

## TOPICAL REVIEW

# BK-type calcium-activated potassium channels: coupling of metal ions and voltage sensing

Jianmin Cui

Department of Biomedical Engineering and Cardiac Bioelectricity and Arrhythmia Center, Washington University, St Louis, MO 63130, USA

Ion channels and lipid phosphatases adopt a transmembrane voltage sensor domain (VSD) that moves in response to physiological variations of the membrane potential to control their activities. However, the VSD movements and coupling to the channel or phosphatase activities may differ depending on various interactions between the VSD and its host molecules. BK-type voltage,  $\text{Ca}^{2+}$  and  $\text{Mg}^{2+}$  activated  $\text{K}^+$  channels contain the VSD and a large cytosolic domain (CTD) that binds  $\text{Ca}^{2+}$  and  $\text{Mg}^{2+}$ . VSD movements are coupled to BK channel opening with a unique allosteric mechanism and are modulated by  $\text{Ca}^{2+}$  and  $\text{Mg}^{2+}$  binding via the interactions among the channel pore, VSD and CTD. These properties are energetically advantageous for the pore to be controlled by multiple stimuli, revealing the adaptability of the VSD to its host molecules and showing the potential for intracellular signals to affect the VSD in order to modulate the function of its host molecules.

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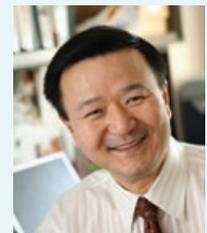
Corresponding author Email: jcui@biomed.wustl.edu

## Introduction

BK channels are large conductance voltage and  $\text{Ca}^{2+}$  activated  $\text{K}^+$  channels encoded by the *Slo1* gene (Atkinson *et al.* 1991). Similar to other  $\text{K}^+$  channels, BK channels contain a central ionic pore comprising transmembrane  $\alpha$  helices S5 and S6 from four Slo1 subunits and the selectivity filter for  $\text{K}^+$  permeation (Atkinson *et al.* 1991; Adelman *et al.* 1992; Butler *et al.* 1993; Shen *et al.* 1994; Tseng-Crank *et al.* 1994; Doyle *et al.* 1998). The opening of BK channels repolarizes the membrane potential and reduces  $\text{Ca}^{2+}$  entry into the cell by closing voltage-dependent  $\text{Ca}^{2+}$  channels, which regulate various physiological processes including neurotransmitter release in synapses (Robitaille & Charlton, 1992; Raffaelli *et al.* 2004; Wang, 2008), contraction of smooth muscle cells in airway and blood vessels (Brayden & Nelson, 1992; Kotlikoff, 1993; Brenner *et al.* 2000), circadian pacemaker output in central nerve systems (Meredith *et al.* 2006; Pitts *et al.* 2006; Kent & Meridith, 2008) and electric tuning by hair cells in vertebrates (Art & Fettiplace, 1987; Fuchs *et al.* 1988; Hudspeth & Lewis, 1988). Each Slo1 subunit has a voltage sensor domain (VSD) formed by transmembrane segments S1–S4 and a large cytosolic domain (CTD) containing  $\text{Ca}^{2+}$  binding sites (Fig. 1). Voltage sensor movements in response to depolarization of the membrane potential and the binding of intracellular  $\text{Ca}^{2+}$  are separately coupled to the pore to activate the channel.

Similar molecular mechanisms may underlie the coupling between voltage sensor movements and pore opening in voltage gated  $\text{K}^+$  ( $\text{K}_V$ ) channels and BK channels (Cui *et al.* 2009). However, while in some  $\text{K}_V$  channels, such as Shaker, the coupling between the voltage sensor and pore seems to be tight or even obligatory, i.e. the pore opens when and only when all the voltage sensors are activated, the coupling in BK channels is allosteric, i.e. the pore can open when the voltage sensor is either activated or at rest but the opening is favoured when the voltage sensor is activated. In addition, in BK channels the CTD is located close to the membrane-spanning voltage

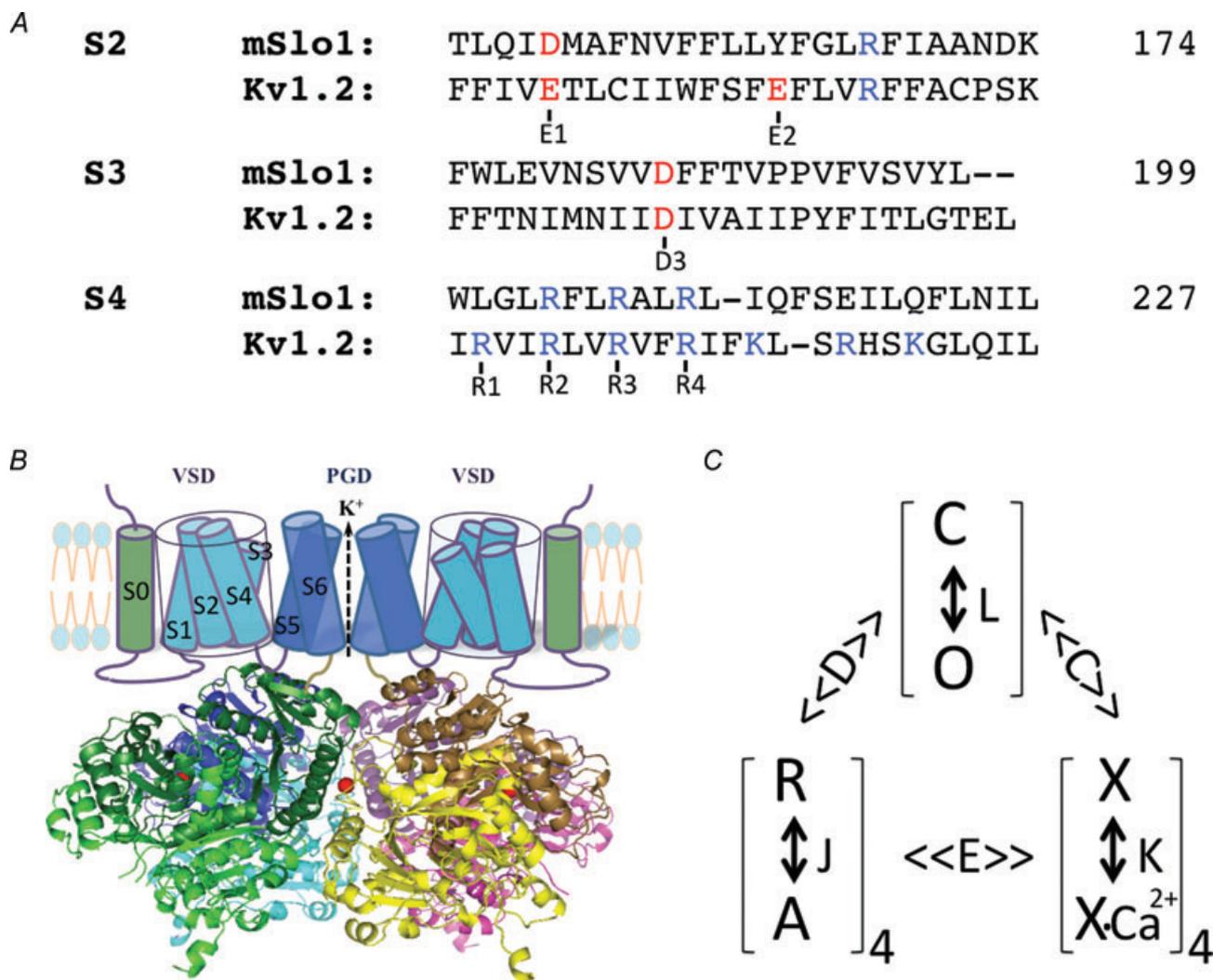
**Jianmin Cui** is the Spencer T. Olin Professor of Biomedical Engineering at Washington University in St Louis. He received a PhD in Physiology and Biophysics from State University of New York at Stony Brook and a post-doctoral training at Stanford University. He was an assistant professor of Biomedical Engineering at Case Western Reserve University before moving to St Louis. His research interests are on membrane permeation to ions, drugs and genes, including the molecular mechanisms of ion channel function and ultrasound-mediated drug/gene delivery. He is a recipient of the Established Investigator Award from the American Heart Association.



sensor and pore domains. This structural feature allows intracellular  $Mg^{2+}$  to bind closely to the cytosolic side of the voltage sensor and directly affect voltage sensor movements via an electrostatic interaction (Yang *et al.* 2007). This article will review the special properties of voltage sensing and the influence by  $Ca^{2+}$  and  $Mg^{2+}$  ions in BK channels.

### Allosteric mechanisms of voltage- and $Ca^{2+}$ -dependent activation of BK channels

The VSDs of BK channels contain charged amino acid residues that are highly conserved among  $K_V$  channels (Fig. 1A). The arginine residues at positions 207, 210 and 213 in the S4 segment (all residue numbers are based



**Figure 1. Voltage- and  $Ca^{2+}$ -dependent activation of BK channels**

A, sequence alignment of S2–S4 in the voltage sensor domain of mSlo1 and  $K_V1.2$  channels, highlighting the conserved charged residues (red and blue) that are important in voltage-dependent activation of both BK and  $K_V$  channels. The position of the last amino acid in each segment of mSlo1 is given (numbers). B, structure model of BK channels. The cartoon of the membrane-spanning domains only shows two subunits for clarity. The cytosolic domain is the tetrameric Slo1 structure (Yuan *et al.* 2010). The subunits are represented by different colours and the dark and light shades of each colour show the RCK1 and RCK2 structural domains within the same subunit (RCK: regulator for conductance of  $K^+$ ). A bound  $Ca^{2+}$  ion to the  $Ca^{2+}$  bowl in each subunit is shown as a red sphere. C, model of allosteric gating of BK channels by voltage and  $Ca^{2+}$ . The channel can open and close (O and C) with an equilibrium constant L, which is modulated by the activation of four voltage sensors (R and A, with an equilibrium constant J) and the binding of four  $Ca^{2+}$  ions (a simplification from actual binding of eight  $Ca^{2+}$  ions, X and  $X-Ca^{2+}$ , with a dissociation constant K) with the allosteric factor D and C, respectively. The voltage sensor activation and  $Ca^{2+}$  binding also affect each other with a factor E. The model was proposed by Horrigan & Aldrich (2002).

on the mbr5 sequence of mouse Slo1 (mSlo1) subunit, GenBank accession number, GI: 347143) (Butler *et al.* 1993) correspond to R2, R3 and R4 in  $K_V$  channels (Gandhi & Isacoff, 2002), Asp153 in S2 corresponds to E1/D1 in  $K_V$  channels (Silverman *et al.* 2003; Wu *et al.* 2010), and Asp186 in S3 corresponds to D3 in  $K_V$  channels (Fig. 1A). The S2 segment does not contain the conserved E2 residue of  $K_V$  channels at the cytosolic side, but instead a positively charged Arg167 that is located four residues below the E2 position. Although the mutations of each of the three arginine residues in S4 alter voltage dependence of BK channel opening (Diaz *et al.* 1998; Cui & Aldrich, 2000), it was shown that only Arg213 contributes to voltage sensing (Ma *et al.* 2006). In addition, the charged residues Asp153 and Arg167 in S2 and Asp186 in S3 also contribute to voltage sensing. Neutralization of each of these residues in the four mSlo1 subunits reduced effective gating charge by 1.20, 0.92, 0.48 and 0.88, respectively, from 2.32 of the WT channel (Ma *et al.* 2006). The role of these charges is somewhat different from that of the equivalent charged residues in  $K_V$  channels, where the arginine residues in S4 serve as the primary gating charges, while the negatively charged residues in S2 and S3 interact with S4 arginines to stabilize channel protein (Papazian *et al.* 1995; Tiwari-Woodruff *et al.* 1997; Long *et al.* 2005) and steer S4 movements during activation (Wu *et al.* 2010); only E2 in the S2 segment of the Shaker  $K^+$  channel had been shown to serve as a gating charge (Seoh *et al.* 1996). Since in BK channels the charged residues in S2, S3 and S4 contribute similarly to voltage sensing, the movements of the VSD in BK and  $K_V$  channels may differ during activation. A model proposes that in BK channels the VSD undergoes a global conformation change that results in a repacking of all membrane-spanning segments in the VSD during activation, as opposed to a large movement of S4 alone in  $K_V$  channels (Ma *et al.* 2006; Pantazis *et al.* 2010).

In BK channels, each Slo1 subunit contains an additional transmembrane segment S0 at the N-terminus of the VSD (Wallner *et al.* 1996) (Fig. 1B). In the BK channel structure, S0 is located at the periphery of VSD close to S2 (Liu *et al.* 2008; Wang & Sigworth, 2009). S0 is important for modulation of BK channels by auxiliary  $\beta$  subunits; it is also essential for the function of the Slo1 protein since the truncated Slo1 lacking S0 could not express any currents but the coexpression of the truncated Slo1 with a separate S0 segment restored channel function (Wallner *et al.* 1996). Mutations of S0 affect voltage-dependent activation of BK channels (Koval *et al.* 2007). Nevertheless, it is not clear how S0 contributes to the movements of the VSD during voltage dependent activation.

Voltage sensor activation in BK channels facilitates channel opening. In the absence of intracellular  $Ca^{2+}$ , the channel can be activated by voltage; the macroscopic conductance and activation rate increase

with depolarization (Meera *et al.* 1996; Cui *et al.* 1997). Likewise, the open probability of single BK channels increases with voltage (Nimigeon & Magleby, 2000; Rothberg & Magleby, 2000; Talukder & Aldrich, 2000). Voltage sensor movements can be measured by gating currents (Stefani *et al.* 1997; Horrigan & Aldrich, 1999) and fluorescence changes from the fluorophore labelling the S3–S4 linker (Savalli *et al.* 2006) prior to channel opening. Mutations of charged amino acid residues in the VSD reduce the steepness of the voltage dependence of channel opening (Diaz *et al.* 1998; Cui & Aldrich, 2000; Ma *et al.* 2006). Finally, intracellular  $Mg^{2+}$  (Yang *et al.* 2007) and the auxiliary  $\beta$  subunits of BK channels (Bao & Cox, 2005; Orío & Latorre, 2005; Savalli *et al.* 2007; Yang *et al.* 2008b) modulate channel opening by altering voltage sensor function.

While it is clear that BK channels are activated by voltage, the channel can open even when the voltage sensors are not activated; conversely, the channel may remain closed when the voltage sensors are activated. For an ion channel that opens only after the voltage sensors are activated, the probability of channel opening ( $P_o$ ) depends on voltage with an exponential function at negative voltages, where the voltage sensors are at the resting state and the probability of their traverse to the activated state is low (Almers, 1978; Sigworth, 1994; Sigg & Bezanilla, 1997). For instance, the Shaker  $K^+$  channel shows such a  $P_o$ -voltage relation at negative voltages where  $P_o$  is as low as  $10^{-7}$ , suggesting the tight or even obligatory relation between voltage sensor activation and channel opening (Islas & Sigworth, 1999). However, for BK channels, the  $P_o$ - $V$  relation becomes flat as voltage decreases below 0 mV, deviating from the exponential relationship at  $P_o \approx 10^{-5}$  and is much shallower at  $P_o \leq 10^{-6}$  (Horrigan *et al.* 1999; Cui & Aldrich, 2000; Yang *et al.* 2010). This result indicates that BK channels can open spontaneously and independently of voltage sensor movements at negative voltages. Another piece of evidence for the non-obligatory relation between voltage sensor movements and channel opening is that the On-gating current elicited by a depolarizing voltage pulse from a negative voltage has a much faster time course than channel opening. The ionic current does not turn on until the On-gating current completes relaxation to 0 (Fig. 2B) (Horrigan & Aldrich, 1999), suggesting that the gating charges have moved to the activated state before the channels open. Thus, the voltage sensors can activate while the channel is closed and the activation of the voltage sensors facilitates channel opening. Based on these results and the voltage dependence of ionic and gating currents, an allosteric model was proposed for voltage-dependent activation of BK channels (Horrigan *et al.* 1999). In this model the voltage sensors can activate when the channel is either at the closed or open state but in open channels the activation of the voltage sensors is more favoured;

reciprocally, the activation of the voltage sensors promotes channel opening (Fig. 1C).

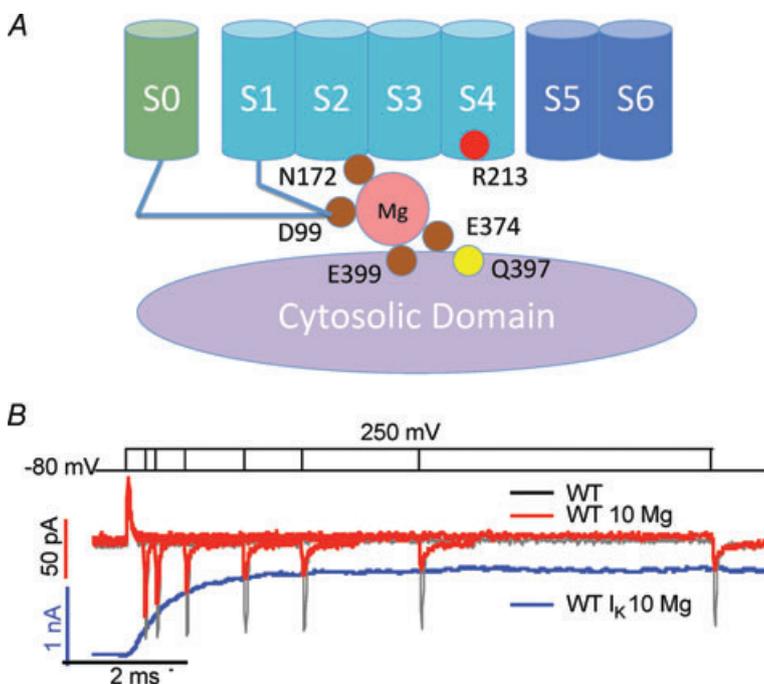
The allosteric coupling between voltage sensor movements and channel opening in BK channels is particularly suitable for  $\text{Ca}^{2+}$ -dependent activation. At voltages below  $-150$  mV, where the voltage sensors are at the resting state, an increase of intracellular  $\text{Ca}^{2+}$  concentration enhances  $P_o$  by  $\sim 10^4$  (Horrigan & Aldrich, 2002; Sweet & Cox, 2008; Yang *et al.* 2010). Therefore,  $\text{Ca}^{2+}$  binding can open the channel without voltage sensor activation, which is energetically advantageous for  $\text{Ca}^{2+}$  to activate the channel, especially at negative voltages.

Two  $\text{Ca}^{2+}$  binding sites have been identified in the CTD of the Slo1 subunit by functional studies and mutagenesis. One site is located in the  $\text{Ca}^{2+}$  bowl that contains consecutive Asp residues at positions 895–901 (Moss *et al.* 1996a,b; Schreiber & Salkoff, 1997; Schreiber *et al.* 1999; Bian *et al.* 2001; Braun & Sy, 2001; Bao *et al.* 2004), and the other resides at Asp367 (Xia *et al.* 2002). Recently, an X-ray crystallographic structure of the CTD of Slo1 shows a  $\text{Ca}^{2+}$  ion binding to the  $\text{Ca}^{2+}$  bowl that is coordinated by the side chains from Asp898 and Asp900 and main chain carbonyl from Gln892 and Asp895 (Yuan *et al.* 2010), but the second  $\text{Ca}^{2+}$  binding site was not identified in the structure.  $\text{Ca}^{2+}$ -dependent activation of BK channels can also be described by an allosteric model (McManus & Magleby, 1991; Cox *et al.* 1997), in which  $\text{Ca}^{2+}$  can bind to the channel in both the closed and open states but with a higher affinity when the channel is open, and therefore  $\text{Ca}^{2+}$  binding shifts the equilibrium between the closed and open states toward open (Fig. 1C). Because both voltage and  $\text{Ca}^{2+}$  promote BK channel

opening and because both voltage sensor activation and  $\text{Ca}^{2+}$  binding are favoured when the channel is open,  $\text{Ca}^{2+}$  binding shifts voltage dependence of channel opening toward less positive voltages, while depolarization enhances the apparent affinity and cooperativity of  $\text{Ca}^{2+}$  binding (Cox *et al.* 1997; Cui *et al.* 1997). Thus, voltage sensor movements are modulated by  $\text{Ca}^{2+}$ , and vice versa, through an allosteric connection via channel opening. In addition,  $\text{Ca}^{2+}$ -dependent activation and voltage sensor movements also affect each other more directly (Horrigan & Aldrich, 2002; Sweet & Cox, 2008), possibly via the interaction between the VSD and CTD (see below).

### $\text{Mg}^{2+}$ activates BK channels through an electrostatic interaction with the VSD

In addition to the sites at the  $\text{Ca}^{2+}$  bowl and Asp367 that bind  $\text{Ca}^{2+}$  with affinities in the micromolar range, BK channels contain another site that binds both  $\text{Ca}^{2+}$  and  $\text{Mg}^{2+}$  with affinities in the millimolar range (Golowasch *et al.* 1986; Oberhauser *et al.* 1988; Shi & Cui, 2001; Zhang *et al.* 2001). Because this site binds metal ions at the physiological concentration of  $\text{Mg}^{2+}$ , it is known as the  $\text{Mg}^{2+}$  binding site. This site is unique in that  $\text{Mg}^{2+}$  is coordinated by residues from both the CTD (Glu374 and Glu399) (Shi *et al.* 2002; Xia *et al.* 2002) and the VSD (Asp99 and Asn172) (Yang *et al.* 2008a) (Fig. 2A). It shows that the CTD in the BK channel is spatially very close to the VSD. Consistent with this result, it was shown that two Cys mutations that substitute for Asp99 in the VSD and Gln397 in the CTD can form a disulfide bond (Yang *et al.* 2008a) that is 2.9–4.6 Å in distance  $\text{C}\beta\text{--C}\beta$



**Figure 2.  $\text{Mg}^{2+}$  modulation of the voltage sensor function in BK channels**

A, cartoon showing amino acid residues involved in  $\text{Mg}^{2+}$  binding and interaction with the voltage sensor domain. B, gating currents in the absence and presence of 10 mM  $\text{Mg}^{2+}$  (black and red traces) in response to various lengths of voltage pulses from  $-80$  mV to  $+250$  mV (top). The ionic current (blue trace) in response to a  $-80$  to  $+250$  mV voltage pulse in 10 mM  $\text{Mg}^{2+}$  is aligned with gating currents to show the correlation in time courses between channel opening and the decay in Off-gating current amplitude in 10 mM  $\text{Mg}^{2+}$ . Adapted from Yang *et al.* (2007), ©2007 National Academy of Sciences of the USA.

of the two Cys residues (Hazes & Dijkstra, 1988). The close association between the VSD and CTD was also demonstrated in the electron cryomicroscopy structure of BK channels (Wang & Sigworth, 2009). Recently, it was shown that X-ray crystallographic structure of the Slo1 CTD fits well with the cytosolic face of the  $K_V$  channel VSD (Yuan *et al.* 2010).

The  $Mg^{2+}$  ion bound to the BK channel is positioned close to the VSD, allowing it to affect the VSD through an electrostatic interaction. Evidence for the involvement of the VSD in  $Mg^{2+}$ -dependent activation of BK channels was first shown in a study in which the mutation of Arg213 to Gln (R213Q) abolished  $Mg^{2+}$  sensitivity of BK channel activation (Hu *et al.* 2003). Since Arg213 is the only residue in S4 contributing to gating charge (Ma *et al.* 2006), this result suggested that  $Mg^{2+}$  might alter voltage sensor movements by an electrostatic interaction. The evidence supporting this idea was shown in a number of experiments (Yang *et al.* 2007). First,  $Mg^{2+}$ -dependent activation depends on the ionic strength of the intracellular solution, and the effect of  $Mg^{2+}$  diminishes with increasing ionic strength, consistent with the idea that  $Mg^{2+}$  activates the channel with an electrostatic interaction. Second, a positive charge at residue 213 is necessary and sufficient for  $Mg^{2+}$ -dependent activation. Neutralization of the charge of Arg213 with mutations R213Q or R213C abolishes  $Mg^{2+}$ -dependent activation, while a covalent modification of R213C by a positively charged [2-(trimethylammonium)ethyl]methane thio-sulfonate (MTSET<sup>+</sup>) restores  $Mg^{2+}$  sensitivity. This result suggests that Arg213 is the target of the electrostatic interaction with  $Mg^{2+}$ . The third experiment supporting the electrostatic interaction is that positive charges added to Gln397 by either mutation or chemical modification can mimic the effect of  $Mg^{2+}$  to activate the channel. Gln397 is located close to the  $Mg^{2+}$  binding site, only two residues away from one of the  $Mg^{2+}$  coordinator Glu399 (Shi *et al.* 2002; Xia *et al.* 2002), and mutations of Gln397 to charged residues alter  $Mg^{2+}$  binding by electrostatic interactions (Yang *et al.* 2007). These results indicate that the electric field of the charges at 397 overlaps with that of the bound  $Mg^{2+}$  ion. Therefore, both charges can interact with Arg213 through the electric fields.

The electrostatic repulsion between  $Mg^{2+}$  and Arg213 in S4 slows down the return of the voltage sensor from the activated state to the resting state. This is supported by measuring the effect of  $Mg^{2+}$  on gating currents of BK channels (Yang *et al.* 2007). Interestingly, the effect of  $Mg^{2+}$  on gating currents depends on whether the channel is in the closed or open state. In BK channels, the voltage sensor is allosterically coupled to the activation gate and can activate in both the closed and open state (Horrigan & Aldrich, 1999). The On-gating current in response to a depolarization pulse decays rapidly before the onset of the ionic current (Fig. 2B), representing voltage sensor

activation in the closed state. It was found that  $Mg^{2+}$  has little effect on the On-gating current but prolongs and reduces the amplitude of the Off-gating current upon membrane repolarization (Yang *et al.* 2007; Horrigan & Ma, 2008). The changes in the Off-gating current by  $Mg^{2+}$  are increasingly apparent with longer depolarization pulses and the time course matches with the onset of the ionic current elicited by the same voltage (Fig. 2B), indicating that the returning of the voltage sensor to the resting state as measured by the Off-gating current is slowed down by  $Mg^{2+}$  more when the channel is open than when the channel is closed. These results suggest that channel opening enhances the interaction between  $Mg^{2+}$  and S4, possibly by a conformational change that moves the bound  $Mg^{2+}$  closer to S4.

### Summary

BK channels are activated by voltage and intracellular  $Ca^{2+}$  and  $Mg^{2+}$ . The BK channel contains the transmembrane VSD as well as  $Ca^{2+}$  and  $Mg^{2+}$  binding sites in a large CTD. The voltage and metal sensors all control the opening of the same ionic pore in response to various physiological signals. While the architecture of the VSD in BK channels is similar to that of other voltage-dependent channels, the movements of the VSD during channel activation and the coupling between the VSD and the activation gate in BK channels are unique, and as such the charged residues in S2, S3 and S4 make similar contributions to voltage sensing and the VSD activation promotes channel opening through an allosteric mechanism. These properties are well suited to the function of BK channels to sense multiple stimuli and may be derived from the interactions of the VSD not only with the pore but also with the unique S0 segment and the CTD in BK channels. These interactions among different structural domains also mediate the modulation of VSD movements by  $Ca^{2+}$  via allosteric mechanisms and by  $Mg^{2+}$  via electrostatic interactions.

The movements of the VSD in response to depolarization can either open or close ion channels (Perozo *et al.* 1992; Mannikko *et al.* 2002) and can activate lipid phosphatases (Murata *et al.* 2005; Hossain *et al.* 2008), indicating that the VSD has the ability to adapt to various environments in its host protein molecules and functional requirements. The VSD may also be modulated by intracellular factors such as ligand binding, post-translational modification, and the interactions with cytosolic domains, accessory proteins or subunits to alter the function of voltage-dependent ion channels (Perozo & Bezanilla, 1990; Jones *et al.* 1997; Terlau *et al.* 1997; Bao & Cox, 2005). The study of voltage-dependent activation of BK channels may provide insights into the principles of the VSD versatility and modulation that affect the function of VSD containing molecules.

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