

THEMED SECTION: ENDOTHELIUM IN PHARMACOLOGY

RESEARCH PAPER

Ca²⁺/calcineurin regulation of cloned vascular K_{ATP} channels: crosstalk with the protein kinase A pathway

NN Ori¹, AM Thomas¹, BA Perrino², A Tinker¹ and LH Clapp¹

¹BHF Laboratories, Department of Medicine, University College, London, UK, and ²Department of Physiology and Cell Biology, University of Nevada School of Medicine, Reno, NV, USA

Background and purpose: Vascular ATP-sensitive potassium (K_{ATP}) channels are activated by cyclic AMP elevating vasodilators through protein kinase A (PKA). Direct channel phosphorylation is a critical mechanism, though the phosphatase opposing these effects is unknown. Previously, we reported that calcineurin, a Ca²⁺-dependent phosphatase, inhibits K_{ATP} channels, though neither the site nor the calcineurin isoform involved is established. Given that the type-2 regulatory (RII) subunit of PKA is a substrate for calcineurin we considered whether calcineurin regulates channel activity through interacting with PKA.

Experimental approach: Whole-cell recordings were made in HEK-293 cells stably expressing the vascular K_{ATP} channel (K_{IR6.1}/SUR2B). The effect of intracellular Ca²⁺ and modulators of the calcineurin and PKA pathway on glibenclamide-sensitive currents were examined.

Key results: Constitutively active calcineurin A α but not A β significantly attenuated K_{ATP} currents activated by low intracellular Ca²⁺, whereas calcineurin inhibitors had the opposite effect. PKA inhibitors reduced basal K_{ATP} currents and responses to calcineurin inhibitors, consistent with the notion that some calcineurin action involves inhibition of PKA. However, raising intracellular Ca²⁺ (equivalent to increasing calcineurin activity), almost completely inhibited K_{ATP} channel activation induced by the catalytic subunit of PKA, whose enzymatic activity is independent of the RII subunit. *In vitro* phosphorylation experiments showed calcineurin could directly dephosphorylate a site in Kir6.1 that was previously phosphorylated by PKA.

Conclusions and implications: Calcineurin A α regulates K_{IR6.1}/SUR2B by inhibiting PKA-dependent phosphorylation of the channel as well as PKA itself. Such a mechanism is likely to directly oppose the action of vasodilators on the K_{ATP} channel.

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Keywords: Calcineurin; ATP-sensitive potassium channels; K_{IR6.1}/SUR2B; vascular; protein kinase A; whole-cell patch clamp

Abbreviations: [Ca²⁺]_i, intracellular Ca²⁺; CAP, calcineurin auto inhibitory peptide; CnA, catalytic subunit A of calcineurin; CnB, regulatory subunit B of calcineurin; CsA, Cyclosporin A; H-89, N-[2-(p-Bromocinnamylamino)ethyl]-5-isoquinolinesulphonamide-2HCl; HEK-293, human embryonic kidney 293; K_{ATP}, ATP-sensitive potassium; KCOs, K⁺ channel opening drugs; MBP, maltose-binding protein; NFAT, nuclear factor of activated T-cells; NDPs, nucleotide diphosphates; PKA, protein kinase A; PKA_{cat}, catalytic subunit of protein kinase A; Rp-cAMPS, Rp-2'-O-monobutyl-5-isoquinolinesulphonamide; SUR, sulphonylurea receptor; RII, type-2 regulatory subunit

Introduction

Vascular ATP-sensitive K⁺ (K_{ATP}) channels contribute to the maintenance of resting membrane potential and local blood

flow. They modulate vascular tone because of the steep relationship between membrane potential and Ca²⁺ influx through voltage-dependent Ca²⁺ channels (Quayle *et al.*, 1997). Vasoconstrictor and vasodilator hormones will influence this relationship through opposing effects on the channel (Quayle *et al.*, 1997; Buckley *et al.*, 2006). The structure of the K_{ATP} channel is an octomeric complex composed of a pore-forming subunit (K_{IR6.x}) to which ATP binds and a sulphonylurea receptor (SUR), the primary target for

Correspondence: Professor Lucie H Clapp, BHF Laboratories, Department of Medicine, UCL, Rayne Building, 5 University Street, London WC1E 6JF, UK. E-mail: l.clapp@ucl.ac.uk
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sulphonylureas, K⁺ channel opening drugs (KCOs) and nucleotide diphosphates (NDPs) (Seino and Miki, 2003). Based on subunit expression, channel characteristics and gene deletion experiments, Kir6.1/SUR2B almost certainly constitutes the vascular K_{ATP} channel (termed K_{NDP}) that is relatively insensitive to ATP, activated by NDPs and inhibited by glibenclamide (Chutkow *et al.*, 2002; Cui *et al.*, 2002; Miki *et al.*, 2002; Li *et al.*, 2003). The consequence of deleting either the SUR2 or Kir6.1 gene, is to produce coronary vasospasm, sudden death and a markedly reduced vasodilatory response to KCOs (Chutkow *et al.*, 2002; Miki *et al.*, 2002; Kane *et al.*, 2006). In addition, SUR2^{-/-} mice are hypertensive while Kir6.1^{-/-} mice have increased mortality towards endotoxin, and do not display the classic hypotensive response to this bacterial toxin (Kane *et al.*, 2006).

K_{ATP} channels represent an important target for vasodilators that elevate cyclic AMP. These include hormones such as calcitonin-gene related peptide, adenosine, prostacyclin and vasoactive intestinal peptide, whose effects of relaxation and hypotension are sensitive to glibenclamide (Quayle *et al.*, 1997; Buckley *et al.*, 2006; Yang *et al.*, 2008). Recently, the mechanism of channel activation was shown to involve direct phosphorylation by protein kinase A (PKA) of Ser and Thr residues located on SUR2B and Kir6.1 (Quinn *et al.*, 2004; Shi *et al.*, 2007; 2008a,b). While, the nature of the phosphatase opposing such phosphorylation is unknown, we have shown that the Ca²⁺-dependent phosphatase, calcineurin can regulate K_{ATP} channels both *in vitro* (Wilson *et al.*, 2000) and *in vivo* (Singer *et al.*, 2005). Such a mechanism allows the channel to sense changes in intracellular Ca²⁺ ([Ca²⁺]_i), being activated at resting levels and inhibited as [Ca²⁺]_i approaches micromolar levels (Wilson *et al.*, 2000). Thus, hormonal regulation of K_{ATP} channels is likely to be influenced by calcineurin, though this has yet to be demonstrated. Moreover, the mechanism by which this phosphatase inhibits the channel is unknown.

Calcineurin is a highly conserved, Ca²⁺/calmodulin-dependent Ser/Thr phosphatase with a catalytic subunit A (calcineurin A or CnA) that binds calmodulin and a regulatory subunit B (calcineurin B or CnB) that binds Ca²⁺ (Rusnak and Mertz, 2000). The CnA subunit is encoded by three separate genes, which give rise to CnA α , CnA β and CnA γ isoforms (Herzig and Neumann, 2000; Rusnak and Mertz, 2000). While CnA α and CnA β are co-expressed in most tissues, expression of the γ isoform is restricted to the testis and discrete regions of the brain (Herzig and Neumann, 2000; Eastwood *et al.*, 2005). Calcineurin is activated in response to a sustained elevation of cytoplasmic Ca²⁺ and is best known for its role in the Ca²⁺-dependent regulation of nuclear factor of activated T-cells (NFAT) which controls T-cell activation (Crabtree, 2001). In addition, this phosphatase is a major regulator of ion channel function, inhibiting or activating voltage-dependent Ca²⁺ channels (Schuhmann *et al.*, 1997), Ca²⁺-activated Cl⁻ channels (Greenwood *et al.*, 2004) and a variety of K⁺ channels (Czirjak and Enyedi, 2006; Loane *et al.*, 2006; Park *et al.*, 2006). While several different mechanisms may underlie the effects of calcineurin on ion channels, the type-2 regulatory (RII) subunit of PKA is a well-known cellular substrate for calcineurin (Perrino *et al.*, 2002) and has been shown to

co-localize with calcineurin in cardiac cells (Santana *et al.*, 2002). This raises the possibility that calcineurin may oppose PKA activation of the K_{ATP} channel through inhibiting PKA activity. Alternatively, it may directly dephosphorylate the channel itself. We therefore investigated Ca²⁺/calcineurin and PKA regulation of Kir6.1/SUR2B channels stably expressed in HEK-293 cells.

Methods

Stable lines were generated in human embryonic kidney 293 (HEK-293) cells containing Kir6.1 with SUR2B as described previously (Cui *et al.*, 2001).

Whole-cell recording

Membrane currents were recorded under voltage-clamp in the whole-cell recording configuration of the patch-clamp technique using an Axopatch 200B amplifier (Axon Instruments, Foster City, CA). Currents were filtered at 1 kHz and sampled at 2 kHz via a Digidata 1322A (Axon Instruments) interface. Data were acquired and analysed using pClamp8 computer software (Axon Instruments). Patch pipettes were made from thin walled (OD 1.5 mm) borosilicate glass capillaries (Harvard Apparatus, Edenbridge, Kent), which were pulled and fire-polished using a DMZ-Universal puller (Zeitz-Instruments, München, Germany) to give resistances of 2–4 M Ω . Electrode capacitance was reduced by coating tips with a parafilm/mineral oil suspension and was compensated electronically. Series resistance was compensated to 70% using the amplifier. Whole-cell bath solutions (pH 7.4) contained the following (in mmol·L⁻¹): 140 KCl, 5 HEPES, 1.2 MgCl₂ and 2.6 CaCl₂. Pipette solutions (pH 7.2) contained the following (in mmol·L⁻¹): 140 KCl, 5 HEPES, 1.2 MgCl₂, 10 EGTA, 3 MgATP, 0.5 Na₂GDP and 0–1.78 CaCl₂. CaCl₂ was added in an amount to yield 0, 18 and 36 nmol·L⁻¹ intracellular free Ca²⁺, as calculated with the CaBuf computer program (G Droogmans, Physiology Laboratory, KU Leuven, Belgium), which takes into account the buffering capacity of ATP, NDPs and Mg²⁺. All measurements of basal or drug-activated currents were made at least 10–15 min after 'break-in'. The magnitude of K_{ATP} current was assessed by sensitivity to 10 μ mol·L⁻¹ glibenclamide (I_{Glib}). Drugs were applied either through the pipette or in the bath solutions.

Synthesis of constitutively active calcineurin

Constitutively active calcineurin isoforms were created by introducing stop codons into the cDNA for CnA, causing the translated CnA subunits to truncate immediately to the C-terminal of the calmodulin-binding domain and delete the auto-inhibitory domain. Methodologies for cDNA manipulation, baculovirus screening and purification of CnA constructs using cultures of Sf21 cells have been described previously (Perrino *et al.*, 2002).

In vitro phosphorylation assay

Cloning, expression and purification of the maltose-binding protein (MBP) and the C-terminus of Kir6.1 (MBP-Kir6.1C)

were carried out as previously described (Quinn *et al.*, 2003). *In vitro* phosphorylation with the catalytic subunit of PKA was carried out as previously described (Quinn *et al.*, 2004) except that 2 h after phosphorylation, samples were washed 3 times with 1 mL of HEPES buffer followed by a single wash with 1 mL of calcineurin reaction buffer (50 mmol·L⁻¹ HEPES, pH 7.4, 18 nmol·L⁻¹ CaCl₂, 100 mmol·L⁻¹ NaCl, 6 mmol·L⁻¹ MgCl₂, 1 mmol·L⁻¹ DTT, 1% Triton X-100, EDTA-free complete protease inhibitor cocktail). Samples were then resuspended in 50 µL of calcineurin reaction buffer and a 10 µL aliquot removed for SDS analysis. Samples were subsequently centrifuged and all of the supernatant removed before addition of 9 µmol·L⁻¹ calmodulin ± 100 nmol·L⁻¹ of human recombinant CnA α subunit in calcineurin reaction buffer. Samples were incubated for 1 h at 37°C and then washed 4 times with 1 mL of calcineurin reaction buffer. The protein was subsequently eluted with 2 \times Laemmli gel loading buffer, run on a 10% SDS-PAGE gel and subjected to autoradiography.

Statistical analysis

Data were analysed using Clampfit 8.2 programme (Axon Instruments) and Graphpad Prism 4 (San Diego, CA). Values are given as means ± standard error of the mean (SEM) of glibenclamide-sensitive current (I_{glib}) densities (pA/pF), and n indicates the number of cells. Statistical significance was assessed using a paired or unpaired Student's t -test or one-way analysis of variance (ANOVA) with correction for multiple comparisons between different groups of cells. P -values < 0.05 were considered to be statistically significant.

Materials

Glibenclamide, Mg²⁺-adenosine triphosphate, Na₂ guanosine 5' diphosphate (GDP), forskolin, levcromakalim and the catalytic subunit of PKA (PKA_{cat}) were obtained from Sigma Chemical Co. (Poole, Dorset, UK). Cyclosporin A (CsA), calcineurin auto inhibitory peptide (CAP), okadaic acid, N-[2-(p-Bromocinnamylamino)ethyl]-5-isoquinolinesulphonamide·2HCl (H-89) Rp-2'-O-monobutryryl-cAMPS (Rp-cAMPS) and human recombinant CnA α subunit were all from Biomol (Exeter, UK). Gö6976 and calmodulin were bought from Calbiochem (San Diego, USA). Forskolin, H-89, Gö6976, levcromakalim and glibenclamide were dissolved in dimethyl sulphoxide but final concentrations after dilution were less than 0.1%, which had no effect on whole-cell currents. PKA_{cat} was prepared as a stock of 1 or 10 U·µL⁻¹ of a solution containing 6 mg·mL⁻¹ of DTT. All other stock solutions were made up in distilled water.

Nomenclature of molecular targets, including receptors and ion channels, is in accordance with the Guide to Receptor and Channels (Alexander *et al.*, 2008).

Results

Effect of Ca²⁺ on whole-cell K_{ATP} currents

We have previously demonstrated that increasing [Ca²⁺]_i inhibited K_{ATP} currents in freshly isolated vascular smooth muscle cells (Wilson *et al.*, 2000). We therefore wished to

establish if Kir6.1/SUR2B channels stably expressed in HEK-293 cells were similarly regulated by [Ca²⁺]_i. When cells were dialysed with solutions containing 0, 18 or 36 nmol·L⁻¹ free Ca²⁺ concentrations, basal currents evoked at all potentials were reduced as [Ca²⁺]_i was increased (Figure 1A). Likewise, the magnitude of glibenclamide-sensitive current (I_{glib}) was largest in the 0 Ca²⁺ pipette solution and smallest in cells dialysed with 36 nmol·L⁻¹ Ca²⁺, as shown by the average current-voltage (I - V) curves in Figure 1B. Thus, the magnitude of current measured at -80 mV dropped by 33% with 18 nmol·L⁻¹ Ca²⁺ (P < 0.05) and by 71% with 36 nmol·L⁻¹ Ca²⁺ (P < 0.001) compared with 0 nmol·L⁻¹ Ca²⁺ (Figure 1C). These results confirm that Ca²⁺ regulates Kir6.1/SUR2B.

Role of calcineurin

Having confirmed that Ca²⁺ regulates the channel, we investigated whether this involved signalling through calcineurin. We used two chemically unrelated inhibitors, calcineurin auto-inhibitory peptide (CAP; 100 µmol·L⁻¹) and the immunophilin, CsA (Cyclo A; 10 µmol·L⁻¹). Representative time-dependent plots comparing the magnitude of currents at -80 mV with and without 100 µmol·L⁻¹ CAP in the pipette solution is shown in Figure 2A. In the presence of CAP, currents were noticeably larger, and gave rise to bigger glibenclamide-sensitive currents (I_{Glib}). In a series of experiments, CAP completely reversed the Ca²⁺-dependent inhibition of the channel, doubling the magnitude of I_{Glib} seen with 18 nmol·L⁻¹ intracellular free Ca²⁺ (Figure 2B,C) and increasing it ~3.5 fold in cells dialysed with 36 nmol·L⁻¹ Ca²⁺ (Figure 2E,F). Likewise, I_{Glib} in the presence of 10 µmol·L⁻¹ Cyclo A was significantly greater (P < 0.05; Figure 2C) than that observed in control cells.

Although our data indicated the involvement of calcineurin in the Ca²⁺-dependent regulation of Kir6.1/SUR2B, we wished to know whether regulation involved protein phosphatase type 1 (PP1), whose activity can be enhanced by calcineurin-dependent dephosphorylation of the PP1 inhibitory peptide inhibitor-1 (Perrino and Soderling, 1998; Herzig and Neumann, 2000). Therefore, the observed effects of CAP and CsA could in part reflect a decrease in PP1 activation. To examine this possibility, we internally applied the PP1 inhibitor, okadaic acid (1 µmol·L⁻¹) in cells dialysed with 18 nmol·L⁻¹ free Ca²⁺. This failed to change the magnitude of I_{glib} recorded at any potential (Figure 2C,D) leading us to conclude that PP1 is neither involved in the Ca²⁺-dependent regulation of Kir6.1/SUR2B nor has direct effects on the K_{ATP} channel itself. We also investigated the possibility that Ca²⁺ inhibits K_{ATP} current through activation of protein kinase C (PKC), which is known to inhibit both native and cloned vascular K_{ATP} channels (Quayle *et al.*, 1997; Thorneloe *et al.*, 2002; Quinn *et al.*, 2003). We found that 1 µmol·L⁻¹ Gö6976, an inhibitor of Ca²⁺-dependent isoforms of PKC, had no effect on K_{ATP} current magnitude at any potential whether cells were dialysed with 18 or 36 nmol·L⁻¹ Ca²⁺ (Figure 2C-E).

To further confirm that calcineurin could indeed inhibit K_{ATP} channels, cells were dialysed with constitutively active recombinant isoforms of the catalytic subunit, CNA α or CNA β (Perrino *et al.*, 2002). These were added to the patch

pipette at a concentration of $100 \text{ nmol}\cdot\text{L}^{-1}$ and experiments carried out in $0 \text{ nmol}\cdot\text{L}^{-1} \text{ Ca}^{2+}$ to ensure minimal contribution from endogenous calcineurin. Figure 3 shows that $\text{CNA}\alpha$ but not $\text{CNA}\beta$ inhibited K_{ATP} current, giving rise to basal currents ($35 \pm 9.6 \text{ pA/pF}$, $n = 6$) of similar magnitude to that observed in $36 \text{ nmol}\cdot\text{L}^{-1} \text{ Ca}^{2+}$ ($27.0 \pm 7.5 \text{ pA/pF}$, $n = 16$). These results therefore suggest that $\text{CNA}\alpha$ is the isoform likely to be involved in channel regulation.

Modulation by PKA

As PKA phosphorylation of Kir6.1/SUR2B can lead to channel opening (Quinn *et al.*, 2004) and calcineurin is known to dephosphorylate and inhibit the RII subunit of PKA (Perrino *et al.*, 2002), we investigated whether calcineurin/ Ca^{2+} effects on the channel involved inhibition of the PKA pathway. As a first step, we sought to confirm that PKA regulated basal channel activity by using two inhibitors, H-89 ($10 \mu\text{mol}\cdot\text{L}^{-1}$) and Rp-cAMPs ($100 \mu\text{mol}\cdot\text{L}^{-1}$). We found that both these agents significantly attenuated the magnitude of K_{ATP} current measured in the presence of $18 \text{ nmol}\cdot\text{L}^{-1} \text{ Ca}^{2+}$ (Figure 4A,B) by 56% and 70% respectively. Next, we examined the combined effect of H-89 ($10 \mu\text{mol}\cdot\text{L}^{-1}$) and CAP ($100 \mu\text{mol}\cdot\text{L}^{-1}$) included in the same patch pipette. Using this approach, the magnitude of the I_{Glib} , while greater than that observed with H-89 alone (Figure 4B), was still significantly less ($P < 0.05$) than that observed if cells were dialysed with CAP alone under the same experimental conditions (Figure 2C). These results are consistent with the notion that a portion of calcineurin action involved inhibition of PKA.

Likewise, the effects of the adenylate cyclase activator, forskolin, whose stimulatory action on K_{ATP} channels in HEK-293 cells is completely dependent on PKA (Quinn *et al.*, 2004), were essentially abolished when Ca^{2+} was raised from 18 to $36 \text{ nmol}\cdot\text{L}^{-1}$ (Figure 5A,B). Again, this could be partially restored ($P < 0.001$, $n = 8$) when the pipette also contained $100 \mu\text{mol}\cdot\text{L}^{-1} \text{ CAP}$ (Figure 5B), suggesting that suppression of forskolin activation of the channel by Ca^{2+} involved calcineurin either acting on PKA or on the channel itself. To distinguish between these possibilities, two experimental approaches were used. First, we examined channel regulation using the constitutively active catalytic subunit of PKA (PKA_{cat}), whose activity can no longer be inhibited by calcineurin because it lacks the RII subunit. Dialysis of cells with PKA_{cat} ($10 \text{ U}\cdot\text{mL}^{-1}$) produced K_{ATP} currents that were 3 times larger than in control cells (Figure 5C,D). However, on raising the intracellular free Ca^{2+} to $36 \text{ nmol}\cdot\text{L}^{-1}$ (equivalent to increasing calcineurin activity), the same concentration of PKA_{cat} failed to have any effect on whole-cell currents. Some activation was achieved when the concentration was raised to $100 \text{ U}\cdot\text{mL}^{-1}$, though this failed to reach significance. These results strongly suggest that PKA and calcineurin can compete for the same phosphorylation sites on the K_{ATP} channel. This contrasts with responses to the classical K_{ATP} channel opener, levcromakalim ($10 \mu\text{mol}\cdot\text{L}^{-1}$) which activated currents of similar magnitude in both 18 and $36 \text{ nmol}\cdot\text{L}^{-1}$ free Ca^{2+} (Figure 5E). The other experimental approach was to examine the ability of calcineurin to dephosphorylate MBP-K_{IR}6.1C, a C-terminal domain of Kir6.1 that contains a serine site (S385) capable of being

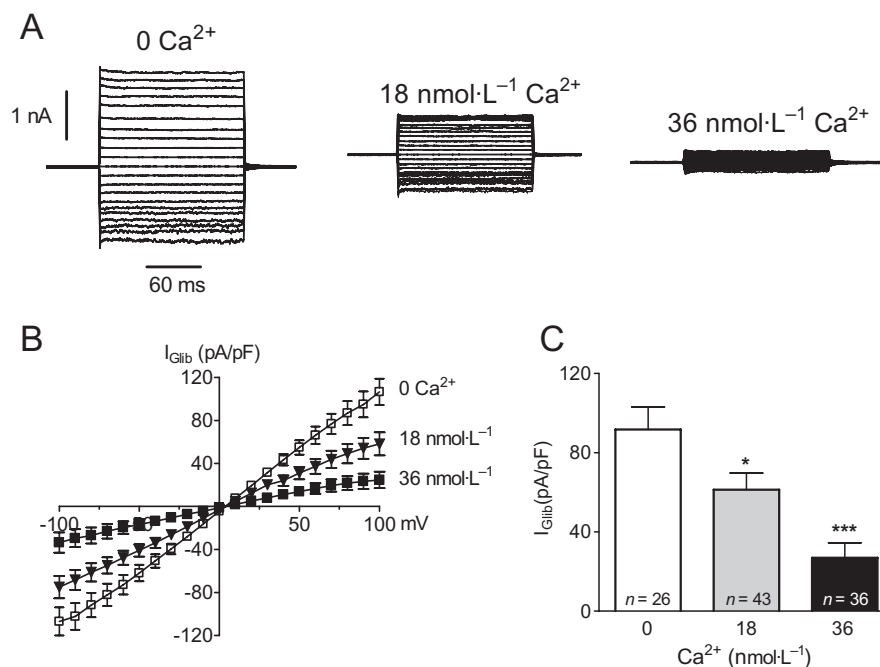


Figure 1 Intracellular Ca^{2+} inhibits whole-cell K_{ATP} currents in HEK-293 cells stably expressing Kir6.1/SUR2B . (A) Recordings of membrane currents from three separate cells dialysed with a pipette solution containing 0, 18 or $36 \text{ nmol}\cdot\text{L}^{-1}$ free Ca^{2+} . Currents were evoked from a holding potential of 0 mV by stepping the voltage for 150 ms in 10 mV increments from -100 mV to $+100 \text{ mV}$. (B) Mean current-voltage (I - V) relationships of steady-state current recorded under the three different $[\text{Ca}^{2+}]$ conditions shown in A. Data have been plotted as glibenclamide-sensitive (I_{Glib}) current (control current minus that in the presence of $10 \mu\text{mol}\cdot\text{L}^{-1}$ glibenclamide) and currents normalized to cell capacitance. (C) Mean I_{Glib} evoked at -80 mV taken from data in B. * $P < 0.05$, *** $P < 0.001$ when compared with 0 Ca^{2+} .

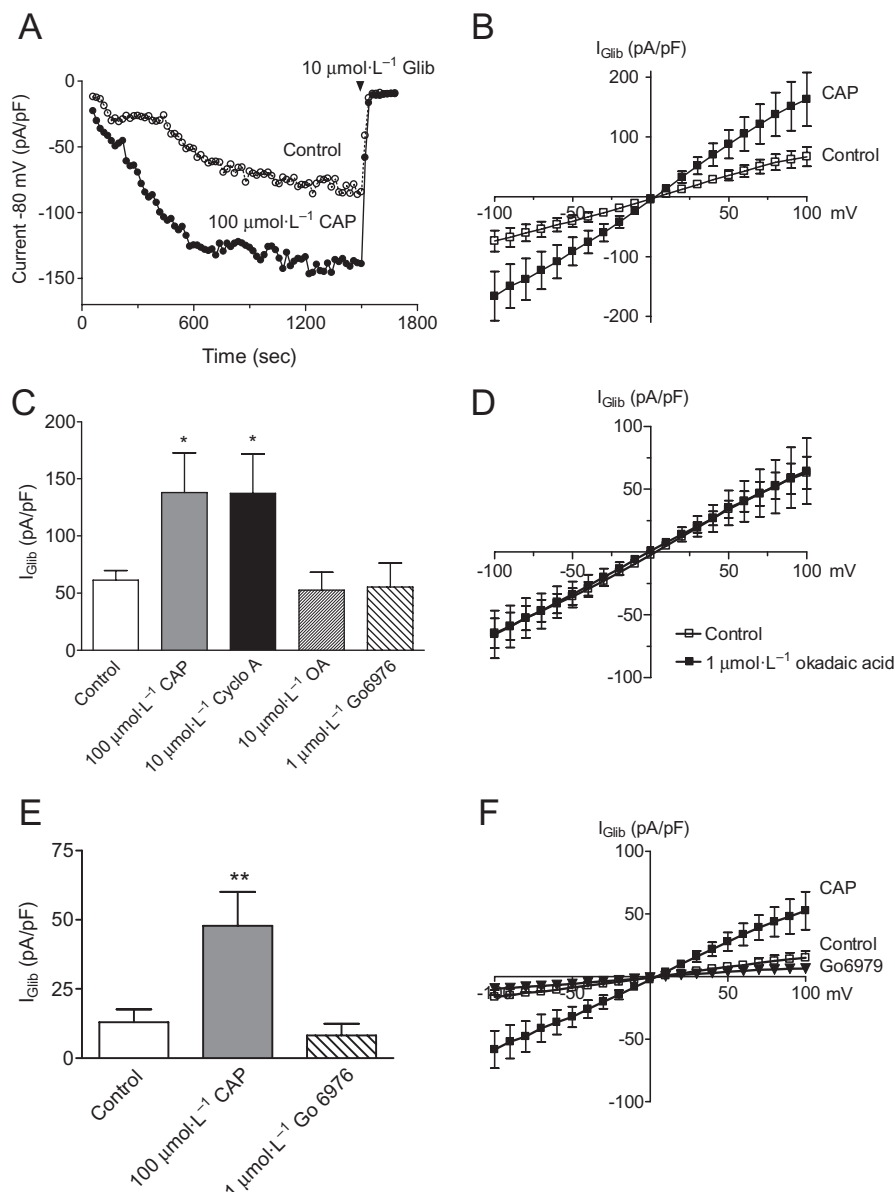


Figure 2 Calcineurin but not PP1 or PKC inhibitors increase Kir6.1/SUR2B currents. (A) Time-course of currents recorded from two cells dialysed in the absence (control) and presence of calcineurin auto-inhibitory peptide (CAP; $100 \mu\text{mol}\cdot\text{L}^{-1}$) in the pipette. Currents were evoked by voltage steps (150 ms duration) applied from a holding potential of 0 mV to -80 mV and repeated every 15 s. Time 0 represents the onset of recording, and glibenclamide ($10 \mu\text{mol}\cdot\text{L}^{-1}$) was given at 25 min to assess the size of basal K_{ATP} current. (B) Mean I_{Glib} plotted at different potentials in the absence (control, $n = 19$) or presence of $100 \mu\text{mol}\cdot\text{L}^{-1}$ CAP ($n = 8$). Currents were elicited by 150 ms voltage steps from -100 to $+100$ mV. (C) Mean I_{Glib} evoked at -80 mV in control cells ($n = 43$) compared with cells dialysed with CAP ($n = 8$), cyclosporin A (Cyclo A, $n = 10$), Gö6976 ($n = 17$) or okadaic acid (OA, $n = 4$). (D) I–V plots generated as in B in the absence (control) or presence of OA. (E) Mean I_{Glib} evoked at -80 mV in cells dialysed with $36 \text{ nmol}\cdot\text{L}^{-1}$ under control conditions ($n = 16$) or in the presence of CAP ($n = 8$) or Gö6976 ($n = 5$) in the pipette. (F) I–V plots generated using the protocol described in B in the absence (control) or presence of CAP or Gö6976 in pipette solutions containing $36 \text{ nmol}\cdot\text{L}^{-1}$ Ca^{2+} . In all other cases (A–D), the pipette solution contained $18 \text{ nmol}\cdot\text{L}^{-1}$ Ca^{2+} . * $P < 0.05$, ** $P < 0.01$ when compared with control.

phosphorylated by PKA (Quinn *et al.*, 2004). Figure 6A shows a Coomassie-stained SDS-PAGE gel loaded with either MBP or MBP-K_{IR}6.1C in the absence or presence of PKA_{cat} and CnA α . In essence, the gel shows roughly equivalent loading of MBP-K_{IR}6.1C (doublet between 60 and 75 kD) in the phosphorylation assay shown in Figure 6B. As described previously, MBP-K_{IR}6.1C was a substrate for PKA-mediated phosphorylation, an effect that was subsequently reduced in the presence of CnA α (Figure 6B). In contrast, PKA_{cat} does not

phosphorylate the MBP control protein nor does CnA α have any effect. Therefore, these results strongly suggest that the pore of K_{IR}6.1 can act as a direct substrate for CnA α mediated dephosphorylation.

Discussion

Previously, we demonstrated that Ca^{2+} regulates basal K_{ATP} channel activity in vascular smooth muscle cells isolated from

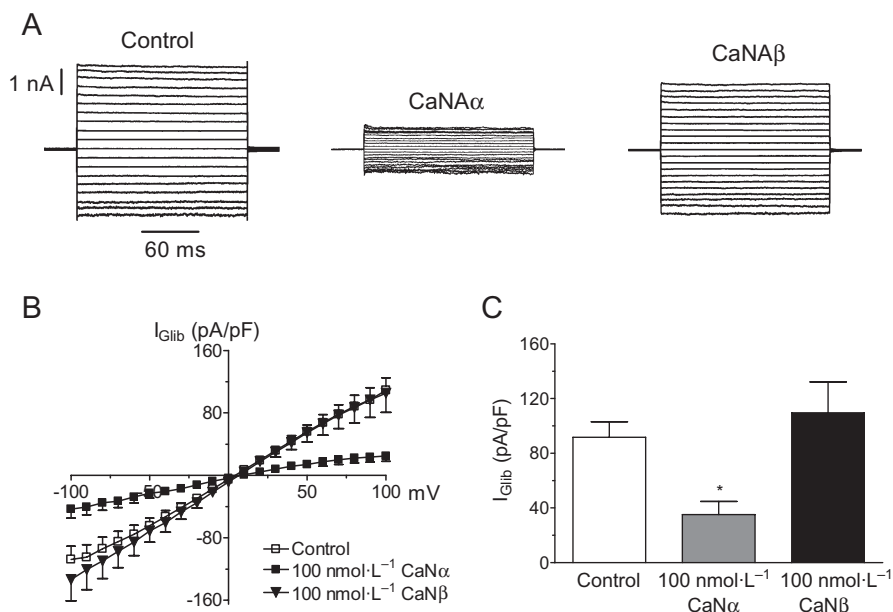


Figure 3 CnA α but not CnA β inhibits whole-cell K_{ATP} currents. (A) Recordings of membrane currents from three separate cells dialysed with a pipette solution containing 0 added Ca^{2+} in the absence (left panel) or presence of constitutively active CnA α (middle panel) and CnA β (right panel). Currents were evoked from a holding potential of 0 mV by stepping the voltage in 10 mV increments from -100 mV to +100 mV. (B) Mean I_{Glib} plotted at different potentials in the absence (control, $n = 19$) or presence of either 100 nmol·L⁻¹ CnA α ($n = 6$) or CnA β ($n = 7$) in the pipette. (C) Mean I_{Glib} evoked at -80 mV in control cells compared with cells dialysed with constitutively active CnA β or CnA α . * $P < 0.05$ when compared with CnA β .

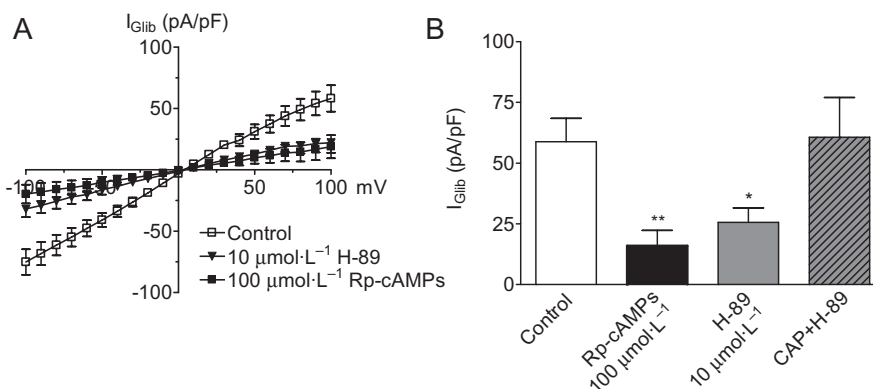


Figure 4 Ca^{2+} /calcineurin regulation of K_{ATP} channels is intrinsically linked to PKA. (A) Mean I_{Glib} plotted at different potentials in the absence (control, $n = 15$) or presence of either H-89 ($n = 8$) or Rp-cAMPS ($n = 9$) in the pipette. (B) Mean I_{Glib} evoked at -80 mV in control cells compared with cells dialysed with either Rp-cAMPS, H-89 or H-89 with 100 μ mol·L⁻¹ CAP ($n = 8$). In both A & B the pipette contained 18 nmol·L⁻¹ Ca^{2+} . * $P < 0.05$, ** $P < 0.01$ when compared with control.

rat aorta (Wilson *et al.*, 2000). In the present study, we investigated the mechanism underlying this regulation in HEK-293 cells stably expressing Kir6.1/SUR2B, a model system where many of the properties of native vascular K_{ATP} channels are faithfully reconstituted (Thorneloe *et al.*, 2002; Quinn *et al.*, 2003; 2004). Our data revealed that calcineurin is likely to be the main mediator of Ca^{2+} -dependent inhibition of Kir6.1/SUR2B and that this phosphatase inhibited channel function by opposing PKA-dependent phosphorylation. This conclusion is based on a number of observations. First, there was selective reversal of Ca^{2+} -induced channel inhibition by two chemically unrelated inhibitors of calcineurin, CsA and CAP. The latter corresponds to a C-terminal domain (residues 457–

482) of the calmodulin-binding domain of calcineurin making it a highly specific inhibitor with little effect on PP1, PP2A or CaM kinase II activity. Likewise, it was concluded that all Ca^{2+} -dependent regulation of $K_V2.1$ was attributable to calcineurin activity in HEK-293 cells (Park *et al.*, 2006). Second, constitutively active CnA α was able to inhibit K_{ATP} currents activated under conditions of low endogenous calcineurin activity (no added Ca^{2+} in the pipette), confirming our cyclosporin and CAP data. Third, PKA inhibitors substantially reduced basal K_{ATP} currents and responses to calcineurin inhibitors, suggesting crosstalk between these two signalling pathways (Figure 7). Fourthly, raising $[Ca^{2+}]_i$ to 36 nmol·L⁻¹ essentially abolished the activating effects of both PKA_{cat} and

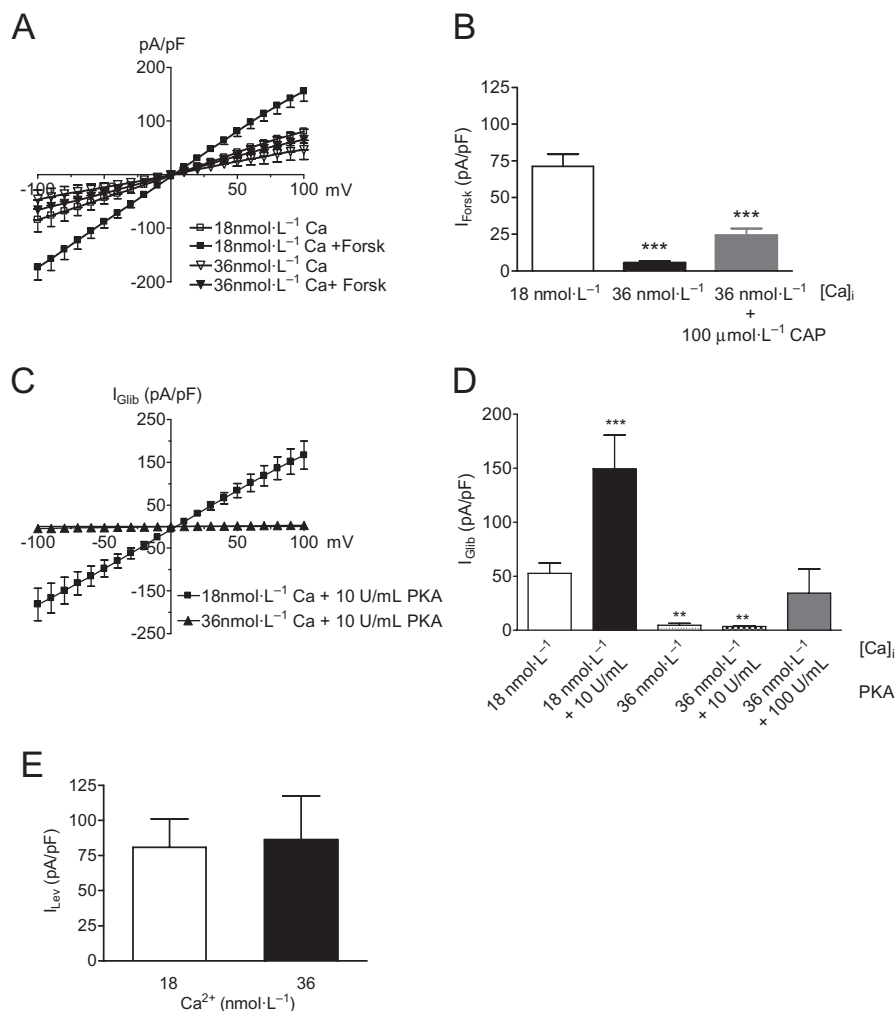


Figure 5 Ca²⁺/calcineurin disrupts PKA but not levcromakalim activation of K_{ATP} currents. (A) Mean current (pA/pF) plotted at different potentials before and 5 min after the application of 10 μmol·L⁻¹ forskolin at two different internal Ca²⁺ concentrations of 18 (*n* = 12) and 36 nmol·L⁻¹ (*n* = 9). (B) Average forskolin-induced current (I_{Forsk}) at -80 mV in cells in the absence or presence of 100 μmol·L⁻¹ CAP (*n* = 8). (C) Mean I_{Glib} induced by the catalytic subunit of PKA in the presence of either 18 (*n* = 11) or 36 (*n* = 17) nmol·L⁻¹ free Ca²⁺ in the pipette. (D) Column graph of currents evoked at -80 mV in the absence and presence of activated PKA in the patch pipette. (E) Mean levcromakalim current (I_{lev}) induced at -80 mV with either 18 (*n* = 5) or 36 nmol·L⁻¹ (*n* = 7) free Ca²⁺ in the pipette. Currents in the presence of 10 μmol·L⁻¹ levcromakalim were subtracted from those obtained before application. ***P* < 0.01, ****P* < 0.001 when compared with control.

forskolin. As PKA_{cat} is constitutively active and therefore not subject to deactivation by calcineurin, this failure could only result from dephosphorylation of sites normally phosphorylated by PKA. Lastly, consistent with direct channel effects, CnAα dephosphorylated MBP-K_{IR}6.1C, a C-terminal domain of K_{IR}6.1 which contains a serine site (S385) directly phosphorylated by PKA and critical for channel activation by either forskolin or PKA_{cat} (Quinn *et al.*, 2004). In other studies, direct binding of calcineurin to K⁺ channels has been demonstrated, though the interaction sites are different, involving CnB subunit for the Ca²⁺-activated K⁺ channel (Loane *et al.*, 2006) or the NFAT binding site on CnAα for TRESK (Czirjak and Enyedi, 2006).

One of the significant findings in the present study was the demonstration that CnAα but not CnAβ regulated K_{IR}6.1/SUR2B, with only the former significantly inhibiting whole-cell currents. Such an isoform-selective effect of CnAα has previously been described for Ca²⁺-activated Cl⁻ channels in

pulmonary artery (Greenwood *et al.*, 2004). So far this is the only isoform reported to regulate either Ca²⁺-activated K⁺ channels (Loane *et al.*, 2006) or TRESK, a member of the twin pore family of K⁺ channels (Czirjak and Enyedi, 2006). Moreover, CnAα has the appropriate cytosolic cellular distribution to regulate plasmalemmal ion channels in rat aortic smooth muscle whereas CnAβ does not, being largely confined to the perinuclear region, where it is differentially translocated into the nucleus in response to smooth muscle mitogens (Jabr *et al.*, 2007). In pulmonary artery, the cellular distribution of the two isoforms was found to be more homogeneous, though translocation to the membrane was only observed with CnAα, but not CnAβ when [Ca²⁺]_i was elevated to 500 nmol·L⁻¹ (Greenwood *et al.*, 2004). Despite an 85% homology between CnAα and CnAβ, different substrate affinities and catalytic activities are displayed by these two isoforms (Perrino *et al.*, 2002), which may contribute to their selective cellular function.

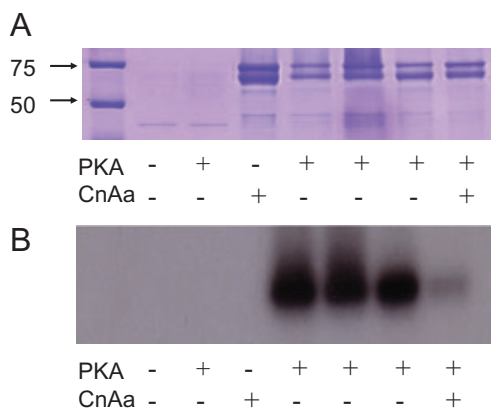


Figure 6 (A) 10% SDS-PAGE stained with Coomassie showing 2 μg of either MBP (lanes 2–3) or MBPKir6.1C (lanes 4–8) ± protein kinase A (PKA) and ± calcineurin α (CnAα). Marker sizes in kDa are indicated. B. Autoradiograph of the gel in A showing phosphorylation via PKA following a 2 h incubation and subsequent dephosphorylation in the presence of CnAα after 1 h. Presence or absence of PKA and CnAα are indicated.

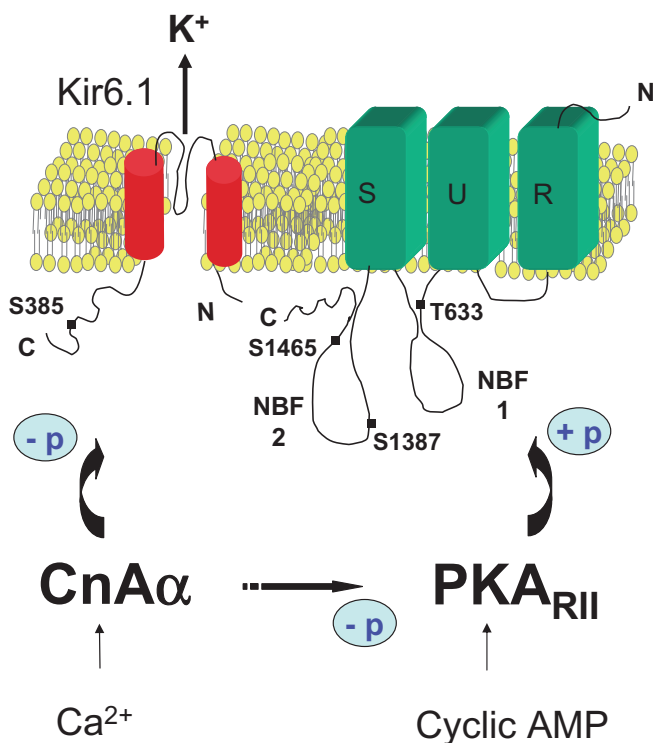


Figure 7 Diagram of possible mechanisms of calcineurin regulation of the vascular K_{ATP} channel. The putative sites involved in PKA and calcineurin regulation are shown on Kir_{6.1} and the nucleotide binding domains 1 (NB1) and 2 (NB2) in SUR2B.

In our study we found no evidence that either PP2A or PP1 regulated SUR2B/Kir6.1 under basal conditions. This conclusion is based on no discernible effect of okadaic acid, which blocks either phosphatase in the low to high nanomolar range respectively (Herzig and Neumann, 2000). Lack of involvement from PP1 was contrary to our expectation because calcineurin is known to activate PP1 through dephosphorylating either PP1 and/or its endogenous inhibitors

(DARPP32, inhibitor-1), thus making its activity potentially dependent on Ca²⁺/calcineurin in the intact cell (Hubbard and Cohen, 1989; Halpain *et al.*, 1990; Mulkey *et al.*, 1994). Given that HEK-293 cells do express endogenous PP1 (Morimoto *et al.*, 2004; Huang *et al.*, 2005), this might suggest that either the cellular localization of this phosphatase is not in the vicinity of the K_{ATP} channel or that it cannot oppose PKA phosphorylation of functional channel sites. Little is known about the role of phosphatases in vascular smooth muscle, though in rat mesenteric artery, PP1 and PP2A inhibitors either had no effect or slightly increased basal K_{ATP} currents while actually decreasing them in guinea-pig mesenteric artery (Firth *et al.*, 2000; Hayabuchi *et al.*, 2001). In contrast, PKC-mediated inhibition of K_{ATP} channels was significantly enhanced by okadaic acid in both vascular and non-vascular smooth muscle (Firth *et al.*, 2000) suggesting Ca²⁺-independent phosphatases may play a more prominent role in opposing PKC signalling. In tissues expressing other K_{ATP} channel subtypes, PP2A inhibits channel activity, promotes rundown and opposes the effects of PKC (Kubokawa *et al.*, 1995; Kwak *et al.*, 1996; Light *et al.*, 1996), while calcineurin appears a major regulator of skeletal muscle channels in humans (Singer *et al.*, 2005), but not in rat renal tubule cells (Kubokawa *et al.*, 1995).

We also considered the possibility that Ca²⁺ might inhibit the K_{ATP} channel through activation of Ca²⁺-dependent isoforms of PKC. We think this unlikely based on lack of effect of Gö6976, an inhibitor of conventional PKC isoforms (α,β,γ) which are activated by both Ca²⁺ and diacylglycerol. This finding is also consistent with previous observations showing angiotensin II inhibition of vascular K_{ATP} channels involves PKCε, a Ca²⁺-insensitive isoform of PKC (Hayabuchi *et al.*, 2001; Sampson *et al.*, 2007). Moreover, the same isoform is responsible for PKC-mediated inhibition of Kir6.1/SUR2B induced by phorbol esters (Quinn *et al.*, 2003).

We sought to investigate whether calcineurin could act by modulating the degree of phosphorylation of residues located on either the channel itself or a regulatory protein such as PKA. Our results support a mechanism whereby calcineurin and PKA can directly compete for the same residues on the K_{ATP} channel, with the degree of channel activation depending on the relative activities of PKA and calcineurin (Figure 7). Based on our phosphorylation data, we would surmise that S385 in Kir6.1 is one of the sites involved in this reciprocal regulation. This is consistent with the corresponding site in Kir6.2 (S372) being the major site of phosphorylation promoted by G_s-coupled receptors or direct stimulators of PKA (Beguin *et al.*, 1999). We believe however that additional PKA sites (Quinn *et al.*, 2004; Shi *et al.*, 2007; 2008a) contribute to the action of calcineurin. We have previously shown two other sites in SUR2B (T633, S1465) to be functionally involved in the activation of Kir6.1/SUR2B by cyclic AMP elevating agents (Quinn *et al.*, 2004; Shi *et al.*, 2007), while in other studies S1387, a site in the second nucleotide-binding domain of SUR2B, is reported to be the main site (Shi *et al.*, 2007; 2008a). It is also possible other serine sites (S1571 in SUR1) might contribute to basal channel activation (Beguin *et al.*, 1999). Taken together, it is likely that multiple sites are dephosphorylated by calcineurin. Indeed K_v2.1 channels expressed in HEK cells are

extensively phosphorylated at several serine sites under basal conditions (Park *et al.*, 2006). These sites can be variably dephosphorylated by calcineurin to give graded regulation of channel activity, a mechanism which allows for the fine tuning of neuronal action potential firing (Park *et al.*, 2006).

A multisite phosphorylation mechanism for PKA activation of the vascular K_{ATP} channel is not strongly supported by the data of Shi and colleagues. Based on systematic mutational analysis of all putative PKA sites, it was concluded that the main phosphorylation site responsible for the functional effects of forskolin or the β_2 -receptor agonist isoprenaline is S1387 (Shi *et al.*, 2007; 2008a). Moreover, on making the exact mutations they were unable to repeat the observations in our original paper (Quinn *et al.*, 2004). How, therefore, do we reconcile these seemingly conflicting data? The most notable difference in their studies is that experiments were performed in the presence of EGTA in the bath or pipette solutions with no added Ca^{2+} (Shi *et al.*, 2007; 2008a). Thus, based on our results presented here, we would predict that in 0 Ca^{2+} , more PKA sites (or the relative proportion of an individual site) would become phosphorylated due to the low calcineurin activity. The latter would also promote higher basal PKA activity, due to increased phosphorylation of the RII subunit (Figure 7). This may explain why forskolin-induced currents relative to activation by the K_{ATP} channel opener were smaller than we observed in the present study in 18 $nmol \cdot L^{-1}$ Ca^{2+} (40% vs. 100% respectively). Thus, it is likely that both groups are investigating contributions from different PKA sites under the different calcium levels. Furthermore, Shi and colleagues could not completely rule out a role for S385, as mutating this site to glutamate instead of alanine, did cause about a 30% reduction in the forskolin-induced activation of Kir6.1/SUR2B (Shi *et al.*, 2007). Lastly, contributions from other unidentified sites could not be ruled out given the moderate activation by PKA_{cat} observed in their S1387 mutant (Shi *et al.*, 2007).

What might be the functional significance of calcineurin regulation of the K_{ATP} channel? We postulate that under conditions where Ca^{2+} levels are elevated, for example, in the presence of a vasoconstrictor, this would promote channel closure. Indeed angiotensin II inhibits K_{ATP} channels in part through inhibiting basal PKA activity (Hayabuchi *et al.*, 2001), which we would argue results from calcineurin opposing PKA phosphorylation. The other consequence might be to limit and/or impair channel opening by vasodilator agents known to activate PKA. Interestingly, K_{ATP} channels do not contribute to forskolin-induced relaxation of phenylephrine contractions unless tissues are first incubated with endotoxin (Wilson and Clapp, 2002). In this context, opening may result from prolonged NO production and/or superoxide generation inhibiting calcineurin activity (Sommer *et al.*, 2002). The net effect would be to promote hyperphosphorylation of the K_{ATP} channel even at relatively high Ca^{2+} levels. A parallel might be drawn in patients on cyclosporin therapy who develop life-threatening hyperkalemia (raised plasma K^+ levels), which can successfully be treated with glibenclamide (Singer *et al.*, 2005). Thus low calcineurin activity is likely to drive channel opening under physiological and pathophysiological conditions.

In conclusion, this study reveals that calcineurin $A\alpha$ is the main mediator of the inhibitory action of Ca^{2+} on Kir6.1/SUR2B and that the mechanism involves dephosphorylation of PKA-phosphorylated sites on the channel. Such a mechanism may oppose the action of vasodilator hormones known to signal through PKA.

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Conflict of interest

None.

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