLOD3*. Certainly, for the improvement of such a therapeutic strategy for clinical application, a suitable vector system is necessary. We also need to explore the effects of *PLOD3* knockdown on normal tissues in future before clinical application of this strategy can be contemplated. We demonstrated that the level of activated PKC δ or PKC α was increased upon treatment with siP $LOD3$ and that the siP $LOD3$ -induced apoptosis was decreased by inhibiting the expression of *PLOD3* and PKC δ or PKC α . Thus, we are further planning to identify the functional relationship between *PLOD3* and PKC δ or PKC α ; the cellular events or functional mechanisms underlying these also need to be elucidated.

In summary, the *PLOD3* siRNA used in this study are potent tools to modulate *PLOD3* expression and they might ultimately be developed into attractive anti-tumor therapeutics. Further, *PLOD3* might represent a prognostic biomarker and a target for reversing cisplatin resistance in lung cancer.

Materials and methods

Cell culture and treatment

Human lung cancer cell lines (H460 and A549) were purchased from ATCC (Manassas, Virginia, USA), and we established a radioresistant H460 (R-H460) cell line derived from parental radiosensitive H460 lung cancer cells treated cumulatively with 2 Gy radiation twice a week for 20 weeks⁵. H460, A549, and R-H460 cells were cultured in RPMI-1640 medium supplemented with 10% fetal bovine serum. Cells were irradiated using a 137 Cs-ray source (Atomic Energy of Canada) at a dose rate of 3.81 Gy/min. Where indicated, cells were treated with etoposide, hydroxyurea, doxorubicin, and cisplatin to induce apoptosis. Caspase inhibitors [Z-VAD-FMK (Adipogen), Z-VDVAD-FMK, Z-DEVD-FMK, and Z-

YVAD-FMK (R&D Systems)] were used to block caspase activation. Inhibitor APY29 was used to block IRE1 α activation. Where indicated, cells were treated with 3-methyladenine to block autophagy.

PLOD3 siRNA transfections

The following human *PLOD3*-specific siRNAs, synthesized by Genolution, were used: #1; 5'-GGUAAAAGA AGGAAAUGGAUU-3', #2; 5'-GGAAGUACAAGG AUGAUGAUGACGACGA-3', #3; 5'-AUAUGAUCauc AUGUUUGUUU-3', #4; 5'-GCCUAAAUCUGGAUCA UAAUU-3'. The siRNA duplexes were transfected into cells using Lipofectamine[®] RNAiMAX Reagent according to the manufacturer's guidelines.

Western blot analysis

Western blot analyzes were performed as described previously¹⁶ using primary antibodies targeting the following proteins: *PLOD3* (Proteintech Group, Chicago, IL, USA); p-PERK, PERK, p-eIF2 α , HA, GFP, lamin A/C, α -tubulin, PKC α , PKC δ (Santa Cruz Biotechnology); cleaved-PARP (Asp214), cleaved caspase-3, eIF2 α , IRE1 α , γ H2AX, and p-PKC δ (Cell Signaling Technology); p-IRE1 α (Novous); ATF6, caspase-2 and caspase-4 (Abcam); and β -actin (Sigma) was used as a loading control.

Quantitative reverse transcription-polymerase chain reaction (qRT-PCR)

Total RNA was isolated using an RNeasy Mini kit (Qiagen). qPCR was performed in triplicate using a PIKOREAL 96 (Thermo) and SYBR Premix Ex Taq (Takara Bio, Shiga, Japan). A two-temperature thermocycling program was used, with 42 cycles of 95 °C (denaturation) and 55 °C (annealing). Target-gene amplification signal was normalized to that of *GAPDH* in the same reaction.

Cell viability assay

Cells were seeded at 5000 cells/well in a 96-well plate and incubated for 24 h in accordance with the indicated experimental conditions. For quantification of cell viability, an equal volume of culture medium containing Cyto XTM Reagent (LPS solution) was added to the cells and the plate was incubated for 4 h. Cell viability was then measured using Multiskan EX (Thermo Fisher Scientific, Germany) at 450 nm.

Fluorescence-activated cell sorting (FACS)

Cells (3×10^5) were seeded in a 60-mm dish and exposed to the indicated experimental conditions. For quantification of cell death, cells were trypsinized, washed in phosphate-buffered saline (PBS), and then resuspended in 1 \times binding buffer. Cells were incubated with annexin V (AV) and propidium iodide (PI) for

15 min, and analyzed with a FACScan flow cytometer (Becton Dickinson).

Clonogenic assay

Cells were seeded in triplicate into 60-mm tissue culture dishes at 2, 4, 8, 16, and 32×10^2 cells/dish and exposed to 0, 2, 4, 6, and 8 Gy, respectively. Cells were exposed once to different doses of radiation. After 14 days, colonies arising from surviving cells were stained with trypan blue and counted using a colony counter (Imaging Products).

Tumor xenograft animal model

Studies with the tumor xenograft animal model were performed as described previously⁴³ using control siRNA or PLOD3 siRNA #2 (KIRAMS 2015–0070).

Immunofluorescence confocal microscopy

Immunofluorescence staining was performed as described previously⁴³ using primary antibodies against PLOD3 (Proteintech), PKC α , PKC δ , and GRP94 (Santa Cruz Biotechnology). Cell nuclei were counter-stained with DAPI. Images were acquired using a confocal laser-scanning microscope (model LSM 710, Carl Zeiss) and were processed with ZEN 2009 Light Edition (Carl Zeiss).

Data mining using Kaplan–Meier plotter

Kaplan–Meier survival curves in relation to *PLOD3* expression were generated for lung cancer, using KM Plotter (<http://kmpplot.com/>). Cancer type was defined as lung cancer and data type as mRNA, whereas analysis type was defined as cancer vs. normal.

Caspase activity assay

Caspase activities were measured using caspase family activity assay kits (Abcam) according to the manufacturer's recommendations. Data were collected using a Multiskan EX at 405 nm.

Proteasome activity

Proteasome activities were measured using proteasome-GloTM chymotrypsin-like, trypsin-like, and caspase-like cell-based assays (Promega) according to the manufacturer's recommendations. Data were collected using a Victor X2 multi-label reader (Perkin Elmer).

ROS assay

R-H460 cells were treated with siRNA under the indicated experimental conditions. The cells were incubated with 10 nM 2',7'-dichlorofluorescein diacetate (Molecular Probes) in the dark at 37 °C for 30 min. Cell staining was examined with a laser-scanning confocal microscope (model LSM 710, Carl Zeiss) equipped with an argon laser tuned to an excitation wavelength of 488 nm, LP505 emission filter (515–540 nm), and Zeiss Axiovert -100X

objective lens. Two groups of cells were randomly selected from each sample.

In situ proximity ligation assay

Paraformaldehyde-fixed R-H460 cells were permeabilized with 0.2% Triton X-100, washed, and blocked with blocking solution (Olink Bioscience). Mouse monoclonal anti-PLOD3 antibody (Proteintech Group) together with rabbit polyclonal antibodies against PKC α , PKC β I, PKC β II, PKC γ , PKC δ , and PKC ζ (Santa Cruz Biotechnology) were used for the proximity ligation reaction. The assay was conducted using the Duolink Detection Kit (Sigma) according to the manufacturer's protocol.

Plasmid construction and transfection

Plasmids were constructed by standard cloning techniques and were verified by DNA sequencing. Human PLOD3 cDNA (wild type) purchased from Origene (Cat. No. SC324563) was PCR-amplified and cloned into the pcDNA3.1 vector. Wild type PLOD3 was amplified with primers 5'-GATGGATCCATGACCTCCTCGGGGCTGGA-3' (BamHI) and 5'-TCTCGAGTCAGGGGTTCGACAAAGGAC-3' (XhoI), and the amplicon was inserted between BamHI and XhoI in the pHA vector. PKC α and GFP-PKC δ expression vectors were purchased from the Korea Human Gene Bank. Plasmids were transfected using Mirus-2020 Reagent according to the manufacturer's guidelines.

Immunoprecipitation

R-H460 cells were transfected with vectors under the indicated experimental conditions for 48 h. The cells were washed twice with PBS, harvested, and lysed for 30 min in NP-40 buffer [50 mM Tris-HCl (pH 8), 150 mM NaCl, 1% NP-40, and 100X protease and phosphatase inhibitor cocktail]. Samples were diluted to 500 μ g of protein in 800 μ l of buffer and pre-cleared for 1 h at 4 °C with 50 μ l of a 50% slurry of protein A/G-Sepharose beads (GE Healthcare). After brief centrifugation to remove pre-cleared beads, 1 μ g of antibody against HA (Santa Cruz Biotechnology) was added to each sample and incubated on a rocking platform at 4 °C overnight. The immune complex was precipitated by incubation with 40 μ l of protein A/G-Sepharose beads at 4 °C for 4 h. The beads were washed thrice with immunoprecipitation buffer and then boiled with sample buffer [0.1 M Tris-HCl (pH 6.8), 4% SDS, 40 mM EDTA, 20% glycerol, and β -mercaptoethanol].

Subcellular fractionation

Cells were lysed using subcellular fractionation buffer (250 mM sucrose, 20 mM HEPES (pH 7.4), 10 mM KCl, 1.5 mM MgCl₂, 1 mM EDTA, 1 mM EGTA, 1 mM DTT, and protease inhibitors), and the lysate was passed

through a 25-gauge needle 10 times using a 1-ml syringe. The lysate was placed on ice for 20 min and centrifuged at 3000 rpm for 5 min to obtain nuclear pellet. The pellet was washed with fractionation buffer 11 times, and the supernatant was centrifuged at 8000 rpm. The supernatant here refers to the cytosolic and membrane fraction.

Statistical analysis

Cell culture experiments were performed at least in triplicate. All data are expressed as mean \pm standard deviation values. Statistical differences between groups were assessed using Student's *t*-test (two-tailed) analysis. *P*-values were interpreted as follows: not significant (n.s.), **P* < 0.05, ***P* < 0.01, and ****P* < 0.001.

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Author contributions

S.H. and E.K. conceived the idea, designed the experiment, and wrote the manuscript. J.B., H.Y., G.K., J.L., J.K., Y.J., J.C., C.L., J.S., J.A., and J.K. performed the experiments. S.H. and E.K. analyzed the data. All authors read and approved the final version of the manuscript.

Conflict of interest

The authors declare that they have no conflict of interest.

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