

Protein Kinase C- δ Negatively Regulates T Cell Receptor-induced NF- κ B Activation by Inhibiting the Assembly of CARMA1 Signalosome^{*[5]}

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Background: TCR-induced NF- κ B activation is crucial for T cell activation.

Results: PKC δ inhibited TCR-induced NF- κ B activation by a mechanism independent of its kinase activity.

Conclusion: PKC δ negatively regulates T cell activation by inhibiting the assembly of CARMA1 signalosome.

Significance: It provides a novel mechanism on negative regulation of T cell activation.

T-cell receptor (TCR)-induced T-cell activation is a critical event in adaptive immune responses. The engagement of TCR complex by antigen along with the activation of the costimulatory receptors trigger a cascade of intracellular signaling, in which caspase recruitment domain-containing membrane-associated guanylate kinase 1 (CARMA1) is a crucial scaffold protein. Upon stimulation, CARMA1 recruits downstream molecules including B-cell CLL/lymphoma 10 (Bcl10), mucosa-associated lymphoid tissue lymphoma translocation gene 1 (MALT1), and TRAF6 to assemble a specific TCR-induced signalosome that triggers NF- κ B and JNK activation. In this report, we identified protein kinase C δ (PKC δ) as a CARMA1-associated protein by a biochemical affinity purification approach. PKC δ interacted with CARMA1 in TCR stimulation-dependent manner in Jurkat T cells. Overexpression of PKC δ inhibited CARMA1-mediated NF- κ B activation, whereas knockdown of PKC δ potentiated TCR-triggered NF- κ B activation and IL-2 secretion in Jurkat T cells. Reconstitution experiments with PKC δ kinase-dead mutant indicated that the kinase activity of PKC δ was dispensable for its ability to inhibit TCR-triggered NF- κ B activation. Furthermore, we found that PKC δ inhibited the interaction between MALT1 and TRAF6, but not the association of CARMA1 with PKC θ , Bcl10, or MALT1. These observations suggest that PKC δ is a negative regulator in T cell activation through inhibiting the assembly of CARMA1 signalosome.

TCR-induced activation of T cells plays a central role in adaptive immune responses. Activated T cells undergo clonal expansion, differentiation, and begin to execute their immunological functions either by producing cytokines for regulating

immune responses or by killing target cells infected with pathogens directly. The tightly controlled activation of transcription factors of the nuclear factor- κ B (NF- κ B)⁴ family plays a crucial role in these processes (1).

TCR-induced activation of NF- κ B signaling pathway is initiated from the interaction of TCR with specific antigen in the context of MHC and costimulation of CD28 by its ligands on antigen-presenting cells. Upon stimulation, a cascade of tyrosine phosphorylation of signaling components is successively triggered, which results in the recruitment and activation of a number of signaling proteins, including phosphoinositide-dependent kinase 1 (PDK1) (2). PDK1 phosphorylates PKC θ and recruits it to lipid rafts in the plasma membrane (3). Concurrently PDK1 recruits CARMA1 to lipid rafts. Phosphorylated PKC θ phosphorylates the close-by CARMA1 at specific serine residues, and leads to the conformational changes of CARMA1 from an inactive form to an active one (4–6). Active CARMA1 assembles a CBM complex by associating with Bcl10, which in turn recruits MALT1 (7, 8). The CBM complex subsequently recruits downstream signal proteins, such as TRAF6, caspase 8, and TAK1, to collaboratively activate the inhibitor of κ B kinase (IKK) complex. IKK is then phosphorylated and undergoes degradation, which leads to the activation of NF- κ B.

Among the CBM signalosome, the scaffold protein CARMA1 is crucial for the recruitment of downstream signal proteins, as well as their plasma membrane localization (9–11). Its activity requires fine tuning to elicit proper activation of NF- κ B. Post-transcriptional modifications of CARMA1, such as phosphorylation and ubiquitination, determine the outcome of TCR stimulation. For examples, kinases such as PKC θ , IKK β , Ca²⁺/calmodulin-dependent protein kinase II, and hematopoietic progenitor kinase 1 can phosphorylate CARMA1 at different serine residues in the PKC-regulated domain (PRD), and subsequently lead to NF- κ B activation (4, 5, 12–14). However, casein kinase 1 α phosphorylates CARMA1 at S608, which impairs its ability to activate NF- κ B (15). In addition, monoubiquitination of CARMA1 catalyzed by Cbl-b disrupts the for-

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⁴ The abbreviations used are: NF- κ B, nuclear factor- κ B; TCR, T-cell receptor; IKK, inhibitor of κ B kinase; CARMA, caspase recruitment domain-containing membrane-associated guanylate kinase; PDK, phosphoinositide-dependent kinase.

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mation of CBM complex (16). Although much has been reported, the subtle regulation of CARMA1 signalosome remains not so clear thus far.

To identify potential regulators of CARMA1-mediated NF- κ B activation, we performed biochemical affinity purification with CARMA1 as a bait protein, and identified PKC δ as a protein specifically associated with CARMA1. Knockdown of PKC δ accelerated TCR-induced NF- κ B activation and IL-2 secretion, suggesting an inhibitory role of PKC δ in TCR-mediated NF- κ B activation. Furthermore, we demonstrated that the kinase activity of PKC δ was not required for its inhibitory role. Instead, PKC δ inhibited TCR-induced NF- κ B activation by steric hindrance of recruiting TRAF6 to CARMA1 signalosome.

EXPERIMENTAL PROCEDURES

Reagents—The Abs against the following antigens were purchased from the indicated companies: human CD3, human CD28, and mouse IgG1 (BD Biosciences); human I κ B α and phospho-I κ B α (S32, S36) (Cell Signaling Technology); human PKC δ (Santa Cruz Biotechnology); human β -actin, Flag, and hemagglutinin (HA) epitopes (Sigma-Aldrich). Mouse antisera were raised against recombinant human CARMA1 (aa 1–468), Bcl10 or MALT1 (aa 348–813) respectively. Human IL-2 ELISA kit (Dake Biotechnology), dual-luciferase reporter assay kit (Promega), and Gamma Bind G Plus-Sepharose (GE Healthcare Life Sciences) were purchased from the indicated companies.

Constructs—NF- κ B luciferase reporter plasmid was provided by Dr. Gary Johnson. Mammalian expression plasmids for human CARMA1 and its mutants, PKC δ and its mutants, PKC θ , Bcl10, or MALT1, fused with Flag-, HA-, or RFP-epitope, were constructed by standard molecular biology techniques. Double-stranded oligonucleotides for RNA interference corresponding to the target sequences were inserted into pSUPER. Retro vector (Oligoengine) according to protocols recommended by the manufacturer. The target sequence for human PKC δ -siRNA #1 is 5'-AAACACTGGTGCAGAAGAA-3'; for PKC δ -siRNA #3 is 5'-GCAGCAAGTGCAACATCAA-3'.

Protein Purification and Mass Spectrometry Analysis—HEK293 cells (5×10^8) stably transfected with pCTAP-CARMA1 (Stratagene) were collected and the cell lysates were subjected to tandem affinity purification procedures. The purified CARMA1-associated proteins were digested by trypsin in solution. The tryptic peptides were analyzed by HPLC-ESI/MS/MS with a Thermo Finnigan LTQ adapted for nanospray ionization. The tandem spectra were searched against *Homo sapiens* National Center for Biotechnology Information reference data base using SEQUEST. Results was filtered by Xcorr +1 > 1.9, +2 > 2.2, +3 > 3.5, sp > 500, Deltcn > 0.1, Rsp < = 5.

Transfection and Reporter Assays—HEK293 cells were transfected with mammalian expression plasmids by standard calcium phosphate precipitation. Jurkat T cells were transfected with siRNAs or expression plasmids by retroviral transduction. For luciferase reporter assays, HEK293 cells (1×10^5) were seeded on 24-well dishes and transfected the next day. In the

same experiment, we added empty control plasmid to ensure that each transfection received the same amount of total DNA. To normalize the transfection efficiency, 0.01 μ g pRL-TK (*Renilla* luciferase) reporter plasmid was added to each transfection. Luciferase assays were performed with a dual-specific luciferase assay kit. Firefly luciferase activities were normalized on the basis of *Renilla* luciferase activities. All reporter assays were repeated for at least three times. Data shown were average values \pm S.D. from one representative experiment.

Cell Sorting by FACS—Constructs of MIGR-GFP-PKC δ -WT (wild type) or MIGR-GFP-PKC δ -KD (kinase-dead mutant, K378A) were transfected into PKC δ -knockdown Jurkat T cells (18). Forty-eight hours later, GFP-positive cells were isolated based on fluorescence intensity using the MoFlo XDP Sorter (Beckman Coulter). Approximately 1×10^5 cells of each sample with >95% purity were obtained and cultured for further experiments.

Coimmunoprecipitation and Immunoblot Analysis—For transient transfection and coimmunoprecipitation experiments, HEK293 cells (2×10^6) were transfected for 20 h. The transfected cells were lysed in Nonidet P-40 lysis buffer (20 mM Tris, 150 mM NaCl, 1% Nonidet P-40, 1 mM EDTA, pH 7.5) supplemented with proteases inhibitors (10 μ g/ml aprotinin, 10 μ g/ml leupeptin, 1 mM phenylmethylsulfonyl fluoride). For each immunoprecipitation, 0.4 ml aliquot of lysates was incubated with 0.5 μ g of antibody or control IgG and 25 μ l of GammaBind G Plus-Sepharose at 4 $^{\circ}$ C for 2 h. Sepharose beads were washed three times with 1 ml lysis buffer containing 0.5 M NaCl. The precipitates were analyzed by standard immunoblotting procedures.

For endogenous coimmunoprecipitation experiments, Jurkat T cells (5×10^7) were costimulated by mouse anti-human CD3 (10 μ g/ml), mouse anti-human CD28 (10 μ g/ml), and 10 μ g/ml goat anti-mouse IgG for crosslinking at 37 $^{\circ}$ C for the indicated times. Cells were then lysed and subjected to coimmunoprecipitation and immunoblot analysis as described above.

ELISA—IL-2 production in the culture medium of Jurkat cells was measured after CD3/CD28 crosslinking for 48 h using a human IL-2 ELISA kit.

RESULTS

Identification of PKC δ as a CARMA1-associated Protein—CARMA1 plays a central role in signaling to NF- κ B activation by TCR or BCR engagement. For a better understanding of signaling events in this pathway, we purified CARMA1-associated proteins by a tandem affinity purification system, and identified the eluted proteins by a shotgun mass spectrometry analysis method. By comparing with other unrelated purifications using the same method, we identified several candidates, including PKC δ , as specific CARMA1-associated proteins. Coimmunoprecipitation results confirmed the interaction between ectopic PKC δ and CARMA1 in HEK293 cells (Fig. 1A). Besides, PKC δ also interacted with MALT1 and TRAF6, but not Bcl10 (Fig. 1B).

To determine whether CARMA1 and PKC δ interact in untransfected cells, we stimulated Jurkat T cells by crosslinking CD3 and CD28 with antibodies, and performed endogenous

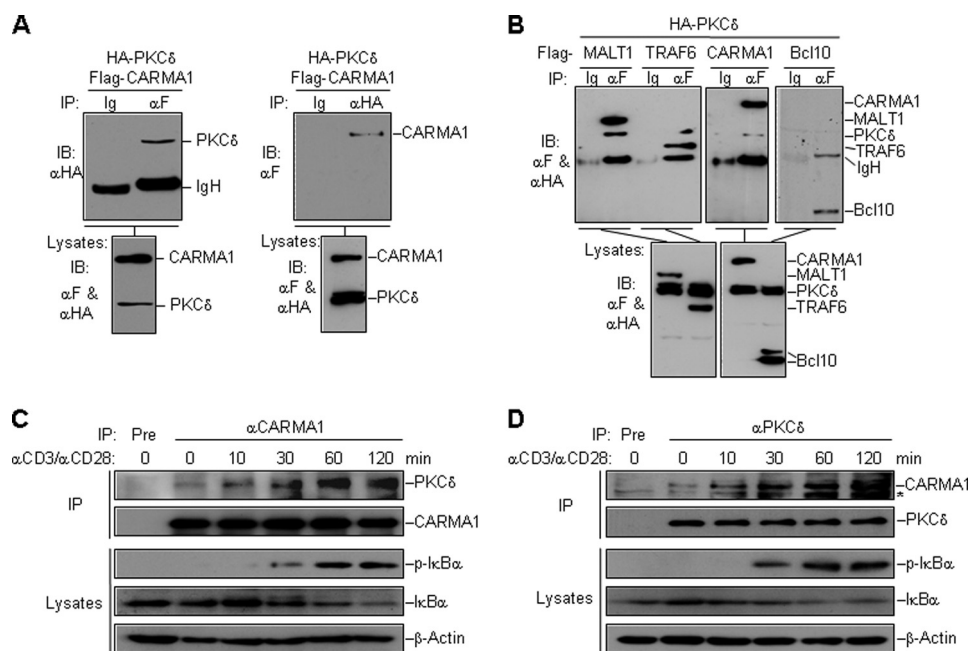


FIGURE 1. Identification of PKC δ as a CARMA1-associated protein. A, interaction of overexpressed HA-PKC δ and Flag-CARMA1. Expression plasmids encoding HA-PKC δ (10 μ g) and Flag-CARMA1 (4 μ g) were transfected into HEK293 cells. Lysates were immunoprecipitated with antibodies to Flag (left panel), HA (right panel), or control IgG, and then analyzed by immunoblotting with an antibody to HA (upper left panel) or Flag (upper right panel). Expression of the transfected proteins were analyzed by immunoblotting with antibodies to Flag and HA (lower panel). B, PKC δ interacts with MALT1 and TRAF6 but not Bcl10. HEK293 cells were transfected with the indicated plasmids. Coimmunoprecipitation and immunoblot analysis were performed with the indicated antibodies. C and D, CD3/CD28 costimulation induces the association between PKC δ and CARMA1. Jurkat T cells were stimulated by CD3/CD28 crosslinking for the indicated time. Lysates were immunoprecipitated with an antibody to CARMA1 (C) or PKC δ (D). The precipitates and lysates were analyzed by immunoblotting with the indicated antibodies. Ig, control mouse IgG; α F, anti-Flag; Pre, preimmune serum; *, nonspecific band.

coimmunoprecipitation. The results showed that endogenous PKC δ was recruited to CARMA1 after CD3/CD28 costimulation (Fig. 1, C and D). Immunofluorescent confocal microscopy also indicated that a population of PKC δ was translocated to the plasma membrane upon TCR stimulation (supplemental Fig. S1).

PKC δ Negatively Regulates TCR-induced Activation of NF- κ B—To address whether PKC δ is involved in CARMA1-dependent NF- κ B activation, we performed NF- κ B reporter assays. Overexpression of PKC δ alone had no marked effect on the basal level of NF- κ B activation. However, when cotransfected with CARMA1, PKC δ inhibited CARMA1-induced NF- κ B activation in a dose-dependent manner (Fig. 2A).

To evaluate the role of PKC δ in TCR-induced NF- κ B signaling, we used RNAi to knockdown PKC δ in human Jurkat T cells. We made four PKC δ siRNA constructs, and determined their effects on knockdown of PKC δ . As shown in Fig. 2B, the #3 PKC δ -siRNA construct could inhibit the expression of transiently transfected PKC δ to 20% of the control sample ($p < 0.01$). The #1 siRNA constructs could reduce PKC δ level to 60% of the control sample, whereas #2 and #4 siRNA plasmid had minimal effects. We thus chose #3 and #1 siRNA constructs for further experiments.

We generated stable Jurkat T cells expressing PKC δ -siRNAs by retroviral transduction. Comparing with the controls, PKC δ -knockdown Jurkat T cells exhibited enhanced phosphorylation and degradation of I κ B upon the stimulation of CD3/CD28 crosslinking (Fig. 2C). Consistently, knockdown of PKC δ potentiated the production of IL-2, which is a pro-inflammatory cytokine induced by NF- κ B activation (Fig. 2D). The extent

of IL-2 enhancement was correlated with the efficiency of PKC δ knockdown (Fig. 2D). These evidences suggest that PKC δ itself plays a negative role in TCR-induced activation of NF- κ B.

The Kinase Activity of PKC δ Is Not Required for Its Inhibitory Role in TCR-induced NF- κ B Activation—It has been reported that upon TCR ligation, PKC δ is phosphorylated and regulates granule exocytosis in a kinase-dependent manner (17). Therefore, we tested whether the kinase activity of PKC δ was required for TCR-induced NF- κ B activation. We reconstituted PKC δ -knockdown Jurkat T cells with PKC δ -wild type (WT) or PKC δ -kinase dead (KD) mutant (18). The reconstitutions were confirmed by immunoblotting (Fig. 3A). The reconstituted cells, PKC δ -knockdown cells and controls were stimulated by CD3/CD28 crosslinking, and then the culture medium of each sample was collected to measure IL-2 production, while the cell lysates were subjected to immunoblots to detect NF- κ B activation. The results showed that the ectopic expression of both WT and KD PKC δ reversed the promotion of IL-2 production and I κ B phosphorylation and degradation in PKC δ -knockdown cells (Fig. 3, B and C), indicating that the kinase activity was dispensable for the inhibitory role of PKC δ in TCR-induced NF- κ B activation.

PKC δ Is Associated with the MAGUK Region of CARMA1, and Has No Marked Effect on the Interaction between CARMA1 and PKC θ —The N terminus of CARMA1 contains a CARD domain and a coiled-coil domain. The C-terminal region contains a MAGUK-homology structure, which is composed of postsynaptic density 95/dislarge/zona occludens 1 (PDZ), Src homology 3 (SH3), and guanylate kinase (GUK) domains (10).

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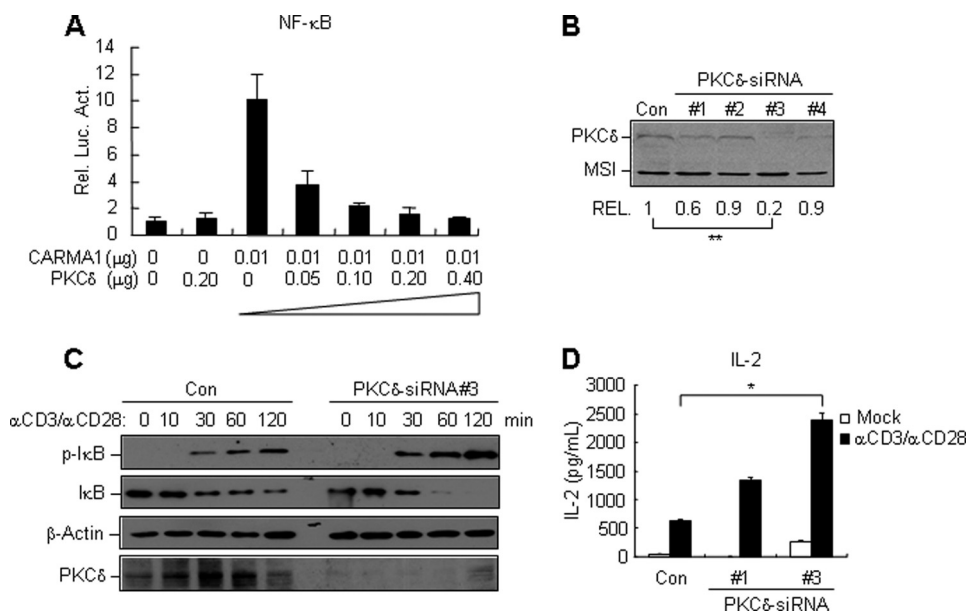


FIGURE 2. PKC δ down-regulates TCR-induced activation of NF- κ B. *A*, PKC δ dose-dependently inhibits CARMA1-induced NF- κ B activation in reporter assays. HEK293 cells were transfected with the indicated mammalian expression plasmids and an NF- κ B reporter plasmid for 18 h, luciferase assays were performed 16 h after transfection. *B*, effects of PKC δ -siRNAs on the expression of PKC δ . The PKC δ -siRNA or control siRNA (green fluorescent protein-siRNA) plasmids, together with Flag-PKC δ and Flag-MSI constructs, were transfected into HEK293 cells. Twenty-four hours after transfection, cell lysates were analyzed by immunoblotting with an anti-Flag antibody. The PKC δ bands from three independent experiments were quantitated using the Bio-Rad Quantity One Program and normalized by levels of the control protein MSI. The average levels of PKC δ are shown at the bottom of the blot. **, $p < 0.01$, $n = 3$. *C*, knockdown of PKC δ potentiates TCR-induced activation of NF- κ B. Jurkat T cells stably transfected with PKC δ -siRNA #3 or control siRNA were stimulated by CD3/CD28 crosslinking for the indicated times. The cell lysates were analyzed by immunoblotting with the indicated antibodies. *D*, knockdown of PKC δ enhances IL-2 secretion upon CD3/CD28 costimulation. Jurkat T cells stably transfected with PKC δ -siRNA #1, #3, or control siRNA were stimulated by CD3/CD28 crosslinking for 48 h. The culture medium was collected and subjected to ELISA. The average levels of IL-2 from three independent experiments are shown as mean \pm S.D., $n = 3$. *, $p < 0.05$.

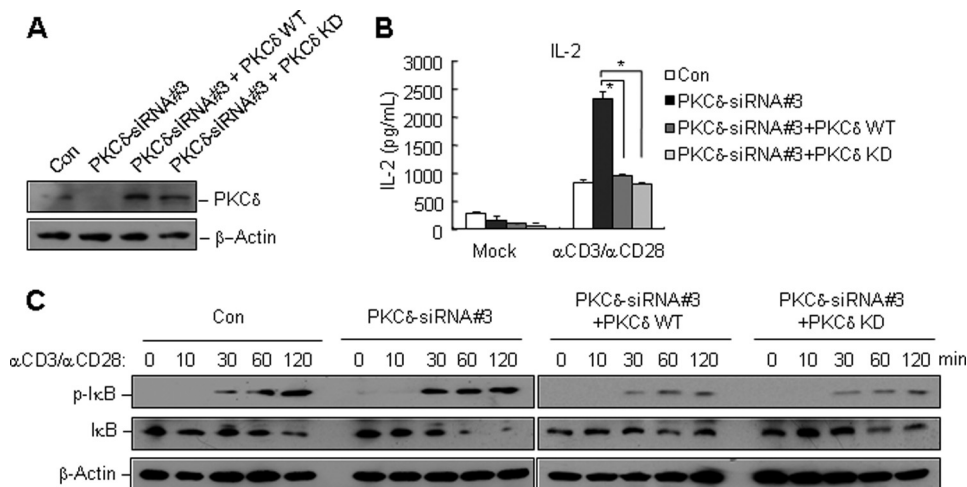


FIGURE 3. The kinase activity of PKC δ is dispensable for its inhibitory role in TCR-induced activation of NF- κ B. *A*, PKC δ -knockdown Jurkat T cells were transfected with MIGR-GFP-PKC δ -WT (wild type) or MIGR-GFP-PKC δ -KD (kinase-dead) mutant by retroviral transduction. Seventy-two hours later, reconstituted cells were sorted by FACS. The efficiency of reconstitution was examined by immunoblotting with an antibody to PKC δ or β -actin. *B* and *C*, PKC δ -knockdown Jurkat T cells, reconstituted Jurkat T cells with PKC δ WT or KD mutants, and control cell lines were stimulated by CD3/CD28 crosslinking. The culture medium was collected and subjected to ELISA for measuring the production of IL-2 (*B*). The cell lysates were applied to immunoblots with indicated antibodies (*C*).

PKC θ , a novel PKC isoform, associates with the PRD region of CARMA1, which links the coiled-coil domain and PDZ domain. PKC θ phosphorylates CARMA1 at Ser-552 and Ser-645 in PRD, which is essential for TCR-induced NF- κ B activation (4, 5, 7). PKC δ , as another novel PKC isoform, shares 85% in catalytic domain and 62.5% in overall protein sequence homology with PKC θ , thus it may interfere with the interaction between CARMA1 and PKC θ through competitive binding.

However, domain-mapping experiments suggested that PKC δ interacted with all the domains consisted in the MAGUK region of CARMA1 but not PRD region (Fig. 4*A*). CARMA1 associated with the catalytic domain (aa 353–668) of PKC δ , but not the regulatory domain (aa 1–352) (Fig. 4*B*). As an unexpected result, PKC δ had no marked effect on the CARMA1-PKC θ interaction, determined by competitive coimmunoprecipitation assays (Fig. 4*C*).

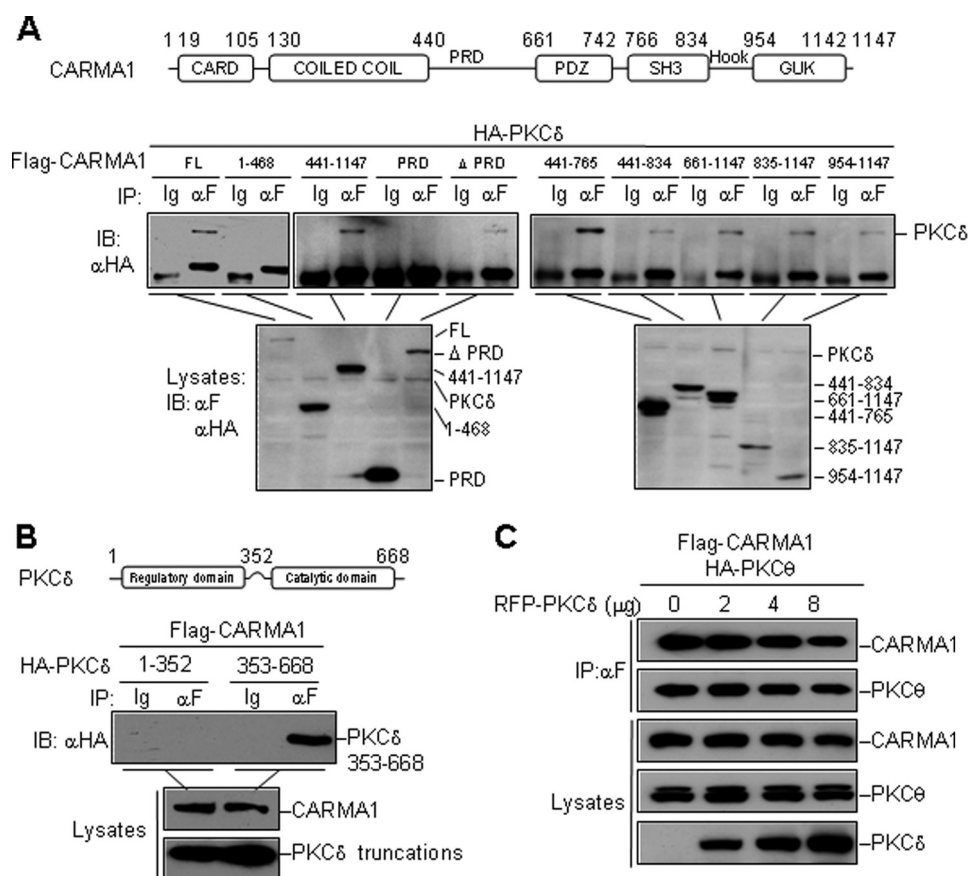


FIGURE 4. PKC δ interacts with MAGUK region of CARMA1 and has no marked effect on CARMA1-PKC θ interaction. *A*, mapping of the minimal PKC δ -interaction region in CARMA1. Transfection, immunoprecipitation and immunoblot assays were performed as described before. *B*, CARMA1 interacts with the catalytic domain of PKC δ (aa 353–668). *C*, PKC δ has no marked effect on CARMA1-PKC θ interaction. Flag-CARMA1 (4 μ g), HA-PKC θ (8 μ g), and RFP-PKC δ (0–8 μ g as indicated) were transfected into HEK293 cells. Lysates were immunoprecipitated with an anti-Flag antibody and then analyzed by immunoblotting with the indicated antibodies (*upper panel*). Expression of the transfected proteins were detected by immunoblotting with antibodies to Flag and HA (*lower panel*).

PKC δ Inhibits the Interaction between MALT1 and TRAF6—We next investigated whether PKC δ acted through inhibiting the assembly of CARMA1 signalosome. The competitive coimmunoprecipitation experiments showed that PKC δ had no marked effect on the interactions between CARMA1-Bcl10, CARMA1-MALT1, or Bcl10-MALT1 (Fig. 5A). In contrast, PKC δ inhibited the association between MALT1 and TRAF6 in a dose-dependent manner (Fig. 5A).

We further performed endogenous coimmunoprecipitation experiments on PKC δ -knockdown Jurkat T cells or control cells to confirm the effect of PKC δ on MALT1-TRAF6 interaction. The results showed that TRAF6 in controls was associated with MALT1 moderately after CD3/CD28 crosslinking, peaked at 30 min, and became weaker after 60 min. While in PKC δ -knockdown cells, obviously more TRAF6 was associated with MALT1 after CD3/CD28 crosslinking. The association peaked at 10 min, and became weaker after 30 min (Fig. 5B). These results suggested that PKC δ inhibited the assembly of CARMA1 signalosome by interfering the association between MALT1 and TRAF6.

DISCUSSION

In this study, we investigated the role of PKC δ in NF- κ B activation in T cells and reported several observations. First, PKC δ participates in the negative control of TCR-induced

NF- κ B activation. Second, the kinase activity of PKC δ is dispensable for the negative regulation. Third, PKC δ is recruited to CARMA1 upon TCR stimulation, and inhibits the subsequent recruitment of TRAF6 by MALT1.

T lymphocytes express multiple PKC isoforms, among which PKC θ is considered as the most important isoform participating in T cell activation (19). It is essential for CARMA1 phosphorylation and further NF- κ B activation. It also phosphorylates RapGEF2 and enhances T cell adhesion (20). Although PKC θ was reported to be the only PKC isoform translocated to immunological synapse after TCR stimulation (21–23), more evidences suggest that other PKCs, such as PKC α , also translocate to the plasma membrane upon TCR stimulation and possibly participate in T cell activation (24). Besides, PKC ζ and PKC ι are PKC θ -associated PKCs in T cells, and possibly assist in PKC θ -mediated signaling (25). In contrast to these PKCs, PKC δ is reported to serve a negative role in TCR-induced IL-2 cytokine production and T cell proliferation (26). However, the mechanism of this negative regulation is not clear so far.

By identifying the CARMA1-associated proteins, we found that PKC δ dynamically interacted with CARMA1 and was translocated to the plasma membrane in a TCR stimulation-dependent manner. It has been shown that following TCR ligation, PKC δ is rapidly phosphorylated at Thr-505 on the activa-

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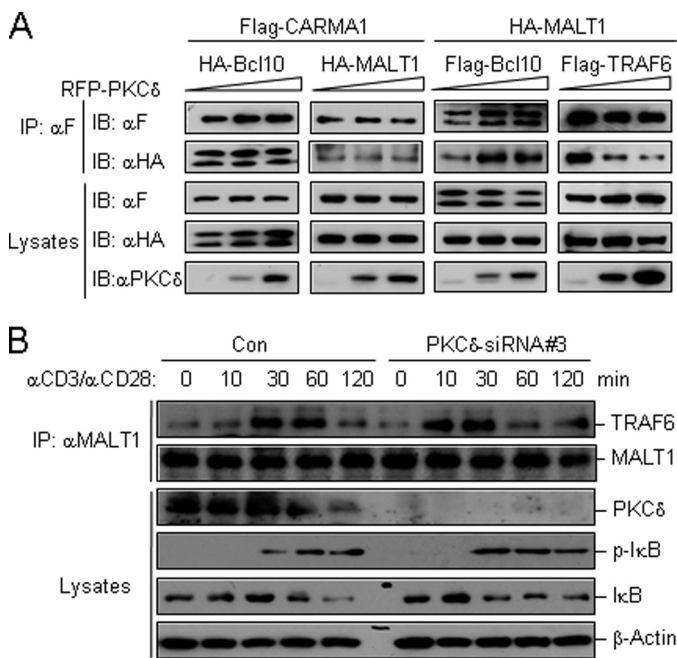


FIGURE 5. PKC δ inhibits the association of MALT1 with TRAF6. *A*, PKC δ disrupts the interaction of MALT1-TRAF6, but not CARMA1-Bcl10, CARMA1-MALT1, and Bcl10-MALT1. The expression plasmids were cotransfected in HEK293 cells as indicated. Immunoprecipitation and immunoblots were performed using the indicated antibodies. *B*, knockdown of PKC δ by siRNA accelerates the recruitment of TRAF6 to MALT1 upon CD3/CD28 costimulation. PKC δ -knockdown Jurkat T cells and controls (5×10^7 each) were stimulated by CD3/CD28 crosslinking for the indicated time. Cell lysates were subjected to immunoprecipitation with an antibody to MALT1. Immunoprecipitates and lysates were immunoblotted using the indicated antibodies.

tion loop (17, 27). This phosphorylation leads to the activation of PKC δ , which regulates lytic granule exocytosis in CTLs in a kinase-dependent manner (17). In this study, however, we found that the kinase activity of PKC δ was not required for its regulation of TCR-induced NF- κ B activation, because the promotion of either I κ B phosphorylation or IL-2 production in PKC δ -knockdown Jurkat T cells could be reversed by reconstitution with kinase-dead mutant of PKC δ , similar to wild-type PKC δ . These surprising findings suggest that negative regulation of TCR-signaling by PKC δ may be mediated through steric hindrance of the assembly of CARMA1 signalosome.

Although PKC δ and PKC θ share high protein sequence homology, they are associated with different region of CARMA1: PKC δ interacts with the MAGUK region, and PKC θ interacts with the PRD region. These findings indicated that PKC δ did not function through competitive binding to CARMA1 with PKC θ , which was confirmed by our competitive binding assays. Upon TCR stimulation, phosphorylated CARMA1 forms a CBM complex with Bcl10 and MALT1. Then MALT1 directly recruits TRAF6 and TAK1 into the CARMA1 signalosome (28, 29). TRAF6 functions as an E3 ligase to catalyze the K63-linked polyubiquitination of MALT1, Bcl10, and IKK γ , and thus promotes signaling to NF- κ B activation (28, 30). Our results showed that PKC δ had no apparent effect on the association of CARMA1 with PKC θ , Bcl10, or MALT1. Instead, PKC δ inhibited the interaction between MALT1 and TRAF6 in a dose-dependent manner, whereas knockdown of PKC δ enhanced the TCR-induced recruitment of endogenous

TRAF6 to MALT1. These results provide an explanation for the inhibitory role of PKC δ in TCR-induced NF- κ B activation: PKC δ is recruited to CARMA1 signalosome upon TCR stimulation and inhibits the recruitment of essential downstream signal components, such as TRAF6.

PKC δ -deficient mice were reported to be susceptible to autoimmune disease, indicating that PKC δ plays a regulatory role in immune systems. Although the development of immune organs in PKC δ -deficient mice appears to be normal, cultured PKC δ -deficient lymphocytes, both B and T cells, display enhanced proliferation in response to various mitogenic stimuli (26, 31). Since signal transduction from antigen receptors in both B and T cells to NF- κ B shares the same downstream signaling events through CARMA1-Bcl10-MALT1 complex, our results might also explain the enhanced proliferation of PKC δ -deficient B cells. Taken together, our findings suggest that PKC δ negatively regulates TCR-induced NF- κ B activation and IL-2 production by inhibiting the assembly of CARMA1 signalosome.

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