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Novel small molecule inhibiting CDCP1-PKC δ pathway reduces tumor metastasis and proliferation

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Key words

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CUB domain-containing protein-1 (CDCP1) is a trans-membrane protein predominantly expressed in various cancer cells and involved in tumor progression. CDCP1 is phosphorylated at tyrosine residues in the intracellular domain by Src family kinases and recruits PKC δ to the plasma membrane through tyrosine phosphorylation-dependent association with the C2 domain of PKC δ , which in turn induces a survival signal in an anchorage-independent condition. In this study, we used our cell-free screening system to identify a small compound, glycoconjugated palladium complex (Pd-Oqn), which significantly inhibited the interaction between the C2 domain of PKC δ and phosphorylated CDCP1. Immunoprecipitation assays demonstrated that Pd-Oqn hindered the intercellular interaction of phosphorylated CDCP1 with PKC δ and also suppressed the phosphorylation of PKCô but not that of ERK or AKT. In addition, Pd-Oqn inhibited the colony formation of gastric adenocarcinoma 44As3 cells in soft agar as well as their invasion. In mouse models, Pd-Oqn markedly reduced the peritoneal dissemination of gastric adenocarcinoma cells and the tumor growth of pancreatic cancer orthotopic xenografts. These results suggest that the novel compound Pd-Oqn reduces tumor metastasis and growth by inhibiting the association between CDCP1 and PKCô, thus potentially representing a promising candidate among therapeutic reagents targeting protein-protein interaction.

UB-domain containing protein 1 (CDCP1), a type I transmembrane glycoprotein with three CUB domains in the extracellular region, was initially reported as being predominantly expressed in colorectal cancers.⁽¹⁾ During the investigation of phosphoproteins associated with anchorage-independent growth of lung adenocarcinoma cells, we previously discovered that CDCP1 protein plays a major role in the induction of anoikis resistance in metastatic tumors as a substrate of Src family kinases (SFKs).⁽²⁾ Specifically, CDCP1 phosphorylation at tyrosines in the intercellular domain upon SFK activation mediates PKCS recruitment to the plasma membrane and its subsequent activation through specific interaction of the PKC δ C2 domain with tyrosine-phosphorylated CDCP1.⁽²⁻⁴⁾ Thus, CDCP1 phosphorylation promotes anchorage-independent cell survival and degradation of the extracellular matrix through the interaction with PKC δ .^(2,5,6) In addition, it has been revealed that CDCP1 suppresses anchorage-independent cell death by inhibiting autophagy,⁽⁷⁾ inhibits the epithelial phenotype in pancreatic cancer cells, and is involved in the trastuzumab resistance of HER2-positive breast cancer.⁽⁸⁻¹⁰⁾ Furthermore, the expression of CDCP1 is induced by activated Ras and correlates with poor prognosis in lung, pancreatic, renal, and ovarian cancer.^(6,11-13) Together, such studies have indicated that the CDCP1-PKCδ signaling pathway is involved

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This is an open access article under the terms of the Creative Commons Attrib ution-NonCommercial License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited and is not used for commercial purposes. in various features of tumor progression such as invasion and metastasis in multiple types of cancers.

Numerous experiments using mouse models have demonstrated that CDCP1 represents a potent therapeutic target for various cancers. For example, the suppression of CDCP1 reduced invasion and the peritoneal dissemination of gastric scirrhous carcinoma,⁽¹⁴⁾ whereas blockage of the CDCP1 pathway using specific antibodies suppressed the growth of lung and breast cancer cells⁽¹⁵⁾ or of ovarian cancer tissues transplanted in mice.⁽¹⁶⁾ Furthermore, a cytotoxin saporin-conjugated anti-CDCP1 antibody reduced the growth and dissemination of subcutaneous tumors in a mouse xenograft model of prostate cancer cells.⁽¹⁷⁾

Therefore, based on the previous work by ourselves and others, we considered that inhibiting the specific interaction between CDCP1 and PKC δ might represent an alternative therapeutic approach for targeting the CDCP1 pathway in addition to the antibody-mediated models that directly target CDCP1 listed above. Accordingly, in the current study we performed *in vitro* screening of compounds that inhibited the association of phosphorylated CDCP1 with the C2 domain of PKC δ to identify a potent small molecule inhibitor. The ability of the candidate molecule to block the CDCP1-PKC δ signaling pathway and inhibit cancer cell proliferation and invasion was

assessed *in vitro*, as was its ability to reduce tumor growth and peritoneal dissemination in a mouse model.

Materials and Methods

Cell culture. The human gastric cancer 44As3 and 58As9 cells have been previously described.⁽¹⁸⁾ BxPC3, MiaPaCa-2 (pancreatic cancer), and A549 (lung cancer) cell lines were obtained from the American Type Culture Collection. The cells were cultured in RPMI1640 (Thermo Fisher Scientific, Waltham, MA, USA) supplemented with 10% fetal bovine serum (FBS) (Thermo Fisher Scientific) at 37°C in a humidified atmosphere containing 5% CO₂. Human normal gastric fibroblast (NF-37) cells were kindly gifted from Dr. Yashiro.⁽¹⁹⁾ NF-37 cells were cultured in DMEM-high glucose (Thermo Fisher Scientific) supplemented with 10% FBS. HEK293T cells were cultured in DMEM supplemented with 10% FBS.

Antibodies. An anti-GFP antibody (#598) was purchased from MBL, Nagoya, Japan. An anti-FLAG M2 antibody conjugated with peroxidase was purchased from Sigma (St. Louis, MO, USA). The anti- β -actin antibody (N21) and horse radish peroxidase (HRP)-conjugated anti-goat IgG antibody were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Anti-Akt (#9272), anti-phospho-Akt (Ser473) (#9271), anti-CDCP1 (#4115), anti-phospho-CDCP1 (Tyr734) (#9050), anti-ERK1/2 (#9102), anti-phospho-ERK1/2 (Thr202/Tyr204) (#9101), anti-PKC δ (#2058), and anti-phospho-PKC δ (Tyr311) (#2055) antibodies were purchased from Cell Signaling Technology (Denvers, MA, USA). Anti-CDCP1 antibody for immunoprecipitation was originally developed as described previously.⁽²⁾

Biding assay for CDCP1-PKC δ interaction and chemical screening. Biotinylated peptide diluted with phosphate buffered saline calcium- and magnesium-free (PBS(-)) at 1 µg/mL was incubated in streptavidin-coated 96-well plates (Thermo Fisher Scientific) at 37°C for 1 h. After washing twice with PBS(-) containing 0.05% Tween-20 (PBS-T), 50 µL C2-FLAG diluted with blocking buffer (3% bovine serum albumin Fraction V [BSA; Roche, Mannheim, Germany] in PBS-T) was added. For chemical screening, a 2 nM concentration of C2-FLAG was used and each compound was added at 20 µM, then the plates were incubated at 37°C for 2 h. After the wells were washed with PBS-T four times, anti-FLAG M2 conjugated with peroxidase (Sigma) diluted at 1:10 000 by blocking buffer was added and incubated at room temperature (RT) for 30 min. After washing with PBS-T four times, the bound antibody was detected by colorimetric reaction using 50 µL 3,3',5,5'-tetramethylbenzidine (TMB) (Wako, Osaka, Japan), then the reaction was stopped by adding the same volume of 1 M HCl. The absorbance of the wells at 450 nm was measured using an iMark microplate reader (BioRad, Hercules, CA, USA). The deposited chemical library was obtained from the Screening Committee of Anticancer Drugs, Japan. The peptide sequence used in the present study is shown in Table S1.

HTRF assay for CDCP1-PKC δ interaction. The biotinylated peptide with phospho-tyrosine (bio-762pY) and a C2-FLAG recombinant protein were used for HTRF assay. Binding was detected using an anti-FLAG antibody conjugated with Eu (anti-FLAG-Eu) (Cisbio, Codolet, France) and streptavidin-d2 (Cisbio). All of the components were diluted with HTRF assay buffer (20 mM sodium-potassium phosphate (pH7.5), 0.5% BSA, 0.25 M potassium fluoride, 0.05% Tween-20). The components of C2-FLAG, bio-762pY, anti-FLAG-Eu, and streptavidin-d2 were mixed in 20 μ L total volume in a 384-well plate (784075; Greiner, Kremsmünster, Austria) and incubated at RT for 1 h. The signal was measured using SynergyH1 with the HTRF filter set (Bio Tek Instruments, Inc., Winooski, VT, USA).

Immunoprecipitation. HEK293T cells were transfected with p3xFLAG-CMV-14-CDCP1, pEGFP-N1-PKC δ , and/or pLNCX-vSrc using Lipofectamine2000 according to the manufacture's manual (Thermo Fisher Scientific). After a day, the cells were harvested in PLC buffer (50 mM HEPES [pH 7.0], 150 mM NaCl, 1.5 mM MgCl₂, 10% glycerol, 1% Triton X-100, and 5 mM ethylene glycol-bis(β -aminoethyl ether)-N,N, N',N'-tetraacetic acid) containing protease (Wako) and phosphatase inhibitor cocktails (Nakalai, Kyoto, Japan) and incubated on ice for 15 min. After centrifugation at 15 000 g, the



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Fig. 1. Establishment of an assay system for the interaction of CDCP1 with PKCδ. (a) Domain structure of PKC δ and the region of the C2 recombinant protein (1-160) that was used for the C2-CDCP1 binding assay. (b) Purification of C2-FLAG. GST-C2-FLAG protein purified from bacterial lysates was proteolyticaly digested (lane 1) and the GST was removed (lane 2). The proteins were subjected to SDS-12.5% PAGE and Coomassie brilliant blue (CBB) staining. (c) Selective binding of the C2 domain to the phosphorylated site at Y762 of CDCP1. Biotinylated peptide, bio-762pY, bio-762, bio-734pY, or bio-734, immobilized on a 96-well plate coated with streptavidin and C2-FLAG (0.5-500 nM) was used for the binding assay. (d) Competitive inhibition in the C2-CDCP1 binding assay by free CP762pY peptide. C2-FLAG (2 nM) and bio-762pY were used for the assay and the binding reaction was performed in the presence of 0.1-10 000 nM CP762pY or CP762 peptide that was not biotinylated.

supernatant was subjected to immunoprecipitation as the input cell lysate. The protein concentration of the lysate was determined using the BCA protein assay (Thermo Fisher Scientific) with BSA as a standard. The cell lysate (2 mg) was mixed and incubated with anti-FLAG agarose (SIGMA) (15 μ L bed volume) at 4°C with rotation for 2 h. The agarose was then washed five times with PLC buffer and boiled in sodium dodecyl sulfate (SDS) sample buffer. Immunoprecipitation of endogenous proteins were described previously.⁽²⁾ Briefly, 44As3 cells treated with DMSO or 10 μ M Pd-Oqn were lysed in PLC buffer. The cell lysate was incubated with Protein G Sepharose (GE Healthcare, Chicago, IL, USA) that was bound with anti-CDCP1 antibody or normal rabbit IgG. The beads were washed with PLC buffer extensively and the bound proteins were finally eluted with SDS-sample buffer.

Western blotting. The cell lysate (20 µg) or immunoprecipitated samples were subjected to SDS-10% polyacrylamide gel electrophoresis (PAGE) and then transferred to a polyvinylidene fluoride membrane (Millipore, Darmstadt, Germany). The membrane was incubated with the Blocking One (Nakalai) and then incubated with primary antibody: anti-GFP (1/5000), anti-FLAG (1/5000), anti-CDCP1 (1/1000), anti-β-actin (1/1000), anti-PKCS (1/1000), anti-p-PKCS (1/1000), anti-p-ERK1/2 (1/ 1000), anti-ERK1/2 (1/1000), anti-p-AKT (1/1000), or anti-AKT (1/1000) antibody at 4°C for 16 h. The bound antibody was detected using Pierce ECL Plus Western Blotting Substrate (Thermo Fisher Scientific) after binding of anti-rabbit or goat IgG antibody conjugated with HRP (GE Healthcare). For immunoprecipitated proteins, TrueBlot anti-rabbit IgG (Rockland Immunochemicals, Limerick, PA, USA) was used as a secondary antibody. The chemiluminescent signals were visualized and analyzed using an Amersham Imager 600 (GE Healthcare).

Cell proliferation assay. Cells were plated onto 96-well plates at 3×10^3 cells per well and cultured for 2 days in the presence or absence of chemical compounds. Viable cells were determined using the Cell Counting Kit (Dojindo, Kumamoto, Japan).

Soft-agar colony formation assay. 44As3 cells $(1 \times 10^4 \text{ cells})$ suspended with the complete medium containing 0.35% agarose were plated on the bottom layer of the complete medium with 0.5% agarose. RPMI1640-10%FBS containing Pd-Oqn or dimethyl sulfoxide (DMSO) was added on top of the soft-agar and then placed in a 37°C incubator. The medium was changed to fresh medium twice a week. After 21 days, the colonies were stained with crystal violet and the numbers of visible colonies per well were counted.

Migration and invasion assay. For the invasion assay, 100 μ L of 0.1 mg/mL ice-cold Matrigel (Corning, Corning, NY, USA) was added onto a transwell chamber (8.0 μ m pore size) (Corning) and dried. 44As3 (1 × 10⁵ cells) suspended in RPMI1640 without serum was added to the top of the chamber. RPMI1640-10% FBS was added in the bottom chamber. After 48 h, the cells on the chamber top were removed using a cotton swab and the invaded or migrated cells were fixed by 4% paraformaldehyde. The cells were stained with DAPI and visualized using a fluorescent microscope (BZ-9000, KEYENCE) with a 20× objective lens. The stained nuclei were counted using ImageJ free software.

Preparation of Pd-Oqn and Pt-Oqn. Pd-Oqn $\{= \text{Chloror}\{N-(hydroxo-quinoline-2-ylmethylidene)-\beta-D-glucosamine}\}$ palladium(II)} and Pt-Oqn $\{= \text{Chloror}\{N-(hydroxo-quinoline-2-ylmethylidene)-b-D-glucosamine}\}$ platinum(II)} were synthesized as described in Yano *et al.*⁽²⁰⁾ Briefly, 8-hydroxy-2-



Fig. 2. Identification of small molecules that inhibit C2-CDCP1 binding. (a) Inhibitory effect of small compounds on the C2-CDCP1 binding assay. The small compounds 5-3B, 5-3C, and 5-3D (1.0–50 μ M) or DMSO were used in the C2-CDCP1 binding assay as inhibitory compounds. (b) Structures of 5-3C and 5-3D, identified as inhibitory compounds on C2-CDCP1 binding. 5-3C; Pd-Oqn. 5-3D; Pt-Oqn. (c) Inhibition kinetics of Pd-Oqn and Pt-Oqn on C2-CDCP1 binding. HTRF assays with Pd-Oqn (Pd) or Pt-Oqn (Pt) were performed and IC50 values were calculated.

quinolinecarbaldehyde was added to a methanol solution of D (+)-glucosamine hydrochloride with *tert*-BuOK at RT to give Schiff base and a stoichiometric amount of Na₂PdCl₄ or Pt (DMSO)₂Cl₂ was added to give Pd-Oqn or Pt-Oqn, individually. Each compound was purified by recrystallization in MeOH/dimethyl formamide. The structures of the final products were confirmed by nuclear magnetic resonance analysis. The compounds were dissolved in DMSO at 100 mM and stored at -20° C as small aliquots until use.

Mouse models. Pd-Oqn was dissolved in DMSO at 100 mg/ mL and then diluted in 0.5% methylcellulose containing 0.1%

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Inhibitor of CDCP1-mediated metastasis



tween-80. 44As3 cells (5×10^5) were inoculated intraperitoneally into 6-week-old BALB/c-nu/nu male mice purchased from Charles River Laboratories Japan, Inc. Pd-Oqn (15 mg/ kg) or the same volume of DMSO was administered by intraperitoneal injection starting at 1 day after the inoculation as described in. At 14 days after inoculation, the mice were sacrificed by cervical dislocation under isoflurane anesthesia and dissected. The number of mesentery nodules visible by eye was determined. For the pancreatic cancer model, Mia-PaCa-2 cells were inoculated in the pancreas of 8-week-old C.B-17/lcr-scid/scid Jcl (SCID) female mice purchased from CLEA Japan, Inc. Pd-Oqn (15 mg/kg) or DMSO was administrated by intraperitoneal injection starting at 1 day after the inoculation as described in. After 23 days, the mice were sacrificed and the pancreatic tumor was removed and weighed. These experiments were approved by the Committee for Ethics of Animal Experimentation and conducted in accordance with the guidelines for Animal Experiments in the National Cancer Center.

Statistical analysis. Statistical analysis was performed using the Student's *t*-test with Excel software. Data are expressed as the means \pm standard deviation (SD).

Results

Establishment of a screening system for CDCP1 pathway inhibitor identification. To identify small molecules that inhibit the CDCP1 signaling pathway, we established an assay system to quantitatively detect the interaction of phosphorylated CDCP1 Fig. 3. Inhibition of the CDCP1-PKC δ pathway by Pd-Oqn. (a) Blockage of the CDCP1-PKC δ interaction by Pd-Oqn in cells. Immunoprecipitation was performed as described in Materials and Methods. Immunoprecipitated proteins and input lysates were subjected to western blotting with the antibodies as indicated at the right side. The top indicates the transfection plasmids and Pd-Oqn addition. (b) Inhibition of the phosphorylation of PKCo by Pd-Oqn. 44As3 cells were treated with Pd-Oqn and the lysates were subjected to western blotting with the antibodies as indicated at the right side. (c) Inhibition of the endogenous CDCP1-PKCo interaction by Pd-O qn. 44As3 cells were treated with DMSO (-) or Pd-Oqn (Pd) for 24 h and the cell lvsates were subjected to immunoprecipitation with anti-CDCP1 antibody or control IgG. The precipitated proteins (IP) and the cell lysate (input) were subjected to western blotting with the antibodies as indicated at right side. CDCP1 bands showed both full-length (FL) and cleaved form (CL) at about 135 and 70 kDa respectively. Molecular weight is indicated at the left.

with PKC δ . It has been reported that the C2 domain of PKC δ binds to the sequence motif containing phosphorylated tyrosine 762 of CDCP1 whereas the SH2 domain of Src binds to CDCP1 phosphorylated tyrosine at 734.⁽⁴⁾ Therefore, biotiny-lated peptides (23 a.a.) derived from the C2-binding motif containing the phosphorylated tyrosine (Y762) of CDCP1 along with several controls such as non-phosphorylated peptides were synthesized and immobilized on streptavidin-coated plates (Table S1). The C2 domain of PKC δ (a.a. 1–160) tagged with FLAG (C2-FLAG) was purified as described in Supplementary Doc. S1 and protein purity was confirmed by SDS-PAGE (Fig. 1a,b).

The association of CDCP1 peptides with the C2-FLAG protein was detected by colorimetric reaction with TMB following binding of an anti-FLAG antibody conjugated with peroxidase (Fig. S1A). To confirm the specificity of the assay system, we first checked whether the C2 domain preferentially bound to a phosphotyrosine-containing CDCP1 peptide corresponding to the PKCS binding site. The results showed that C2-FLAG bound to the CDCP1 peptide phosphorylated at Y762 (bio-762pY) but not obviously to the peptide phosphorylated at Y734 (bio-734pY). The interaction with the non-phosphorylated peptide (bio-762Y) was substantially weaker than bio-762pY (Fig. 1c). In addition, a competitive peptide (11 a.a.) of 762pY (CP762pY) but not 762 (CP762) inhibited the interaction of C2 with the bio-762pY peptide (Fig. 1d). These findings demonstrated that this assay system was capable of detecting specific association of the phosphorylated Y762 of CDCP1 with the C2 domain of PKC δ and exhibited sufficient



Fig. 4. Inhibitory effect of Pd-Oqn on the growth of cancer cells. (a, b) Inhibition of colony formation of 44As3 cells in soft agar by Pd-Oqn and Pt-Oqn. The soft agar colony formation assay was performed as described in Materials and Methods. Representative photos of the colonies in soft agar with DMSO, Pd-Oqn, or Pt-Oqn (5, 25, and 100 μ M) are shown in (a). The bar graph indicates the number of colonies in soft agar with DMSO, 5 μ M Pd-Oqn, or Pt-Oqn as the means \pm SD of three biological replicates. Statistical significance was determined using the standard Student's *t*-test. **P* < 0.05. (d) Inhibition of cell proliferation of various cancer cells by Pd-Oqn. The cells were cultured with Pd-Oqn (1.0–100 μ M) or DMSO (control) and cell viability was determined using the CCK assay after 48 h. The graph indicates the relative value to DMSO treated cells of each cell line as the means \pm SD of three biological replicates.

sensitivity for use in screening molecules that inhibited the interaction of PKC δ with phosphorylated CDCP1.

Identification of an inhibitor of the CDCP1-PKC δ interaction. The first screening library used to identify inhibitory molecules toward the interaction of CDCP1 with PKC δ consisted of 495 deposited chemical compounds. Two compounds (5-3C and 5-3D) were identified based on their outstanding inhibitory effect on the interaction (Fig. S1B). We then separately confirmed that these two compounds significantly inhibited the interaction in a dose-dependent manner (Fig. 2a), compared with the effect of an adjacent unrelated compound (5-3B) as a control. The small compound corresponding to 5-3C was a palladium complex (Pd-Oqn) with glucose and quinolinol and 5-3D was a platinum complex (Pt-Oqn) with the same structure (Figs 2b and S2). In these molecules, each metal atom is surrounded by four donor atoms, two nitrogen atoms, a phenolic oxygen atom, and a chloride ion to form an approximately square planar structure. Next, we proceeded to further screen another chemical library consisting of 10 000 compounds using the same assay system for the purpose of identifying several additional inhibitory compounds that are not as potent as the Pd-Oqn and Pt-Oqn compounds.

We also developed another binding assay based on homogeneous time-resolved fluorescence resonance energy transfer (HTRF) technology to confirm the inhibitory effect of these compounds on the CDCP1 with PKC δ interaction. For this assay, we used C2-FLAG and bio-762pY and binding was detected using anti-FLAG-Eu and Streptavidin-d2 (Fig. S3A). The interaction of C2-FLAG with bio-762pY was detected in a dose-dependent manner using the HTRF assay (Fig. S3B). In addition, the interaction was inhibited by the non-biotinylated CP762pY peptide with an IC50 value calculated as 91 nM (Fig. S3C). The inhibition of the interaction by Pd-Oqn and Pt-Oqn was also detected using this assay with IC50 values calculated as 2.3 and 12.4 μ M, respectively (Fig. 2c). These results suggested that Pd-Oqn was more effective at inhibiting the interaction of phosphorylated CDCP1 with PKC δ .

Inhibitory effect of Pd-Oqn on the CDCP1-PKC δ pathway. We next tested whether the PKC δ -phosphorylated CDCP1 interaction inhibition effected by Pd-Oqn was also observed in cells. Accordingly, HEK293T cells were transfected with expression plasmids of CDCP1-FLAG and PKC δ -GFP with or without v-Src. Immunoprecipitation of HEK293T cells with an anti-FLAG antibody revealed that PKC δ co-precipitated with CDCP1 in cells expressing both proteins but not in control cells that expressed PKC δ without CDCP1. Furthermore, the amount of PKC δ bound with CDCP1 was increased by v-Src expression owing to phosphorylation of CDCP1 (Fig. 3a, lanes 4 and 6). Co-precipitated PKC δ with CDCP1 was decreased by Pd-Oqn treatment in the presence of v-Src (lanes 6 and 7) but was not significantly affected in the absence of v-Src (lanes 4 and 5).

Following interaction with phosphorylated CDCP, PKCS is transferred to the plasma membrane and activated by SFK phosphorylation.⁽⁶⁾ Thus, as expected, inhibition of the PKCδphosphorylated CDCP1 interaction by Pd-Oqn suppressed tyrosine phosphorylation of PKC\delta (Y311) in a dose-dependent manner whereas phosphorylation of AKT and ERK were not affected (Fig. 3b). We further examined whether Pd-Oqn affected on the interaction of endogenous CDCP1 with PKC\delta. Immunoprecipitation of 44As3 cell lysate with anti-CDCP1 antibody revealed that endogenous PKC δ co-precipitated with CDCP1 was decreased by Pd-Oqn treatment (Fig. 3c). In addition, Pd-Oqn did not affect on the cleavage of CDCP1 although it has been recently reported that the cleaved isoform rather than full-length of CDCP1 activates the downstream signaling.⁽²¹⁾ These results demonstrated that Pd-Oqn-induced disruption of PKC^δ interaction with CDCP1 could suppress the downstream signaling in cells.

Effect of Pd-Oqn and Pt-Oqn on the growth and invasion of cancer cells. Next, we tested whether the addition of Pd-Oqn

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Fig. 5. Inhibitory effect of Pd-Oqn on the invasion of cancer cells. 44As3 cells were subjected to invasion (a, b) and migration (c, d) assays in the presence or absence of Pd-Oqn. Representative photos of invaded (a) or migrated (c) cells are shown as nuclear staining with DAPI. The graph indicates the cell number of invaded (b) or migrated (d) cells upon DMSO or Pd-Oqn treatment as the means \pm SD of three biological replicates. Statistical significance was determined using the Student's *t*-test.

and Pt-Oqn affects anchorage-independent growth of 44As3 cells, comprising diffuse-type gastric adenocarcinoma cells in which CDCP1 is highly phosphorylated, using a colony formation assay in soft agar. The number 44As3 cell colonies in soft agar significantly decreased in the presence of 5 µM Pd-Oqn as compared to that reported for vehicle treatment and was inhibited completely at 25 and 100 µM Pd-Oqn (Fig. 4a,b). Furthermore, both Pd-Oqn and Pt-Oqn could inhibit proliferation of A549, BxPC3, MiaPaCa-2, and 44As3 cells in a dosedependent manner under normal culture conditions with the inhibitory effect of Pd-Oqn being significantly higher than that of Pt-Oqn (Fig. S4A). The inhibitory effect of Pd-Oqn on the proliferation of 44As3 cells was most significant among the cell lines examined, while the proliferation of normal fibroblast NF-37 cells was least affected by Pd-Oqn (Fig. 4c). This difference in sensitivity to Pd-Oqn among cell lines may indicate the differences in expression and phosphorylation levels of CDCP1 in individual cell lines (Fig. S4B).

We also examined whether Pd-Oqn could affect cell motility and invasion using a Boyden chamber-based assay. The addition of Pd-Oqn reduced the number of cells that moved to the bottom area through the membrane coated with Matrigel extracellular matrix but not significantly without Matrigel (Fig. 5). These results indicated that Pd-Oqn inhibited the invasion of 44As3 cells, possibly by degradation of the extracellular matrix, a process also regulated by the CDCP1 pathway as reported.^(5,6)

In vivo effect of Pd-Oqn on tumor metastasis and growth. Finally, we examined the effect of Pd-Oqn on mouse models of peritoneal dissemination of gastric cancer and tumor growth of pancreatic cancer. For the metastatic model, 44As3 cells were injected into the peritoneal cavity of BALB/c-nu/nu mice. PdOqn (15 mg/kg) was injected into the mouse peritoneum six times (Fig. 6a). The weights of the mice were not affected by Pd-Oqn treatment (Fig. S5). After 14 days, the mice were sacrificed and the metastatic nodules on the mesenterium were counted. Notably, the metastatic nodules of the mice were markedly decreased or diminished by Pd-Oqn treatment (Fig. 6b,c).

For the pancreatic tumor model, MiaPaCa-2 cells were transplanted into the pancreas of SCID mice. Pd-Oqn (15 mg/kg) was injected into the mouse peritoneum seven times (Fig. 7a). Pancreatic tumors were dissected from the mice after 23 days and the weights of the pancreatic tumor tissues were determined. The pancreatic tumor weights of the mice treated with Pd-Oqn were slightly but significantly lower than those of the control mice were (Fig. 7b,c). These results indicated that Pd-Oqn suppressed peritoneal dissemination as well as tumor growth *in vivo* through inhibition of the CDCP1-PKCδ signaling pathway.

Discussion

In this study, we first identified small molecules that inhibited the interaction of a phosphorylated site of CDCP1 with the C2 domain of PKC δ . The small molecule, Pd-Oqn, inhibited the growth and invasion of cancer cells *in vitro* and also reduced the peritoneal dissemination of gastric adenocarcinoma and pancreatic adenocarcinoma tumor growth in mouse models. The C2 domain of PKC δ has previously been characterized as a binding domain for tyrosine-phosphorylated sites of CDCP1.⁽⁴⁾ In addition, we have reported that the CDCP1-PKC δ signaling pathway plays important roles in tumor progression such as the anchorage-independent growth and metastasis of cancer cells.^(2,6) Our findings in the current study



Fig. 6. Effect of Pd-Oqn on the peritoneal dissemination of gastric adenocarcinoma. (a) Schedule of the experiment for injection of Pd-Oqn or DMSO into the mouse model of peritoneal dissemination using 44As3 cells. (b) Representative photos of the nodules on the mouse mesenterium. (c) The dot plot indicates the number of nodules in each mouse treated with Pd-Oqn (n = 13) or DMSO (n = 10). The bars in the plot indicate the median value. Statistical significance was determined using the Student's t-test.

further showed that Pd-Oqn could reduce tumor metastasis and growth through inhibition of the recruitment of PKC δ to phosphorylated CDCP1 and SFK-mediated phosphorylation of PKC δ (Fig. S6).

Src family kinases are activated in multiple cancer cells and plays a critical role in cancer progression through the regulation of various substrate proteins and pathways.⁽²²⁻²⁴⁾ Accordingly, SFK inhibitors have been developed and tested in clinical trials.^(25,26) However, SFK is also ubiquitously expressed in normal tissues and involved in normal tissue development and cell differentiation through the conditional activation and phosphorylation of various physiological substrates.⁽²⁷⁻²⁹⁾ Therefore, compounds inhibiting the catalytic activity of SFK might naturally show significant sideeffects.⁽³⁰⁾ Conversely, CDCP1, a substrate of SFK, is predominantly expressed in various cancer cells whereas it exhibits limited expression in normal tissues including hematopoietic and neural stem/progenitor cells^(31,32) although its functional roles in these tissues have not yet been elucidated. Furthermore, CDCP1 deficiency in mouse shows no obvious effects



Fig. 7. Effect of Pd-Oqn on pancreatic cancer of orthotopic xenografts. (a) Schedule of the experiment for the injection of Pd-Oqn or DMSO into the orthotopic xenograft model of pancreatic cancer. (b) Photo of the pancreatic tumor from the dissected mouse. (c) The bar graph indicates the weight of the tumor treated with Pd-Oqn (n = 6) or DMSO (n = 5) as the means \pm SD. Statistical significance was determined using the Student's t-test.

on development or adult tissues as compared to wild-type animals (Nakashima K, Sakai R et al, unpubl. data). These findings suggest that inhibition of the CDCP1 pathway by Pd-Oqn may preferentially affect tumor progression but not normal development and cell differentiation. Consistent with this supposition, the growth inhibitory effect of Pd-Oqn was selectively shown in cell lines such as A549 or 44As3 in which the CDCP1 pathway is activated rather than in fibroblasts and *in vivo* Pd-Oqn administration did not significantly affect general condition of mice at doses shown to be effective against cancers.

The PKC family is implicated in cancer pathology including proliferation, metastasis, and angiogenesis.^(33,34) PKC δ in particular plays a critical role in proapoptotic events through its phosphorylation and organelle translocation including to the nucleus and mitochondria although it also can act as an antiapoptotic protein.^(35,36) The function of PKC δ depends on various factors including its localization, phosphorylation, and the presence of signaling molecules.⁽³⁶⁾ However, although PKC inhibitors have been subjected to clinical trials for cancer therapy, they have mostly failed owing to lack of efficacy on tumors and severe damage on normal tissues.⁽³⁷⁾ Alternatively, as previously stated, PKC δ localization and activation on cell

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membranes occurs through interaction with phosphorylated CDCP1 and may contribute to the induction of signaling for cancer progression. Thus, Pd-Oqn may selectively act as a suppressor for cancer progression that is triggered by PKC δ association with CDCP1.

In the current study, we found that Pd-Oqn inhibited the invasion but not the migration of 44As3 cells in a chamber assay. We have previously found that CDCP1 regulates the activation of matrix metallopeptidase (MMP)2 and secretion of MMP9.⁽³⁸⁾ In addition, CDCP1 is an essential regulator of the trafficking and function of MT1-MMP and the invadopodia-mediated invasion of cancer cells.⁽⁵⁾ Here, we showed that in a mouse model, Pd-Oqn reduced the peritoneal dissemination of gastric adenocarcinoma. These results suggested that Pd-Oqn inhibited the invasion and metastasis of cancer cells through suppression of the extracellular matrix degradation that is promoted by the CDCP1 pathway.

Pd-Oqn was found to clearly block the interaction of phosphorylated CDCP1 with PKC δ and inhibit PKC δ phosphorylation. Pd-Oqn is classified under the same category as metal complex compounds such as cisplatin, which induce DNA damage. Pd-Oqn did not, however, induce the phosphorylation of histone H2AX (γ -H2AX), a marker of DNA damage response (Fig. S7). Although Pd-Oqn notably inhibited the interaction of the C2 domain with the 762pY peptide, the mechanism of Pd-Oqn action on the complex remains to be clarified.

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Overall, the results of our study indicate that the novel compound Pd-Oqn reduces tumor metastasis and growth by inhibiting the association between CDCP1 and PKC δ , thus potentially representing a promising candidate among therapeutic reagents targeting protein—protein interaction. Furthermore, whereas previous attempts to identify compounds inhibiting specific protein—protein interactions have not generally been very successful, the application of metal-containing compounds such as Pd-Oqn might provide a means toward overcoming this limitation.

Disclosure Statement

The authors have no conflict of interest.

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Supporting Information

Additional Supporting Information may be found online in the supporting information tab for this article:

Doc. S1. Materials and Methods.

Fig. S1. Chemical Screening.

Fig. S2. Structures of the inhibitory compounds for the C2-CDCP1 interaction.

Fig. S3. HTRF assay.

Fig. S4. Effect of the compounds on proliferation of cancer cells.

Fig. S5. Body weight of the mice treated with Pd-Oqn.

Fig. S6. Model for molecular action of Pd-Oqn.

Fig. S7. No DNA damage response by Pd-Oqn.

Table S1. Peptide sequence.