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G $\beta\gamma$ -mediated activation of protein kinase D exhibits subunit specificity and requires G $\beta\gamma$ -responsive phospholipase C β isoforms

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Abstract

Background: Protein kinase D (PKD) constitutes a novel family of serine/threonine protein kinases implicated in fundamental biological activities including cell proliferation, survival, migration, and immune responses. Activation of PKD in these cellular activities has been linked to many extracellular signals acting through antigen receptor engagement, receptor tyrosine kinases, as well as G protein-coupled receptors. In the latter case, it is generally believed that the G α subunits of the G $_q$ family are highly effective in mediating PKD activation, whereas little is known with regard to the ability of G $\beta\gamma$ dimers and other G α subunits to stimulate PKD. It has been suggested that the interaction between G $\beta\gamma$ and the PH domain of PKD, or the G $\beta\gamma$ -induced PLC β /PKC activity is critical for the induction of PKD activation. However, the relative contribution of these two apparently independent events to G $\beta\gamma$ -mediated PKD activation has yet to be addressed.

Results: In this report, we demonstrate that among various members in the four G protein families, only the G α subunits of the G $_q$ family effectively activate all the three PKD isoforms (PKD1/2/3), while G α subunits of other G protein families (G $_s$, G $_i$, and G $_{12}$) are ineffective. Though the G α subunits of G $_i$ family are unable to stimulate PKD, receptors linked to G $_i$ proteins are capable of triggering PKD activation in cell lines endogenously expressing (HeLa cells and Jurkat T-cells) or exogenously transfected with (HEK293 cells) G $\beta\gamma$ -sensitive PLC $\beta_{2/3}$ isoforms. This indicates that the G $_i$ -mediated PKD activation is dependent on the released G $\beta\gamma$ dimers upon stimulation. Further investigation on individual G $\beta\gamma$ combinations (i.e. G β_1 with G γ_{1-13}) revealed that, even if they can stimulate the PLC β activity in a comparable manner, only those G $\beta_1\gamma$ dimers with γ_2 , γ_3 , γ_4 , γ_5 , γ_7 , and γ_{10} can serve as effective activators of PKD. We also demonstrated that G $_i$ -mediated PKD activation is essential for the SDF-1 α -induced chemotaxis on Jurkat T-cells.

Conclusions: Our current report illustrates that G $\beta\gamma$ dimers from the G $_i$ proteins may activate PKD in a PLC $\beta_{2/3}$ -dependent manner, and the specific identities of G γ components within G $\beta\gamma$ dimers may determine this stimulatory action.

Keywords: G proteins, G α subunits, G $\beta\gamma$ dimers, PLC β , PKD

Background

Protein kinase D (PKD) constitutes a novel family of diacylglycerol (DAG)-responsive serine/threonine protein kinases with different structural, enzymological and regulatory properties from the protein kinase C (PKC)

family members [1-3]. To date, three members of the PKD family have been identified: human PKD1 (its mouse ortholog being PKC μ), and the more recently identified PKD2 and PKD3 (also named PKC ν), among which PKD1 is the most extensively characterized isoform. Emerging studies have revealed that PKDs are implicated in a complex array of fundamental biological activities, including cell survival [4], migration [5], proliferation [6], and immune responses [7]. In addition, growing evidence links PKDs to signal transduction pathways in tumor development and cancer progression.

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In many cases, specific PKD isoforms are dysregulated in cancer cells [8].

All PKDs share a common modular structure, with a tandem repeat of zinc finger-like cysteine-rich motifs at their NH₂ termini that display high affinity for DAG or phorbol ester, a pleckstrin homology domain (PH domain) for negative regulation of kinase activity [9], and a C-terminal catalytic domain containing transphosphorylation and autophosphorylation sites. Activation of PKD isoforms is generally attributed to phosphorylation at a pair of highly conserved serine residues (for human: Ser⁷³⁸ and Ser⁷⁴² in PKD1; Ser⁷⁰⁶ and Ser⁷¹⁰ in PKD2; Ser⁷³¹ and Ser⁷³⁵ in PKD3) in the activation loop of the kinase domain by PKC. As PKC can be activated by many extracellular signals, stimulation of PKD isoforms has been demonstrated by antigen receptor engagement [10], stimulation of receptor tyrosine kinases (RTKs) such as platelet-derived growth factors (PDGF) receptors [11] and vascular endothelial growth factor (VEGF) receptors [12], as well as activation of various G protein-coupled receptors (GPCRs). Among the large GPCR family, receptors with preferential coupling to G_q, including those responsive to bombesin, vasopressin, endothelin, bradykinin [13], cholecystokinin [14], tachykinin [15] and angiotensin II have been demonstrated to activate PKD in a variety of cell types. Other G protein members like G₁₂ and G₁₃ have also been proposed to activate PKD3 in a PKC- and Rac-dependent manner [16]. In addition, it has been reported that G_q, G_i and G_{12/13} may cooperate in LPA-induced PKD activation [17], but the relative contribution of specific G protein subunits (e.g. G_{αi} versus G_{βγ}) to PKD activation remains undefined.

The functional specificity of G proteins was originally accredited to the G_α subunits, with the G_{βγ} dimers being viewed as negative regulators of G protein signaling. Yet, there is growing evidence that G_{βγ} dimers can also act as active mediators in signal transduction, thus conferring an additional level of signal specificity [18-20]. The G_β identity in the G_{βγ} dimer imparts selectivity on its interaction with effectors like phospholipase C_β [21], as well as in the regulation of neutrophil function [22]. Moreover, since the G_γ component is structurally and expression-wise diverse, it imposes additional complexity in signal transduction. For instance, only certain G_{βγ} combinations (mainly those containing γ₂, γ₄, γ₇ or γ₉) are linked to significant STAT3 activation [23]. Functional selectivity of G_γ subunits has also been reported [24-26]; deletion of the *Gng3* gene leads to increased susceptibility to seizures in mice with significant reductions in G_{β2} and G_{α13} subunit levels in certain brain regions [25], whereas knock-out of the *Gng7* gene is associated with reductions in the G_{αolf} subunit content and adenylyl cyclase activity of the murine striatum [24].

These observations demonstrate that members of the G_γ subunit family are not functionally interchangeable.

It has been suggested that the interaction between G_{βγ} and the PH domain of PKD [27], or the G_{βγ}-induced PLC_β/PKC activity is critical for the induction of PKD activation [28]. However, the relative contribution of these two apparently independent events to G_{βγ}-mediated PKD activation has yet to be addressed. Recently, G_{βγ} combinations containing G_{γ2} (i.e. G_{β1γ2} and G_{β3γ2}) have been shown to be effective activators for PKD [29], but the relevant capabilities of other G_{βγ} dimers remain unclear.

In this report, we demonstrated that all family members of the G_q subfamily (G_q, G₁₁, G₁₄, and G₁₆) can induce PKD1, PKD2 and PKD3 activation. G_s cannot elicit a PKD response, whereas G_i members may induce PKD activation in a G_{βγ}-dependent manner. For the G_{βγ}-induced PKD activation, even in the presence of PLC_{β2} or PLC_{β3}, only certain G_{βγ} dimer combinations are capable of activating the kinase effectively. Moreover, we showed that this selective G_{βγ} dimer-mediated PKD activation is accompanied by enhanced interaction between the two components when PLC_{β2/3} is present.

Materials and methods

Materials

HEK293 and Jurkat T-cells were obtained from American Type Culture Collection (Rockville, MD). Pertussis toxin (PTX) was purchased from List Biological Laboratories (Campbell, CA). Cell culture reagents including Dulbecco's phosphate-buffered saline (PBS), trypsin, fetal bovine serum (FBS), penicillin-streptomycin mixture, RPMI 1640 medium, minimum essential medium (MEM), Dulbecco's modified Eagle's medium (DMEM) and Lipofectamine PLUSTM were obtained from Invitrogen (Carlsbad, CA). The cDNAs encoding PLC_{β1}, PLC_{β2} and PLC_{β3} were obtained from Dr. Richard Ye (University of Illinois at Chicago). Flag-tagged human G_{β1} and G_{β2}, HA-tagged human G_{γ1}, G_{γ2}, G_{γ3}, G_{γ4}, G_{γ5}, G_{γ7}, G_{γ8}, G_{γ9}, G_{γ10}, G_{γ11}, G_{γ12} and G_{γ13} cDNA constructs were obtained from UMR cDNA Resource Center (Rolla, MO). Antiserum including anti-Flag and anti-HA were purchased from Roche Molecular Biochemicals (Indianapolis, IN). Cell culture reagents including Lipofectamine PlusTM were obtained from Invitrogen (Carlsbad, CA). Myo-[³H] inositol was purchased from DuPont NEN (Boston, MA). M2 affinity gels and protein A-agarose were obtained from Sigma (St. Louis, MO). HA-PKD1 and FLAG-PKD2 constructs were gifts from Dr. J. Van Lint (Katholieke Universiteit Leuven, Belgium), and Myc-PKD3 constructs were kindly provided by Dr. Q. J. Wang (University of Pittsburgh, PA).

Cell culture and transfection

HEK293 cells were cultured in MEM supplemented with 10% (v/v) FBS, 50 units/ml penicillin, and 50 µg/ml streptomycin. Jurkat T-cells were cultured in RPMI1640 containing 10% (v/v) FBS. For PLC assays and co-immunoprecipitation assays, HEK293 cells were seeded at 60% confluency into 12-well plates or 6-well plates, respectively. Transfection was performed on the following day using Lipofectamine PLUSTM reagents. For the establishment of stable cell lines (293/BK₂R, 293/β₂AR and 293/fMLPR), exponentially growing HEK293 cells were transfected with cDNA of BK₂R, β₂AR or fMLPR in pcDNA3.1-zeo using Lipofectamine PLUSTM. The cells were then selected with Zeocin (50 µg/mL). 293/fMLPR-Gα₁₆ cells were established by transient transfection of 293/fMLPR stable cell lines with Gα₁₆ in pcDNA3.

In vitro PKD Assay

Twenty-four hours after transfection, HEK293 cells were serum-starved overnight and then treated with 500 µl (per well) of ice-cold detergent-containing lysis buffer (50 mM Tris-HCl, pH 7.5, 100 mM NaCl, 5 mM EDTA, 40 mM Na₄P₂O₇, 1% Triton X-100, 1 mM dithiothreitol, 200 µM Na₃VO₄, 100 µM phenylmethylsulfonyl fluoride, 2 µg/ml leupeptin, 4 µg/ml aprotinin, and 0.7 µg/ml pepstatin). Lysates obtained were subjected to *in vitro* PKD kinase assay. Fifty µl of each supernatant was used for the detection of PKD isoform expression and stimulatory phosphorylation, and the remaining lysate (450 µl) was incubated overnight at 4°C with specific affinity gels to immune-precipitate the corresponding PKD isoform (anti-HA for HA-PKD1; anti-FLAG for FLAG-PKD2; and anti-Myc for Myc-PKD3). The resulting immunoprecipitates were washed twice with lysis buffer and twice with kinase assay buffer (30 mM Tris-HCl, pH 7.4, 10 mM MgCl, and 1 mM DTT). Washed immunoprecipitates were resuspended in 40 µl of kinase assay buffer containing 2.5 mg/ml of Syntide-2 (PLARTLSVAGLPGKK), and the kinase reactions were initiated by the addition of 10 µl of ATP buffer containing 1 µCi of [γ-³²P]-ATP per sample. After 10-min incubation at 30°C with occasional shaking, the reactions were terminated by adding 100 µl of 75 mM H₃PO₄ and spotting 75 µl of the reaction mix onto P-81 phosphocellulose paper. Free [γ-³²P]-ATP was separated from the labelled substrate by washing the P-81 paper four times (5 min each) in 75 mM H₃PO₄. The papers were dried and the radioactivity incorporated into Syntide-2 was determined by scintillation counting.

Electroporation

The knock down of PKD1, PKD2 and PKD3 was performed by introducing the corresponding PKD isoform-specific siRNA from Invitrogen (Carlsbad, CA,

USA) using Nucleofector[®] Kit V from Lonza (Basel, Switzerland). Briefly, 1×10⁶ cells per sample were resuspended in Nucleofector[®] Solution and supplement provided at room temperature. siRNA against PKD1, PKD2 or PKD3 (200 pmol each) was added to the samples and then electroporated using the Nucleofector[®]. Electroporated cells were then incubated at room temperature for 10 min before transferring them into the 12-well plate with culture medium. The knock down of PLCβ₁, PLCβ₂ and PLCβ₃ was performed in similar manner, with the corresponding isoform-specific siRNA obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

Western blotting analysis

Cells in 12-well plate were lysed in 300 µl of ice-cold lysis buffer (50 mM Tris-HCl, pH 7.5, 100 mM NaCl, 5 mM EDTA, 40 mM NaP₂O₇, 1% Triton X-100, 1 mM dithiothreitol, 200 µM Na₃VO₄, 100 µM phenylmethylsulfonyl fluoride, 2 µg/ml leupeptin, 4 µg/ml aprotinin and 0.7 µg/ml pepstatin). Clarified lysates were resolved on 1 µ2% SDS-polyacrylamide gels and then transferred to nitrocellulose membranes (Westborough, MA). Stimulatory phosphorylation of PKD1, PKD2, ERK and CREB were detected by their corresponding antisera and horseradish peroxidase-conjugated secondary antisera. The immunoblots were visualized by chemiluminescence with the ECL kit (Amersham Biosciences). Antibodies sources are as follows: anti-phospho-PKD1-Ser^{744/748}, anti-phospho-PKD1-Ser⁹¹⁶ (also recognize human PKD1-Ser^{738/742} and Ser⁹¹⁰, respectively), anti-phospho-ERK-Thr²⁰²/Tyr²⁰⁴, anti-PKD1 were obtained from Cell Signaling Technology (Beverly, MA). Anti-phospho-PKD2-Ser⁸⁷⁶ and anti-PKD2 were purchased from R & D Systems (Minneapolis, MN). Anti-PKD3 was obtained from Bethyl Laboratories (Montgomery, TX).

Measurement of intracellular Ca²⁺ transient by FLIPR[®]

Jurkat T-cells were serum-starved overnight in the absence or presence of PTX (10 ng/ml) and then washed with Hank's balanced salt solution (HBSS). Washed cells (1×10⁶ cells/ml) were preloaded with Fluo-4 (AM) followed by incubation at 37°C for 1 h. These labeled cells were then transferred to a black-walled and clear-bottomed 96-well plate (1×10⁵ cells/well) placed in the Fluorometric Imaging Plate Reader (FLIPR), and 50 µl of HBSS (with or without agonists) was added to each well. The resulting fluorescent signals that reflect the intracellular Ca²⁺ transients were monitored by an excitation wavelength of 488 nm and detection with the emission wavelength from 510 to 570 nm.

Co-immunoprecipitation assay

Transfected cells were lysed in the lysis buffer as described before. Cell lysates were centrifuged (12000 g, 4°C, 5 min) to remove cellular debris. Lysates were incubated at 4°C overnight with M2 affinity gels (20 µl/ sample) for the binding with Flag-tagged Gβ subunits. The resulting immunoprecipitates were collected by centrifugation at 1,000 g, 4°C, for 3 min and then washed three times with 500 µl lysis buffer. Bound proteins were eluted by 50 µl of lysis buffer and 10 µl of 6× SDS-containing sample buffer, and boiled for 5 min prior to separation by 12% SDS-polyacrylamide gel electrophoresis (PAGE). Flag-tagged Gβ, HA-tagged Gγ subunits, PLCβ₂ and PKD1 in the immunoprecipitates were detected by their corresponding antisera followed with horseradish peroxidase-conjugated secondary antisera in Western blotting analysis.

Chemotactic assay

The chemotactic ability of Jurkat T cells was evaluated using transwell plates (Costar, Cambridge, MA) with polycarbonate inserts with 5-µm pores (Costar 3421). Lower chambers were loaded with 600 µl of migration media alone or containing SDF-1α at the concentration of 100 nM. Cells (0.1 ml) at 1×10^6 /ml were added to the top chamber of a 24-well transwell (6.5-µm diameter, 5-µm pore size) and incubated for 4 h at 37°C. The cells which passed through the membranes and migrated to the lower chambers were quantified under microscopy.

Statistics

The values shown in each figure represent mean ± SEM from at least three individual experiments. Statistical analyses were performed by ANOVA, followed by the Bonferroni's post test. Differences with a value of $P < 0.05$ were considered statistically significant.

Results

Previous studies on Gα subunit-induced activation of PKD isoforms were primarily performed on the PKD1 prototype with Gα_q [30,31], leaving the activation profile of the PKD family rather incomplete. Most of these studies employed aluminum tetrafluoride (AlF₄⁻) to elicit G protein-mediated activation of PKD. Although AlF₄⁻ can selectively stimulate heterotrimeric G proteins over monomeric GTPases [32,33], AlF₄⁻ activates multiple heterotrimeric G proteins simultaneously and thus cannot be used to identify the specific G proteins involved in the activation of PKD. On the basis of these considerations, we aimed to firstly define the role of different Gα subunits in promoting the activation of all three PKD isoforms. We performed screening on Gα subunit-mediated PKD1 phosphorylation. HEK293 cells were transfected with wild-type (WT) or constitutively active

(RC/QL) Gα subunits (Gα_q, Gα₁₁, Gα₁₄, Gα₁₆, Gα₁₂, Gα₁₃, Gα₁₁, Gα₁₂, Gα₁₃, Gα_z and Gα_s) and then assayed for PKD phosphorylation by phospho-PKD-specific antibodies. HEK293 cells have previously been shown to express all three PKD isoforms [34].

The phosphorylation of a pair of highly conserved serine residues in the activation loop (Ser⁷³⁸ and Ser⁷⁴² in PKD1; Ser⁷⁰⁶ and Ser⁷¹⁰ in PKD2; Ser⁷³¹ and Ser⁷³⁵ in PKD3) plays a crucial role in human PKD activity [35]. Some early studies on PKD targeted the autophosphorylation sites (Ser⁹¹⁶ in PKD1 and Ser⁸⁷⁶ in PKD2) as surrogate markers of mouse PKD activity, though a recent report has demonstrated that this site is not required for activation [36]. Therefore, anti-phospho-PKD1 Ser^{744/748} and Ser⁹¹⁶ antibodies (also recognize human PKD1 at Ser^{738/742} and Ser⁹¹⁰, respectively) were both adopted for the evaluation of PKD1 activation. As shown in Figure 1, expression of WT Gα subunits did not induce significant PKD1 phosphorylation as compared to the vector control, although expression of Gα₁₁ or Gα₁₄ slightly enhanced the basal PKD phosphorylation. Conversely, prominent phosphorylation of PKD1 was observed in cells expressing one of the constitutively active mutants from the Gα_q subfamily (Gα_q, Gα₁₁, Gα₁₄, or Gα₁₆). Western blot analysis verified that the expression levels of PKD1 were similar and that both WT and constitutively active Gα subunits were expressed at comparable levels (Figure 1). In contrast, there was no detectable phosphorylation of PKD1 by constitutively active mutants from G_i, G_s, or G₁₂ subfamilies (Figure 1). This is consistent with earlier studies demonstrating that the constitutively active mutants of Gα₁₂ and Gα₁₃ did not induce PKD activation in COS-7 cells [30].

To examine whether Gα subunits from the G_q subfamily are all capable of inducing activation of all three isoforms of PKD, HEK293/HA-PKD1, HEK293/FLAG-PKD2 and HEK293/Myc-PKD3 stable cell lines were established and then transiently transfected with WT or the RC/QL mutants of Gα subunits (Gα_q, Gα₁₁, Gα₁₄, Gα₁₆), followed by *in vitro* [³²P]-kinase assays using syntide-2 as an exogenous substrate for PKD. As shown in Figure 2A, PKD isoforms isolated from all three stable cell lines transfected with vector control or plasmids encoding the WT Gα subunits exhibited low catalytic activity. In contrast, those immunoprecipitated from stable cell lines overexpressing a constitutively active mutant displayed marked increase in PKD kinase activity. Comparable expressions of Gα subunits and PKD isoforms in the various transfectants were confirmed by Western blot analyses (Figure 2B). We also examined the phosphorylation of specific PKD isoforms in the same samples. Since anti-phospho-PKD1^{738/742} exhibits some cross-reactivity with PKD2 and PKD3, anti-phospho-PKD1⁹¹⁰ was also employed to detect PKD1

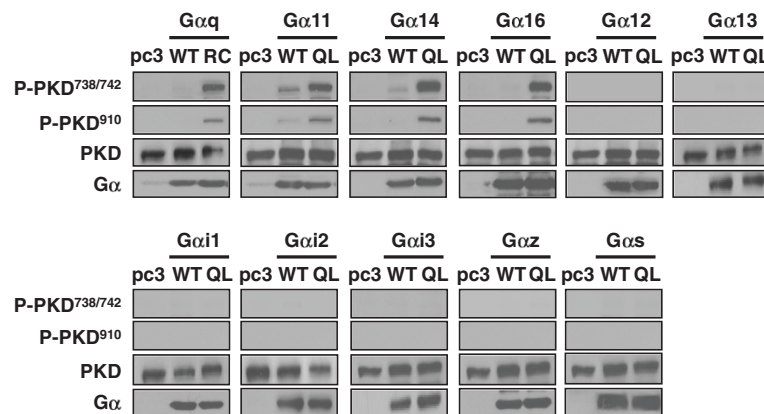


Figure 1 Constitutively active mutants of G_q family members stimulate PKD1 phosphorylation. HEK293 cells were transfected with pcDNA3, wild type (WT) or constitutively active (RC or QL) G_a subunits of G_q , G_{12} , G_i and G_s families. Transfectants were then lysed, and proteins were subjected to SDS-PAGE and immunoblotting with antibodies against phospho-PKD1, PKD1, and specific G_a subunits.

phosphorylation. Likewise, anti-phospho-PKD2⁸⁷⁶ was used for PKD2. As PKD3 lacks the phosphorylation site equivalent to phospho-PKD1⁹¹⁰, only the phosphorylation at PKD3^{731/735} was monitored. In agreement with the results from the *in vitro* kinase assay, stimulatory PKD phosphorylation for all three PKD isoforms was enhanced in the presence of constitutively active G_a mutants from the G_q subfamily (Figure 2B). Unlike members of the G_q subfamily, constitutively active $G_{\alpha_{i1}}$ failed to stimulate the kinase activity of all three forms of PKD (Figure 2A) or elevate their level of phosphorylation (Figure 2B). Similar results were obtained with other members of the G_i ($G_{\alpha_{i2}}$, $G_{\alpha_{i3}}$, $G_{\alpha_{oA}}$, $G_{\alpha_{z}}$, $G_{\alpha_{t1}}$, and $G_{\alpha_{t2}}$), G_s ($G_{\alpha_{sL}}$ and $G_{\alpha_{oIf}}$) and G_{12} (G_{12} and G_{13}) families (Additional file 1: Figure S1 and Additional file 2: Figure S2). Collectively, these results demonstrated that PKD1, PKD2 and PKD3 can be specifically activated by the constitutively active G_a subunits from the G_q family, but not by those of G_i , G_s or G_{12} families.

The preceding experiments suggest that the G_a subunits from the G_q family contribute to elevated PKD phosphorylation. To examine in more detail the stimulation of PKD by G protein signaling, we tested different G_q -, G_s - and G_i -coupled receptors for their ability to activate PKD1 in HEK 293 cells. HEK293 cells were transfected with the G_q -coupled bradykinin BK₂ receptor (Figure 3A), G_s -coupled β_2 -adrenergic receptor (Figure 3B) or G_i -coupled fMLP receptor (Figure 3C), and the transfectants subsequently examined for agonist-induced PKD1 activation. Phosphorylation of CREB or ERK was simultaneously monitored as positive controls of G_s - and G_i -signaling, respectively. In line with the data in Figures 1 and 2, only bradykinin (which stimulates the G_q -coupled BK₂ receptor) rapidly and potently stimulated PKD1 phosphorylation (Figure 3A), while isoproterenol and fMLP failed to induce any detectable

PKD activation despite obvious phosphorylation of CREB or ERK (Figure 3B and C). Since many G_i -coupled receptors including the fMLP receptor are capable of interacting with $G_{\alpha_{16}}$ [37], it is expected that co-expression of $G_{\alpha_{16}}$ would turn on G_q -related signals, thus allowing effective stimulation of PKD1 phosphorylation. As illustrated in Figure 3D, prominent fMLP-induced PKD1 phosphorylations at both Ser^{738/742} and Ser⁹¹⁰ were observed in HEK293 cells co-expressing the G_i -coupled fMLP receptor and $G_{\alpha_{16}}$ (Figure 3D); the fMLP-induced response was readily detected by 2 min and was maintained up to 30 min. These results further confirmed the specificity of G_{α_q} -mediated PKD activation and implied that many GPCRs are capable of regulating the function of PKD through members of the G_q subfamily. This may have particular relevance to hematopoietic cells since the promiscuous $G_{\alpha_{16}}$ and $G_{\alpha_{14}}$ are mainly expressed in immune cells and are capable of recognizing a large number of GPCRs [38,39].

Next, we investigated whether PKD phosphorylation can be induced upon activation of G_q -coupled receptors that are endogenously expressed in HeLa cells. Serum starved HeLa cells were treated with various agonists targeting G_q -, G_i - and G_s -coupled receptors for various durations, and PKD1 phosphorylation was determined by Western blot analysis. As expected, bradykinin and histamine acting on G_q -coupled receptors effectively induced a marked increase in PKD phosphorylation at the activation loop (Figure 4A). Agonists that act on G_s -coupled β -adrenergic receptor (isoproterenol) and GLP receptor (glucagon-like peptide) failed to activate PKD, even when stimulatory phosphorylation of ERK was clearly detected (Figure 4B). Unexpectedly, stimulation of G_i -coupled α_2 -adrenergic receptor (by UK14304) and CXCR₄ receptor (by SDF-1 α) led to observable PKD activation. This is in contrast to the result presented in

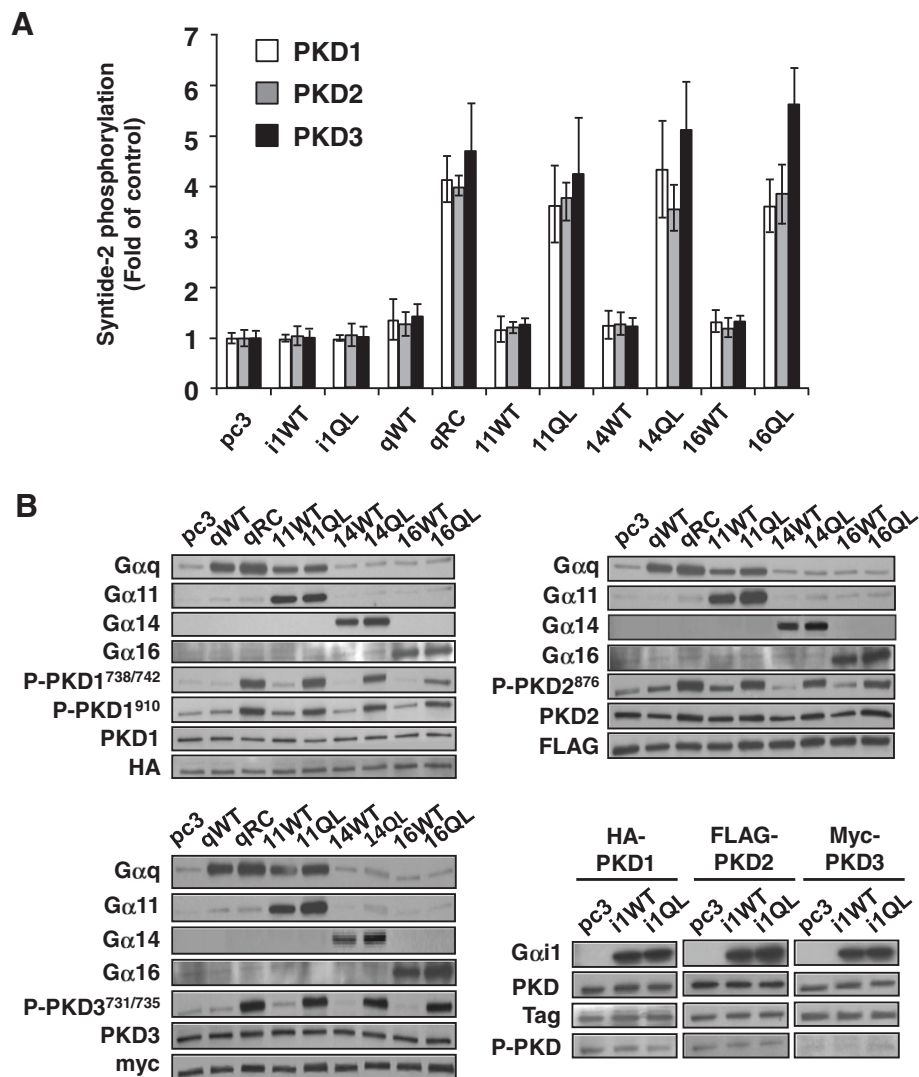


Figure 2 All Gα subunits from the G_q subfamily can activate PKD1, PKD2 and PKD3. (A) Stably expressed and epitope-tagged PKD isoforms were immunoprecipitated from HEK293 cells transiently co-transfected with Gα subunits, and syntide-2 phosphorylation assays were carried out as described under "Materials and Methods". Results are the average of at least three independent experiments, and presented as fold of control (±S.E.M.). (B) Cell lysates from HEK293 transfectants as described in (A) were subject to Western blot analysis using antibodies against specific Gα subunits (except anti-Gα_q which cross reacts with Gα₁₁), phosphorylated PKD isoforms, PKD1, 2, 3 as well as their corresponding tag. Similar results were obtained in three separate experiments.

Figure 3C where stimulation of the G_i-coupled fMLP receptor in HEK293 cells failed to promote PKD activation.

The ability of G_i-coupled receptors to stimulate PKD phosphorylation in HeLa cells was contrary to the results obtained with either Gα_iQL (Figures 1 and 2) or the G_i-coupled fMLP receptor in HEK293 cells (Figure 3C). Given that Gα_q-induced activation of PKD is known to be mediated via PLCβ/PKC [30], and that Gα_i apparently could not activate PKD, we hypothesized that PKD activation by the G_i-coupled receptors in HeLa cells was mediated by the Gβγ subunits, presumably via Gβγ-sensitive PLCβ₂ or PLCβ₃. Gβγ-induced activation of

PKD in HeLa cells have indeed been reported [27]. To test this hypothesis, we first examined the endogenous expression of PLCβ₂ and PLCβ₃ in both HEK293 and HeLa cells. Western blot analysis revealed that HEK293 cells expressed barely detectable levels of PLCβ₂ and PLCβ₃, whereas PLCβ₃ (but not PLCβ₂) was abundantly expressed in HeLa cells (Figure 5A).

To determine the importance of Gβγ-sensitive PLCβ_{2/3} in Gβγ-mediated PKD activation, HEK293/Gγ₂ stable cells were transiently transfected with FLAG-Gβ₁₋₂ in the absence or presence of PLCβ_{2/3}. Because consistent expression of Gγ subunits (~6 kDa) is more difficult to achieve in transient transfections, HEK293 cells stably

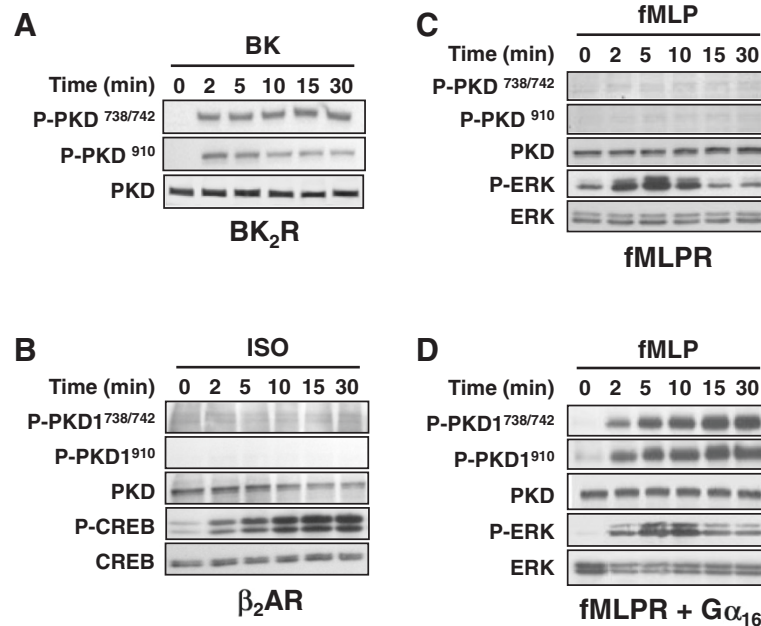


Figure 3 G_q, but not G_s or G_i signaling is linked to stimulatory phosphorylation of PKD in HEK293 cells. HEK293 cells were stably transfected with BK₂R (A), β₂AR (B) or fMLPR, in the absence (C) or presence of Gα₁₆ (D). Transfectants were serum starved for 4 h prior to treatment with 100 nM bradykinin (BK), 10 μM isoproterenol (ISO) or 300 nM N-formyl-methionyl-leucyl-phenylalanine (fMLP) for the indicated durations. Cell lysates were resolved in SDS-PAGE, and the presence of ERK, PKD and CREB phosphorylation was detected by their respective anti-phospho or anti-total antisera. Activation of PKD was observed only for BK₂R and fMLPR/Gα₁₆ stable cells. CREB activation served as a positive control for β₂AR stable cells.

expressing G_γ₂ were employed in these assays. As expected, co-expression of various combinations of Gβ_γ alone did not induce any stimulatory phosphorylation as compared to the vector control in HEK293 cells (Additional file 3: Figure S3A). Upon co-expression with PLCβ₃, however, both Gβ₁γ₂ and Gβ₂γ₂ markedly enhanced the level of PKD phosphorylation; the expression of PLCβ₃ alone had no significant effect on PKD phosphorylation (Figure 5B). Likewise, co-expression of Gβ₁γ₂ or Gβ₂γ₂ with PLCβ₂ induced significant PKD phosphorylation (Figure 5C). These results not only suggest the crucial role of PLCβ_{2/3} in Gβ_γ-mediated PKD activation, but also help to explain the differences in G_i-mediated PKD phosphorylation in HEK293 and HeLa cells.

Since the G_γ subunit identity has been shown to affect signaling specificity [24–26], we determined whether other Gβ₁γ dimer combinations can effectively induce PKD1 activity in the presence of PLCβ_{2/3}. Hence, HEK293 cells were transfected with pcDNA3 (vector control) and one of the twelve combinations of Gβ₁γ_x dimer, with or without PLCβ₂. As shown in Figure 5D (lower panel), transfection of Gβ_γ dimers alone did not significantly enhance the phosphorylation of PKD1 beyond the vector control. Among all of the Gβ₁γ_x combinations tested, Gβ₁γ₂, Gβ₁γ₃, Gβ₁γ₄, Gβ₁γ₅, Gβ₁γ₇ and Gβ₁γ₁₀ consistently triggered strong and significant

PKD1 phosphorylation upon co-expression with PLCβ₂, however, there was no significant change in PKD1 phosphorylation in other Gβ₁γ_x/PLCβ₂-overexpressing cells (Figure 5D, lower panel). Comparable expressions of all Gβ₁γ_x combinations and PLCβ₂ were detected in the transfectants (data not shown), resulting in elevated levels of IP₃ formation (Figure 5D, upper panel) as reported previously [21]. We also tested whether selected Gβ₁γ_x/PLCβ₂ combinations can induce *in vitro* kinase activity of the different PKD isoforms (PKD1–3). In agreement with the Gβ₁γ_x/PLCβ₂-induced PKD1 phosphorylation profile, Gβ₁γ₂/PLCβ₂ and Gβ₁γ₇/PLCβ₂ induced significant PKD kinase activity with all three PKD isoforms, while Gβ₁γ₉/PLCβ₂ failed to do so (Additional file 3: Figure S3B). Similar Gβ₁γ_x-mediated PKD activation profile was obtained with PLCβ₃ (data not shown). As expected, Gβ₁γ_x failed to induce PKD phosphorylation with PLCβ₁ which is insensitive to Gβ_γ (Additional file 3: Figure S3C).

Having demonstrated that certain Gβ₁γ_x/PLCβ_{2/3} combinations were more effective in triggering PKD activity in HEK293 cells, we asked if similar Gβ₁γ_x selectivity for PKD phosphorylation could be observed in HeLa cells that endogenously express high level of Gβ_γ-sensitive PLCβ₃ (Figure 5A). Due to the relatively low levels of endogenously expressed PKD1 [34], HeLa cells were transiently co-transfected with cDNAs encoding

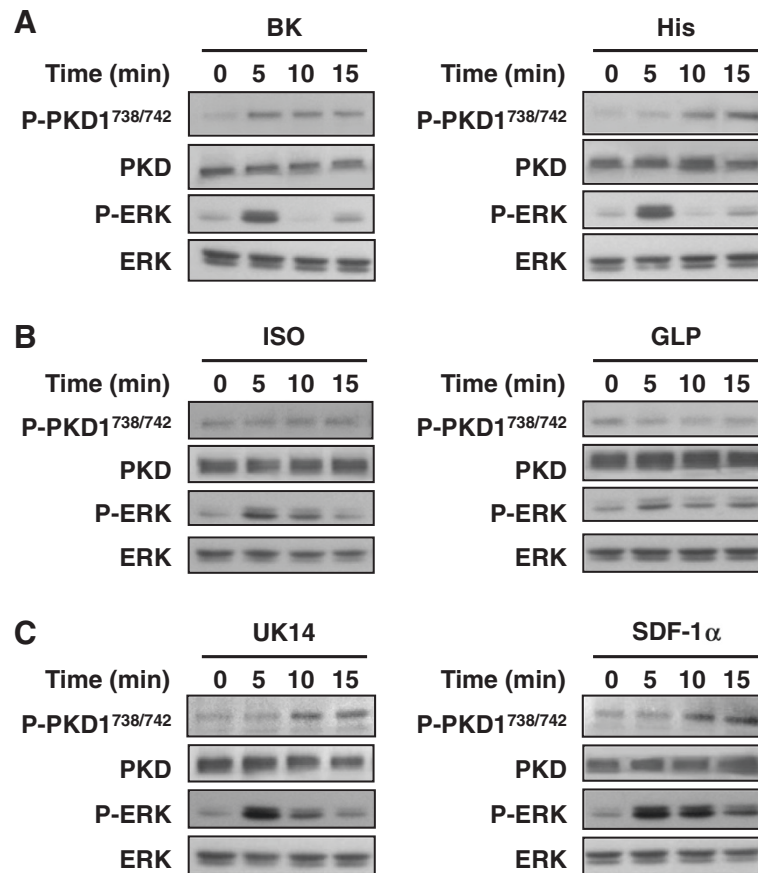


Figure 4 Stimulation of Gq/Gi-coupled receptors activates PKD1 in HeLa cells. HeLa cells were serum starved, followed by treatment with (A) 100 nM bradykinin (BK) or 10 μ M histamine (His); (B) 10 μ M isoproterenol (ISO) or 100 nM glucagon-like peptide (GLP); (C) 10 μ M UK14304 (UK14) or 100 nM stromal cell-derived factor-1 (SDF-1) for the indicated times. Cell lysates were detected for phospho-PKD, PKD, phospho-ERK and ERK with their respective anti-sera.

PKD1 and G $\beta_1\gamma_2$, G $\beta_1\gamma_7$ or G $\beta_1\gamma_9$, followed by serum starvation and subsequent immuno-detection of stimulatory phosphorylated PKD. The results obtained with endogenous PLC β_3 -expressing HeLa cells (Figure 5E) were essentially similar to those obtained from the PLC $\beta_{2/3}$ -transfected HEK293 cellular background (Figure 5D, lower panel). This further indicates that the identity of the G γ subunit may confer specificity to G $\beta\gamma$ -mediated PKD phosphorylation.

It has previously been suggested that G $\beta\gamma$ activates PKD through direct interaction at its PH domain [27]. However, overexpression of G $\beta\gamma$ dimers failed to stimulate PKD phosphorylation in HEK293 cells (Figures 5B-D and Additional file 3: Figure S3A-B) unless G $\beta\gamma$ -responsive PLC $\beta_{2/3}$ was co-expressed (Figures 5D and Additional file 3: Figure S3B-C). Despite the fact that all of the functional G $\beta_1\gamma_x$ dimers tested are capable of stimulating PLC β activity [21], only certain G $\beta_1\gamma_x$ dimers (e.g. G $\beta_1\gamma_2$) effectively stimulated PKD phosphorylation in the presence of PLC $\beta_{2/3}$ (Figure 5D, lower panel). Hence, we hypothesized that the presence of PLC $\beta_{2/3}$

may allow specific G $\beta\gamma$ to associate with PKD. For this, HEK293 cells were transiently transfected with pcDNA3 (vector control), G $\beta_1\gamma_x$ (G $\beta_1\gamma_7$, G $\beta_1\gamma_9$) with or without PLC β_2 . FLAG-tagged G β_1 was immunoprecipitated from the lysates of the transfectants, and the immune complexes were subjected to SDS-PAGE, followed by Western blotting for any PKD co-immunoprecipitated with G β_1 . As shown in Figure 6, phosphorylated PKD1 was clearly detectable in the immunoprecipitates prepared from transfectants expressing both G $\beta_1\gamma_7$ dimer and PLC β_2 , but not when PLC β_2 was absent. Despite comparable expressions of the various constructs (Figure 6, right panel), hardly any PKD1 was pulled down by the FLAG-tagged G β_1 in cells expressing G $\beta_1\gamma_9$ with or without PLC β_2 (Figure 6, left panel). It should be noted that both G $\beta_1\gamma_7$ and G $\beta_1\gamma_9$ were able to interact with PLC β_2 in a comparable manner because the latter was detected in the immunoprecipitates (Figure 6, left panel). As the current data showed that G $\beta\gamma$ dimers alone are ineffective in the co-immunoprecipitation with PKD, hence, our findings not only demonstrate the

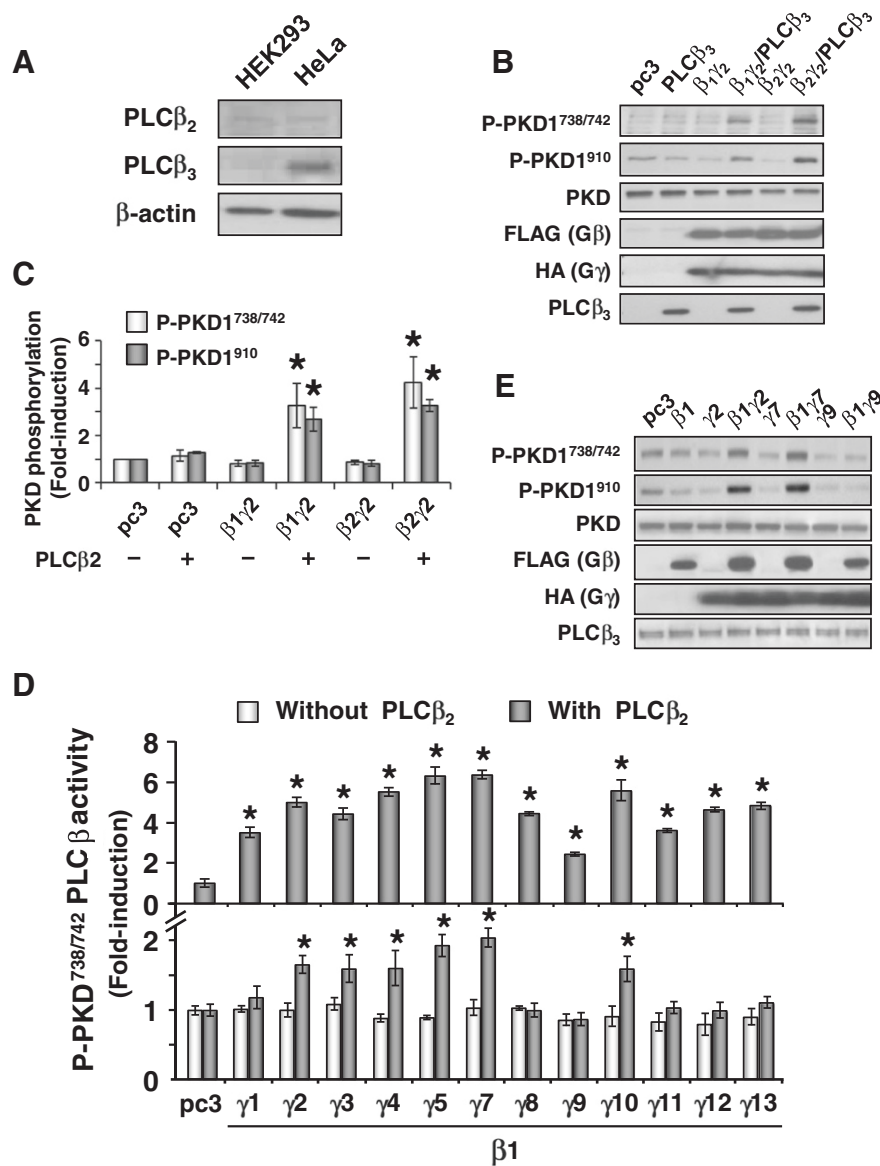


Figure 5 PLC $\beta_{2/3}$ is required for G $\beta\gamma$ dimer-induced PKD activation. (A), Expression of PLC β_2 and PLC β_3 in HEK293 and HeLa cells was determined with their respective anti-sera. HEK293 cells were transfected with pcDNA3 and G $\beta\gamma$ dimers with or without PLC β_3 (B) or PLC β_2 (C). PKD activation was detected by specific phospho-antibodies. Results shown are the mean \pm S.E.M. of at least three independent experiments. (D) HEK293 cells were transfected with pcDNA3, PLC β_2 , G $\beta\gamma$ with or without PLC β_2 . Transfectants were lysed and the extracts analyzed by PLC assay or SDS-PAGE together with Western blot using antibodies against phosphorylated PKD1. Band intensity was quantified by Image J software (National Institute of Health, Bethesda, MD, USA) and depicted in graphical form, and presented as a fold-induction of the pcDNA3 control. Results shown are the mean \pm S.E.M. of at least three independent experiments. (E) HeLa cells were transiently transfected with HA-tagged PKD1 together with vector control or G $\beta\gamma$ dimers. The expression of PKD1, FLAG-G β , HA-G γ and endogenous PLC β_3 were detected with their specific antibodies.

crucial role of PLC β_2 for the effective binding between G $\beta\gamma$ dimers and PKD, but also implicate that only specific G $\beta\gamma$ dimers are capable of interacting and activating PKD in the presence of PLC β_2 .

Having established that PKD1-3 activation is promoted by ectopic expression of certain G $\beta\gamma$ complexes, we investigated whether G $\beta\gamma$ -mediated PKD activation was implicated in G $_i$ -linked biological function. Cell migration [34]

and invasion [40] represent some of the known cellular functions of PKD. Since Jurkat T-cells express the G $_i$ -coupled receptor CXCR $_4$ and it is responsive to stromal cell-derived factor 1 α (SDF-1 α) for chemotaxis [41], it appears to be a good cellular system for this investigation. First of all, we examined whether PLC β_2 and PLC β_3 are endogenously expressed in Jurkat T cells. Indeed, Jurkat T-cells endogenously express both PLC β_2 and PLC β_3

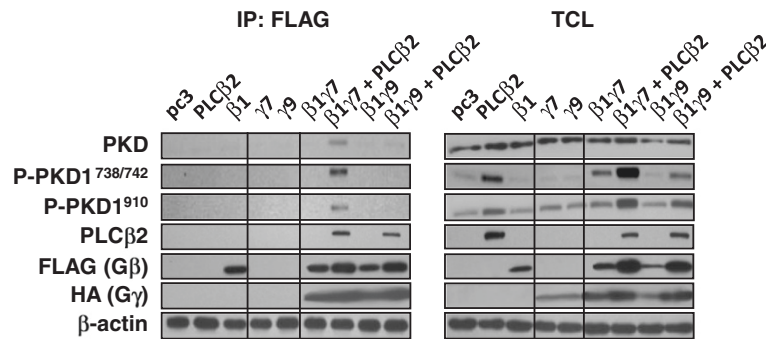


Figure 6 $G\beta_1$ interacts with PKD1 in the presence of specific $G\gamma$ subunits and $PLC\beta_2$. HEK293 cells were transiently transfected with pcDNA3, $PLC\beta_2$, and different $G\beta\gamma$ combinations as indicated, followed by serum starvation. Cell lysates were subjected to immunoprecipitation by FLAG-tagged affinity gel and SDS-PAGE to resolve for proteins bound to FLAG- $G\beta_1$. Black lines represent positions at which images from the same blot but on different lanes were merged.

isoforms, with the former being more abundant (Figure 7A). Next, we used PTX (which ADP-ribosylates G_i proteins) to confirm that SDF-1 α -induced signaling and chemotaxis in Jurkat T-cells are mediated via G_i proteins. Both SDF-1 α -induced intracellular Ca^{2+} mobilization (Figure 7B) and chemotaxis (Figure 7C) in Jurkat T-cells were completely abolished upon PTX pretreatment. These results imply that CXCR₄ utilizes G_i proteins to stimulate chemotaxis and $PLC\beta$ -mediated Ca^{2+} mobilization in Jurkat T cells. The latter response was presumably mediated by $G\beta\gamma$ dimers released from activated G_i proteins [42,43].

To determine whether PKD contributed to SDF-1 α -induced chemotaxis in Jurkat T cells, we asked if this chemotactic response can be inhibited by the PKD inhibitor, G66976. We were able to demonstrate that SDF-1 α -induced chemotaxis could be suppressed by pretreatment with G66976 (Figure 7D). In agreement with a previous report [41], the PI3K inhibitor wortmannin (Figure 7D) also inhibited the SDF-1 α -stimulated chemotaxis. Next, we assessed if PKD can be activated by the G_i -coupled CXCR4. Jurkat T-cells were pretreated with or without PTX, followed by SDF-1 α stimulation. Since Jurkat T-cells predominantly express PKD2 [44], only PKD2 phosphorylation was determined. SDF-1 α stimulated PKD2 phosphorylation became evident within 10 min and peaked at 15 min after agonist addition (Figure 7E). The response was effectively abolished by PTX pretreatment of Jurkat T-cells. As a control, phospho-ERK was similarly monitored; SDF-1 α also stimulated ERK phosphorylation in a PTX-sensitive manner (Figure 7E).

To substantiate that SDF-1 α -induced chemotaxis in Jurkat T-cells is PKD2-dependent, we used specific validated siRNA oligonucleotides to knock down the expression of PKD2. As shown in Figure 7F, control and scrambled siRNAs had no effect on PKD2 expression, while silencing of PKD2 led to a remarkable reduction in PKD2 expression; siRNAs targeting either PKD1 or

PKD3 did not affect the expression of PKD2. The siRNA-mediated knockdown of PKD2 effectively inhibited the SDF-1 α -induced chemotaxis, whereas the controls and siRNAs targeting PKD1 and PKD3 did not significantly suppress chemotaxis (Figure 7F). Furthermore, silencing of $PLC\beta_{2/3}$ but not $PLC\beta_1$ resulted in the suppression of SDF-1 α -induced chemotaxis in Jurkat T-cells, illustrating the importance of $G\beta\gamma$ -responsive $PLC\beta$ isoforms in this activity (Figure 8A). As SDF-1 α also acts on G_i -coupled CXCR4 receptor in HeLa cells for PKD activation (Figure 4C), we then performed similar knockdown treatment to verify the possible $PLC\beta_{2/3}$ -dependency. Our result demonstrated that this G_i -induced signaling also required the $G\beta\gamma$ -responsive $PLC\beta_{2/3}$ isoforms to stimulate the PKD activation (Figure 8B).

Discussion

Extending from prior reports on the regulation of PKD1 by G_q [30,45], the present study demonstrates unequivocally that each and every member of the G_q subfamily (i.e., $G\alpha_q$, $G\alpha_{11}$, $G\alpha_{14}$ and $G\alpha_{16}$) are capable of inducing the kinase activity of all PKD isoforms (Figure 2). The ability to β stimulate PKD activity is apparently unique to the $G\alpha_q$ members because other $G\alpha$ subunits belonging to the G_i , G_s , or G_{12} subfamilies all failed to induce PKD phosphorylation or kinase activity (Figures 1, 2, Additional file 1: Figure S1 and Additional file 2: Figure S2). However, it should be noted that addition of AlF_4^- to cells co-expressing PKD and wild type $G\alpha_{13}$ can lead to PKD activation [46]. Such an observation is confounded by the fact that AlF_4^- may activate multiple G proteins simultaneously. The lack of effect on PKD by the constitutively active mutant of $G\alpha_{13}$ has in fact been reported [30]. Hence, it is reasonable to conclude that only members of the G_q subfamily are efficiently linked to PKD activation.

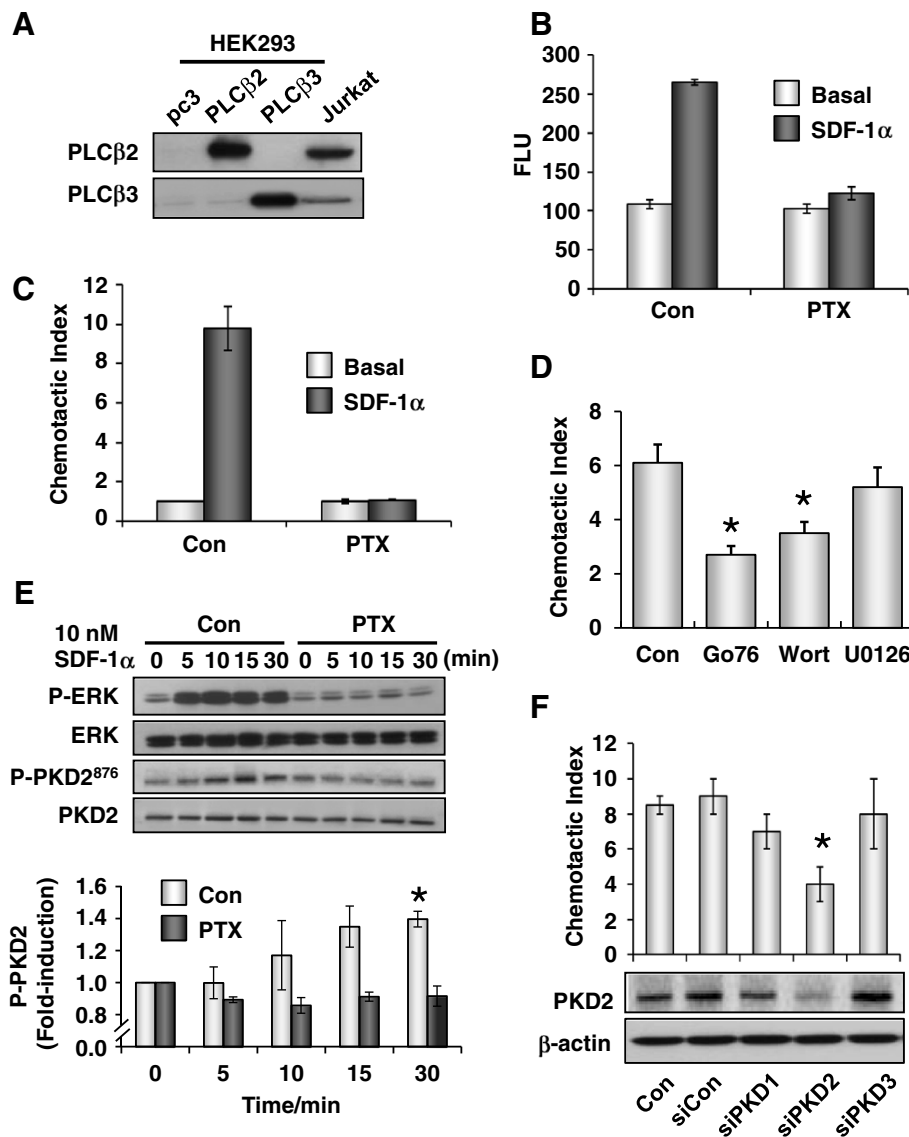


Figure 7 SDF-1 α -induced chemoataxis in Jurkat T cells is dependent on PTX-sensitive G_i proteins and PKD2. (A) Expression of PLC $\beta_{2/3}$ in Jurkat T cells were examined alongside with parental HEK293 cells or HEK293 cells overexpressing PLC $\beta_{2/3}$. (B) Bar diagram showing maximum SDF-1 α -induced Ca²⁺ mobilization in Jurkat T cells with or without PTX pretreatment. (C) Jurkat T cells pretreated with or without PTX were subjected to SDF-1 α -induced chemotactic assay. The chemotactic index was expressed as the ratio of the numbers of cells found in the lower compartments, between the agonist-stimulated and the unstimulated groups. (D) Jurkat T cells were pretreated with or without specific inhibitors for PKD (100 nM Gö6976), PI3K (100 nM wortmannin), and MEK/ERK (10 μ M U0126) and then subjected to the chemotactic assay in the presence of SDF-1 α . (E) Jurkat T cells pretreated with or without PTX were stimulated with 10 nM SDF-1 α for the indicated durations. Cell lysates were subjected to immunoblotting for ERK and PKD2 phosphorylation. (F) Jurkat T cells were transfected with vehicle (RNase free water), scrambled siRNA (siCon), siPKD1, siPKD2 or siPKD3 oligonucleotides for 72 h prior to chemotactic assay. Cells were also harvested and lysates were subjected to Western blot analysis with specific antibody against PKD2. β -actin was used as loading control.

Despite the preponderance of G_q in mediating GPCR-induced activation of PKD, stimulation of G_i-coupled receptors in HeLa cells resulted in PKD phosphorylation (Figure 4). This may be explained by the observation that HeLa cells endogenously express G $\beta\gamma$ -responsive PLC $\beta_{2/3}$ [28], thereby allowing G $\beta\gamma$ released from activated heterotrimeric G_i proteins to mediate PKD activation through the G $\beta\gamma$ /PLC/PKC axis. One would expect

that stimulation of G_i-coupled receptors will result in PKD activation in cells endowed with PLC $\beta_{2/3}$. However, if the endogenous PLC $\beta_{2/3}$ is responsive to G $\beta\gamma$ dimers and all active G protein heterotrimers liberate free G $\beta\gamma$ dimers, then it remains puzzling why stimulation of G_s-coupled receptors cannot activate PKD via PLC $\beta_{2/3}$ (Figure 4B). A recent report has revealed that differential dissociation may exist among different G proteins [47],

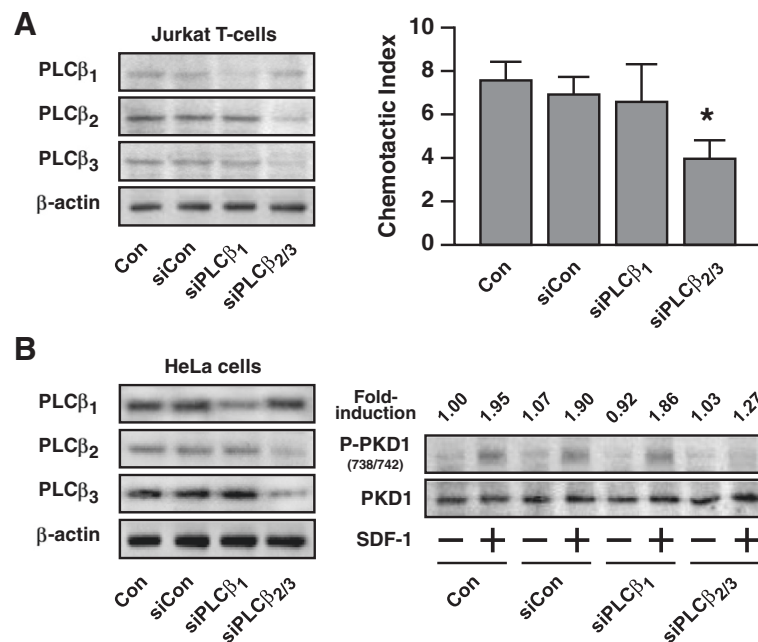


Figure 8 Requirement of Gβγ-responsive PLCβ_{2/3} isoforms in SDF-1α-induced signaling activities in Jurkat T-cells and HeLa cells.

(A) Jurkat T-cells were transfected with the vehicle control (RNase free water), scrambled siRNA (siCon), siPLCβ₁, or siPLCβ₂ and siPLCβ₃ oligonucleotides for 72 h prior to chemotactic assay upon SDF-1α treatment. Cells lysates were subjected to Western blot analysis with specific antibody against various PLCβ isoforms. β-actin was used as loading control. *Simultaneous knockdown the Gβγ-responsive PLCβ₂ and PLCβ₃ isoforms resulted in significant suppression of SDF-1α-induced chemotaxis. (B) HeLa cells subjected to similar PLCβ knockdown profiles were examined for SDF-1α-induced PKD activation, the levels of phospho-PKD1 in cell lysates were detected by Western blot analysis. The fold-induction represents the ratio of band intensities (phospho-PKD1) with respect to the basal (1.00) of the control group. Similar results were obtained in two independent experiments.

though it has long been thought that active G protein heterotrimers readily dissociate into Gα-GTP subunits and Gβγ dimers [48]. Activated G_{oA} heterotrimers can seemingly dissociate more readily than activated G_s heterotrimers, and this may account for Gα-specific activation of Gβγ-sensitive effectors [47]. Alternatively, the lack of G_s-induced PKD activation may be attributed to insufficient release of Gβγ dimers as most Gβγ-dependent signaling appeared to require substantial amounts of free Gβγ, which is most often achieved by stimulating the more abundantly expressed G_i proteins [42,43].

Another interesting observation in the present study pertains to the requirement of PLCβ_{2/3} for Gβγ-induced PKD activation (Figure 5). At first sight, our finding seems to suggest a concept different from the previous belief that Gβγ dimers alone can activate PKD through interaction with the PH domain [27]. However, since the cellular model (i.e. HeLa cells) used in Jamora's report expresses significant amount of Gβγ-sensitive PLCβ_{2/3}, it is possible that the presence of PLCβ_{2/3} enables specific Gβγ dimers to act on the PH domain of PKD. It has been demonstrated that Gγ prenylation is one of the important factors for Gβγ interaction with PLC isoforms, as the presence of farnesyl lipid motif in Gγ₁, Gγ₉ and

Gγ₁₁ may lead to a weaker PLC activation as compared to Gβγ dimers containing other Gγ components with geranylgeranyl lipid motif [49]. Indeed, we have observed that Gβ₁γ₁, Gβ₁γ₉ and Gβ₁γ₁₁ are associated with a weaker PLC activation and all of them are incapable of effectively stimulating PKD (Figure 5D and 5E). Hence, the possible influence of Gγ prenylation status cannot be neglected. However, Gβ₁γ₂ and Gβ₁γ₃ induce PLC activities of similar magnitude as those of Gβ₁γ₁₂ and Gβ₁γ₁₃, but only the former two are capable of stimulating PKD. As Gγ₂, Gγ₃, Gγ₁₂, and Gγ₁₃ are commonly incorporated with the geranylgeranyl lipid motif, factors other than Gγ prenylation and PLC activity may also be important for governing the specificity of Gβγ-mediated PKD activation. It can be observed that only certain Gβ₁γ dimers (i.e., those containing γ₂, γ₃, γ₄, γ₅, γ₇, and γ₁₀) but not others (i.e., those containing γ₁, γ₈, γ₉, γ₁₁, γ₁₂, and γ₁₃) could effectively activate PKD in the presence of PLCβ_{2/3} (Figure 5D, lower panel). Yet, all combinations of Gβ₁γ_x dimers are capable of activating PLCβ₂ [21]. The differential ability of various Gβ₁γ dimers to stimulate PKD is thus unlikely to solely depend on their PLCβ activity alone. It can also be observed that the expression levels of Gβ₁γ₄, Gβ₁γ₇, Gβ₁γ₉, Gβ₁γ₁₁ and Gβ₁γ₁₂ appear

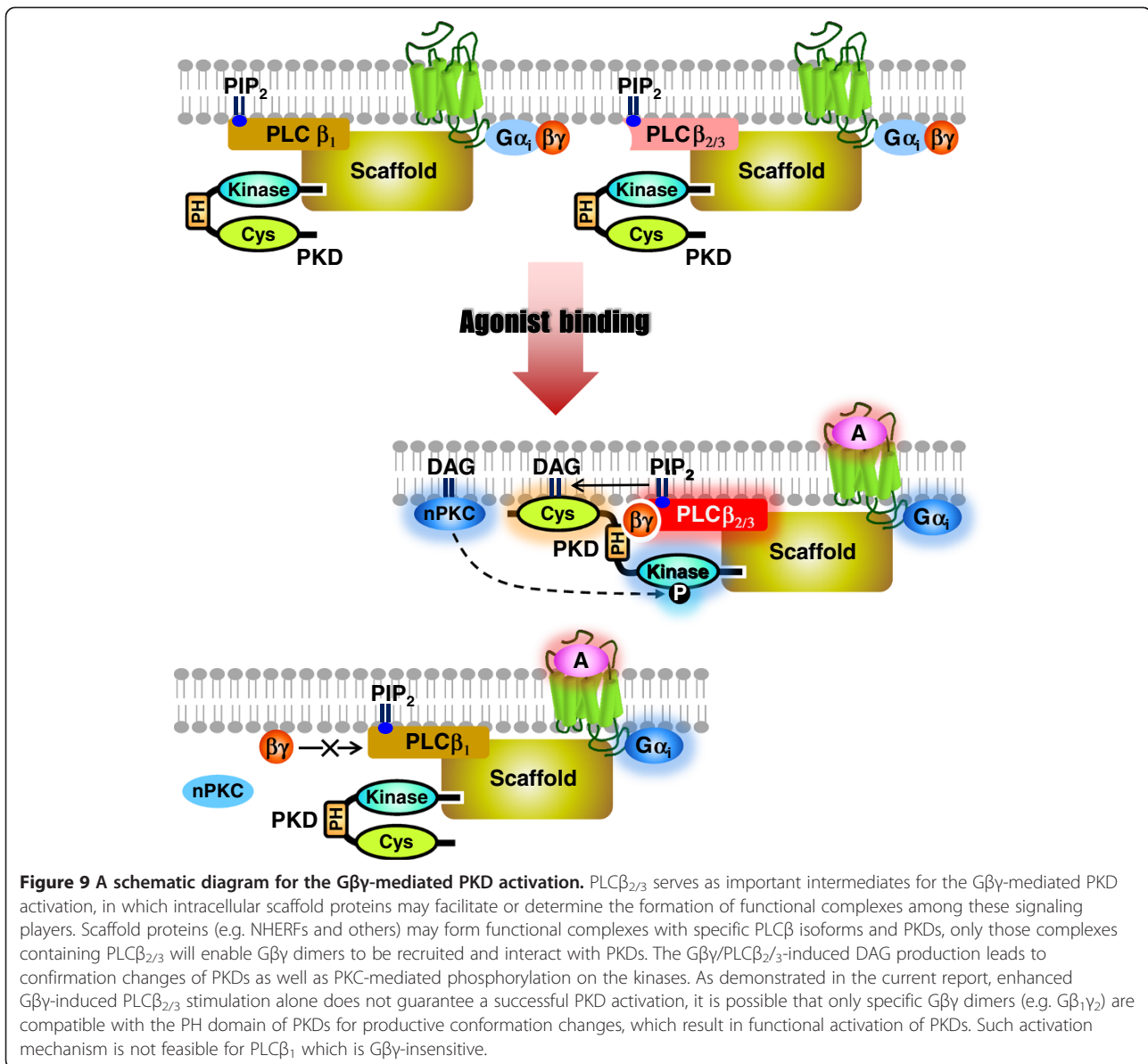
to be increased upon PLC β_2 co-expression (Additional file 4: Figure S4). However, such increased G $\beta\gamma$ expression is not necessarily related to the subsequent PKD activation, as increased G $\beta_1\gamma_9$, G $\beta_1\gamma_{11}$ and G $\beta_1\gamma_{12}$ expressions do not effectively stimulate PKD in the presence of PLC β_2 , whereas G $\beta_1\gamma_2$, G $\beta_1\gamma_3$, G $\beta_1\gamma_5$, and G $\beta_1\gamma_{10}$ trigger the kinase activation without increased levels of subunit expressions (Additional file 4: Figure S4). Hence, G $\beta\gamma$ -mediated PKD activation seems to be a specific function in response to unique G $\beta\gamma$ combinations.

In fact, the ability of specific G $\beta\gamma$ dimers to stimulate PKD phosphorylation may depend on their ability to form a complex with PKD, since only those G $\beta\gamma$ dimers that can stimulate PKD (e.g., G $\beta_1\gamma_7$) could be immunoprecipitated with PKD (Figure 6). The requirement of PLC $\beta_{2/3}$ in G $\beta\gamma$ -mediated PKD signaling might be explained if PLC $\beta_{2/3}$ is an essential component of the signaling complex that stabilizes the interaction between G $\beta\gamma$ and PKD. The possible existence of a G $\beta\gamma$ /PLC $\beta_{2/3}$ /PKD signaling complex is supported by the fact that G $\beta\gamma$ dimers serve as direct activators for PLC $\beta_{2/3}$ [50], probably through the binding of G $\beta\gamma$ to the PH domain of PLC $\beta_{2/3}$ [51], while G $\beta\gamma$ /PKD-mediated Golgi fragmentation can be inhibited by a sequester peptide with identical sequence of the G $\beta\gamma$ -binding PH domain in PKD [27]. Indeed, we have preliminary data suggesting that PLC β_2 can be co-immunoprecipitated with all three PKD isoforms, while PLC β_1 fails to do so (Additional file 5: Figure S5). Apparently the reported capabilities of G $\beta\gamma$ to interact with PLC $\beta_{2/3}$ and PKD seem to support the notion for the formation of a G $\beta\gamma$ /PLC $\beta_{2/3}$ /PKD signaling complex. However, it is unclear as to whether a single G $\beta\gamma$ dimer binds to the PH domains of PLC $\beta_{2/3}$ and PKD sequentially or simultaneously. Similarly, we cannot rule out the possibility that there may be different pools of G $\beta\gamma$ dimers for G $\beta\gamma$ -PLC β and G $\beta\gamma$ -PKD interactions, respectively, and that they may subsequently cooperate with each other to stimulate PKD. Further studies are required to examine the precise interactions between G $\beta\gamma$, PLC $\beta_{2/3}$ and PKD.

The assembly of a G $\beta\gamma$ /PLC $\beta_{2/3}$ /PKD signaling complex may require the participation of scaffolding proteins. In this regard PKD isoforms have been shown to interact with the PDZ domains of a scaffolding protein family named NHERF [52]. Coincidentally, PLC $\beta_{2/3}$ can also interact with different NHERF members [53,54]. Hence, NHERF, as well as other similar scaffold proteins, may act as a nexus for G $\beta\gamma$ /PLC β /PKD signaling (Figure 9), in which intracellular scaffold may facilitate or determine the formation of functional complexes among the signaling players. Scaffolding proteins (e.g. NHERFs and others) may form functional complexes with specific PLC β isoforms and PKDs, and perhaps only those complexes containing PLC $\beta_{2/3}$ will enable G $\beta\gamma$

dimers to be recruited for interaction with PKDs. Such activation mechanism is not feasible for PLC β_1 which is G $\beta\gamma$ -insensitive. The G $\beta\gamma$ /PLC $\beta_{2/3}$ -induced DAG production leads to confirmation changes of PKDs as well as PKC-mediated phosphorylation on the kinases. As demonstrated in the current report, enhanced G $\beta\gamma$ -induced PLC $\beta_{2/3}$ stimulation alone does not guarantee a successful PKD activation, it is possible that only specific G $\beta\gamma$ dimers (e.g. G $\beta_1\gamma_2$) are compatible with the PH domain of PKDs for productive conformation changes, which result in functional activation of PKDs. In fact, our unpublished data showed that PKD activation triggered by G $_i$ -coupled receptors is sensitive to inhibitors for PLC β (e.g. U73122) as well as to G $\beta\gamma$ subunit scavengers (e.g. transducin). Since only specific G $\beta\gamma$ dimers are capable of stimulating PKD in the presence of PLC $\beta_{2/3}$, our results actually suggest a dual requirement of functional PLC β activity and compatible G $\beta\gamma$ dimers for G $_i$ -mediated PKD activation. It remains unclear if all the members in the G $_q$ family (i.e. G α_q , G α_{11} , G α_{14} , and G α_{16}) also activate PKD in a similar manner. However, it should be noted that another scaffold protein named PAR3 have been suggested as a G $_q$ -specific signaling component with selective recruitment of PLC β_1 , while PLC $\beta_{2/3}$ isoforms may have high preferences towards NHERF members in G $_i$ -mediated signaling [53,54]. The involvement of different scaffold proteins may also explain the differential observation that, G α subunits of the G $_q$ family (much stronger activators for PLC β isoforms as compared to G $\beta\gamma$ dimers) are capable of stimulating PKD in a G $\beta\gamma$ -independent manner.

PKD mediates a diverse array of normal biological functions and pathological activities, including cell proliferation and differentiation, cell motility, regulation of cell vesicle trafficking, secretion, and polarity, inflammatory responses, cardiac hypertrophy and cancer [55]. Therein, the transport of protein from the Golgi to plasma membrane is regulated via G $\beta\gamma$ signaling [27,28,56]. From our results, it is postulated that stimulation of G $_i$ -coupled receptor leads to the liberation of free G $\beta\gamma$ dimers, which then interact with PLC $\beta_{2/3}$ and activate PKD. This may help to elucidate part of the mechanism regarding secretory activities regulated by receptor-induced G $\beta\gamma$ translocation between the Golgi and plasma membrane [57], and the characteristic of Golgi as one of the major cellular locations for activated PKD [58]. Indeed, G $\beta\gamma$ dimers are known to mediate many cellular responses and signaling pathways involved in multiple aspects of cellular function. Previous studies have reported that SDF-1 α -induced activation of CXCR4 receptor induces chemotaxis in Jurkat T cells [41]. Here, our results showed that this G $_i$ -coupled chemotactic response may be mediated by the G $\beta\gamma$ /PLC β /PKD axis (Figure 7). However, further investigations are needed to



determine whether these components act in concert. The activation of STAT3, which is an important transcription factor, is also regulated by $G\beta\gamma$ -mediated signaling [23]. Similar to PKD, only distinct combinations of $G\beta\gamma$ can effectively activate STAT3. Nevertheless, the panel of STAT3-activating $G\beta\gamma$ dimers is not identical to the PKD-stimulatory $G\beta\gamma$ complexes; only $G\beta_1\gamma_4$ and $G\beta_1\gamma_7$ are effective activators for both pathways. Taken together, our results suggested that PKD may be implicated in diverse cellular activities, including those mediated by $G\beta\gamma$.

Functional redundancy is a common feature among isoforms of biological molecules. However, it is not always the case. Though the three PKD isoforms are highly

conserved and our results showed that all three PKD isoforms (PKD1, PKD2 and PKD3) are activated equally well by $G\alpha$ subunits from the G_q family, as well as by specific $G\beta_1\gamma_x$ with $PLC\beta_{2/3}$, they may have unique functions. For example, PKD1 plays a non-redundant role in pathological cardiac remodeling, and the homozygous germline deletion of PKD1 causes embryonic lethality [59]. As for PKD2, it has a unique role in endothelial cells [6], lymphoid cells [7], and monocytes [34]. Recent studies have revealed the essential role of PKD3 in the progression of prostate cancer [60] and insulin-independent basal glucose uptake in L6 skeletal muscle cells [61]. Further studies are necessary to elucidate the mechanisms behind GPCR-mediated activation of the three PKD isoforms.

Conclusion

Collectively, among various members of G proteins, only the G_{α} subunits of the G_q family effectively activate all three PKD isoforms (PKD1/2/3), while G_{α} subunits of other G protein families (G_s , G_i , and G_{12}) are inefficient in these kinase activations. However, receptors linked to G_i proteins are capable of triggering PKD activation in cell lines endogenously expressing (HeLa cells and Jurkat T-cells) or exogenously transfected with (HEK293 cells) $G\beta\gamma$ -sensitive PLC $\beta_{2/3}$ isoforms, indicating the involvement of $G\beta\gamma$ dimers for the G_i -mediated PKD activation. Although the presence of PLC $\beta_{2/3}$ is highly important, only those $G\beta_1\gamma$ dimers with γ_2 , γ_3 , γ_4 , γ_5 , γ_7 , and γ_{10} are effective activators of PKD, and the specific interaction between $G\beta\gamma$, PKD and PLC $\beta_{2/3}$ may play a pivotal role in this $G\beta\gamma$ -mediated PKD signaling pathway. Furthermore, the biological significance of G_i -mediated PKD activation is illustrated by SDF-1 α -induced chemotaxis on Jurkat T-cells, in which the chemotactic activity is abolished by pretreatment with PTX and knockdown of PKD. Taken together, our current report illustrates that $G\beta\gamma$ dimers from G_i proteins may activate PKD in a PLC $\beta_{2/3}$ -dependent manner, and the identity of $G\gamma$ of the $G\beta\gamma$ dimer being a determinant.

Additional files

Additional file 1: Figure S1. Constitutively active G_{α} subunits from the G_i subfamily failed to induce PKD activation. (A) HEK293 cells were transiently transfected with pcDNA3 or WT/QL forms of G_{α} subunits from the G_i subfamily. Cell lysates were subjected to SDS-PAGE. G_{α} subunits, phospho-PKD1^{738/742}, phospho-PKD1⁹¹⁰, total PKD1, tag of PKD1 (HA) were analyzed by Western blotting using respective specific antibody. (B) HA-PKD1, FLAG-PKD2 and Myc-PKD3 were immunoprecipitated from cell lysates described in (A), and subjected to *in vitro* PKD kinase assays in terms of syntide-2 phosphorylation. Results are the average of at least three independent experiments, and represented as fold increase over pcDNA3 control (\pm S.E.M.).

Additional file 2: Figure S2. Constitutively active G_{α} subunits from the G_s and G_{12} subfamilies failed to induce PKD activation. (A) HEK293 cells were transiently transfected with pcDNA3 or WT/QL forms of G_{α} subunits from the G_s and G_{12} subfamilies. Cell lysates were subjected to SDS-PAGE. G_{α} subunits, phospho-PKD1^{738/742}, phospho-PKD1⁹¹⁰, total PKD1, tag of PKD1 (HA) were analyzed by Western blotting using respective specific antibody. (B) HA-PKD1, FLAG-PKD2 and Myc-PKD3 were immunoprecipitated from cell lysates described in (A), and subjected to *in vitro* PKD kinase assays. Results are the average of at least three independent experiments, and represented as fold increase over pcDNA3 control (\pm S.E.M.).

Additional file 3: Figure S3. PLC β_2 and specific $G\gamma$ subunits are required in $G\beta\gamma$ -induced PKD activation in HEK293 cells. (A) HEK293 cells stably transfected with pcDNA3 or HA- G_{γ_2} were transfected with pcDNA3, FLAG- $G\beta_1$ or FLAG- $G\beta_2$. Cell lysates were subjected to SDS-PAGE. FLAG- $G\beta$, HA- $G\gamma$, phospho-PKD1^{738/742}, phospho-PKD1⁹¹⁰ and total PKD1 were analyzed by Western blotting using respective specific antibody. (B) HEK293 cells were transiently transfected with vector control, PLC β_2 , various $G\beta\gamma$ dimers and tagged PKD isoforms (HA-PKD1, FLAG-PKD2 and Myc-PKD3). The cultures were then lysed, and the tagged PKD isoforms were immunoprecipitated for *in vitro* PKD kinase assay. (C) HEK293 cells transiently transfected with pcDNA3, $G\beta$, $G\gamma_w$ $G\beta\gamma$ combinations with or

without PLC β_1 or PLC β_2 were lysed, and analyzed by Western blotting for PKD1 phosphorylation.

Additional file 4: Figure S4. The expression profiles of $G\beta\gamma$ dimers and the corresponding PKD activation in the presence of PLC β_2 . HEK293 cells were transfected with pcDNA3, PLC β_2 , various combinations of $G\beta\gamma$ with or without PLC β_2 . Transfectants were lysed, and the lysates were subjected to Western blotting using antibodies against phosphorylated PKD1, PKD, PLC β_2 , Flag-tagged $G\beta_1$ and HA-tagged $G\gamma$ subunits.

Additional file 5: Figure S5. PLC β_2 , but not PLC β_1 , can be co-immunoprecipitated with various PKD isoforms. HEK293 cells were transiently transfected with pcDNA3, PLC $\beta_{1/2}$ with tagged PKD1/2/3 as indicated. HA-PKD1, FLAG-PKD2 and Myc-PKD3 were immunoprecipitated from cell lysates with their respective affinity gels and further analyzed by Western blotting for the possible interaction with PLC $\beta_{1/2}$.

Abbreviations

BK₂R: Bradykinin type II receptor; β_2 AR: β_2 -adrenergic receptor; DAG: Diacylglycerol; fMLPR: N-formyl-methionyl-leucyl-phenylalanine receptor; GPCR: G protein-coupled receptor; PLC β : Phospholipase C β ; PKD: Protein kinase D; PTX: Pertussis toxin.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

WWL and ASLC carried out the experiments participated in the design of the study and wrote the manuscript. LSWP and JZ carried out some of the experiments. YHW participated in the design of the study and revised the manuscript. All authors read and approved the final manuscript.

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