

Protein Kinase A Activation Phosphorylates the Rat CIC-2 Cl⁻ Channel but Does Not Change Activity

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Abstract. Phosphorylation-dependent events have been shown to modulate the activity of several members of the mammalian CLC Cl⁻ channel gene family, including the inward rectifier CIC-2. In the present study we investigated the regulation of rat CIC-2 expressed in the TSA-201 cell line (a transformed HEK293 cell line that stably expresses the SV40 T-antigen) by protein kinases. Protein kinase A activation phosphorylated CIC-2 in vivo, whereas stimulation of protein kinase C with phorbol 12-myristate 13-acetate did not. In vitro labeling studies confirmed that protein kinase A could directly phosphorylate CIC-2, and that protein kinase C and Ca²⁺/calmodulin-dependent protein kinase II did not. Nevertheless, protein kinase A-dependent phosphorylation of CLC-2 failed to regulate either the magnitude or the kinetics of the hyperpolarization-activated Cl⁻ currents. Considered together, we demonstrate that protein kinase A activation results in the phosphorylation of rat CIC-2 in vivo, but this event is independent of Cl⁻ channel activity.

Key words: CIC-2 — Hyperpolarization-activated Cl⁻ Channel — Phosphorylation — Protein kinase A — Regulation

Introduction

The ubiquitously expressed rat CIC-2 channel, a member of the CLC voltage-activated chloride channel gene family (Jentsch et al., 1995, 1999), exhibits an inward rectifying steady-state current-voltage relation and time-

dependent activation at physiological voltages when expressed in *Xenopus* oocytes (Thiemann et al., 1992), DRG neurons (Staley et al., 1996) and HEK293 cells (Park et al., 1998). Similar hyperpolarization-activated Cl⁻ currents have been associated with the expression of CIC-2 in many different types of cells (Arreola et al., 1996; Carew & Thorn, 1996; Clark et al., 1998; Dinudom, Young & Cook, 1993; Enz, Ross & Cutting, 1999; Ferroni et al., 1997; Huber et al., 1998; Joo et al., 1999; Kajita, Omori & Matsuda, 2000; Murray et al., 1995; Park et al., 1998). Indeed, targeted disruption of the *Clcn2* gene confirmed that hyperpolarization-activated Cl⁻ currents in Leydig and Sertoli cells are mediated by CIC-2 (Bösl et al., 2001). Other members of the CLC mammalian gene family include CIC-1 that regulates the resting membrane potential in muscle cells (Steinmeyer et al., 1994), the intracellular Cl⁻ channels CIC-3 (Stobrawa et al., 2001) and CIC-7 (Kornak et al., 2001), isoforms important for renal function—CIC-5, CIC-K (K1 and K2, or Ka and Kb, depending on the species) (Gunther et al., 1998; Kieferle et al., 1994), and isoforms for which the functions are unknown—CIC-4 and CIC-6 (Brandt & Jentsch, 1995).

Regulation of CIC-2 channel activity is multifaceted. The voltage-dependent CIC-2 current is enhanced by low extracellular pH (Jordt & Jentsch, 1997; Sherry et al., 1997; Tewari et al., 2000), is sensitive to the intracellular and extracellular Cl⁻ concentration (Dinudom et al., 1993; Jordt & Jentsch, 1997), and can be either activated (Furukawa et al., 1998; Thiemann et al., 1992; Xiong et al., 1999) or inhibited (Komwatana et al., 1995) by cell swelling. CIC-2 currents are also apparently modulated by protein kinases. Lipid bilayers fused with membranes from *Xenopus* oocytes injected with rabbit CIC-2 cRNA expressed Cl⁻ channel activity enhanced by a cAMP-dependent kinase (PKA)-dependent process

(Sherry et al., 1997). Similarly, the inward-rectifying Cl^- currents in mouse choroid plexus epithelial cells were activated by PKA (Kibble et al., 1997). In contrast, activation of either PKA or protein kinase C (PKC) decreased the amplitude of the CIC-2 inward rectifying currents in human intestinal T84 cells (Fritsch & Edelman, 1996). Human CIC-2 is apparently activated by PKA when expressed in HEK 293 cells (Tewari et al., 2000), whereas, no effect of PKA on this channel was observed in the human bronchial cell line IB3-1 (Schwiebert et al., 1998) or when expressed in *Xenopus* oocytes (Jordt & Jentsch, 1997).

Analysis of the primary amino acid sequence of rat CIC-2 revealed three potential phosphorylation sites for both PKA and Ca^{2+} /calmodulin-dependent protein kinase II (CaMKII). The three potential PKA phosphorylation sites in rat CIC-2 (Ser82, Ser710 and Thr898) are conserved in mouse, rabbit and human CIC-2 protein. All three sites are predicted to be cytoplasmic on either the amino or carboxy termini of CIC-2, and thus are likely to be accessible to intracellular kinases. Therefore, the modulation of channel activity by kinases may involve direct phosphorylation of the CIC-2 protein, although this has not been previously reported. In the present study, we demonstrate that PKC and CaMKII do not phosphorylate rat CIC-2. In contrast, PKA activation phosphorylated CIC-2 *in vivo* and *in vitro*; yet, CIC-2 currents were not modulated by PKA under conditions where the cAMP-dependent Cl^- channel CFTR was activated.

Materials and Methods

SUBCLONING OF CIC-2 AND CFTR INTO GREEN FLUORESCENT PROTEIN EXPRESSION VECTORS

To facilitate identification of transfected cells in patch clamp studies, rat CIC-2 and human CFTR were subcloned into eukaryotic green fluorescent protein (GFP) expression vectors. Cells expressing the channel constructs were identified by epi-fluorescence with an Endow GFP bandpass filter (Chroma Technology, Brattleboro, VT). Rat CIC-2 was subcloned from pCRII-CIC-2 (Park et al., 1998) into the bicistronic expression vector pIRES2-EGFP (Clontech, Palo Alto, CA) using the *Bam*H1 and *Nsi*I restriction sites for removal from pCRII (Invitrogen, San Diego, CA) and from the *Bam*H1 and *Pst*I (compatible cohesive end with *Nsi*I) sites for insertion into pIRES-EGFP. A rat CIC-2-egfp fusion construct was made using a PCR-based technique to introduce an *Age*I restriction enzyme site and to remove the stop codon as described before (Park et al., 1998). Briefly, the open reading frame of rat CIC-2 cDNA was amplified using sense (5'-AAGCAAGAGGAGGCAAGAGGAC-3') and nonsense primers (5'-GGTCACCGTAACTTGTCA-3') with pCRII-CIC-2 as the template, followed by *Age*I digestion. The rat CIC-2 cDNA insert was then subcloned into pEGFP-N1 (Clontech) which had been linearized by *Sma*I and *Age*I to construct pCIC-2-egfp. The inserts in the bicistronic pCIC-2/egfp and fusion pCIC-2-egfp constructs were confirmed by sequencing (MJResearch autosequencer).

The open reading frame of human CFTR was amplified from a cDNA in pBluescript-KS(+) provided by Jack R. Riordan (Mayo Clinic Scottsdale, S.C. Johnson Medical Research Center, Scottsdale, AZ). Primers were designed to remove the stop codon and to generate restriction endonuclease sites for subcloning the PCR product into the expression vector pEGFP-N1. The sense primer was 5'-aaactgcaGTAGGTCTTTGGCATTAGGAGCTTG-3' (lower-case letters indicate a *Pst*I endonuclease site) and the nonsense primer was 5'-acaccggTAGCCTTGATCTT-3' (lower-case letters indicate an *Age*I endonuclease site). The PCR product containing the open reading frame of CFTR was digested with *Pst*I and *Age*I and subcloned into pEGFP-N1 previously digested with *Pst*I and *Age*I. The pCFTR-egfp construct was sequenced to verify in-frame translation of EGFP fused to the carboxyl terminus of CFTR.

TRANSIENT EXPRESSION OF RAT CIC-2 AND HUMAN CFTR-egfp

The TSA-201 cell line (a transformed HEK293 cell line that stably expresses the SV40 T-antigen—kindly provided by Ronald Li and Eduardo Marban, Johns Hopkins University) was used for transient expression of channel proteins. Cells were maintained as previously described (Park et al., 1998) and transiently transfected with CIC-2/egfp bicistronic or CFTR-egfp fusion constructs at 10 μg DNA/100 mm plate using Superfect (Qiagen, Hilden, Germany) and incubated for 48 hours before phosphorylation and electrophysiological studies. In some electrophysiological experiments, HEK293 cells and the CIC-2-egfp fusion construct were used. In this case, cells with positive GFP signals were sorted by their fluorescent intensity using a B-D Vantage cell sorter (Becton Dickinson, Palo Alto, CA). Single cell suspensions at concentration of $1-3 \times 10^6/\text{ml}$ were prepared in PBS containing 1% FBS and analyzed at a rate of about 1,000 cells/sec using an argon ion laser operating at 488 nm wavelength and 200 milliwatts excitation. Two-parameter gating window (forward scattering vs. fluorescent intensity) was used to define the sorted cell populations. GFP positive cells were collected into a 15-ml conical centrifuge tube filled with 10 ml Dulbecco's modified Eagle's medium containing 20% FBS, 2 mM L-glutamine, penicillin (100 units/ml) and streptomycin (100 $\mu\text{g}/\text{ml}$).

IN VIVO METABOLIC $^{32}\text{P}_i$ LABELING OF RAT CIC-2

To examine whether the CIC-2 is phosphorylated *in vivo*, TSA-201 cells transiently transfected with the CIC-2/egfp bicistronic construct were incubated for 4 hr in a phosphate-free Dulbecco's modified Eagle's medium (Gibco/BRL, Grand Island, NY) containing 50 μM sodium orthovanadate and 250 $\mu\text{Ci}/\text{ml}$ $^{32}\text{P}_i$. After labeling, the cells were treated for 15 min with either: (1) cAMP cocktail [20 μM forskolin + 100 μM 3-isobutyl-1-methylxanthine + 500 μM 8-(4-chlorophenylthio) cAMP], (2) 0.5 μM Phorbol 12-myristate 13-acetate (PMA), or (3) vehicle. The reaction was stopped by the addition of ice-cold PBS.

IMMUNOPRECIPITATION AND BIOCHEMICAL ANALYSIS OF CIC-2

Following 15 min *in vivo* treatment with either a cAMP cocktail, PMA or vehicle, CIC-2 transfected TSA-201 cells were washed twice with ice-cold PBS, lysed by incubation at 4°C for 20 min in a modified RIPA buffer [in mM: 50 Tris-HCl (pH 7.4), 500 NaCl, 1 PMSF, and 1 sodium orthovanadate], 0.1% SDS, 1% Nonidet P-40, 0.5% sodium deoxycholate, and centrifuged at $10,000 \times g$ for 15 min. CIC-2 protein was immunoprecipitated by incubation of the supernatants at 4°C with

anti CIC-2 antibody (Park et al., 1998) for 4 hr and then treated overnight with protein A-agarose beads (Pierce, Rockford, IL). The immunoprecipitates were resolved by 8.5% SDS/PAGE electrophoresis. After allowing for the decay of ^{32}P , western blot analysis was performed as we described previously (Park et al., 1998) to verify the amount of CIC-2 loaded in each lane.

For *in vitro* phosphorylation studies of CIC-2, immunoprecipitates from TSA-201 cells transiently transfected with the CIC-2/egfp bicistronic construct were incubated in either a protein kinase A (PKA), protein kinase C (PKC), or calmodulin-dependent protein kinase (CaMKII) assay buffer at 30°C for 60 min in the presence of 10 μCi of [γ - ^{32}P]ATP. The reactions were stopped by washing the protein A-agarose beads twice with RIPA buffer followed by two washes with RIPA buffer containing 1 M NaCl. The CIC-2 proteins were recovered by incubation of the beads in $2 \times$ Laemmli SDS sample buffer at 55°C for 15 min and analyzed by 8.5% SDS/PAGE. The fractionated proteins were then electro-transferred to a polyvinylidene difluoride (PVDF) membrane (Bio-Rad, Hercules, CA) and exposed to Kodak Biomax MS film. Western blot analysis was then performed as we described previously (Park et al., 1998). In the PKA reaction, immunoprecipitates were incubated with 1 ng of PKA catalytic subunit (Promega, Madison, WI) in 50 μl reaction buffer containing (in mM) 50 Tris-HCl (pH 7.4), 10 MgCl_2 , 0.1 ATP, 5 μM cAMP and 100 μg BSA/ml. In the PKC reaction, immunoprecipitates were incubated with 125 ng of PKC (Promega, Madison, WI) in 50 μl reaction buffer containing (in mM) 0.25 EGTA, 0.4 CaCl_2 , 20 Tris-HCl (pH 7.5), 10 MgCl_2 , 0.1 ATP, 100 μg BSA/ml, 320 μg phosphatidylserine/ml, and 32 μg diacylglycerol/ml. In the CaMKII reaction, immunoprecipitates were incubated with 100 ng of CaMKII (New England Biolabs, Beverly, MA) in 50 μl reaction buffer containing (in mM) 20 Tris-HCl (pH 7.5), 10 MgCl_2 , 5 DTT, 0.1 ATP, 0.1 EDTA, 2 CaCl_2 , and 2.4 μM calmodulin. Kinase activity was normalized to give the same incorporation of phosphate using enzyme-specific peptide substrates for the PKA, PKC, and CaMKII reactions: Kemptide (Promega, Madison, WI), Neurogranin (Promega, Madison, WI), and KRQSFDFL oligopeptide (New England Biolabs, Beverly, MA), respectively (*data not shown*).

ELECTROPHYSIOLOGICAL ANALYSIS

Whole-cell patch clamp recordings were done at room temperature (20–22°C) using an Axopatch 1-D amplifier (Axon Instruments, Foster City, CA). The measured junction potentials for the solutions used were less than 1 mV and so no correction for these was applied. The holding potential was 0 mV. Data acquisition was performed using a 12-bit analog/digital converter controlled by a personal computer.

Currents through expressed Cl^- channels were measured with an external solution in which the impermeant cation *N*-methyl-D-glucamine (NMDG) was used. The composition of this solution was (in mM): 140 NMDG-Cl, 2 CaCl_2 , 2 MgCl_2 , 20 HEPES, pH 7.1 (with NMDG). The internal solution used for these recordings contained (mM): 60 Cl, 60 glutamic acid, 120 NMDG, 3 Mg-ATP, 10 EGTA, 10 HEPES, pH of 7.2 (with NMDG). Very shortly after achieving whole-cell mode with these solutions, an outward rectifying current developed in both untransfected and transfected cells. This was likely the volume-sensitive organic osmolyte/anion channel (VSOAC—e.g., (Cannon, Basavappa & Strange, 1998)). This current was transient and disappeared within 10–15 min. The elimination of this current was accelerated by addition of 30 mM mannitol to the external solution. The magnitude and kinetics of expressed CIC-2 and CFTR-egfp were independent of the presence or absence of 30 mM mannitol in the external solution.

Results

IN VIVO AND IN VITRO PHOSPHORYLATION OF RAT CIC-2 BY PROTEIN KINASE A

The primary amino acid sequence of CIC-2 was analyzed for putative phosphorylation sites. Potential cAMP protein kinase phosphorylation sites were identified using a Prosite consensus sequence [RK]2-x-[S/T] (Taylor, 1989), or [RK]2-x-x-[S/T], as described (Feramisco, Glass & Krebs, 1980). As there are a number of exceptions to these rules, an 80% match was used as a threshold value. Rat CIC-2 contains three potential phosphorylation sites for both PKA and Ca^{2+} /calmodulin-dependent protein kinase II (CaMKII; at Ser82, Ser710 and Thr898), and 10 potential phosphorylation sites for PKC. All PKA sites are conserved in mouse, rabbit and human CIC-2 protein and the localization of these sites to the cytoplasmic amino and carboxy termini of CIC-2 indicates that they are accessible to intracellular kinases.

There is considerable evidence that CIC-2 is regulated by protein kinases (Fritsch & Edelman, 1996; Kibble et al., 1997; Sherry et al., 1997; Tewari et al., 2000); however, the mechanism of regulation has never been directly addressed. To determine whether rat CIC-2 is a substrate for protein kinases *in vivo*, TSA-201 cells were metabolically labeled with $^{32}\text{P}_i$ and then stimulated with either a PKA-activating cocktail or the PKC activator PMA (*see Methods*). Panel A of Fig. 1 shows that PKA activation leads to CIC-2 phosphorylation *in vivo*, but PKC produced little, if any, phosphorylation. To demonstrate the relative amount of CIC-2 protein added to each lane, a western blot of the membrane used in panel A was performed using a CIC-2-specific antibody (Fig. 1, panel B).

Figure 1 shows that activation of PKA results in phosphorylation of CIC-2 *in vivo*. However, this experiment does not differentiate between direct phosphorylation of CIC-2 by PKA and phosphorylation of CIC-2 as the consequence of PKA acting through another kinase. To address this issue we immunoprecipitated CIC-2 expressed in TSA-201 cells and then determined whether PKA, PKC, or CaMKII can phosphorylate CIC-2 *in vitro*. Panel A of Fig. 2 shows that *in vitro* PKA phosphorylated the expressed CIC-2 protein, but PKC and CaMKII failed to do so. To determine the relative amount of CIC-2 loaded in each lane, western blot analysis was performed on the same membrane (panel B of Fig. 2). The protein indicated by the solid arrows in panels A and B is the predicted size for CIC-2 (~100 kDa). The broken arrows point out a much smaller protein (~75 kDa) that is very strongly phosphorylated by PKA (panel A) and is labeled by the CIC-2 antibody (panel B). The identity of this latter protein is unknown, but likely represents a cleavage product of

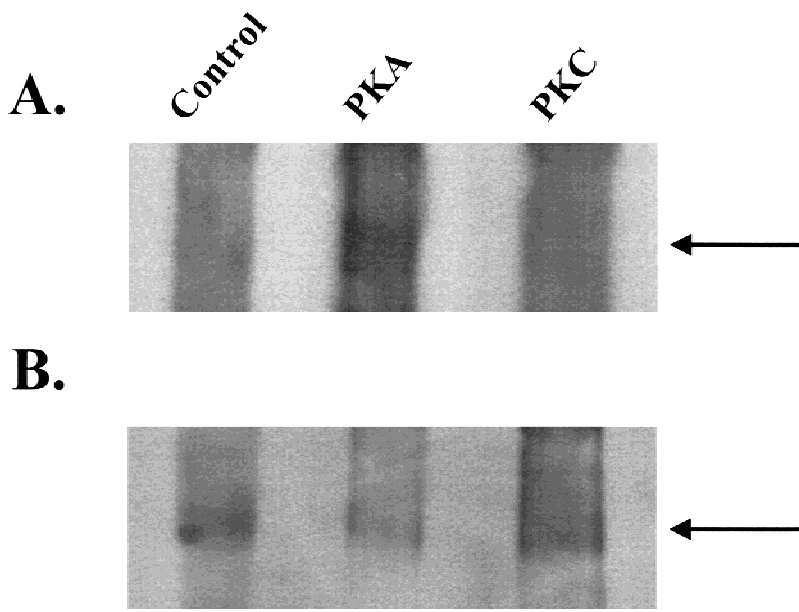


Fig. 1. PKA-dependent phosphorylation of CIC-2 in vivo. Rat CIC-2 was transiently expressed in TSA-201 cells and labeled with ^{32}P . Panel A. After labeling, the cells were treated for 15 min with either a PKA-stimulation cocktail [20 μM forskolin + 100 μM 3-isobutyl-1-methylxanthine + 500 μM 8-(4-chlorophenylthio) cAMP], or a PKC activator [0.5 μM Phorbol 12-myristate 13-acetate (PMA)], or vehicle (*Control*). The immunoprecipitates were resolved by 8.5% SDS/PAGE electrophoresis. Panel B. To determine the relative amount of CIC-2 loaded in each lane, western blot analysis was performed after allowing for the decay of ^{32}P . The arrows in panels A and B indicate the position of CIC-2 (~100 kDa).

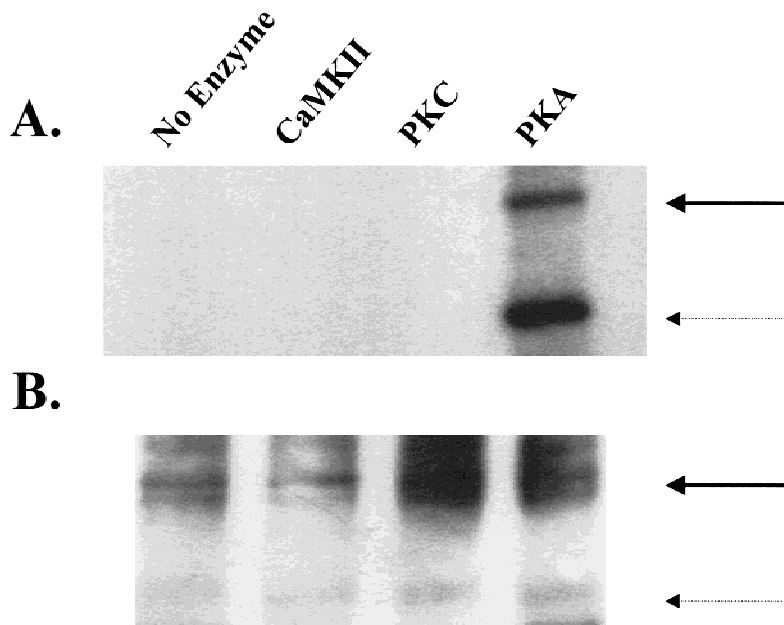


Fig. 2. In vitro labeling of CIC-2 by PKA. Rat CIC-2 was transiently expressed in TSA-201 cells and immunoprecipitated for in vitro labeling studies. Panel A. Immunoprecipitated CIC-2 was incubated in the presence of either PKA, PKC, CaMKII or no enzyme and 10 μCi of [γ - ^{32}P]ATP, analyzed by 8.5% SDS/PAGE, and transferred to PVDF membrane. Panel B. To determine the relative amount of CIC-2 loaded in each lane, western blot analysis was performed after allowing for the decay of ^{32}P . The solid arrows in panels A and B indicate the position of CIC-2 (~100 kDa). The broken arrow in panels A and B indicate the position of an ~75 kDa protein.

CIC-2. Moreover, this protein cannot represent the slightly smaller (~97–99 kDa) truncated form of CIC-2 (Cid et al., 2000; Loewen et al., 2000). Together, these experiments indicate that PKA activation phosphorylates CIC-2 and this in vivo event may be direct.

PROTEIN KINASE A ACTIVATION DOES NOT REGULATE RAT CIC-2 Channel Activity

To determine whether modulation of CIC-2 activity correlates with the phosphorylation induced by increased cAMP, chloride currents were measured using patch-

clamp techniques. The CIC-2/egfp bicistronic and CIC-2-egfp fusion constructs were used to facilitate identification of transfected cells expressing CIC-2. Figure 3 shows that treatment with a cAMP cocktail (the same cocktail used for in vivo labeling studies) for 20 min had no significant effect on the magnitude or the activation kinetics (Fig. 3, *inset*) of CIC-2. The average fraction change of the CIC-2 current at -120 mV was 0.99 ± 0.17 after 20 min stimulation ($n = 8$). The properties of the CIC-2 channel-egfp fusion protein ($n = 5$) were comparable to those without the GFP fusion ($n = 3$).

To verify that the stimulation protocol was sufficient to activate channels in TSA-201 cells, a CFTR-egfp fu-

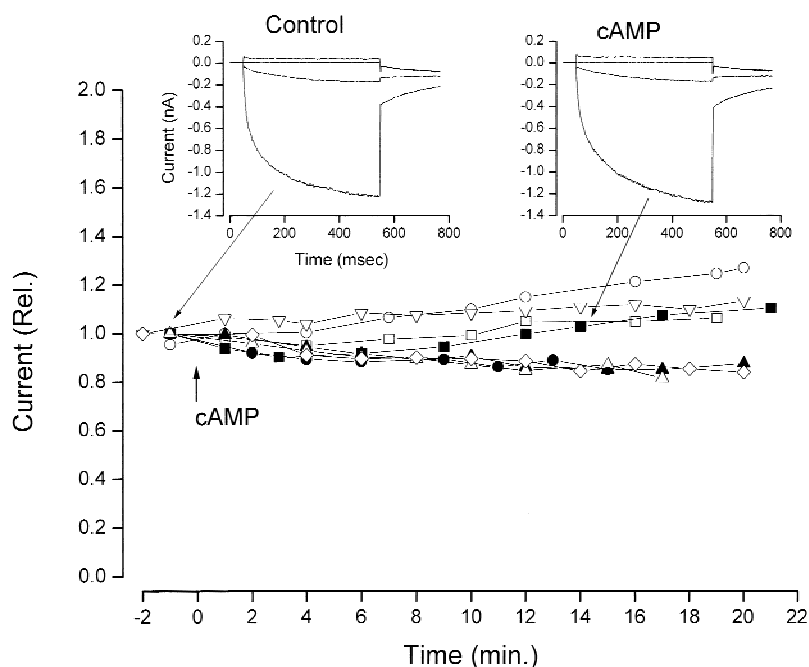


Fig. 3. CIC-2 currents in phosphorylating conditions. Currents at the end of hyperpolarizing pulses to -120 mV (except for Δ , which was -100 mV) as a function of time of the experiment. Currents normalized to the values recorded 2 min prior to application of the phosphorylating solution. The cAMP phosphorylating solution (*see Methods*) was added at zero time. Each symbol represents a different cell. Currents from cells represented by closed symbols were recorded with a solution of normal tonicity; 30 mM mannitol was added for the cells indicated by open symbols. Experiments represented by Δ , ∇ , and \diamond were done without the GFP fusion, whereas those represented by \blacktriangle , \square , \blacksquare , \circ , and \bullet were done with the CIC-2-egfp fusion. Insets: currents recorded at -120 , -60 , and $+60$ mV from the 0 mV holding potential. Tails recorded at -50 mV following the test pulse. Currents recorded from the indicated cell at the indicated times.

sion construct was used to express human CFTR. Figure 4 shows that the maximal response to the cAMP cocktail in most cells occurred in less than 5 min—the slowest responding cell showed effects within 5 min but the maximal response wasn't until 10 min. The average current at -80 mV before cAMP addition was -0.029 ± 0.006 nA in CFTR expressing cells. In contrast, the average current after cAMP addition was -6.1 ± 0.85 nA (SEM, $n = 6$), a greater than 200-fold increase in the current amplitude.

Discussion

CIC-2 is a voltage activated Cl^- channel that has wide tissue distribution suggesting that it may play an important functional housekeeping role in many different cell types. In the stomach, CIC-2 appears to be activated by low external pH, and thus could be involved in HCl secretion (Malinowska et al., 1995; Sherry et al., 1997; Stroffekova et al., 1998). CIC-2 is highly expressed in the brain (Thiemann et al., 1992), where it has been hypothesized to regulate the intracellular Cl^- concentration of neurons (Staley et al., 1996). According to this model, CIC-2 expression would produce a large negative shift in the Cl^- equilibrium potential and influence the production of action potentials. In the lung, CIC-2 expression is developmentally regulated, expression levels decreasing dramatically shortly after birth (Blaisdell et al., 1999; Chu et al., 1999; Murray, Chu & Zeitlin, 1996; Murray et al., 1995). Its targeting to the apical membrane of lung cells suggests that it may play an important role in Cl^- and water secretion necessary for the devel-

opment of this organ (Blaisdell et al., 2000). However, CIC-2-deficient mice were normal for these proposed functions; targeted disruption of the *Clcn2* gene leads to severe degeneration of male germ cells and photoreceptors (Bösl et al., 2001).

Understanding the acute regulation of this protein may have important clinical implications. In the present study, we show that rat CIC-2 is phosphorylated following PKA activation *in vivo*, and is directly phosphorylated by PKA *in vitro*. Nevertheless, the currents generated by these channels are not affected by this PKA-induced modification. In contrast, PKC (neither *in vivo* nor *in vitro*) and CaMKII (*in vitro*) failed to phosphorylate rat CIC-2. However, we cannot rule out the possibility that PKC and/or CaMKII activation might indirectly regulate CIC-2 channel activity. Our results are consistent with those of Schwiebert et al. (1998) and Jordt and Jentsch (1997) where no effect of PKA on the human CIC-2 channel was observed. Conversely, activation of PKA decreased the amplitude of the CIC-2 inward rectifying currents in human intestinal T84 cells (Fritsch & Edelman, 1996). However, CIC-2-like currents in mouse choroid plexus epithelial cells (Kibble et al., 1997) and human CIC-2 currents expressed in HEK293 cells (Tewari et al., 2000) are apparently activated by PKA. The simplest explanations for these differences are that PKA-dependent regulation of CIC-2 is cell-type or species-specific. In Fig. 3 we show the rat CIC-2 currents expressed in HEK293 and TSA-102 cells. In both cases, PKA activation failed to have an effect on CIC-2 current amplitude or activation kinetics. Thus, the cell type used for expression of CIC-2 cannot explain the difference in this case. Moreover, currents produced by

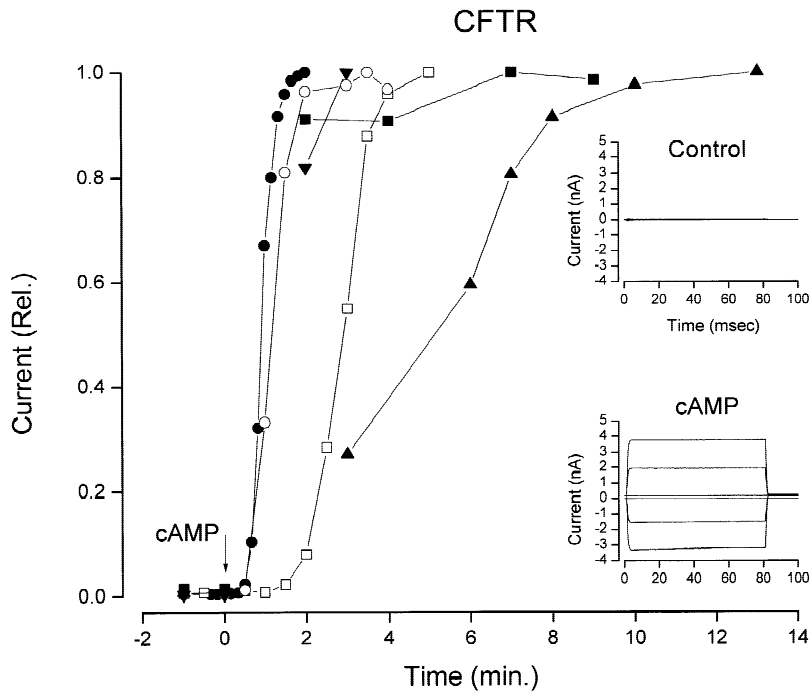


Fig. 4. CFTR currents in phosphorylating conditions. Currents at the end of hyperpolarizing pulses to -80 mV as a function of time of the experiment. Currents normalized by the largest recorded value. The cAMP phosphorylating solution (see Methods) was added at zero time. Each symbol represents a different cell. Insets: currents recorded at -60 , -30 , $+30$, and $+60$ mV from the 0 mV holding potential. Currents recorded from the cell identified by \blacktriangle before and after 10 min in the phosphorylating solution.

human CIC-2 when expressed in HEK293 cells (Tewari et al., 2000) show properties quite different from those seen for human CIC-2 when expressed in other cell types or for CIC-2 from other species. Specifically, both “trademarks” of CIC-2, the inward rectification and the time-dependent activation, are absent (Tewari et al., 2000).

In conclusion, we show protein kinase A activation phosphorylates rat CIC-2 in vivo, but this activation is independent of Cl^- channel activity. This has important implications as to the mechanism by which CIC-2 activity is acutely regulated.

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