

EFFECTS OF CALCIUM CHANNEL ANTAGONISTS ON THE PHOSPHORYLATION OF MAJOR PROTEIN KINASE C SUBSTRATES IN THE RAT HIPPOCAMPUS

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Abstract—K⁺-induced depolarization of rat hippocampal slices resulted in significant increases in the phosphorylation state of myristoylated, alanine-rich C kinase substrate (MARCKS; also known as 87K, pp80) and neuromodulin [also known as growth associated protein 43 (GAP43), B50, F1] as determined by back-phosphorylation using protein kinase C. The effect of organic and inorganic Ca²⁺ antagonists on the phosphorylation of these major protein kinase C substrates in the rat hippocampus was studied to determine whether Ca²⁺ influx through L- or N-type voltage-sensitive Ca²⁺ channels was required for the phosphorylation changes observed. The depolarization-induced changes appeared to be dependent on extracellular Ca²⁺, based on evidence indicating that the chelation of extracellular Ca²⁺ with ethylene glycol-bis (β-amino-ethyl ether)-N,N,N',N'-tetraacetic acid (EGTA) inhibited these changes. In addition, pretreatment of the slices with 500 μM Cd²⁺, but not 300 nM nimodipine, 10 μM ω-conotoxin GVIA or 10 μM MK-801, blocked the K⁺-induced change in phosphorylation. These results suggest that K⁺-induced changes in the phosphorylation of MARCKS and neuromodulin are mediated by Ca²⁺-dependent mechanisms other than, or in addition to, those sensitive to the organic Ca²⁺ channel antagonists employed.

Nimodipine, a 1,4-dihydropyridine Ca²⁺ channel antagonist, has a variety of effects on the brain that include neuroprotection, accelerated neural regeneration and cognition enhancement [1,2]. In neuronal systems, Ca²⁺ and various protein kinases are implicated in mechanisms of ischemic cell damage, neuroplasticity and in electrophysiological models of imprinting [3–8]. In the present study, the effects of nimodipine and other Ca²⁺ blockers were investigated to determine whether these treatments modulated depolarization- and extracellular Ca²⁺-dependent phosphorylation of hippocampal proteins by protein kinase C.

Different methods have been used to study drug effects on Ca²⁺-dependent protein phosphorylation. The use of tissue homogenates incubated with drug, cofactors and [γ -³²P]ATP relies on endogenous kinase and phosphatase activity that may not be representative of the *in situ* state. An alternative *in vivo* method labels the intracellular ATP pool of intact tissue or cells with radioactive ³²PO₄ prior to drug treatment. In both methods the substrates of interest are isolated using immunoprecipitation techniques or electrophoretic separation. As critically discussed by Nestler and Greengard [9], the *in vivo* studies are dependent upon the consistency of labeling the ATP pool, and the high background,

due to the use of millicurie amounts of ³²P, may make direct quantitation difficult. In using either method, the comigration on gel electrophoresis of numerous phosphorylated proteins may make identification difficult. Back-phosphorylation techniques, properly conducted, can determine the state of phosphorylation of discrete substrates for a specific kinase [9]. We report here for the first time a back-phosphorylation procedure using protein kinase C to study the mechanism of action of Ca²⁺ channel antagonists.

MATERIALS AND METHODS

Materials. Drugs, chemicals and protein kinase C were obtained as follows: nimodipine (Miles Inc.); ω-conotoxin GVIA (Peninsula Laboratories Inc.); (±) MK-801 (Research Biochemicals Inc.); Na₂HPO₄, KH₂PO₄, NaOH, KCl, CdCl₂, NaCl, CaCl₂ (Baker); acrylamide, bis-acrylamide, piperazine diacrylamide (PDA⁺), urea, sodium dodecyl sulfate (SDS, Ultrapure grade), glycine, 2-mercaptoethanol, N,N,N',N'-tetra-methylethyldiamine (Bio-Rad); N-(2-hydroxyethyl)piperazine-N'-2-ethanesulfonic acid (HEPES) (Calbiochem); phosphoric acid (H₃PO₄) (EM Science); ampholines (40%) pH 5–7, pH 5–10 (LKB Pharmacia); [γ -³²P]ATP (10–30 Ci/mmol) (New England Nuclear); aprotinin, leupeptin, dithiothreitol, Nonidet P-40, sodium phosphatidylserine, 1,2-dioleoyl-rac-glycerol, magnesium acetate, citric acid, ethylene glycol-bis-(β-amino-ethyl ether)-N,N,N',N'-tetraacetic acid (EGTA), CaCl₂, Trizma Base and prestained molecular weight markers "SDS 7B" (Sigma Chemical).

The protein kinase C was purified from rat

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† Abbreviations: EGTA, ethylene glycol-bis (β-amino-ethyl ether)-N,N,N',N'-tetraacetic acid; HEPES, N-(2-hydroxyethyl)piperazine-N'-2-ethanesulfonic acid; MARCKS, myristoylated, alanine-rich C kinase substrate; PDA, piperazine diacrylamide; and SDS, sodium dodecyl sulfate.

forebrain [10] and was supplied by Drs. A. C. Nairn and P. Greengard of Rockefeller University, New York, NY.

Equipment. The Bio-Rad model 175 tube cell and Protean II 20-cm multicell 2D Vertical Electrophoresis Systems were used for protein separation and the AMBIS Mark II Radioanalytic Imaging System for quantitative 2D gel analysis.

Methods. Phosphorylation changes were determined using a modification of the direct back-phosphorylation technique of Guitart and Nestler [11], which measures the dephospho form of the protein. The general procedure was as follows: (1) treatment of hippocampal slices; (2) homogenization and extraction of slices with acidification (pH 2.9) to inactivate endogenous kinases and phosphatases, thereby maintaining the phosphorylation state of the proteins; (3) centrifugation of homogenates resulting in an acid extract composed of soluble and membrane-bound proteins; (4) neutralization of the extract followed by phosphorylation with purified protein kinase C and [γ - 32 P]ATP; and (5) electrophoretic separation of the phosphorylated proteins using two-dimensional polyacrylamide gel electrophoresis. Since only the dephospho form of the protein is phosphorylated in this system, a decrease in phosphorylation generally reflects an increase in the phosphorylation state of the protein [11, 12].

Experiments were performed *ex vivo* using hippocampal slices (400 μ m) obtained from 3- to 9-month-old male Long-Evans Hooded rats. The slices were placed, two per tube, in 5 mL of a low Ca^{2+} Krebs-bicarbonate buffer (pH 7.33) of the following composition (in mM): NaCl, 124; KCl, 3.33; CaCl_2 , 0.01; KH_2PO_4 , 1.25; MgSO_4 , 1.33; NaHCO_3 , 25.7; D-glucose, 10; HEPES, 20. The slices were equilibrated for 60 min at 33–34° with water-saturated oxygenation (95% O_2 , 5% CO_2). After 60 min the solution was replaced with Krebs buffer containing a physiological level of Ca^{2+} (1.3 mM) and incubated for an additional 90 min [13]. After the pre-equilibration period, the slices were pretreated for 15 min with either vehicle or drug. The tissue was depolarized by increasing the K^+ concentration 50 mM by the addition of 250 μ L of a 1 M KCl solution, and the slices were incubated for an additional 5 min. The buffer was removed and the slices sonicated in 300 μ L of acidified homogenization buffer (pH 2.9) containing 20 mM citric acid, 0.1% Nonidet P-40, 10 mM Tris, 1 mM EGTA, 1 mM dithiothreitol, 0.1 trypsin inhibitor units/mL aprotinin, and 10 μ g/mL leupeptin [modified from Ref. 11].

Acidified homogenates (pH 2.9) were centrifuged at 10,000 *g* for 15 min at 4°. Aliquots of the supernatant, containing acid-soluble proteins, were neutralized with the addition of 200 mM Na_2HPO_4 (20%, v/v) [11, 14]. Protein content was determined by the method of Bradford [15], using bovine serum albumin as the standard. An aliquot (\approx 15–30 μ L) containing 10 μ g protein was back-phosphorylated in a reaction solution (total volume 80 μ L) containing 50 mM Tris, pH 7.4, 10 mM magnesium acetate, 1 mM EGTA, 10 mM dithiothreitol, 0.5% Nonidet P-40, 10 μ g phosphatidylserine, 0.4 μ g dioleoyl-rac-glycerol, 1.8 mM Ca^{2+} , 0.01% bovine serum albumin,

10 μ g leupeptin, 60 ng protein kinase C and 8 μ M [γ - 32 P]ATP (10–30 Ci/mmol) [modified from Ref. 16]. The reaction was initiated by the addition of [γ - 32 P]ATP and incubated for 30 min at 30°. The reaction was terminated and proteins were denatured with 47 mg urea and 8 μ L stop solution [final concentration: 9 M urea, 2% Nonidet P-40, 2% ampholines (ratio of pH 3–10 to pH 5–7 1:1), 5% 2-mercaptoethanol] [11].

Samples were subjected to two-dimensional electrophoretic separation using a modification of the method of O'Farrell [17]. Isoelectric focusing tube gels (2 mM) containing 9 M urea, 4% acrylamide (ratio of acrylamide to PDA, a cross-linker, was 25:1), 2% Nonidet P-40, and 4% ampholines (ratio of pH 3–10 to pH 5–7 was 1:1) were prepared and polymerized for 4–6 hr prior to use. Eighty microliters of the reaction mixture was loaded onto the tube gels and overlaid with 20 μ L of sample overlay buffer (6.5 M urea, 0.4% ampholine, pH 3–10). Tube gels were electrophoresed for 20–24 hr at 400 V (constant). The anode solution was 10 mM H_3PO_4 and the cathode solution was 20 mM NaOH. The tube gels were extruded and soaked in 3 mL SDS stop solution (50 mM Tris, pH 6.7, 4% glycerol, 2% SDS, 2% 2-mercaptoethanol, 0.01% bromophenol blue) for 40–60 min at room temperature.

Second-dimensional gels were run by layering the tube gel on an SDS-polyacrylamide slab gel [18]. The acrylamide concentration of the upper stacking gel was 3.6% and that of the lower resolving gel was 7.5% with 4% *N,N'*-methylene-bis-acrylamide. Gels were electrophoresed at 50 V (constant) overnight until the dye front had traveled 11–12 cm from the bottom of the stacking gel. The electrophoretic separation was calibrated for molecular weight by use of prestained markers run in the second dimension (Sigma "SDS 7B"), and the pH gradient of the focusing gel was determined by homogenizing 1-cm sections of an equilibrated tube gel in 200 μ L of deionized water and measuring the pH. The gradient was linear from pH 3.8 to 7.0.

The two-dimensional gels were dried under vacuum at 80° and major proteins were directly quantitated in counts per minute on an AMBIS Radioanalytic Imaging System (San Diego, CA). Instrument sensitivity and linearity were verified for ^{32}P from 30 to 30,000 dpm (\approx 30% efficiency) using a 1-hr scan. Autoradiographic exposures on Kodak X-OMAT 8 \times 10 film were used for comparative purposes. Statistical analysis was performed using a two-way analysis of variance with a Bonferroni *t*-test to determine the level of significance.

RESULTS

Substrates for protein kinase C were analyzed in rat hippocampal CA1 slices. Approximately ten major phosphoproteins were detected by back-phosphorylation and separation using two-dimensional electrophoretic techniques. Two of the prominent phosphoproteins were identified as myristolated, alanine-rich C kinase substrate (MARCKS) and neuromodulin based on their electrophoretic properties (Fig. 1). Using increasing

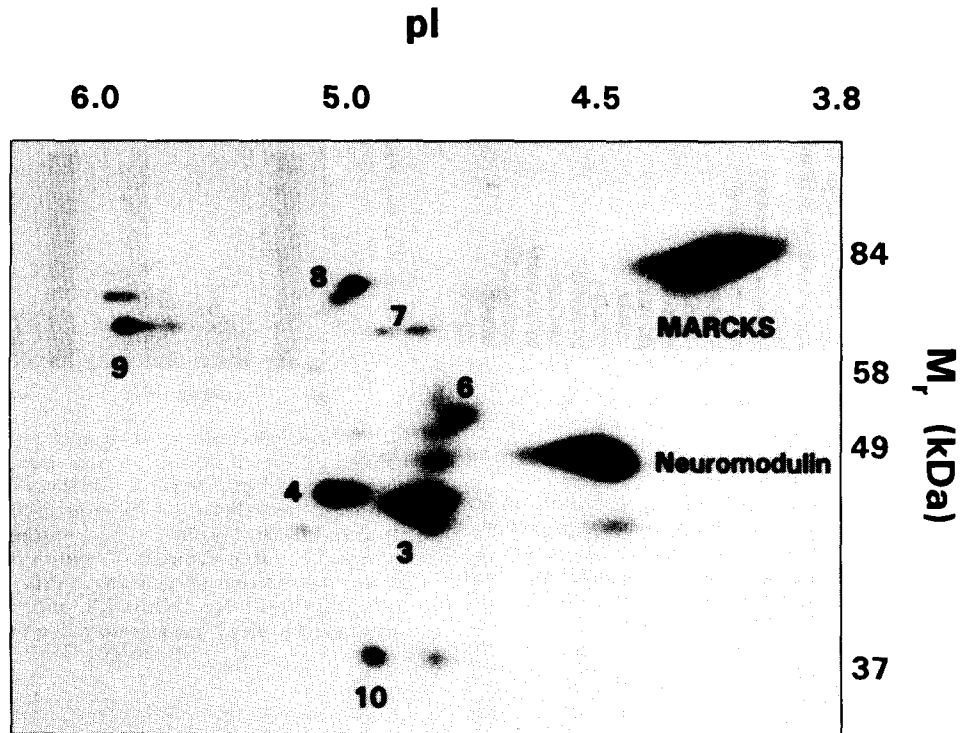


Fig. 1. Autoradiogram of a two-dimensional gel showing ^{32}P incorporation into hippocampal slice proteins back-phosphorylated by protein kinase C. Aliquots of neutralized acid extracts containing $10\text{ }\mu\text{g}$ protein were incubated at 30° with 60 ng protein kinase C (sp. act. $0.3\text{ nmol phosphate/min}/\mu\text{g}$ using histone III S as substrate), $8\text{ }\mu\text{M}$ $[\gamma\text{-}^{32}\text{P}]\text{ATP}$ in a volume of $80\text{ }\mu\text{L}$.

concentrations of protein kinase C, it was demonstrated that 60 ng was necessary to achieve maximal phosphorylation (Fig. 2A) of MARCKS and neuromodulin. Analysis of the time course of phosphorylation indicated that 30 min was an appropriate incubation time for back-phos-

phorylation (Fig. 2B). The acid extraction procedure recovered $>95\%$ of MARCKS, $\approx 80\%$ of neuromodulin and $\approx 50\%$ of the remaining phosphoproteins as determined by quantitation of the total yield of each phosphoprotein present in the supernatant and pelleted material.

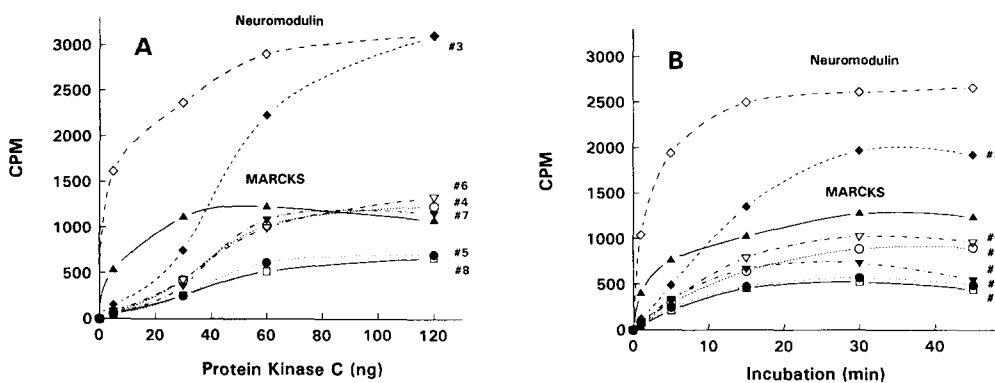


Fig. 2. (A) Effect of increasing protein kinase C concentration on the back-phosphorylation of protein substrates in acid extracts from hippocampal CA1 slices. Samples were incubated for 30 min at 30° . (B) Time course of phosphorylation of protein substrates in acid extracts from the CA1 region of rat hippocampus. Aliquots of neutralized acid extracts were incubated at 30° with 60 ng protein kinase C. Incorporation was terminated at selected time points and radioactivity was quantitated as described in Materials and Methods.

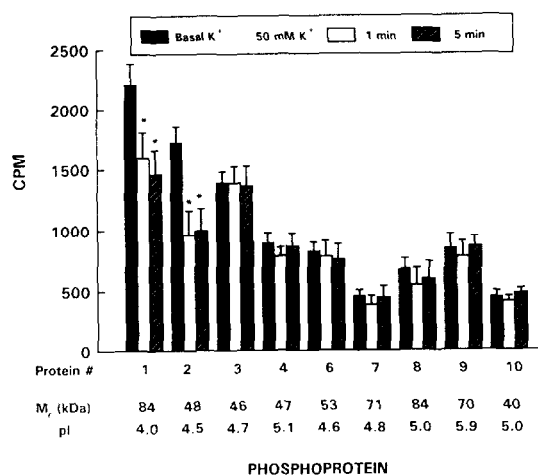


Fig. 3. Effects of K^+ -induced depolarization of intact hippocampal slices on the phosphorylation of proteins by protein kinase C. Results are expressed as mean cpm \pm SEM per protein; $N = 4$ per group. Key: (*) significantly different from control, $P < 0.05$.

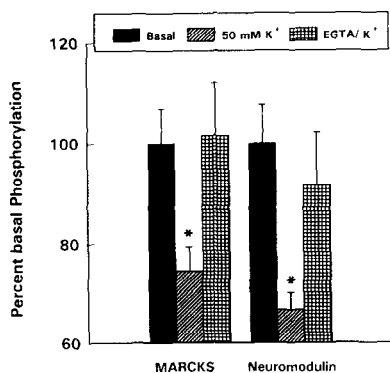


Fig. 4. Effect of low extracellular Ca^{2+} on K^+ -induced depolarization changes. Hippocampal slices were pretreated for 15 min in low Ca^{2+} ($10 \mu M$) Krebs buffer with $100 \mu M$ EGTA prior to K^+ depolarization for 5 min. Data represent the mean per cent \pm SEM of 5–6 determinations obtained in two independent experiments. Basal values ($N \approx 5$) in cpm were 2130 ± 130 for MARCKS and 1150 ± 90 for neuromodulin. Key: (*) significantly different from control, $P < 0.05$.

Depolarization of intact hippocampal slices with $50 mM K^+$ for 5 min produced a significant change in phosphorylation of the two major protein kinase C substrates identified as MARCKS (#1) and neuromodulin (#2) (Fig. 3). The addition of EGTA ($100 \mu M$) using low Ca^{2+} ($10 \mu M$) buffer prevented the depolarization-induced changes (Fig. 4). Pretreatment with nimodipine ($300 nM$) and/or ω -conotoxin GVIA ($10 \mu M$), or MK-801 ($10 \mu M$) for 15 min prior to depolarization had no effect on the K^+ -induced phosphorylation change. Cd^{2+} ($500 \mu M$), a non-specific inhibitor of cation channels [19],

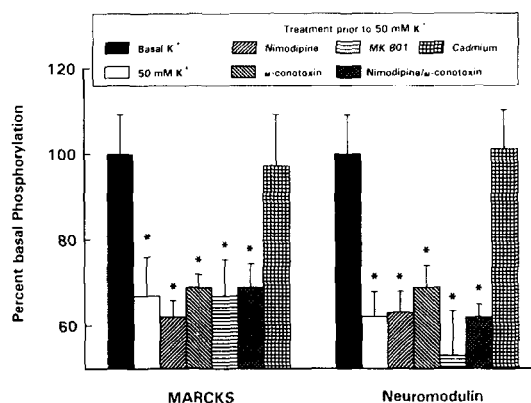


Fig. 5. Acute effect of Ca^{2+} antagonists on depolarization-induced changes in MARCKS and neuromodulin phosphorylation in hippocampal slices. Slices were pretreated with either nimodipine ($300 nM$) or ω -conotoxin GVIA ($10 \mu M$), MK-801 ($10 \mu M$), or Cd^{2+} ($500 \mu M$). Results are expressed as mean per cent of control SEM \pm ($N = 4$). Basal values in cpm for cumulative experiments ($N = 7$) were 2820 ± 410 for MARCKS and 1940 ± 190 for neuromodulin. Key (*) significantly different from control, $P < 0.05$.

abolished the depolarization-induced changes in phosphorylation (Fig. 5).

DISCUSSION

This study demonstrates for the first time the application of a back-phosphorylation procedure using purified protein kinase C to determine changes in the phosphorylation state of major endogenous substrate proteins. Previous investigations employing back-phosphorylation techniques have studied substrates of the cyclic AMP-dependent kinase in neuronal preparations using the purified catalytic subunit [11, 14]. Procedures to examine substrates of protein kinase C have relied upon front or endogenous phosphorylation techniques to quantitate changes to proteins.

The method used here quantitated phosphorylation changes in MARCKS and neuromodulin induced by K^+ depolarization of hippocampal slices with results similar to those obtained with immunoprecipitation techniques used in the study of neuromodulin [5]. These changes were dependent on extracellular Ca^{2+} as shown by experiments using low Ca^{2+} conditions. This result is consistent with those of other studies [5, 20]. The finding that Cd^{2+} ($500 \mu M$) inhibited depolarization-induced changes in phosphorylation has been reported recently in studies of neuromodulin in synaptosomes [21].

Neuromodulin is localized presynaptically and its phosphorylation is correlated with neurotransmitter release in hippocampal slices [3, 5]. The lack of effect of combined treatment with nimodipine ($300 nM$) and ω -conotoxin ($10 \mu M$) in the present study may indicate that Ca^{2+} influx through neither L- nor N-type channels is rate-limiting for phosphorylation changes and/or neurotransmitter

release induced by K^+ depolarization. Electrophysiological studies in hippocampal CA1 neurons indicate that there is a large component ($\approx 50\%$) of voltage-sensitive calcium current that is insensitive to both nimodipine and ω -conotoxin, but blocked by Cd^{2+} [22]. Ca^{2+} influx through this "P-type" Ca^{2+} channel [23] may contribute to the depolarization-induced changes in phosphorylation observed in the present study. In addition, the lack of effect of MK-801 (10 μ M), a noncompetitive N-methyl-D-aspartate (NMDA) receptor antagonist [24], suggests that calcium influx through the NMDA ion channel is not solely responsible for these changes in phosphorylation. It is likely that multiple pathways contribute to the increase in intracellular Ca^{2+} that activates protein kinase C under the conditions of these experiments.

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