

Original Article

Protein kinase C α inhibits myocardin-induced cardiomyocyte hypertrophy through the promotion of myocardin phosphorylation

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Abstract

Myocardin plays a key role in the development of cardiac hypertrophy. However, the upstream signals that control the stability and transactivity of myocardin remain to be fully understood. The expression of protein kinase C α (PKC α) also induces cardiac hypertrophy. An essential downstream molecule of PKC α , extracellular signal-regulated kinase 1/2, was reported to negatively regulate the activities of myocardin. But, the effect of cooperation between PKC α and myocardin and the potential molecular mechanism by which PKC α regulates myocardin-mediated cardiac hypertrophy are unclear. In this study, a luciferase assay was performed using H9C2 cells transfected with expression plasmids for PKC α and myocardin. Surprisingly, the results showed that PKC α inhibited the transcriptional activity of myocardin. PKC α inhibited myocardin-induced cardiomyocyte hypertrophy, demonstrated by the decrease in cell surface area and fetal gene expression, in cardiomyocyte cells overexpressing PKC α and myocardin. The potential mechanism underlying the inhibition effect of PKC α on the function of myocardin is further explored. PKC α directly promoted the basal phosphorylation of endogenous myocardin at serine and threonine residues. In myocardin-overexpressing cardiomyocyte cells, PKC α induced the excessive phosphorylation of myocardin, resulting in the degradation of myocardin and a transcriptional suppression of hypertrophic genes. These results demonstrated that PKC α inhibits myocardin-induced cardiomyocyte hypertrophy through the promotion of myocardin phosphorylation.

Key words: PKC α , myocardin, cardiomyocyte hypertrophy, phosphorylation

Introduction

Cardiac hypertrophy, an essential adaptive or maladaptive response of myocytes to diverse pathophysiological stimuli, is characterized by increased cardiomyocyte size, elevated protein synthesis, and altered gene expression [1–3]. Numerous transcription factors, such as GATA-binding protein 4, myocyte enhancer factor 2, nuclear factor

of activated T cells, and serum response factor (SRF), have been identified as transcriptional regulators that regulate the expressions of a series of hypertrophic genes through the recruitment of cofactors [4,5].

Myocardin, an SRF coactivator belonging to the secreted aspartyl proteinase superfamily, is specifically expressed in cardiomyocytes and smooth muscle cells. Myocardin potently transactivates CArG

box (CC[A/T]₆GG)-containing cardiac and smooth muscle target genes through an association with SRF [6–10]. Previous studies have confirmed that the overexpression of myocardin induces hypertrophy and fetal cardiac gene expression [9,11,12]. However, the upstream signals that control the stability and transactivity of myocardin remain to be fully understood.

Protein kinase C (PKC), a serine/threonine kinase, plays a key role in the regulation of cardiac contraction, hypertrophy, injury, failure, etc. [13–15]. It has been shown that PKC α is a key regulator of cardiomyocyte hypertrophic growth [16]. Using adenovirus-mediated transfection of wild-type or dominant inhibitory forms of PKC α , - β 2, - δ , and - ϵ in neonatal rat cardiomyocytes, Braz *et al.* [17] demonstrated that the inhibition of PKC α reduces agonist-induced hypertrophy. Similarly, dominant negative PKC α suppresses agonist-induced cardiac hypertrophy [16].

As an essential downstream molecule of PKC α , extracellular signal-regulated kinase 1/2 (ERK1/2) was reported to negatively regulate the activities of myocardin [18]. But, the effect of cooperation between PKC α and myocardin on cardiac hypertrophy and the exact molecular mechanism by which PKC α regulates myocardin-mediated cardiac hypertrophy are unclear. In this study, the relationship between myocardin and PKC α was explored. Our results showed that PKC α inhibits myocardin-induced cardiomyocyte hypertrophy in H9C2 cells overexpressing PKC α and myocardin. Furthermore, PKC α promotes the myocardin phosphorylation, resulting in myocardin degradation and transcriptional activity decrease.

Materials and Methods

Cell culture

Neonatal rat cardiomyocytes were isolated from day 1 to day 3 Sprague-Dawley rat pups, as described previously [19]. The isolated cells were cultured in Dulbecco's modified Eagle's medium/F-12 (DMEM/F-12; Gibco, Gaithersburg, USA) supplemented with 20% fetal bovine serum (FBS) at 37°C in humidified air with 5% CO₂.

The rat cardiomyocyte-derived cell line H9C2 was obtained from American Type Culture Collection (Manassas, USA) and cultured in high glucose DMEM supplemented with 10% FBS.

Cell transfection and luciferase reporter assay

The pcDNA3.1-myocardin expression plasmid encoding mouse myocardin was constructed, as described previously [19]. PKC α cDNA was amplified by polymerase chain reaction (PCR) and subsequently cloned into the pCMV vector. The promoter regions of atrial natriuretic factor (ANF) (–957/+37) were amplified by PCR and cloned into pGL3 luciferase reporter vector. The primers used to generate PKC α -flag and ANF-luc are shown as follows: ANF: forward 5'-AGG AACGCGTGCCTTCACGGATCACTTC-3', reverse 5'-TCTGCTCG AGGCTGTCTCGGCTCACTCT-3' and PKC α : forward 5'-GGTACT CGAGATGGCTGACGTTTCCCG-3', reverse 5'-CCGCGGTACC CATACTGCACTCTGTAAGAT-3'. The luciferase reporter gene plasmid containing the 4 \times CarG box was a gift from Professor Eric N. Olson of the Department of Molecular Biology at the University of Texas Southwestern Medical Center (Dallas, USA).

For the transfection experiments, neonatal rat cardiomyocytes were transfected using TurboFect transfection reagent (Thermo, Waltham, USA), and H9C2 cells were transfected using Lipofectamine 2000 transfection reagent (Invitrogen, Carlsbad, USA). After 6 h of incubation, the medium was replaced with normal culture medium. The luciferase reporter assay was performed in 24-well plates. Twenty-four hours after transfection, the cells were harvested, and luciferase

activity was measured using a luciferase assay system (Promega, Madison, USA).

Immunofluorescence

The cardiomyocytes were fixed with 4% paraformaldehyde for 15 min and treated with 0.25% Triton X-100 for 20 min at room temperature, followed by blocking with normal goat serum for 20 min at room temperature. After overnight incubation with mouse anti- α -actinin (Sigma, St Louis, USA), the cells were subsequently incubated with the fluorescein isothiocyanate-conjugated goat anti-mouse IgG secondary antibody (Santa Cruz Biotechnology, Santa Cruz, USA) for 30 min at 37°C. After washing with phosphate-buffered saline (PBS), the samples were observed under a laser scanning confocal microscope (OLYMPUS, Tokyo, Japan). The fluorescent stain 4',6'-diamidino-2-phenylindole was used to stain the nuclei, and the relative area was analyzed using iPP 6.0 software.

Reverse transcriptase-PCR and quantitative real-time PCR

Reverse transcriptase (RT)-PCR and quantitative real-time PCR (qRT-PCR) analyses were carried out as described previously [19]. qRT-PCR was performed using Biosystems StepOne™ Real-Time PCR system (Applied Biosystems, Foster City, USA), according to the manufacturer's instructions. Fast SYBR® Green Master Mix was obtained from Applied Biosystems. The data were expressed as the relative expression levels after normalization to GAPDH. The PCR primers used in the experiment are as follows: ANF for RT-PCR: forward 5'-GAAGTCAACCCGTCTCA-3', reverse 5'-ATCCTGTCA ATCCTACCC-3'; brain natriuretic peptide (BNP) for RT-PCR: forward 5'-TTCTGCTCCTGCTTTTCCTT-3', reverse 5'-CGGTCTAT CTTCTGCCAAA-3'; GAPDH for RT-PCR and qRT-PCR: forward 5'-ATTCAACGGCACAGTCAAGG-3', reverse 5'-GCAGAA GGGGCGGAGTGA-3'; and qRT-PCR primers: ANF for qRT-PCR: forward 5'-GGGCTTCTTCTCTTCTCTG-3', reverse 5'-CGCTTCA TCGGTCTGCTC-3' and BNP for qRT-PCR: forward 5'-TTCTGC TCCTGCTTTTCC-3', reverse 5'-CTTTTGTAGGGCCTTGGT-3'.

Western blot analysis

After treatment, cells were washed twice with ice-cold PBS and lysed in RIPA buffer (CW BioTech, Beijing, China). The proteins were subsequently electrophoresed on a 12% sodium dodecyl sulfate-polyacrylamide gel (SDS-PAGE), followed by transferring onto a nitrocellulose (NC) membrane and blocking with PBS containing 5% skimmed milk powder (w/v) for 60 min at room temperature. The membranes were immunoblotted with mouse anti-myocardin (Sigma), rabbit anti-PKC α (Beyotime Biotechnology, Haimen, China), rabbit anti-ANF (Abcam, Cambridge, UK), rabbit anti-BNP (Abcam), anti-phospho-ERK1/2 (Beyotime Biotechnology), anti-ERK1/2 (Cell Signaling, Beverly, USA), and anti- β -actin (Santa Cruz) antibodies overnight at 4°C, followed by incubation with horseradish peroxidase (HRP)-conjugated anti-mouse or anti-rabbit secondary antibodies (Li-COR Biosciences, Lincoln, USA) for 1 h at room temperature. The specific proteins were visualized using the Odyssey Infrared Imaging System (Gene Company Limited, Beijing, China). The relative protein expression levels were analyzed using ImageJ software.

Total protein/DNA content ratio analysis

Protein and DNA from cardiomyocytes were measured with a published method described previously [20]. Myocytes were collected

using trypsin and then split into two equal aliquots for protein and DNA measurements. For protein assays, one aliquot was resuspended in protein lysis buffer (50 mM Tris-HCl, pH 7.4, 2 mM ethylenediaminetetraacetic acid (EDTA), 1% NP-40, 100 mM NaCl, and 0.1 mM phenylmethylsulfonyl fluoride). The total cell protein was measured using a BCA protein assay kit (Solarbio, Beijing, China) using bovine serum albumin as the standard. The other aliquot was resuspended in Tris-HCl, EDTA, NaCl (TEN) buffer (10 mM Tris-HCl, pH 7.4, 1 mM EDTA, pH 8.0, and 150 mM NaCl). DNA was measured by using Hoechst 33258 (Solarbio) using Salmon sperm DNA (Solarbio) as the standard. The DNA was analyzed at 365 nm excitation and 460 nm emission using a Synergy4 system (BioTek, Winooski, USAS). Protein-to-DNA ratio was calculated in the extracts obtained from individual wells of each culture tray.

Immunoprecipitation and co-immunoprecipitation analyses

The immunoprecipitation (IP) experiments were performed to detect the phosphorylation of myocardin in H9C2 cells. The protein extracts were isolated from cells using RIPA buffer. The tagged proteins were immunoprecipitated overnight at 4°C using an anti-Myc antibody, followed by the addition of protein A/G agarose (CW0349) and further incubation for 3 h. All complexes were pelleted at 1400 g for 3 min. Proteins bound to the beads were fractionated through 12% SDS-PAGE and transferred to an NC membrane. The membranes were immunoblotted with anti-Myc-tag (Proteintech, Chicago, USA), anti-phospho-threonine (Cell Signaling), anti-phospho-serine (IMMUNECHEM, Burnaby, Canada), and anti- β -actin (Santa Cruz) antibodies overnight at 4°C, followed by incubation with corresponding HRP-conjugated secondary antibodies (Li-COR Biosciences) for 1 h at room temperature. The specific proteins were visualized using the Odyssey Infrared Imaging System (Gene Company Limited).

For Co-IP assay, a similar protocol was followed using COS7 cells to confirm the interaction between myocardin and PKC α . IP was performed using anti-Myc or anti-flag antibodies (Proteintech). The myc- or flag-precipitated proteins were immunoblotted using anti-flag and anti-myc antibodies, respectively.

Statistical analysis

Data were expressed as the mean \pm SEM, and the number of experiments performed independently was also shown. Data were analyzed using Student's *t*-test. Differences at *P* < 0.05 were considered statistically significant.

Results

PKC α decreases the transcriptional activity of myocardin

To determine the relationship between myocardin and PKC α , H9C2 cells were transfected with expression plasmids for myocardin, ANF-luc, and increasing amounts of PKC α , and then luciferase assays were performed. The overexpression of myocardin significantly increased ANF promoter transcriptional activity, whereas co-transfection with expression plasmids for myocardin and PKC α decreased the transcriptional levels of ANF in a dose-dependent manner (Fig. 1A,B).

To further determine whether PKC α represses the transcriptional activity of myocardin, H9C2 cells were transfected with expression plasmids for myocardin and ANF-luc/4 \times CArG-luc and then stimulated with increasing amounts of PMA, an activator of PKC α , followed by analysis using luciferase assay. The stimulation of myocardin-transfected H9C2 cells with increasing amounts of PMA for 12 h

decreased ANF promoter activity in a dose-dependent manner (Fig. 1C). As shown in Fig. 1D,E, the co-transfection of PKC α and stimulation with 50 nM PMA significantly inhibited the transcriptional activity of myocardin. These results demonstrated that PKC α negatively regulates the transcriptional activity of myocardin.

PKC α inhibits myocardin-induced cardiomyocyte hypertrophy

To investigate the role of PKC α in myocardin-induced cardiac hypertrophy, neonatal rat cardiomyocytes were transfected with expression plasmids for myocardin and/or PKC α , followed by starvation for 6 h and stimulation with or without 50 nM PMA for an additional 24 h. The overexpression of myocardin or PKC α increased the cell surface area, whereas co-transfection with the expression plasmids for myocardin and PKC α strongly decreased the cell size (Fig. 2A). The protein/DNA ratio was determined to demonstrate cellular hypertrophy (Fig. 2B). The overexpression of myocardin or PKC α resulted in a significant increase in the protein/DNA ratio. Myocardin-induced increase in the protein/DNA ratio was attenuated by the overexpression of PKC α . As shown in Fig. 2B, myocardin-activated ANF and BNP mRNA levels were significantly inhibited by PKC α . PKC α also markedly reduced myocardin-stimulated ANF and BNP expressions (Fig. 2C, D). Interestingly, co-transfection with the expression plasmid for PKC α also suppressed the myocardin level (Fig. 2E,F). Thus, these results showed that PKC α inhibits myocardin-induced cardiomyocyte hypertrophy.

PKC α inhibitor recovers the PKC α -mediated repression of myocardin transcriptional activity and enhances the expression of the myocardin target genes ANF and BNP

To further determine whether PKC α inhibits myocardin-induced cardiomyocyte hypertrophy, H9c2 cells were co-transfected with expression plasmids for myocardin and PKC α . In addition, cells were pre-treated for 1 h with 100 nM Calphostin C, an inhibitor of PKC α , followed by stimulation with 50 nM PMA for 24 h. As shown in Fig. 3A,B, compared with cardiomyocytes overexpressing both myocardin and PKC α , cells pre-treated with the PKC α inhibitor showed significantly increased surface area and protein/DNA ratio with a hypertrophic growth phenotype. Furthermore, PKC α strongly inhibited myocardin-mediated ANF and BNP expressions, whereas pre-treatment with the PKC α inhibitor dramatically increased the levels of these markers in the presence of myocardin and PKC α (Fig. 3C,D). The PKC α inhibitor also partially restored the myocardin levels (Fig. 3E,F). When pre-treated with Calphostin C for 1 h followed by stimulation with PMA for 12 h, co-transfected cardiomyocytes showed a significant increase in ANF and 4 \times CArG promoter activities (Fig. 3G).

PKC α promotes phosphorylation of myocardin

To confirm the effect of Calphostin C on the function of PKC α , the nuclear translocation of PKC α was tested by immunofluorescent assays. As shown in Fig. 4A, PMA promoted the nuclear translocation of PKC α , whereas Calphostin C inhibited the nuclear translocation of PKC α . To confirm the effect of the ERK inhibitor, U0126, on the phosphorylation of ERK1/2, the phosphorylation of ERK1/2 was detected with anti-phos-ERK1/2 antibodies, and the total expression of ERK1/2 was detected with anti-ERK1/2 antibodies by western blot analysis. As shown in Fig. 4B, U0126 completely counteracted PMA-induced ERK phosphorylation.

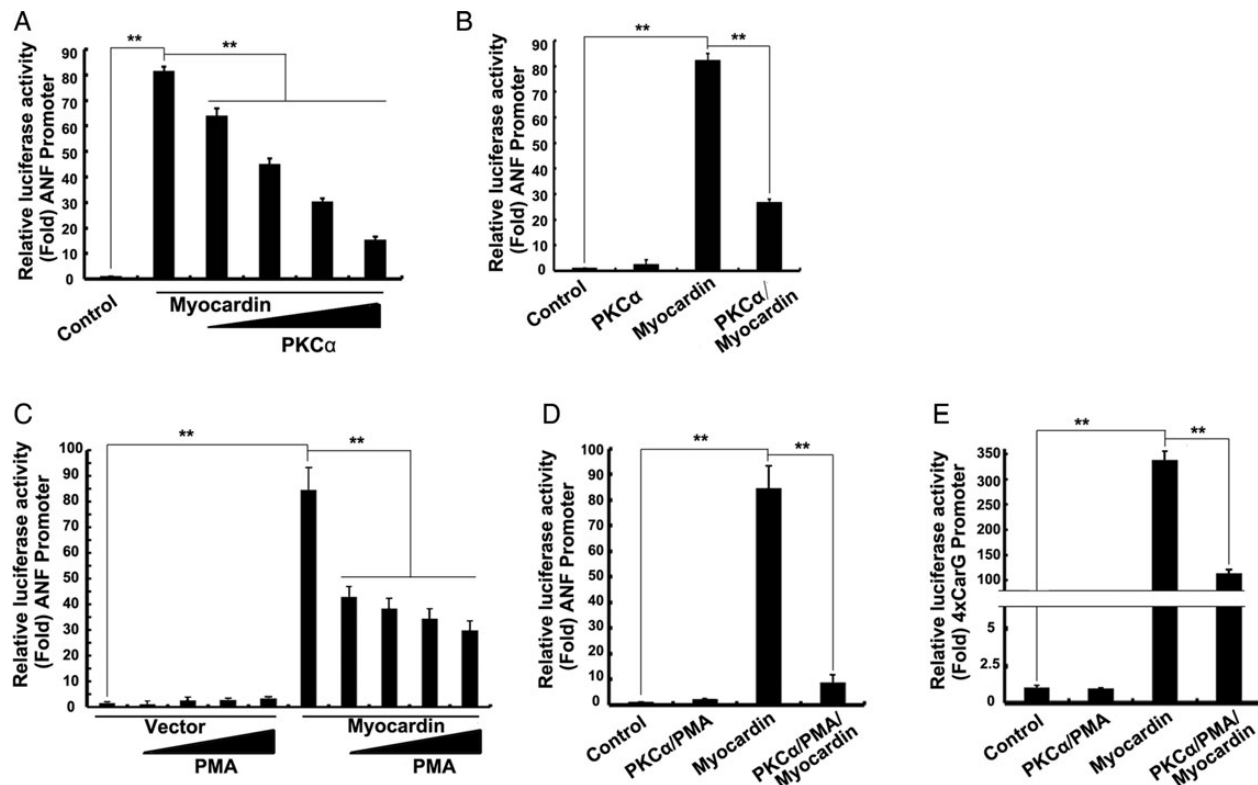


Figure 1. PKC α decreases the transcriptional activity of myocardin (A,B) H9C2 cells were transfected with expression plasmids for myocardin, ANF-luc, and increasing amounts of PKC α (0.3, 0.4, 0.5, and 0.6 μ g). Subsequently, luciferase assays were performed. (C) H9C2 cells were transfected with expression plasmids for myocardin and ANF-luc, starved for 6 h, and stimulated with increasing amounts of PMA (0, 50, 100, 150, and 200 nM) for 12 h, followed by analysis using luciferase assays. (D,E) After co-transfection with expression plasmids for myocardin and/or PKC α , the starved cells were treated with 50 nM PMA for 12 h. The promoter activities of ANF and 4xCarG were analyzed in H9C2 cells. ** $P < 0.01$, $n = 3$.

Myocardin is phosphorylated through either ERK1/2 or glycogen synthase kinase 3 β (GSK-3 β), and phosphorylation reduces the transcriptional activity of this protein, resulting in the significant attenuation of cardiac hypertrophy [21,22]. In this study, PKC α was also found to inhibit myocardin transcriptional activity. Thus, we investigated whether PKC α promotes myocardin phosphorylation. First, the phosphorylation of endogenous myocardin was examined in the presence of PMA. As shown in Fig. 4C, endogenous myocardin was phosphorylated at serine and threonine residues, whereas the inhibition of PKC α /MAPK/ERK through Calphostin C or U0126 impaired myocardin phosphorylation. Next, the phosphorylation of myocardin was detected in H9C2 cells overexpressing myocardin-myc and PKC α -flag. As shown in Fig. 4D, PKC α induced excessive myocardin phosphorylation in cardiomyocytes overexpressing myocardin, and this increased phosphorylation was inhibited by the PKC α -specific inhibitor Calphostin C. Interestingly, U0126, which almost completely inhibits the PMA-induced phosphorylation of myocardin, partially suppressed PKC α -induced phosphorylation at serine residues, but was ineffective for the inhibition of phosphorylation at threonine residues (Fig. 4D). This result suggested that myocardin can be phosphorylated by PKC α directly and by ERK1/2 as well. To further determine whether PKC α interacts with myocardin, H9C2 cells were transfected with expression plasmids for PKC α -flag and/or myocardin-myc, followed by the Co-IP assay. The nuclear proteins were immunoprecipitated using anti-flag or anti-myc antibody. Western blot analysis of the Co-IP experiments showed that PKC α directly interacts with myocardin (Fig. 4E).

Phosphorylation of myocardin promotes the degradation of myocardin through PKC α

It has been reported that myocardin phosphorylation leads to the degradation of myocardin [23]. We also found that PKC α activation decreased the levels of myocardin protein in cells co-transfected with expression plasmids for myocardin and PKC α in the presence of PMA. However, the myocardin mRNA levels remained unaffected, as shown in Fig. 2B,D. Thus, we hypothesized that myocardin phosphorylation results in the degradation of myocardin through PKC α . To confirm this hypothesis, cardiomyocytes were co-transfected with expression plasmids for myocardin and PKC α , followed by treatment with the proteasome inhibitor MG-132 (1 μ M) and 50 nM PMA for 24 h. As shown in Fig. 5A,B, PKC α overexpression did not influence myocardin mRNA levels in H9C2 cells, but decreased myocardin protein levels. When co-transfected cells were incubated with the proteasome inhibitor MG-132, the myocardin protein levels were strongly up-regulated, and myocardin degradation was significantly inhibited (Fig. 5C). These data suggested that PKC α mediates the degradation of myocardin, thereby inhibiting the expression of myocardin.

Discussion

As a coactivator of SRF, myocardin is required to induce cardiac hypertrophy and fetal cardiac gene program [11]. In addition, PKC α regulates the hypertrophic growth of cardiomyocytes through ERK1/2 [18]. Consistent with the results of previous studies [11,24], we also

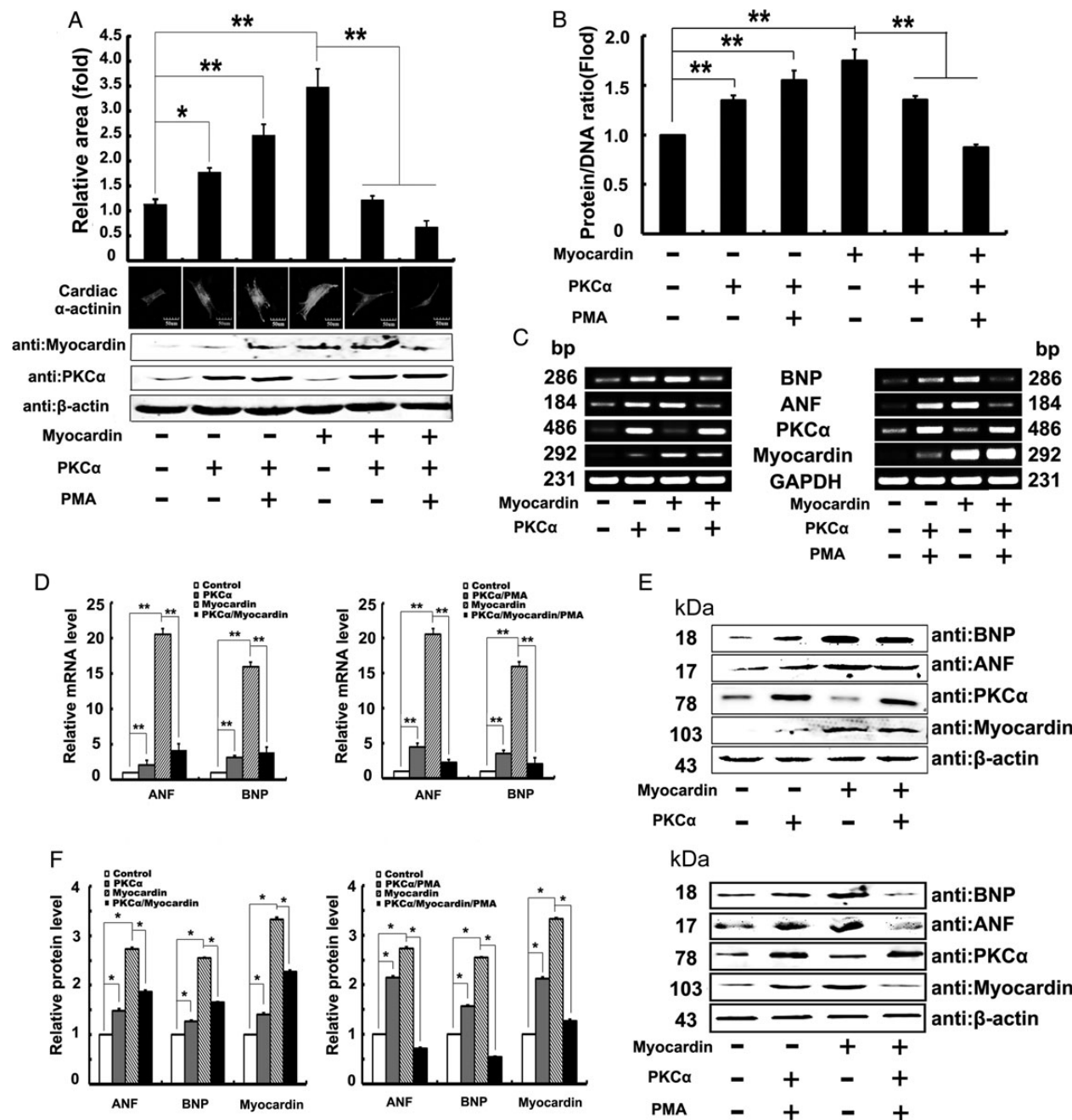


Figure 2. PKCα inhibits myocardin-induced cardiomyocyte hypertrophy (A) Neonatal rat cardiomyocytes were transfected with myocardin and/or PKCα expression plasmids, starved for 6 h, and subsequently stimulated with or without 50 nM PMA for 24 h. The morphological changes of cardiomyocytes were observed under a phase-contrast microscope. The relative area of cardiomyocytes was analyzed using iPP 6.0 software, and the results were normalized to the control. ** $P < 0.01$, $n = 3$. (B) Cellular hypertrophy was determined by measuring the protein/DNA ratio. (C,D) The left panel shows the results of the qRT-PCR analysis of ANF and BNP mRNA levels in H9C2 cells co-transfected with expression plasmids for myocardin and/or PKCα. The right panel shows the results of the qRT-PCR analysis of ANF and BNP mRNA levels in co-transfected H9C2 cells with PMA stimulation. ANF and BNP mRNA levels were analyzed using ImageJ software, and the results were normalized to GAPDH. ** $P < 0.01$, $n = 3$. (E,F) Western blot analysis of BNP, ANF, PKCα, and myocardin expressions after co-transfection with expression plasmids for myocardin and PKCα with (upper panel) or without PMA stimulation (lower panel). The relative protein expression levels were analyzed using ImageJ software, and the results were normalized to β-actin. * $P < 0.05$; ** $P < 0.01$, $n = 3$.

found that the overexpression of either myocardin or PKCα increased the cell surface area and induced cardiac hypertrophy. Furthermore, we examined the relationship between two transcription factors, myocardin and PKCα, in the regulation of cardiomyocyte hypertrophy. Co-transfection with expression plasmids for PKCα strongly decreased the cell size and reduced cardiac hypertrophy induced by

myocardin. These findings suggested that PKCα can inhibit myocardin function. Luciferase reporter assays also confirmed the inhibitory effect of PKCα on the transcriptional activity of myocardin using a luciferase reporter gene plasmid containing 4×CARG box. The results showed that PKCα overexpression suppressed the transcriptional activity of myocardin (Fig. 1E). Furthermore, the PKCα inhibitor

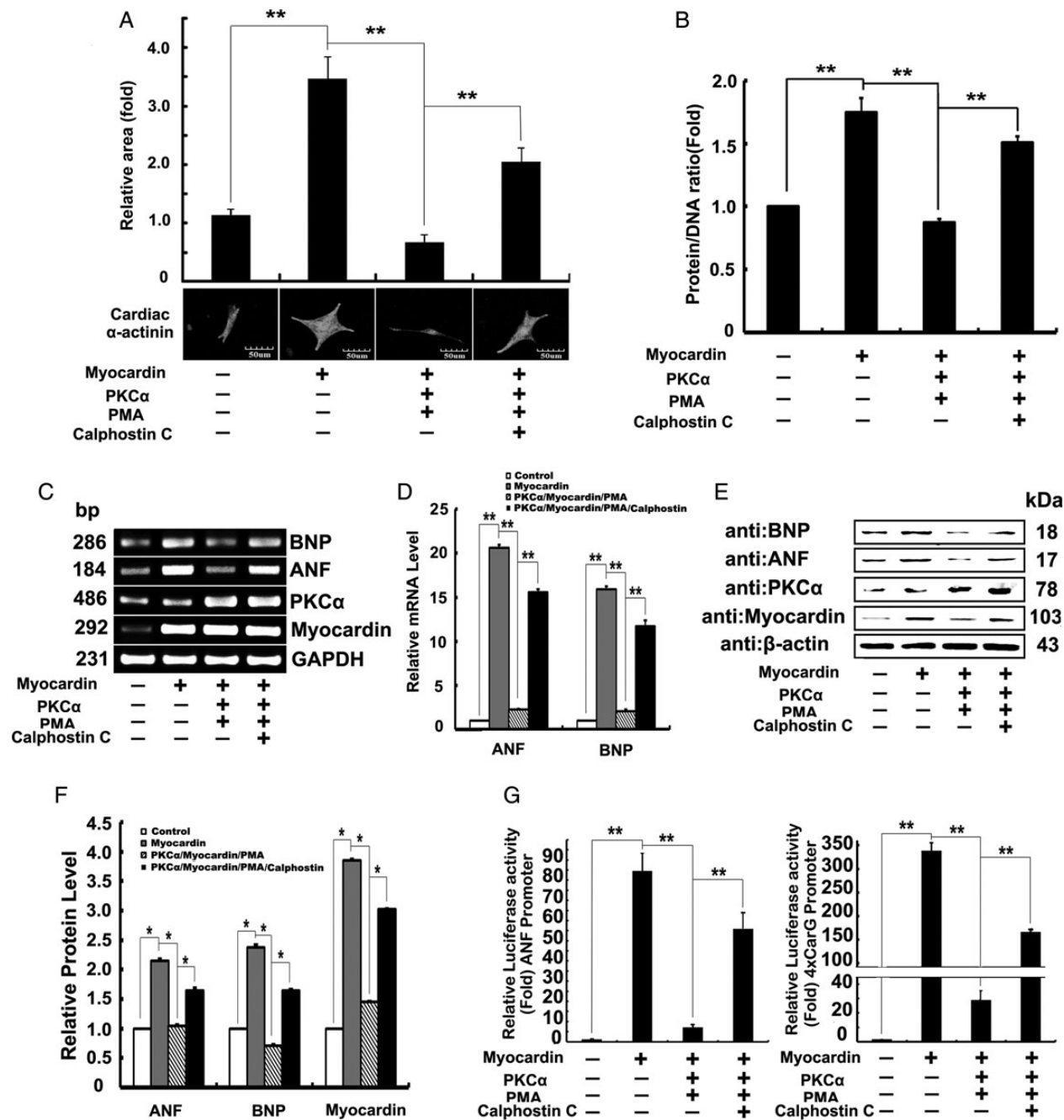


Figure 3. PKCα inhibitor recovers the PKCα-mediated repression of myocardin transcriptional activity and enhances the expression of myocardin target genes ANF and BNP (A) Neonatal cardiomyocytes were co-transfected with expression plasmids for myocardin and/or PKCα, starved for 6 h, pre-treated with 100 nM Calphostin C for 1 h, and subsequently incubated with 50 nM PMA for 24 h. The morphological changes of cardiomyocytes were observed under a phase-contrast microscope. The relative area of cardiomyocytes was analyzed using iPP 6.0 software, and the results were normalized to the control. $**P < 0.01$, $n = 3$. (B) Cellular hypertrophy was determined by measuring the protein/DNA ratio. $**P < 0.01$, $n = 3$. (C,D) The mRNA levels of ANF and BNP were examined in H9C2 cells using qRT-PCR. ANF and BNP mRNA levels were analyzed using ImageJ software, and the results were normalized to GAPDH. $**P < 0.01$, $n = 3$. (E,F) Western blot analysis of myocardin, PKCα, ANF, and BNP expressions. The relative protein expression levels were analyzed using ImageJ software, and the results were normalized to β-actin. $*P < 0.05$, $n = 3$. (G) A luciferase assay was performed after transfection with ANF-luc (left panel) or 4xCarG-luc (right panel) and indicated combinations of expression vectors and stimulatory factors as described in (A). $**P < 0.01$, $n = 3$.

recovered the PKCα-mediated repression of myocardin transcriptional activity and enhanced the expression of the myocardin target genes ANF and BNP (Fig. 3). In addition, myocardin overexpression did not affect the expression of PKCα in cardiomyocytes (Figs. 2D and 3D). These results suggested that the overexpression of PKCα reduced myocardin-induced cardiac hypertrophy.

The impaired myocardin function induced by PKCα was restored after pre-treatment with Calphostin C. Similarly, the activities of myocardin and myocardin-related transcription factors (MRTFs) are also negatively regulated through ERK1/2, an essential downstream molecule of PKCα [21,22]. Purcell *et al.* [25] reported that p65, which could promote the development of cardiac hypertrophy, inhibited

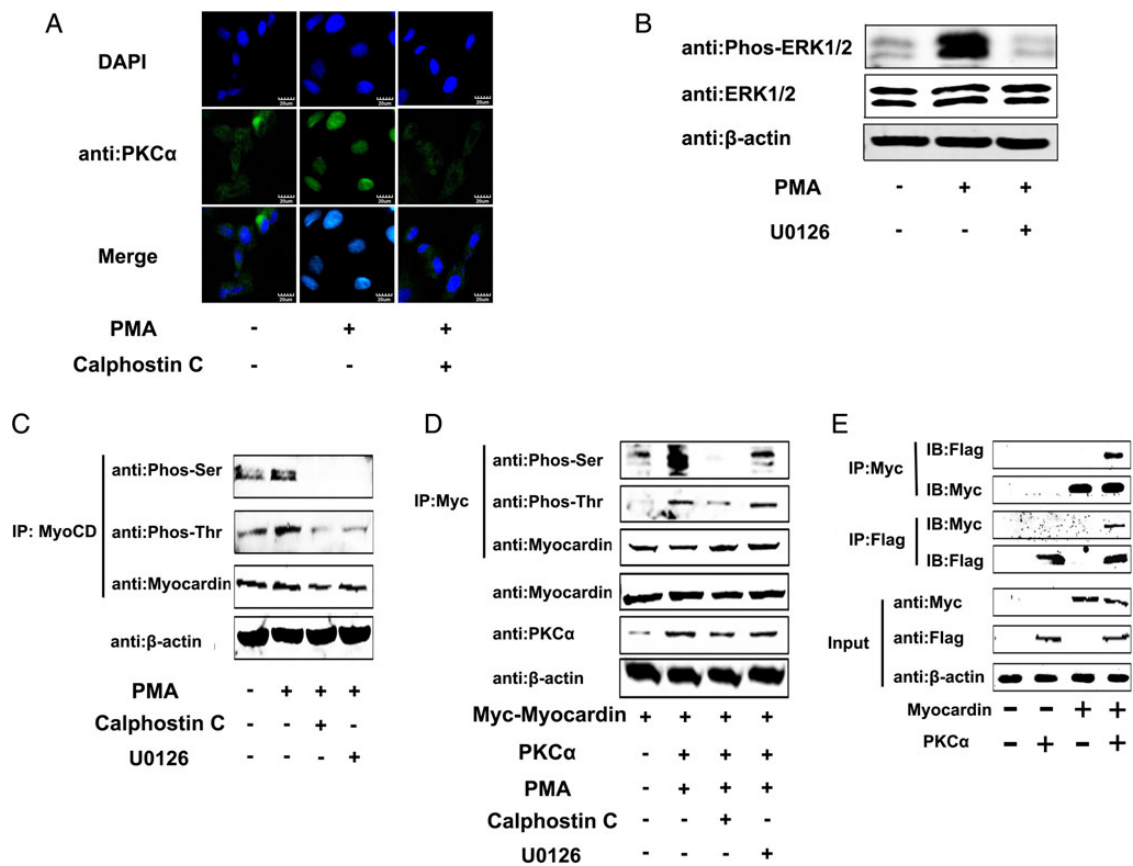


Figure 4. PKCα promotes phosphorylation of myocardin (A,B) H9C2 cells were pre-treated with 100 nM Calphostin C or 10 μM U0126 for 1 h and subsequently stimulated with 50 nM PMA for 24 h. The nuclear translocation of PKCα was detected by immunofluorescent assay and the phosphorylation of ERK1/2 was detected by western blot analysis, respectively. (C) H9C2 cells were pre-treated with 100 nM Calphostin C or 10 μM U0126 for 1 h and then stimulated with 50 nM PMA for 24 h. The proteins were immunoprecipitated with the anti-myocardin antibody, and myocardin phosphorylation was detected by western blot analysis using anti-phos-Ser or anti-phos-Thr antibodies. (D) H9C2 cells were co-transfected with expression plasmids for myocardin-myc and PKCα-flag, starved for 6 h, pre-treated with 100 nM Calphostin C or 10 μM U0126 for 1 h, and subsequently stimulated with 50 nM PMA for 24 h; then myocardin phosphorylation was detected. (E) COS7 cells were transfected with expression plasmids for PKCα-flag and/or myocardin-myc, followed by the Co-IP assay.

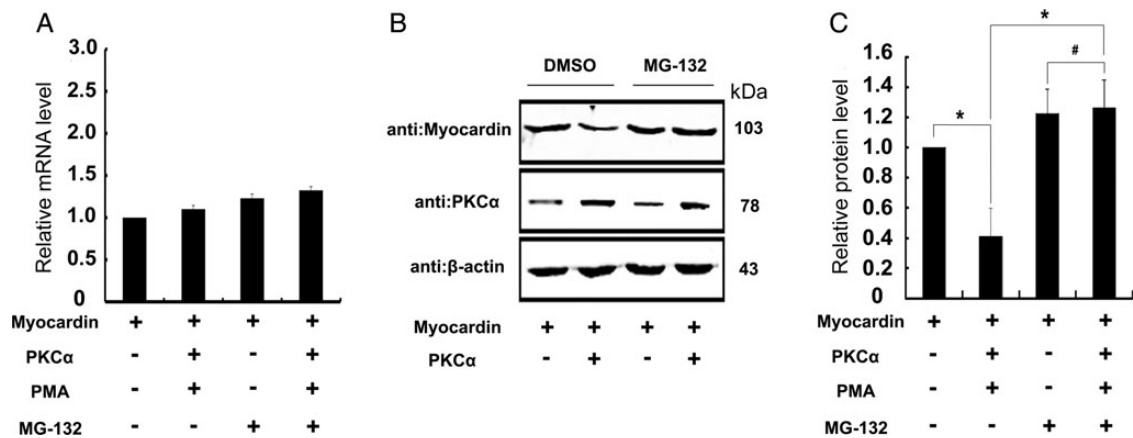


Figure 5. Phosphorylation of myocardin through PKCα promotes the degradation of myocardin (A,B) H9C2 cells were co-transfected with expression plasmids for myocardin and PKCα, and subsequently the cells were treated with a combination of 50 nM PMA and 1 μM MG-132 for 24 h. Myocardin expression was analyzed by qRT-PCR and western blot analysis. (C) The relative levels of myocardin protein were analyzed using ImageJ software, and the results were normalized to β-actin. * $P < 0.05$, # $P > 0.05$, $n = 3$.

myocardin transactivity by physically interacting with it. Liao *et al.* [19] showed that p65 inhibited myocardin-mediated cardiomyocyte hypertrophy by epigenetic modifications.

The activity of myocardin is regulated through post-translational modifications such as phosphorylation, ubiquitination, etc. For example, myocardin phosphorylation inhibits its transactivity.

The phosphorylation of myocardin through ERK1/2 reduces smooth muscle gene transcription [18], and the phosphorylation of myocardin through GSK-3 β suppresses cardiac hypertrophy through the reduced transcriptional activation of myocardin [26]. MRTF-A is phosphorylated through ERK1/2, leading to the inhibition of MRTF-A activity [21,22]. In this study, we demonstrated that endogenous myocardin is slightly phosphorylated through PMA. PKC α might promote myocardin phosphorylation through the activation of ERK1/2 [18]. Consistent with the results of previous studies [21,22], we demonstrated that specific inhibitors of PKC α or ERK1/2 reduced the phosphorylation of endogenous myocardin (Fig. 4A). Similarly, co-transfection with expression plasmids for PKC α in cardiomyocytes significantly increased myocardin serine phosphorylation. This increased phosphorylation is almost completely inhibited by Calphostin C, but partially inhibited by U0126. The effect of U0126 suggested that myocardin can be phosphorylated by PKC α directly and by ERK1/2 as well. Zhang *et al.* [27] have revealed that PKC α can be transferred to the nucleus after being activated by PMA. Thus PKC α may be activated and located into the nucleus, where PKC α can directly phosphorylate myocardin. Our Co-IP assay results showed that PKC α directly interacts with myocardin. These results suggested that PKC α directly suppresses the transcriptional activity of myocardin through promoting the phosphorylation of myocardin.

Furthermore, the phosphorylation of myocardin through GSK-3 β was observed, and E3 ligase C terminus of Hsc70-interacting protein (CHIP) revealed the ubiquitin-mediated degradation of myocardin in the proteasome [23]. ERK1/2 phosphorylates myocardin, but does not cause myocardin degradation [18]. The results obtained in the present study showed that PKC α phosphorylates and promotes the degradation of myocardin, suggesting that PKC α and ERK1/2 phosphorylate myocardin at different sites. All these data suggested that phosphorylation through PKC α attenuates the transcriptional activity and promotes the degradation of myocardin.

Yoshida *et al.* [28] reported the role of Kruppel-like factor 4 (KLF4) in myocardin-induced cardiomyocyte hypertrophy using KLF4 knockout mice and cell line H9C2. Overexpression of KLF4 inhibits myocardin-induced cellular enlargement by decreasing the expression and activity of myocardin in cardiac cells. In this study, our results showed that PKC α could inhibit myocardin-induced cardiomyocyte hypertrophy by directly promoting myocardin phosphorylation and degradation.

In summary, these results demonstrated that PKC α inhibits myocardin-induced cardiomyocyte hypertrophy. PKC α directly phosphorylates and promotes the degradation of myocardin, thereby decreasing its transcriptional activity. These findings provide new insight into the interaction between PKC α and myocardin in cardiac hypertrophy.

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