20-Hydroxyeicosa-Tetraenoic Acid (20 HETE) Activates Protein Kinase C

Role in Regulation of Rat Renal Na⁺,K⁺-ATPase

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Abstract

It is well documented that the activity of Na⁺,K⁺-ATPase can be inhibited by the arachidonic acid metabolite, 20hydroxyeicosa-tetraenoic acid (20 HETE). Evidence is presented here that this effect is mediated by protein kinase C (PKC). PKC inhibitors abolished 20 HETE inhibition of rat Na⁺,K⁺-ATPase in renal tubular cells. 20 HETE caused translocation of PKC α from cytoplasm to membrane in COS cells. It also inhibited Na⁺,K⁺-ATPase activity in COS cells transfected with rat wild-type renal Na⁺, K⁺-ATPase α_1 subunit, but not in cells transfected with Na⁺, K⁺-ATPase α 1, where the PKC phosphorylation site, serine 23, had been mutated to alanine. PKC-induced phosphorylation of rat renal Na⁺,K⁺-ATPase, as well as of histone was strongly enhanced by 20 HETE at the physiologic calcium concentration of 1.3 µM, but not at the calcium concentration of 200 μ M. The results indicate that phospholipase A₂-arachidonic acid-20 HETE pathway can exert important biological effects via activation of PKC and that this effect may occur in the absence of a rise in intracellular calcium. (J. Clin. Invest. 1997. 99:1224-1230.) Key words: cytochrome P-450 • phosphorylation • protein kinase C inhibitors • protein kinase C translocation • proximal tubules

Introduction

Arachidonic acid is released from cell membranes in response to receptor-dependent activation of phospholipase $A_2 (PLA_2)^1$

J. Clin. Invest. © The American Society for Clinical Investigation, Inc. 0021-9738/97/03/1224/07 \$2.00 Volume 99, Number 6, March 1997, 1224–1230 and can be metabolized via at least three different pathways: the cyclooxygenase, the lipoxygenase and the cytochrome P-450 monooxygenase pathways (P-450) (for review, see references 1, 2). The P-450 metabolites of arachidonic acid appear to be of special importance for the regulation of ion pumps and ion channels (3–5). The ω -hydroxylation product 20-hydroxyeicosa-tetraenoic acid (20-HETE) constitutes one of the major arachidonic acid metabolites in the kidney. This metabolite, which is mainly produced in the proximal tubular cells, has been shown to inhibit the activity of Na⁺,K⁺-ATPase, the electrogenic ion pump that catalyzes the transcellular transport of sodium in each tubular segment (5-7). The molecular mechanisms responsible for this effect are unknown. Na⁺,K⁺-ATPase is a substrate for protein kinase C (PKC) and phosphorylation of Na⁺,K⁺-ATPase by PKC has been shown to cause inhibition of the activity of the pump (8-10).

In the present study, we have examined the possibility that 20 HETE may regulate rat renal Na⁺,K⁺-ATPase via a pathway that involves PKC activation. In support of this possibility, we found that 20 HETE–induced inhibition of rat renal Na⁺,K⁺-ATPase in intact cells was abolished not only by specific PKC inhibitors, but also by mutation of the PKC regulatory site, which for the rat Na⁺,K⁺-ATPase α 1 subunit is Ser 23 of the α subunit. Furthermore, it was found that phosphorylation of rat renal Na⁺,K⁺-ATPase α subunit, as well as of histone, was significantly enhanced by 20 HETE. The results indicate that 20 HETE, one of the major renal PLA₂ pathway metabolites, can, at physiological calcium ion concentrations, activate PKC and via phosphorylation reaction(s) inhibit the activity of Na⁺,K⁺-ATPase.

Methods

Microdissection of single tubular segments and determination of Na^+,K^+ -ATPase activity. Kidney proximal tubule segments (length 0.5–1 mm) were dissected from collagenase (0.05%)-perfused Sprague Dawley rats (B&K Universal, Sollentuna, Sweden). After dissection, Na⁺,K⁺ATPase activity was determined in each tubule segment after incubation for 30 min at room temperature in oxygenated modified Hank's solution (MHS) containing (mM): 137 NaCl, 5 KCl, 0.8 MgSO₄, 0.33 Na₂HPO₄, 0.44 KH₂PO₄, 0.25 CaCl₂, 1 MgCl₂, 10 Tris-HCl. 10⁻³ M butyrate was added to the solution to optimize mitochondrial respiration. Tubules were incubated in MHS alone (control tubules) or MHS plus drugs (experimental). The following drugs were used: 20 HETE (10⁻⁶ M); GF109203x (10⁻⁶ M); RO318220 (10⁻⁷ M). Na⁺,K⁺-ATPase activity was measured as described (8).

Purification of rat renal Na⁺, K⁺-ATPase. Na⁺, K⁺-ATPase was purified from rat renal cortex as described (11). The preparation was analyzed by SDS–polyacrylamide gel electrophoresis, stained with Coomassie brillant blue. Densitometric analysis of the stained gels indicated that Na⁺, K⁺-ATPase α subunit was \sim 55% of total protein.

Portions of this work have been presented at the 28th Annual Meeting of the American Society of Nephrology in San Diego, CA on 5 November 1995. Portions of this work have appeared in abstract form (1995. J. Am. Soc. Nephrol. 6:S536).

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^{1.} *Abbreviations used in this paper:* 20 HETE, 20-hydroxyeicosa-tetraenoic acid; OAG, 1-oleoyl-2-acetoyl-*sn*-glycerol; P-450, cytochrome P-450 monooxygenase; PDBu, phorbol 12,13-dibutyrate; Pi, inorganic phosphate; PKC, calcium- and phospholipid-dependent protein kinase; PLA₂, phospholipase A₂.

The activity of the purified enzyme was measured (as described in reference 12) as the rate of formation of inorganic ³²P from $[\gamma^{-32}P]ATP$ in the absence and presence of ouabain.

Phosphorylation of purified Na^+, K^+ -ATPase by PKC. Assays were carried out at 24°C in a reaction volume of 20 µl containing: 50 mM Hepes (pH 7.5), 10 mM MgCl₂, 1 mM EGTA, 10 nM PKC, and 1 µg of purified Na⁺,K⁺-ATPase. Calcium and 20 HETE concentrations are described below for each experiment. Reactions were initiated by the addition of ATP (final concentration 0.1 mM), with a trace of $[\gamma^{-32}P]$ ATP (2,000–3,000 cpm/pmol). Unless specified, the incubation time was 2 min. Reactions were stopped by the addition of SDS sample buffer. Samples were analyzed by electrophoresis using a 7.5% SDS-PAGE. Gels were stained with Coomassie brillant blue, destained, dried, and analyzed by radioautography or phosphorimaging. In accordance with what had been found in previous studies (13), phosphate was only incorporated into the catalytic a subunit of Na⁺,K⁺-ATPase. Radioactivity was quantified by scanning the phosphorimages using a Molecular Analyst program (Bio-Rad Laboratories, Hercules, CA).

The stoichiometry of the phosphorylation reaction was quantified by measuring the amount of protein in the α subunit of Na⁺,K⁺-ATPase, isolated by gel electrophoresis, as described (14). The amino acid composition determined for the protein blotted onto the polyvinylidene fluoride (PVDF) membranes showed good agreement with the expected composition for Na⁺,K⁺-ATPase.

Free calcium concentrations were calculated as described (15), taking into account the multiple equilibria of Ca^{2+} , Mg^{2+} , ATP, and EGTA, and the pH.

Phosphorylation of histone by PKC. Phosphorylation was performed under linear assay conditions using 10 μM histone (type III from calf thymus) as substrate and the assay conditions described above. 10 μg/ml L-α-phosphatidylserine and 1 μM 1-oleoyl-2-acetoyl*sn*-glycerol (OAG) were added to the reaction mediums indicated. Reactions were stopped by the addition of 30% acetic acid and aliquots were spotted on phosphocellulose filters (P81; Whatman Inc., Clifton, NJ). Filters were washed in water for 10 min and then counted in a liquid scintillation counter.

Two-dimensional phosphopeptide mapping. For phosphopeptide mapping of Na⁺,K⁺-ATPase, gel pieces containing labeled Na⁺,K⁺-ATPase α_1 subunit were excised from dried gels, washed, incubated with trypsin, and processed as described (16). Briefly, aliquots were spotted in the middle of the plate, 4 cm from the bottom, and initially separated by horizontal electrophoresis at pH 3.5 in 10% acetic acid/ 1% pyridine until the dye front migrated 7 cm. Chromatography was performed in the vertical dimension in 1-butanol/acetic acid/water/ pyridine (15:3:12:10 vol/vol, respectively). Phosphopeptides were visualized by autoradiography.

Cell culture. COS cells were maintained at 37° C in a humified atmosphere of 95% O₂ and 5% CO₂, in DMEM supplemented with 10% fetal serum. Cells were grown to confluence for 4–5 d in 10-cm dishes.

Analysis of PKC distribution in COS cells. (a) Subcellular fractionation. After a 30-min incubation with DMEM alone (control group) or with the addition of experimental drugs, cell monolayers were washed with ice-cold PBS, scraped in 1 ml homogenization buffer (HB) (20 mM Tris-HCl, 2 mM EGTA, 2 mM EDTA, 1 mM phenylmethylsulfonylfluoride, 10 mM ß-mercaptoethanol, and 100 KIE/ml aprotinin), homogenized and pelleted by centrifugation at 100,000 g for 1 h at 4°C. Membrane fractions were solubilized in HB plus 0.1% triton ×100. Protein concentrations were determined according to Bradford (17). (b) PKC immunoblotting. Cytosolic and membrane samples containing equal amounts of protein (10 µg) were dissolved in loading buffer (1% SDS, 0.25 mM Tris-HCl, 10% glycerol, 5% β-mercaptoethanol, and trace amounts of pyronine), and separated by SDS-PAGE (8% acrylamide), transferred to nitrocellulose and immunoblotted with a PKC α isozyme-specific antipeptide antibody (Santa Cruz Biotechnology, Santa Cruz, CA) as described (9). After washing, filters were incubated with peroxidase-conjugated anti–rabbit polyclonal antibodies (Amersham International, Little Chalfont, UK) and detected using an enhanced chemiluminescence method (Amersham International).

Site-directed mutagenesis and DNA transfection into COS cells. Ser23 of rat α_1 Na⁺,K⁺-ATPase was mutated to alanine using the methods described in detail elsewhere (18). In brief, site-directed mutagenesis was carried out by PCR using as template rat Na⁺,K⁺-ATPase α_1 cDNA cloned in the pCMV eukaryotic expression vector (PharMingen, San Diego, CA). The presence of the correct mutation was confirmed by direct nucleotide sequencing according to Sanger et al. (19). The entire cDNAs coding for wild-type and mutated α_1 were transfected into a Green Monkey kidney cell line (COS cells) using the CaPO₄/DNA coprecipitation method. Since the endogenous COS Na⁺,K⁺-ATPase has a higher sensitivity to ouabain than the rat α_1 subunit Na⁺,K⁺-ATPase, the cells were grown in a medium containing 10⁻⁵ M ouabain. Untransfected COS cells did not survive in this medium. To avoid problems arising from clonal variation, a total of \sim 100 colonies were selected, pooled, and used in all the studies.

Ouabain-sensitive ⁸⁶Rb⁺ uptake and Na⁺,K⁺-ATPase activity in transfected COS cells. Cell monolayers were washed twice with room temperature PBS and preincubated for 30 min at room temperature with 20 HETE (1 μ M). All measurements were made in the presence of 10⁻⁵ M ouabain to inhibit endogenous Na⁺,K⁺-ATPase activity. Protein content was determined by the method of Bradford (17).

 $^{86}\rm{Rb^+}$ uptake, used as an index of K⁺ uptake (20), was initiated by addition of 0.1 ml/well PBS containing 25 $\mu\rm{Ci/ml}$ $^{86}\rm{Rb^+}$ and was linear for at least 5 min. Recordings were made at 2-min intervals. Reactions were stopped by aspiration of the medium and rinsing the cells three times with 1 ml/well of ice-cold PBS, containing 5 mM BaCl₂. Cells were then lysed twice with 0.55 ml of 1 mM NaOH. Radioactivity in cell lysates was measured by liquid scintillation counting (LKB, Wallac, Sweden). Na⁺,K⁺-ATPase activity was determined as the difference between $^{86}\rm{Rb^+}$ uptake in the absence (total activity) and presence (ouabain-insensitive) of ouabain and was expressed as micromoles per minute per milligram protein.

The activity of Na⁺,K⁺-ATPase was assayed in a crude membrane preparation as described above. To prevent dephosphorylation of Na⁺,K⁺-ATPase, the buffer was complemented with the protein phosphatase inhibitors okadaic acid (250 nM) and FK506 (25 nM). An amount of enzyme was selected so that total ATP hydrolysis did not exceed 20%, and ATP hydrolysis was linear with time. The reaction was stopped by the addition of 700 µl of activated charcoal. The ³²P liberated was determined in the supernatant after centrifugation. For the determination of ouabain-insensitive rat ATPase activity, NaCl and KCl were omitted and 5 mM ouabain was added. Na⁺,K⁺-ATPase activity is expressed as picomoles inorganic phosphate (Pi) per hour per milligram protein.

Chemicals. GF109203x was purchased from Calbiochem Corp. (La Jolla, CA), RO318220 was a kind gift from Roche Products LTD (London, UK). Both of these PKC antagonists were stored as stock solutions in DMSO at -20° C. Further dilutions were made in dissection solution. Control experiments showed that, at the concentrations used, DMSO did not interfere with Na⁺,K⁺-ATPase activity. Both GF109203 and RO318220, which are bisindolyl maleimides, are highly specific inhibitors for PKC with IC₅₀ 30–200 nM and 23 nM, respectively (21, 22). Other kinases are affected at much higher concentrations. The IC₅₀ for PKA is > 10 μ M. All these IC₅₀ values were recorded for purified enzyme. GF109203 was used in the concentration of 1 μ M and RO318220 in the concentrations of 0.1 μ M.

PKC was obtained from Boehringer Mannheim (Mannheim, Germany), 1-oleoyl-2-acetoyl-*sn*-glycerol from Avanti Polar Lipids, Inc. (Alabaster, AL), 20-hydroxyeicosa-tetraenoic acid from Cascade Biochem LTD (Berkshire, UK).

Statistics. Statistical comparisons between two groups were performed by Student's t test for unpaired data. Comparisons between several groups were done by analysis of variance. P < 0.05 were considered significant.

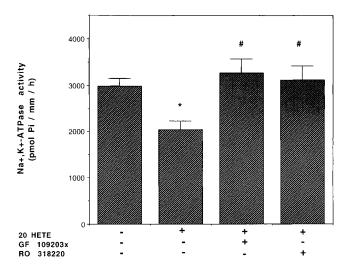


Figure 1. Na⁺,K⁺-ATPase activity in isolated proximal tubules. Individual tubular segments were incubated at room temperature for 30 min in a drug free medium (control, n = 17), or in the presence of the indicated drugs: 20 HETE (10^{-6} M) (n = 13), 20 HETE (10^{-6} M) plus GF109203x (10^{-6} M) (n = 5), 20 HETE (10^{-6} M) plus RO318220 (10^{-7} M) (n = 5). The PKC inhibitors were present 5 min before and throughout the incubation with 20 HETE. Each bar represents the mean±SEM. Statistical comparison between groups was done by one-way analysis of variance followed by Tukey-Kramer test. *P < 0.05 vs. control, *P < 0.05 vs. 20 HETE.

Results

Inhibition of renal tubular Na⁺,K⁺-ATPase by 20 HETE is abolished by PKC inhibitors. Na⁺,K⁺-ATPase activity in proximal tubular segments dissected from rat kidneys was 2,850± 112 pmol Pi mm⁻¹ h⁻¹ (n = 17). 20 HETE (10⁻¹⁰ to 10⁻⁵ M) inhibited Na⁺,K⁺-ATPase activity in a dose-dependent manner in accordance with data already published (5). Inhibition was first observed at 10⁻⁹ M 20 HETE and was maximal (25% inhibition) at 10⁻⁷ M (P < 0.05, n = 3) (data not shown). 20 HETE at concentrations from 10⁻⁹-10⁻⁴ M had no effect on purified rat renal Na⁺,K⁺-ATPase (data not shown), suggesting that the effect of 20 HETE on tubular Na⁺,K⁺-ATPase activity is not due to a direct interaction between the lipid and the pump.

To assess whether the inhibitory effect of 20 HETE was mediated via PKC, proximal tubular segments were first incubated with 20 HETE (10^{-6} M) in the presence of two different PKC inhibitors: GF109203x (10^{-6} M) or RO318220 (10^{-7} M). In this protocol, incubation of tubules with 20 HETE alone resulted in $27\pm4\%$ inhibition of Na⁺,K⁺-ATPase activity (P < 0.05 vs. control, n = 9). The effect of 20 HETE was abolished by GF109203x and RO318220 (P < 0.05 vs. 20 HETE in paired experiments for both drugs) (Fig. 1). The addition of the PKC inhibitors alone had no effect on Na⁺,K⁺-ATPase activity (data not shown).

20 HETE does not inhibit the activity of Na⁺, K⁺-ATPase after mutation of the PKC regulatory site. The site for PKC phosphorylation in rat α_1 subunit of Na⁺, K⁺-ATPase was recently identified as Ser23 (13, 23). cDNAs coding for wild-type rat Na⁺, K⁺-ATPase α_1 subunit or rat α subunit in which Ser23 had been mutated to alanine (S23A) were stably transfected into COS cells. Total ⁸⁶Rb⁺ uptake was similar in cells expressing

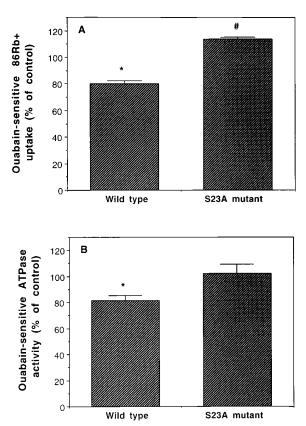


Figure 2. Effect of 20 HETE on ouabain-sensitive ⁸⁶Rb⁺ uptake and Na⁺,K⁺-ATPase activity in transfected COS cells. Cells were transfected with rat wild-type α_1 subunit or with the S23A mutant and preincubated in the presence or absence of 1 μ M 20 HETE for 30 min at room temperature. Ouabain-sensitive ⁸⁶Rb⁺ uptake (*A*) and Na⁺,K⁺-ATPase activity (*B*) were assayed as described in Methods. In control groups (incubation without drugs), ouabain-sensitive ⁸⁶Rb⁺ uptake and Na⁺,K⁺-ATPase activity were 22.9±0.5 pmol/min per mg protein and 4,497±305 pmol Pi/h per mg protein, respectively, for cells expressing wild-type Na⁺,K⁺-ATPase. For cells expressing S23A Na⁺,K⁺-ATPase, the corresponding values were 21.7±1.3 pmol/min per mg protein and 2,780±672 pmol Pi/h per mg protein, respectively. Each bar represents the mean±SEM of four experiments. **P* < 0.01 vs. control by Student's *t* test for unpaired data.

wild-type and S23A Na⁺,K⁺-ATPase. 20 HETE decreased ouabain-sensitive ⁸⁶Rb⁺ uptake by 19.7±1.0% in COS cells expressing wild-type Na⁺, K⁺-ATPase (P < 0.001 vs. control, n = 5). In contrast, 20 HETE caused a small, but significant increase of 14.3±2.0% in ⁸⁶Rb⁺ uptake in cells expressing the S23A mutant (P < 0.02 vs. control, n = 4) (Fig. 2 A). Ouabaininsensitive ⁸⁶Rb⁺ uptake was unaffected by 20 HETE in both cell types (data not shown). In ongoing studies, we have observed that in the presence of phorbol 12,13-dibutyrate (PDBu), ouabain-sensitive ⁸⁶Rb⁺ uptake is decreased in cells expressing rat wild-type Na⁺,K⁺-ATPase, but increased in cells expressing rat S23A Na⁺,K⁺-ATPase. This increase could, in the case of PDBu be attributed to a PKC-mediated stimulation of Na⁺ uptake via the Na⁺-H⁺ exchanger (Belusa, R., Z.-M. Wang, T. Matsubara, B. Sahlgren, I. Dulubova, A.C. Nairn, E. Ruoslahti, P. Greengard, and A. Aperia, manuscript in preparation).

Treatment with 20 HETE induced an $18.3\pm1.7\%$ inhibition of Na⁺,K⁺-ATPase activity in COS cells expressing wild-type

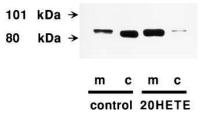


Figure 3. Effect of 20 HETE on intracellular distribution of PKC. Cytosolic (c) and membrane (m) extracts of COS cells that had been incubated with DMEM alone (control group) or 20 HETE (1 μ M) for

30 min at room temperature were prepared as described in Methods, subjected to SDS-PAGE (10 ng protein/lane), transferred to nitrocellulose, and immunoblotted with an antipeptide antibody specific for PKC α .

Na⁺,K⁺-ATPase (P < 0.01 vs. control, n = 4), but had no effect in cells expressing S23A Na⁺,K⁺-ATPase (Fig. 2 *B*). Ouabain-insensitive ATPase activity was unaffected by 20 HETE in both cell types (data not shown).

20 HETE induces translocation of PKC from the cytoplasm to the membrane. The initial step in the activation of many PKC isoforms, including the ubiquitous PKC α , is the translocation of the enzyme from the cytosol to the membrane (24). Incubation of intact COS cells with 20 HETE (1 μ M) for 30 min caused a pronounced decrease in cytosolic PKC α , along with a concomitant, approximately twofold, increase in PKC α associated with the membrane fraction (Fig. 3). Similar results were observed when PDBu, a classical PKC activator, was used (data not shown).

20 HETE enhances phosphorylation of Na^+, K^+ -ATPase by *PKC*. The calcium dependence of the phosphorylation by PKC of the purified rat kidney Na^+, K^+ -ATPase α_1 , was evaluated. As expected, a relationship between calcium and Na^+, K^+ -ATPase α_1 subunit phosphorylation was found with a thresh-

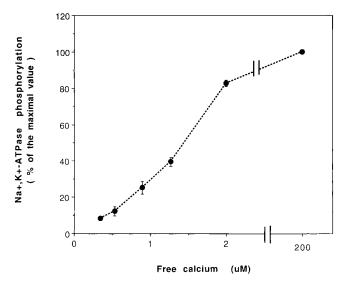


Figure 4. Ca²⁺ dependence of Na⁺,K⁺-ATPase phosphorylation by PKC. 1 µg purified Na⁺,K⁺-ATPase was incubated with PKC at 24°C in a phosphorylation mixture as described in Methods for 20 min at various calcium concentrations. The stoichiometry of phosphorylation measured at 200 µM calcium at steady state conditions was 0.6 mol Pi incorporated per mol of Na⁺,K⁺-ATPase. Data were normalized to the maximal phosphorylation signal from the same experiment (obtained at 200 µM free calcium). Each point represents the mean±SEM from two to four experiments.

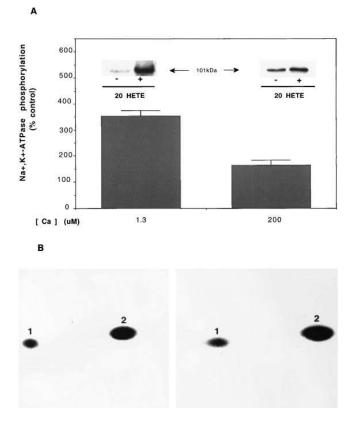


Figure 5. (*A*) Effect of 20 HETE on PKC-dependent Na⁺,K⁺-ATPase phosphorylation. 1 μ g of purified Na⁺,K⁺-ATPase was incubated with PKC at 24°C for 2 min in the presence of the indicated Ca²⁺ concentration, in the absence (control) or presence of 100 μ M 20 HETE. Data are expressed as the increase in the phosphorylation signal in the presence of 20 HETE over the control value for each experiment. Bars represent means ±SEM from three experiments. An autoradiogram from a representative experiment is included in the inset. (*B*) Two-dimensional phosphopeptide mapping of Na⁺,K⁺-ATPase phosphorylated by PKC in the absence (*left*) or presence (*right*) of 100 μ M 20 HETE. Phosphorylation was run for 3 min under the conditions described above. Electrophoresis was performed in the horizontal direction (positive electrode left) and chromatography was performed in the vertical direction.

old calcium concentration of $\sim 0.5~\mu M$ (Fig. 4). Time course studies were performed at both saturating (200 μM) and non-saturating (1.3 μM) calcium concentrations. In both conditions, Na⁺,K⁺-ATPase phosphorylation by PKC was linear during the first 2 min of incubation. 20 HETE (1–170 μM) increased Na⁺,K⁺-ATPase phosphorylation by PKC in a dose-dependent manner, with a threshold concentration of 20 μM (data not shown).

The effect of 20 HETE (100 μ M) on the PKC-dependent phosphorylation of Na⁺,K⁺-ATPase was studied at two different calcium concentrations (1.3 and 200 μ M) under conditions of linearity (Fig. 5 *A*). Phosphorylation of Na⁺,K⁺-ATPase by PKC was enhanced by 20 HETE, mainly at the lower calcium concentration.

Two-dimensional phosphopeptide maps of the phosphorylated α subunit indicated that the same site in the purified enzyme was phosphorylated by PKC, both in control conditions and in the presence of 20 HETE (Fig. 5 *B*).

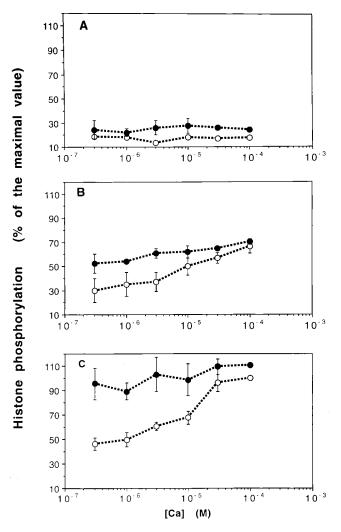


Figure 6. Calcium dependence of PKC activity in the absence and presence of 20 HETE. PKC activity was assayed using histone as the substrate under linear and standard conditions (see Methods) except that various concentrations of $CaCl_2$ were added in the absence (\bigcirc) or presence (\bullet) of 20 HETE (50 μ M). Other lipids were added to the incubation medium as follows: A, none; B, phosphatidylserine (10 µg/ ml); C, phosphatidylserine (10 µg/ml) plus OAG (1 µM). Data were normalized to the phosphorylation signal obtained at 10⁻⁴ M calcium in the presence of phosphatidylserine plus OAG. For each group, histone phosphorylation was also studied under completely calcium free conditions. In these protocols, EGTA (5 mM) was added to the reaction mixture instead of CaCl₂. Under these conditions, histone phosphorylation (percentage of the maximal value) were as follows: control, 17.4±1.5, 23.0±10, and 44.1±7 for A, B, and C, respectively, and in the presence of 20 HETE (50 μ M), 19.4 \pm 2.6, 40.7 \pm 3.6, and 59.7 \pm 14 for A, B, and C, respectively. Each point represents the mean±SEM from two to three experiments.

It is well known that unsaturated fatty acids are extremely difficult to maintain in solution. The major difficulties are related to the adsorption of the fatty acids to tubing and to their poor solubility (25). In most previous in vitro studies on the effect of unsaturated fatty acids on PKC, doses up to 100 μ M have been used (26, 27). The intact cells responded to a lower dose of 20 HETE, namely 1 μ M. It is possible that conditions for the in vitro studies are less favorable for the maintenance

of 20 HETE than the conditions for the intact cell studies. It is also possible that in the intact cell the effect of 20 HETE is amplified for instance by the presence of synergizing fatty acids (28).

20 HETE enhances phosphorylation of histone by PKC. To evaluate if the effect of 20 HETE on phosphorylation of Na⁺,K⁺-ATPase by PKC was due to a direct effect on PKC or on Na⁺,K⁺-ATPase, we also studied the effect of 20 HETE on the phosphorylation of histone by PKC. Histone phosphorylation was slightly enhanced by the presence of 50 µM 20 HETE. This effect was calcium independent (Fig. 6 A). The interaction between 20 HETE, phosphatidylserine, and the diacylglycerol analogue, OAG was also studied. In the presence of phosphatidylserine (10 µg/ml), phosphorylation of histone by PKC was calcium dependent. The concentration dependence for calcium-stimulated phosphorylation of histone by PKC in the presence of phosphatidylserine was markedly shifted to the left by 20 HETE. The maximal response was unaffected by 20 HETE (Fig. 6 B). As expected, the OAG and phosphatidylserine-dependent activation of PKC was also calcium dependent and was greatly enhanced by the addition of 20 HETE (Fig. 6 C).

Discussion

Previous studies have demonstrated that arachidonic acid activates PKC (26-30), that this effect is associated with an increase in the affinity for calcium, and that certain unsaturated fatty acids act synergistically with diacylglycerol and phosphatidylserine to activate PKC (26, 27). In another series of studies, it was shown that certain P-450 metabolites of arachidonic acid regulate the activity of Na⁺,K⁺-ATPase (5–7). The results of the present study indicate that 20 HETE-dependent inhibition of rat renal Na⁺,K⁺-ATPase activity is mediated via PKC. Thus PKC inhibitors abolished the inhibitory effect of 20 HETE on Na⁺,K⁺-ATPase activity and this effect was also abolished in Na⁺,K⁺-ATPase in which the PKC phosphorylation site (S23) was mutated to alanine. Two other lines of evidence indicate that the effect of 20 HETE was due to an effect on PKC and not on the Na⁺,K⁺-ATPase itself; first, 20 HETE had no effect on the activity of purified Na⁺,K⁺-ATPase in the absence of PKC; second, 20 HETE stimulated PKC-dependent phosphorylation of histone.

Rat renal α_1 Na⁺,K⁺-ATPase is a physiological substrate for PKC (8, 13, 14). Many studies, performed either on intact rat renal tubular cells or on purified enzyme, suggest that PKC phosphorylation of rat Na⁺,K⁺-ATPase is associated with a decrease in enzyme activity (8–10, 13, 31), although in one study a stimulatory effect of phorbol ester on enzyme activity was found (32).

The present study suggests that 20 HETE may act as a second messenger, mediating the actions of hormones on phosphorylation of rat $\alpha_1 \text{ Na}^+, \text{K}^+$ -ATPase by PKC. 20 HETE was shown to activate PKC both in the in vitro phosphorylation studies and in intact cells. In the in vitro studies, 20 HETE was found to reduce the concentration of calcium required for the phosphatidylserine-dependent activation of PKC, regardless of whether histone or Na⁺, K⁺-ATPase was used as a substrate. The results are in agreement with the findings that several *cis*unsaturated fatty acids, including arachidonic acid, can enhance the affinity of PKC for calcium (27). In intact cells, 20 HETE facilitated the translocation of the calcium-dependent PKC α from the cytosol to the membrane, a process that is also calcium dependent. 20 HETE mimicked the action of diacylglycerol and phorbol ester both with regard to the effects on histone phosphorylation and with regard to PKC translocation.

Our present results suggest a synergism between 20 HETE, phosphatidylserine, and the diacylglycerol analogue that was more pronounced at physiological intracellular calcium levels. Activation of phospholipase C leads to the release of inositol triphosphate and a transient increase in intracellular free calcium (33). In contrast, activation of PLA₂ is generally not associated with an increase in intracellular calcium. The present results suggest that the calcium-dependent PKC isoforms expressed in the kidney, α and γ (34), can be activated by the PLA₂ pathway at resting intracellular calcium concentrations.

The expression of the arachidonic acid metabolizing P-450 enzymes is highly tissue specific (35). It is important to realize that the PLA₂-PKC signaling pathway may be mediated by arachidonic acid metabolites other than 20 HETE. Thus, a number of biological effects of lipoxygenase metabolites are mediated through the stimulation of PKC. 15(S)-hydroperoxy-5,8,11,13-eicosa-tetraenoic acid (15(S)-HPETE) was, at micromolar concentration, found to induce transmigration of monocytes across an endothelial monolayer. This effect was attenuated in the presence of PKC inhibitors (36). 12(S)-Hydroxyeicosatetraenoic acid (12(S)-HETE) (0.1 μ M) was found to induce the release of cathepsin B. This effect was abolished in the presence of PKC inhibitors (37).

In ongoing studies, we have found that the lipoxygenase metabolite 12(S)-HETE enhances PKC-induced phosphorylation of Na⁺,K⁺-ATPase in vitro and inhibits proximal tubule Na⁺,K⁺-ATPase activity (data not shown).

The results from the present study have important implications for an understanding of the PLA_2 -arachidonic acid signaling pathway in the kidney as well as in other tissues. Both dopamine and parathyroid hormone have been shown to inhibit renal tubular Na⁺,K⁺-ATPase via the PLA_2 -arachidonic acid–20 HETE pathway (5–7). Several other hormones, such as vasopressin and EGF, are known to activate PLA_2 in the kidney (38). It is possible that the effects of these hormones, which can influence tubular transport, renal hemodynamics, and kidney growth and differentiation are also mediated via PKC. In addition to inhibiting tubular Na⁺,K⁺-ATPase activity, 20 HETE has several other biological effects, such as induction of vasoconstriction (39) and inhibition of K⁺ channels (40). The possibility that these effects are also mediated via PKC should be considered.

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