

Dual role of protein kinase C on BK channel regulation

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Large conductance voltage- and Ca²⁺-activated potassium channels (BK channels) are important feedback regulators in excitable cells and are potently regulated by protein kinases. The present study reveals a dual role of protein kinase C (PKC) on BK channel regulation. Phosphorylation of S⁶⁹⁵ by PKC, located between the two regulators of K⁺ conductance (RCK1/2) domains, inhibits BK channel open-state probability. This PKC-dependent inhibition depends on a preceding phosphorylation of S¹¹⁵¹ in the C terminus of the channel α -subunit. Phosphorylation of only one α -subunit at S¹¹⁵¹ and S⁶⁹⁵ within the tetrameric pore is sufficient to inhibit BK channel activity. We further detected that protein phosphatase 1 is associated with the channel, constantly counteracting phosphorylation of S⁶⁹⁵. PKC phosphorylation at S¹¹⁵¹ also influences stimulation of BK channel activity by protein kinase G (PKG) and protein kinase A (PKA). Though the S¹¹⁵¹A mutant channel is activated by PKA only, the phosphorylation of S¹¹⁵¹ by PKC renders the channel responsive to activation by PKG but prevents activation by PKA. Phosphorylation of S⁶⁹⁵ by PKC or introducing a phosphomimetic aspartate at this position (S⁶⁹⁵D) renders BK channels insensitive to the stimulatory effect of PKG or PKA. Therefore, our findings suggest a very dynamic regulation of the channel by the local PKC activity. It is shown that this complex regulation is not only effective in recombinant channels but also in native BK channels from tracheal smooth muscle.

phosphorylation | protein kinase A | protein kinase G | protein phosphatase 1 | tracheal smooth muscle cells

Large conductance Ca²⁺-activated potassium channels (BK channels) are unique in their regulation by both intracellular Ca²⁺ and membrane voltage. They are expressed in many tissues, and are particularly abundant in nerve and smooth muscle, where they play a key role as negative and positive feedback regulators of cell excitability by conducting repolarizing and hyperpolarizing outward currents, as has been impressively demonstrated in mice with targeted deletion of the pore-forming α -subunit (1–5). Alternative splicing of premRNA and protein phosphorylation generates structural and functional diversity of BK channels. Several serine/threonine kinases, such as the cAMP-(PKA)- and cGMP-dependent protein kinase (PKG) and protein kinase C (PKC), potently regulate BK channel activity. Their phosphorylation sites at the pore-forming α -subunit are fully conserved in almost all mammalian alternative splice variants, and mutation of the PKA and PKG phosphorylation sites abolished the kinase effect on channel activity in heterologous expression systems (6–10). In smooth muscle, PKA and PKG predominantly activate BK channels by increasing the apparent voltage and Ca²⁺ sensitivity of the channel, whereas PKC exerts opposite effects (11). Experimental evidence indicates that hormones and drugs that activate PKA or PKG contribute to smooth muscle relaxation by activation of BK channels. In contrast, activators of PKC seem to reinforce contraction by inhibiting BK channels as negative feedback regulators (12). Despite the potential physiological and pathophysiological relevance, the molecular mechanism of BK channel inhibition by PKC has remained elusive. This is surprising because

numerous hormones, neurotransmitters, and drugs bind to G protein-coupled receptors, which signal through pathways leading to activation of PKC.

We show in this study that PKC inhibits the open-state probability of BK channel α -subunits. This inhibition depends on a sequential phosphorylation of two distinct serines in the C terminus of the channel protein. In addition, both PKC phosphorylation sites have important effects on the regulation of BK channels by PKA and PKG in primary cells, e.g., freshly isolated smooth muscle cells of the mouse trachea.

Results

Inhibition of BK Channel Activity by PKC. The PKC-dependent BK channel regulation was investigated with the pore-forming BK channel α -subunit (KCNMA1) cloned from bovine trachea (BK_A, corresponding to the BK_{zero} variant) (9). When inside-out membrane patches were superfused at the cytosolic side with 30 nM of the catalytically active fragment of PKC (PKC_c), a decrease of macroscopic currents was observed at all voltages tested (–100 to +100 mV; Fig. 1A and B). The current decline started 30 s after the application of PKC_c and reached a stable level within 4–5 min. At +80 mV, PKC reduced membrane conductance (G_m) from 13.7 ± 0.7 to 5.9 ± 0.5 nS, i.e., by $57.2 \pm 3.4\%$ ($n = 7$). Reduced G_{max} could not be reversed by increasing voltage or Ca²⁺. The inhibitory PKC effect was abolished when inside-out patches were concurrently superfused for at least 5 min with PKC_c and 5 μ M of the PKC pseudosubstrate inhibitor peptide PKC_{19–31} (Fig. 1A). Representative macroscopic BK currents before (Ctr) and after application of 30 nM PKC_c are shown in Fig. 1B. To avoid effects on channel activity of the storage buffers in which the purified protein kinases were dissolved, freshly excised inside-out patches were superfused with control solutions containing the same amount of storage buffer as the solution with the respective kinase (*SI Materials and Methods*). Experiments carried out on inside-out patches with low channel density in the presence of symmetrically high potassium (140 mM) revealed that PKC decreased channel open probability, NP_o , by shortening channel open time and prolonging the closed state of the channel (Fig. 1C–E). Channel open and closed times were well fitted by two exponentials, corresponding to a short ($\tau_1 \sim 1$ ms) and a longer duration ($\tau_2 \sim 10$ ms for open time and ~ 20 ms for closed time). PKC affected only τ_2 ; it shortened the channel open time from 12.0 ± 5.3 to 7.0 ± 2.9 ms ($n = 9$) and prolonged the closed time from 23.9 ± 3.8 to 47.5 ± 4.7 ms ($n = 7$). The single-channel

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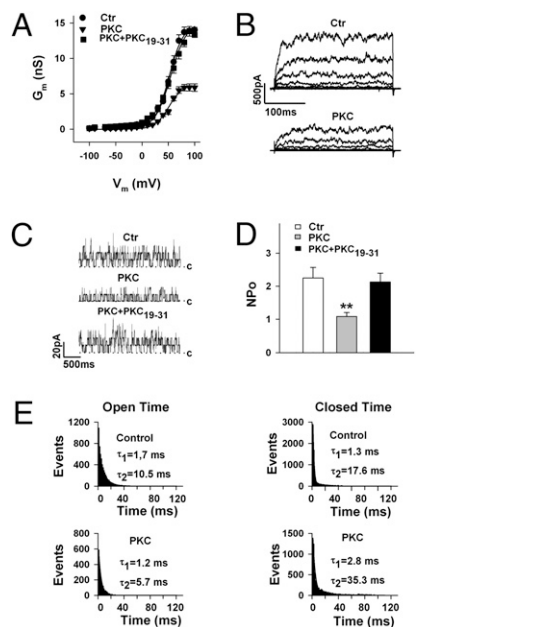


Fig. 1. PKC inhibits BK channel activity by decreasing channel open state probability. (A) Conductance-voltage relationships obtained from inside-out membrane patches of HEK293 cells expressing BK channels. Curves before (Ctr, $n = 15$) and in the presence of 30 nM PKC_c (PKC, $n = 7$) are shown. The PKC pseudosubstrate inhibitor peptide PKC₁₉₋₃₁ (5 μ M; $n = 8$) abolished the PKC effect. (B) Representative macroscopic BK channel currents before (Ctr) and in the presence of 30 nM PKC_c (–40 to +80 mV in 20-mV increments). (C) Decrease of NP_o by 30 nM PKC_c and abolition of this effect by the PKC inhibitor peptide; c indicates the closed state of the channel. Shown are single-channel currents from an inside-out patch with low channel density in the presence of symmetrically high potassium of 140 mM at +40 mV. (D) Summary of NP_o before (Ctr; $n = 11$) and in the presence of PKC_c ($n = 6$), or PKC_c plus PKC₁₉₋₃₁ ($n = 5$). Data are means \pm SEM. (E) PKC shortens the open state and prolongs the closed state of BK channels. Shown are open- and closed-time histograms fitted by two exponentials (τ_1 and τ_2). Recording conditions as in C, except that the data used for the histograms were taken from inside-out patches with only one active channel. $**P < 0.01$ vs. control (Ctr). The intracellular (bath) Ca^{2+} concentration was 1 μ M in A and B, 0.3 μ M in C–E.

conductance, the voltage dependence, and the calcium sensitivity of BK channels were not affected by PKC (Fig. S1).

PKC-Induced Inhibition Depends on Phosphorylation of Ser⁶⁹⁵ and Ser¹¹⁵¹.

By screening the BK channel α subunit for PKC consensus sequences (13), we found in addition to the prominent tandem motif with Ser¹¹⁵¹ and Ser¹¹⁵⁴ seven further serines—Ser³⁹⁰, Ser⁴⁹⁹, Ser⁶²⁵, Ser⁶³⁰, Ser⁶⁹⁵, Ser⁷¹², and Ser¹¹²⁹—as putative PKC phosphorylation sites. Whole-cell I_{BK} was measured in mutants where the respective serines were replaced by alanine. With the exception of the mutants S⁶⁹⁵A and S¹¹⁵¹A, at which the PKC activator phorbol 12-myristate 13 acetate (PMA) had lost its inhibitory effect, the activation of PKC reduced I_{BK} at all potentials by $\sim 50\%$ (Fig. 2A). In contrast, the inactive phorbol ester analog 4 α -PMA (100 nM) had no significant effect on BK channel currents. The positions of the critical residues Ser⁶⁹⁵ and Ser¹¹⁵¹ and the respective flanking regions within the BK_A isoform are shown in Fig. 2B.

To examine whether the negative charge added by phosphorylation of Ser⁶⁹⁵ and Ser¹¹⁵¹ is responsible for BK channel inhibition by PKC, we constructed mutants in which either serine was replaced by the negatively charged amino acid aspartate (S⁶⁹⁵D and S¹¹⁵¹D). Surprisingly, the S¹¹⁵¹D mutant did not significantly differ from the nonmutated BK channel with respect to membrane conductance, the half-maximal activating voltage ($V_{1/2}$) and inhibition of conductance by 30 nM PKC_c (cf. Fig. 2C with Fig. 1A). Introducing a negative charge at position 695 (S⁶⁹⁵D), however,

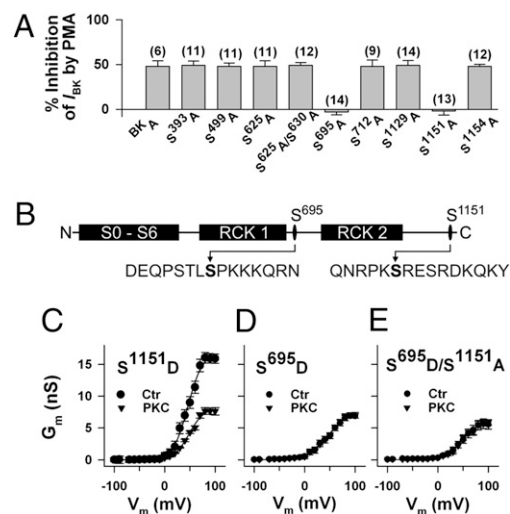


Fig. 2. Inhibition of BK channel activity by PKC depends on serines 695 and 1151. (A) Mutation of the PKC phosphorylation sites Ser⁶⁹⁵ and Ser¹¹⁵¹ to alanine abolished the inhibitory effect of the PKC activator phorbol 12-myristate 13 acetate (100 nM PMA) on BK channel whole-cell currents in transfected HEK293 cells. Percent inhibition of BK currents (I_{BK}) at +80 mV is shown with number of cells in parentheses. The pipette solution contained 0.3 μ M Ca^{2+} . I_{BK} before adding PMA was not significantly different between mutants (range: 121.2 ± 8.0 for S⁶²⁵A and 127.7 ± 12.3 pA pF⁻¹ for S⁴⁹⁹A) and BK_A (125.3 ± 13.1 pA pF⁻¹). (B) Schematic representation of the BK channel with the positions of Ser⁶⁹⁵ and Ser¹¹⁵¹ and their flanking regions, with the first methionine being in the context MANGG according to GenBank accession no. AAK54352.1. (C–E) Conductance-voltage relationships of mutants obtained from inside-out membrane patches before (Ctr) and after application of 30 nM PKC_c (PKC). Means \pm SEM of six (S¹¹⁵¹D) and eight (S⁶⁹⁵D/S¹¹⁵¹A) cells, respectively. Note, G_{max} of channels inhibited by PKC resembles G_{max} of phosphomimetic mutants. The intracellular (bath) Ca^{2+} concentration was 1 μ M.

resulted in channels with a strongly reduced membrane conductance resembling that of nonmutated BK channels in the presence of PKC (cf. Fig. 2D with Fig. 1A). At +80 mV, the mean membrane conductance of the S⁶⁹⁵D mutant was 6.4 ± 0.5 nS ($n = 6$), which is $47.4 \pm 2.1\%$ of the conductance measured in the nonmutated BK channel at the same potential. The mutant was insensitive toward PKC; a $V_{1/2}$ of 47.9 ± 3.6 mV ($n = 6$) vs. 51.6 ± 4.6 mV in seven controls indicates that its voltage-dependent activation was unchanged. Nearly identical results as those shown in Fig. 2D were obtained with the double mutant S⁶⁹⁵D/S¹¹⁵¹D. To determine the contribution of Ser¹¹⁵¹ to the inhibitory effect of PKC, we created the double mutant S⁶⁹⁵D/S¹¹⁵¹A. This mutant showed exactly the same electrophysiological characteristics as the mutants S⁶⁹⁵D and S⁶⁹⁵D/S¹¹⁵¹D, its membrane conductance was strongly reduced with respect to the nonmutated BK channel (by $59.8 \pm 3.9\%$ at +80 mV; $n = 8$), and PKC was ineffective (Fig. 2E). Taken together, these findings indicate that phosphorylation of Ser¹¹⁵¹ is constitutive (or unconditional) in nonmutated BK channels and a prerequisite for PKC-induced BK channel inhibition by phosphorylation of Ser⁶⁹⁵ (conditional phosphorylation). The decrease of membrane conductance depends solely on a negative charge at the amino acid position 695.

Stoichiometry of BK Channel Phosphorylation by PKC. BK channel α -subunits are assembled as tetramers. To investigate the stoichiometry of BK channel tetramer phosphorylation by PKC, we employed a strategy that allowed us to determine electrophysiologically the subunit composition of each BK channel from the single-channel current amplitude by introducing a mutation to the tetraethylammonium (TEA)-sensitive site (10, 14, 15). Introducing the point mutation Y³⁴⁷V in the subunit pore (Fig. 3A) strongly

reduces the sensitivity of BK channels expressed in outside-out patches to extracellular TEA (2 mM), whereas nonmutated channels (WT) were completely blocked (Fig. 3 *B* and *C*). The stoichiometry of mixtures of TEA-sensitive and TEA-insensitive α -subunits could thus be predicted from the single-channel current amplitude in the presence of 2 mM TEA (Fig. 3 *B* and *C*). We then examined in inside-out patches with 2 mM TEA in the patch-pipette channels formed from mixtures of BK α -subunits containing the TEA mutation ($Y^{347}V$, which retains the two PKC phosphorylation sites) coexpressed with TEA-sensitive BK channels in which either Ser⁶⁹⁵ (Fig. 3*A*) or Ser¹¹⁵¹ had been mutated to alanine. The TEA sensitivity of $S^{695}A$ and $S^{1151}A$ homotetramers was not significantly different from nonmutated BK channels. The application of 30 nM PKC_c to the cytosolic side of patches expressing either the homotetrameric $Y^{347}V$ mutant or the heterotetrameric channels, resulted in a robust inhibition of NP_o by ~60% (Fig. 3*D*; note that the data shown here for the mutant $S^{695}A$ are nearly identical with those obtained from the S^{1151} mutant and which are therefore omitted). For comparison, the PKC-induced inhibition of NP_o in homotetrameric nonmutated BK channels (WT), and the lacking PKC effect in homotetrameric $S^{695}A$ mutants, is also shown, albeit in the

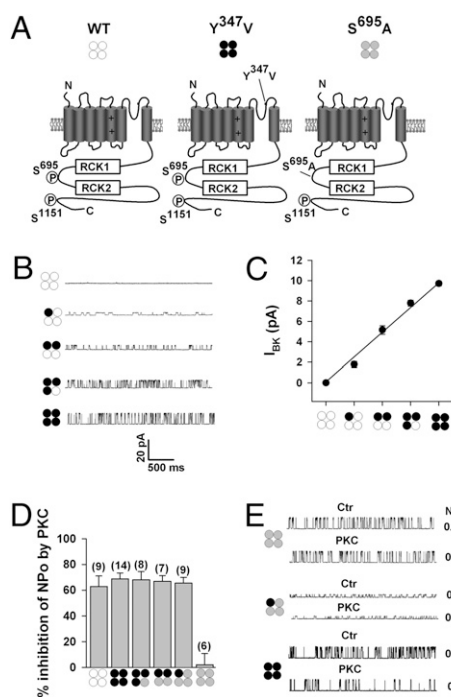


Fig. 3. PKC inhibition of BK channels requires phosphorylation of a single α -subunit within the tetramer. (A) Schematic of the nonmutated BK channel α -subunit (WT, open circles), the TEA-insensitive $Y^{347}V$ α subunit (black circles), and the PKC site mutant $S^{695}A$ (gray circles). The critical PKC phosphorylation sites S^{695} , located between regulator of K^+ conductance (RCK) domains, and S^{1151} are indicated. Representative single-channel traces (B) and summary plot of single-channel amplitude (C) of BK channels in the presence of 2 mM TEA, assembled as heterotetramers of BK and $Y^{347}V$ α -subunits. Pictograms illustrate predicted channel stoichiometry. Recordings were at +40 mV in symmetrically high potassium (140 mM) from outside-out patches of transfected cells, means \pm SEM, 6–14 patches per group. (D) Summary of effects of 30 nM PKC_c on NP_o by using BK- $Y^{347}V$ plus $S^{695}A$ heterotetramers with (3:1), (2:2), and (1:3) stoichiometry. For comparison, the PKC effect on homotetrameric WT-, $Y^{347}V$ -, and $S^{695}A$ - channels are also shown. Data from inside-out patches with symmetrically high potassium (140 mM) at +40 mV. With the exception of WT- and $S^{695}A$ homotetrameric channels, the recording pipette contained 2 mM TEA. Means \pm SEM; n in parentheses. (E) Representative single records before (Ctr) and NP_o after application of 30 nM PKC_c (PKC) to channels composed as indicated by the pictograms. Recordings as in D. Intracellular free Ca^{2+} concentration was 0.3 μ M.

absence of TEA. Taken together, the findings show that only one BK channel α -subunit within the tetramer needs to be phosphorylated for PKC-induced channel inhibition (Fig. 3 *D* and *E*).

PKC Modulates the Effects of PKA and PKG on BK Channels. We have shown earlier that the mutation of the tandem PKC motif in the C terminus to alanine ($S^{1151}A/S^{1154}A$) switches BK channel regulation from PKG to PKA (9). When we applied 300 nM PKG to the cytosolic face of excised inside-out membrane patches expressing the $S^{1151}D$ mutant, we observed a 26 mV shift to the left of the normalized conductance-voltage relation ($V_{1/2}$ before, 51.4 ± 4.2 mV, $n = 12$; $V_{1/2}$ after PKG 25.1 ± 4.4 mV; $n = 6$). Opposite to PKG, the catalytic subunit of PKA (300 nM) had no significant influence on the $S^{1151}D$ mutant ($V_{1/2}$ before 51.4 ± 4.2 mV, $n = 12$; $V_{1/2}$ after PKA 48.5 ± 3.8 mV; $n = 6$; Fig. 4*A*). These findings resemble those obtained earlier with the nonmutated BK channel (9). Inside-out membrane patches expressing the $S^{1151}A$ mutant, however, proved insensitive to stimulation by PKG but were responsive to PKA. $V_{1/2}$ obtained from normalized conductance-voltage relations was 52.1 ± 5.1 mV before (control, $n = 14$) and 50.7 ± 5.8 mV after the application of PKG ($n = 7$; not significant). In contrast, PKA shifted $V_{1/2}$ by 29 mV from 52.1 ± 5.1 ($n = 14$) to 22.7 ± 4.2 mV ($n = 7$; Fig. 4*B*). Nearly identical results as those shown in Fig. 4*A* and *B* were obtained when Ser¹¹⁵¹ was mutated on the $S^{695}A$ background (Fig. S2). Because phosphorylation/dephosphorylation of Ser¹¹⁵¹ seems to determine whether BK channels are regulated by PKG or PKA, we wondered whether BK channels under the influence of 30 nM PKC_c are still regulated by the two cyclic nucleotide-dependent kinases. When in the presence of PKC_c, PKG or PKA was applied to excised inside-out patches expressing the nonmutated BK channel, neither kinase enhanced membrane conductance at any potential (Fig. 4 *C* and *D*). The same result was obtained with the mutant $S^{695}D$ ($n = 6$) and the double mutants $S^{695}D/S^{1151}D$ ($n = 7$) and $S^{695}D/S^{1151}A$ (Fig. 4*E*; $n = 8$). Mutation of Ser⁶⁹⁵ in alanine abolished the inhibitory PKC effect on BK channels but had no influence on the stimulatory effect of additionally applied PKG (Fig. S3). Thus, BK channels in which Ser⁶⁹⁵ is phosphorylated by PKC or contain a negatively charged amino acid at this position are no longer regulated by cyclic nucleotide-dependent protein kinases.

PKC-Dependent Inhibition of BK Channels in Tracheal Smooth Muscle Cells. To verify that the regulation of BK channels described above occurs not only in a heterologous expression system but also under physiological conditions, we investigated the PKC-dependent regulation of BK channel activity in native cells. Recently, we showed that stimulation of muscarinic M_2 receptors by carbamylcholine (CCh) inhibits BK channels in isolated smooth muscle cells of the mouse trachea (TSMCs) by a dual mechanism involving PKC and $G_{\beta\gamma}$ (12). When we elicited whole-cell outward currents (I_{out}) in freshly isolated TSMCs, addition of the specific BK channel-blocking peptide iberiotoxin (300 nM) revealed that more than 90% of I_{out} was conducted by BK channels (Fig. S4). The application of 10 μ M CCh decreased within 5 min I_{out} at all potentials by ~50%. At 80 mV, current densities decreased from 129.9 ± 22.3 to 69.0 ± 8.3 pA pF^{-1} , i.e., by $46.3 \pm 4.7\%$ (Fig. 5*A*; $n = 16$). Additional application of the membrane permeant-specific activators of PKG (300 μ M 8-pCPT-cGMP) or PKA (300 μ M Sp-5,6-DCI-cBIMPS) for 10 min, produced no further significant change of I_{out} (Fig. 5*A*). In contrast, when the protein kinase activators were applied in the absence of CCh, only the PKG activator enhanced I_{out} by $63.1 \pm 5.2\%$ at 80 mV ($n = 6$), whereas the PKA activator or the β -adrenoreceptor agonist isoproterenol (10 μ M) were ineffective (Fig. 5*B*). To demonstrate that the abolition of the PKG effect in the presence of CCh was due to activation of PKC, we superfused TSMCs for 5 min with the PKC inhibitor Ro 31-8220 (1 μ M), which did not significantly affect I_{out} . When 10 μ M CCh was added thereafter, inhibition of I_{out} was not completely prevented because

potentials. At 40 mV, NP_o was 0.42 ± 0.03 before and 0.21 ± 0.02 after the addition of PPI2 (Fig. 5E; $n = 13$). Additional application of 300 nM PKG or PKA for 5 min produced no further significant effect on NP_o (Fig. 5E). When the experiments were repeated in the presence of the PKC inhibitor Ro 31-8220, PPI2 did not inhibit NP_o , and additional application of 300 nM PKG enhanced NP_o 2.6-fold at 40 mV from 0.52 ± 0.06 to 1.34 ± 0.15 (Fig. 5F; $n = 6$). Preferential inhibition of protein phosphatase 2A by 3 nM ocadaic acid for 10 min failed to induce BK channel inhibition and did not abolish PKG-dependent activation of BK channels (Fig. S5).

Discussion

The data reported in this study establish that inhibition of BK channels by PKC depends on the unconditional and conditional phosphorylation of the C-terminal serine residues 1151 and 695, respectively. The inhibition of BK channel conductance by PKC is due to a decrease of channel open probability without changing the unitary current amplitude. From the open- and closed-time histograms it is obvious that the inhibitory effect of PKC is mediated by altering both the opening and closing process to stabilize the non-conducting conformations of the channel. The voltage dependence and the Ca^{2+} sensitivity of the BK channel were not affected by PKC. In addition, the phosphorylation of the BK channel by PKC abolishes the stimulatory effects of PKA and PKG.

We investigated the mechanism underlying the PKC-induced BK channel inhibition using a BK channel isoform that contains no inserts at the alternative splice sites, and has been identified in a variety of species, including man (19, 20). Among nine putative PKC phosphorylation sites, two serines, Ser⁶⁹⁵ and Ser¹¹⁵¹, were identified by site-directed mutagenesis to be critical for PKC-dependent BK channel inhibition. Our experimental evidence indicates that Ser¹¹⁵¹ is constitutively phosphorylated in HEK293 cells. First, the replacement of Ser¹¹⁵¹ by aspartate (S¹¹⁵¹D) to mimic PKC-dependent phosphorylation resulted in mutant BK channels that resembled nonmutated BK channels with respect to G_{max} , $V_{1/2}$, inhibition by PKC, and activation by PKG. Second, the BK channel mutant S¹¹⁵¹A was insensitive to PKC and to stimulation by PKG. It was instead activated by PKA, which is in line with previous data (9) showing that phosphorylation/dephosphorylation of Ser¹¹⁵¹ determines the channel's ability to respond to either PKG or PKA. The phosphorylation of the second critical serine, Ser⁶⁹⁵ by PKC, is conditional. It depends on the preceding phosphorylation of Ser¹¹⁵¹. Most likely, this regulation relies on a conformational change within the C terminus that allows access of the PKC to Ser⁶⁹⁵ only when Ser¹¹⁵¹ is phosphorylated, or when a negatively charged amino acid, like in the mutant S¹¹⁵¹D, mimics the phosphorylation at this site. Ser⁶⁹⁵, which is highly conserved in BK channels of many vertebrates, is located in the linker between the two "regulator of K^+ conductance" (RCK) domains. Both RCK1 and RCK2 domains contain high-affinity Ca^{2+} -binding sites that form parts of a complex functional domain (gating ring) that converts the free energy of Ca^{2+} binding into mechanical work to open the channel (21–23). When Ser⁶⁹⁵ was replaced by aspartate to mimic phosphorylation at this position (S⁶⁹⁵D), BK currents were inhibited like under the influence of PKC. The same reduced channel conductance was also observed in the double mutant S⁶⁹⁵D/S¹¹⁵¹A, in which the constitutive phosphorylation of Ser¹¹⁵¹ was abandoned. The Ser⁶⁹⁵ mutants clearly demonstrate that a PKC-like inhibition of the BK channel depends exclusively on a negatively charged amino acid at position 695. BK channel α -subunits expressing a 59-amino acid stress-regulated exon at splice site 2 (STREX-1) exhibit an additional PKA consensus motif at serine residue 3 within the STREX insert (8, 10). PKA inhibits NP_o of STREX channels via phosphorylation of S3_{STREX}, probably by preventing palmitoylation of the C terminus (24). Interestingly, the S3_{STREX} is located within the RCK1 and RCK2 linker only a few amino acids upstream of Ser⁶⁹⁵. BK channels are assembled as tetramers of pore-forming α -subunits (14), each of which includes a PKC phosphorylation site. We found that phosphorylation of only one BK α -subunit either at Ser¹¹⁵¹ or

Ser⁶⁹⁵ is sufficient for channel inhibition. Thus, the PKC-dependent inhibition obeys a single subunit rule, which is similar to the inhibitory effect of PKA in STREX channels, but different from insertless BK channels where phosphorylation of all four α -subunits is required for channel activation by PKA (10). Whether prevention of palmitoylation of the C terminus also plays a role for PKC-dependent BK channel inhibition is presently unknown. Note, however, that the phosphorylation of S3_{STREX} by PKA produced a ~ 20 -mV rightward shift of the voltage-dependent activation curve (25), whereas PKC reduced the maximal BK channel activity without affecting the apparent voltage sensitivity. A similar observation has been made earlier with PKC on BK channels in pituitary CH₄C₁ cells (26).

The unconditional and conditional phosphorylation of BK channels by PKC implies differential tasks in channel regulation. Besides the direct inhibition of channel activity, PKC phosphorylation apparently modulates PKA- and PKG-mediated effects. In accordance with previously published data (9, 27), the phosphorylation of Ser¹¹⁵¹ alone switches BK channel activation from PKA to PKG, whereas the additional phosphorylation of Ser⁶⁹⁵ renders the BK channel completely insensitive to both cyclic nucleotide-dependent kinases. Such a coregulation is not unique to BK channels but has also been observed in other ion channels. In T lymphocytes the modulation of delayed rectifier potassium channels by PKA is enhanced by PKC phosphorylation (28). Similarly, voltage-gated sodium channels require PKC phosphorylation to allow modulation by PKA (29).

The inhibition of BK channels by phorbol esters and PKC_c has been shown before in isolated smooth muscle cells from different arterial beds (30, 31). In the present study, we used mouse TSMCs to prove that the regulation of BK channel activity by PKC in primary smooth muscle cells is identical to that observed in the recombinant channels. We chose TSMCs for two reasons. First, BK channels in mouse TSMCs represent the zero variant corresponding to the bovine trachea isoform. Second, the muscarinic receptor agonist CCh inhibits BK channels in TSMCs by activating the PKC via M₂ receptors (12). The data obtained in TSMCs are in complete accordance with the regulation observed with the recombinantly expressed BK channels. CCh decreased BK channel activity in TSMCs by $\sim 50\%$ and abolished the stimulating effect of the PKG activator 8-pCPT-cGMP, which was effective in the absence of CCh. The broad-spectrum PKC inhibitor Ro-31-8220 inhibited the effects of CCh. Dephosphorylation of presumably Ser¹¹⁵¹, by pretreating inside-out patches from TSMCs with ATP-free solution in the presence of the PKC pseudosubstrate inhibitor PKC₁₉₋₃₁, resulted in a switch of channel regulation from PKG to PKA, resembling the switch in the S¹¹⁵¹A mutant. The regulation of BK channels by closely associated phosphatases has been shown before (16–18). Therefore, the switch of channel regulation from PKG to PKA was likely due to the activity of a phosphatase that had dephosphorylated the channel at the PKC motif before the channel became responsive to PKA. Applying the selective protein phosphatase 1 peptide inhibitor PPI2 to inside-out patches of TSMCs resulted in a substantial decrease of NP_o and prevented channel activation by either PKG or PKA. This effect was dependent on PKC, because Ro 31-8220 abolished the inhibitory effect of PPI2 and allowed PKG to enhance NP_o . Inhibition of a protein phosphatase 2A by 3 nM ocadaic acid had no influence on BK channel activity. These findings, together with experiments performed in HEK293 cells expressing recombinant channels (Fig. S6), provide strong evidence that BK channels are intimately associated with protein phosphatase 1 and PKC, and that interfering with the balance between kinase and opposing phosphatase activities can lead to the conditional phosphorylation of Ser⁶⁹⁵ with the described consequences on channel regulation. The findings with PPI2 are intriguing because they disprove the concept of an indirect regulation of BK channels by protein kinases. It has been demonstrated that protein phosphatase inhibitors abolish or reverse the stimulatory effect of PKG (17) or of compounds that activate cGMP-dependent pathways, such as the natriuretic peptide ANP (32) or the β -amyloid-

precursor protein β -APP (33), on BK channel activity. These observations had been taken as evidence that BK channels are stimulated by dephosphorylation of the channel protein due to the activation of an associated protein phosphatase. In light of the present findings, however, it is very likely that protein phosphatase inhibitors in TSMCs lead to the phosphorylation of Ser⁶⁹⁵ by an endogenous PKC, which inhibits BK channels and renders them insensitive to cyclic nucleotide-dependent protein kinases.

In summary, we have shown that PKC via unconditional and conditional phosphorylation of the BK channel not only decreases channel open probability but also determines the regulation of the channel by other protein kinases such as PKG and PKA. By inhibiting BK channel activity and blunting the activating effects of PKG and PKA in TSMCs, PKC facilitates contraction and prevents relaxation via NO/cGMP- or cAMP-dependent pathways. The balance between phosphorylation by PKC and dephosphorylation by phosphatase 1 may vary in different cell types and may therefore explain the divergent results reported in the literature on BK channel regulation by PKG and PKA.

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Materials and Methods

HEK293 cells were cultured for 24 h and then transiently transfected with the pcDNA3 plasmid containing the BK channel α -subunit BK_A cloned from a bovine tracheal smooth muscle oligo dT-primed cDNA library (9) or with plasmids encoding the respective BK_A mutant. The GenBank accession numbers for the BK channel isoform are AAK54352.1 (protein) and AY033472 (mRNA; BK_A). Tracheal smooth muscle cells were isolated as described previously (12). Standard patch-clamp recording techniques were used to measure currents in the whole-cell, outside-out, or inside-out patch-clamp configuration. The membrane conductance (G_m) was calculated as $G_m = I/(E_m - E_{rev})$, where I is macroscopic current, E_m is the test membrane potential, and E_{rev} is the reversal potential for potassium. For normalization, G_m values were divided by G_{max} , where G_{max} was defined as the largest G_m value obtained in each experiment. G_m/G_{max} voltage curves were fitted with a Boltzmann equation of the form $G/G_{max} = (1 + \exp((V_{1/2} - V_m)/k))^{-1}$, where $V_{1/2}$ is the membrane potential (V_m) required for half-maximal activation of the channels, and k is the logarithmic voltage sensitivity (i.e., ΔV required for an e-fold increase in activity). Further details of the experimental procedures are given in *SI Materials and Methods*.