Molecular mechanism for depolarization-induced modulation of Kv channel closure

Alain J. Labro,1,2 Jerome J. Lacroix,1 Carlos A. Villalba-Galea,3 Dirk J. Snyders,2 and Francisco Bezanilla1

1Department of Biochemistry and Molecular Biology, University of Chicago, Chicago, IL 60637
2Department of Biomedical Sciences, University of Antwerp, Antwerp 2610, Belgium
3Department of Physiology and Biophysics, Virginia Commonwealth University School of Medicine, Richmond, VA 23298

Voltage-dependent potassium (Kv) channels provide the repolarizing power that shapes the action potential duration and helps control the firing frequency of neurons. The K⁺ permeation through the channel pore is controlled by an intracellularly located bundle-crossing (BC) gate that communicates with the voltage-sensing domains (VSDs). During prolonged membrane depolarizations, most Kv channels display C-type inactivation that halts K⁺ conduction through constriction of the K⁺ selectivity filter. Besides triggering C-type inactivation, we show that in Shaker and Kv1.2 channels (expressed in Xenopus laevis oocytes), prolonged membrane depolarizations also slow down the kinetics of VSD deactivation and BC gate closure during the subsequent membrane repolarization. Measurements of deactivating gating currents (reporting VSD movement) and ionic currents (BC gate status) showed that the kinetics of both slowed down in two distinct phases with increasing duration of the depolarizing prepulse. The biphasic slowing in VSD deactivation and BC gate closure was strongly correlated in time and magnitude. Simultaneous recordings of ionic currents and fluorescence from a probe tracking VSD movement in Shaker directly demonstrated that both processes were synchronized. Whereas the first slowing originates from a stabilization imposed by BC gate opening, the subsequent slowing reflects the rearrangement of the VSD toward its relaxed state (relaxation). The VSD relaxation was observed in the Ciona intestinalis voltage-sensitive phosphatase and in its isolated VSD. Collectively, our results show that the VSD relaxation is not kinetically related to C-type inactivation and is an intrinsic property of the VSD. We propose VSD relaxation as a general mechanism for depolarization-induced slowing of BC gate closure that may enable Kv1.2 channels to modulate the firing frequency of neurons based on the depolarization history.

INTRODUCTION

Voltage-gated potassium (Kv) channels form transmembrane potassium-selective pathways that actively open and close in response to changes in membrane potential. In vivo, these channels are (commonly) responsible for repolarizing the membrane potential back to its resting conditions after an action potential (Hodgkin and Huxley, 1952). The kinetics and abundance of these channels shape the action potential and constitute a critical determining factor of cellular excitability. A typical Kv channel is composed of four individual α subunits (MacKinnon, 1991), each containing six membrane-spanning helices (S1–S6) whereby S5 and S6 segments organize to form a central ion-conducting pore (Doyle et al., 1998; Long et al., 2005). This K⁺ permeation pathway can be sealed off by two separate gates in series: (1) at the level of the K⁺ selectivity filter that collapses and becomes nonpermeable upon C-type inactivation (Liu et al., 1996; Loots and Isacoff, 1998) and (2) at the inner S6 bundle-crossing (BC) gate, which closes or opens depending on the membrane potential (Liu et al., 1997; del Camino and Yellen, 2001).

Control over the BC gate is achieved by the four voltage-sensing domains (VSDs) formed by the S1 through S4 segments that surround the central pore domain. The S4 segment, which contains positively charged amino acids (gating charges), forms the main voltage-sensing element that switches between resting and active conformations upon changes in the membrane potential. This motion is mechanically transmitted via the S4–S5 linker (S4S5L) to the BC gate of the same subunit (Labro et al., 2005; Long et al., 2005), thereby closing or opening the channel’s pore. The dynamics of the VSD motion can be directly monitored by measuring transient gating currents that are produced by the displacement of the gating charges across the membrane electric field (Bezanilla, 2000).
In addition to its resting and active states, the VSD enters a relaxed state upon prolonged depolarizations (Villalba-Galea et al., 2008). In Shaker K⁺ channels, relaxed VSDs return to their resting state twice as slow as from their active state, thus resulting in an apparent shift of the charge (Q) vs. voltage (V) [QV] curves toward negative voltages (Lacroix et al., 2011). Using in-depth biophysical analysis of wild-type (WT) Shaker and Kv1.2 channels, we report here that prolonged depolarizations also produce a twofold slowing down of BC gate closure with a time course synchronized with VSD relaxation. Upon prolonged depolarizations, K⁺ permeation is in several Kv channels terminated via a constriction of the selectivity filter known as C-type inactivation (Loots and Isacoff, 1998; Cuello et al., 2010). As part of these experiments were done in the KcsA channel that lacks a VSD, these results strongly convey the notion that C-type inactivation does not require the presence of a VSD. In agreement with this notion, we provide here compelling evidence that VSD relaxation occurs independently from C-type inactivation and propose that VSD relaxation constitutes a potent physiological molecular mechanism for the modulation of deactivating ionic currents in Kv channels.

MATERIALS AND METHODS

Expression in Xenopus laevis oocytes

The Shaker cDNA used in this study was the fast inactivation removed (IR, Δψ→0) version described previously (Yellen et al., 1991), and the WT human Kv1.2 and Ciona intestinalis voltage-sensitive phosphatase (Ci-VSP) cDNAs were kindly provided by S. Goldstein (Brandeis University, Boston, MA) and Y. Okamura (Osaka University, Osaka, Japan), respectively. cDNAs were cloned into a pbSTα vector modified for enhanced expression into Xenopus oocytes (Starace and Bezanilla, 2001). The Ci-VSP cDNA contains the C363S mutation that inactivates catalytic activity (Murata et al., 2005). cDNAs were linearized with a unique NotI restriction site (New England Biolabs, Inc.) and transcribed into cRNAs using a T7 RNA expression kit (Ambion; Invitrogen). 5–50 ng cRNA was injected into mature Xenopus oocytes 4–24 h after surgical extraction from adult frogs. Injected oocytes were maintained before recordings in a standard oocyte solution containing 100 mM NaCl, 5 mM KCl, 2 mM CaCl₂, and 10 mM HEPES, pH 7.5, supplemented with 50 µg/ml gentamicin for 2–6 d at 16.5 °C.

Electrophysiology and solutions

Ionic and gating current recordings were taken at room temperature with a high performance oocyte clamp amplifier (CA-1B; Dagan Corporation) in the cut-open voltage-clamp setup. Current recordings were digitized with an in-house–built acquisition system based on the a4d4 module in a digital signal processing unit (SB6711; Innovative Integration) at a sampling frequency ranging from 10 to 100 kHz after 5–20-kHz low-pass filtering. Pulse protocols and data acquisition were controlled using in-house–developed software. Recording pipettes with a resistance of 0.8–1.5 MΩ were pulled from 1.5-mm borosilicate glass capillaries (World Precision Instruments) using a vertical heat filament micropipette puller (P87; Sutter Instrument). For gating current recordings, remaining capacitive transient currents after compensation were subtracted online using the −P/4 method for activation protocols (~120-mV holding potential) and +P/4 for deactivation protocols (20-mV holding potential) when possible. For ionic current recordings, the external solution contained 11.5 mM K-methyldisulfonate (MES), 103.5 mM NMDG-MES, 2 mM Ca-MES, and 10 mM HEPES, and the internal solution contained 115 mM K-MES, 2 mM EGTA, and 10 mM HEPES, both adjusted to pH 7.5 using MES. Gating currents were recorded after depleting the oocytes of K⁺ by repetitive activation of the channels. Any remaining ionic current contamination was eliminated by external TEA block, and the external recording solution contained 115 mM TEA-MES, 2 mM Ca-MES, and 10 mM HEPES adjusted to pH 7.5 using MES. The internal solution contained 115 mM NMDG-MES, 2 mM EGTA, and 10 mM HEPES adjusted to pH 7.5.

Site-directed flurometry

To study VSD movement using site-directed voltage-clamp fluorometry, we chose the M356C Shaker mutant because labeling this residue has been shown to give nice time-dependent changes in fluorescence amplitude (ΔF) that follow VSD movements (Cha and Bezanilla, 1997). Oocytes expressing the Shaker mutant M356C were incubated in 0.2 mM dithiothreitol for 30 min before labeling with 20 µM tetramethylrhodamine-5-maleimide (TMRM; Invitrogen) for 20 min on ice. Labeled cells were individually mounted on an epifluorescence cut-open oocyte setup and illuminated with a tungsten halogen lamp as previously described (Cha and Bezanilla, 1997). TMRM fluorescence was measured using an excitation filter (535DF55), a 570-nm dichroic mirror, and an emission filter (595AF60; Omega Filters).

Data analysis

Details of pulse protocols used to elicit gating or ionic currents were adjusted to determine the biophysical properties of each construct adequately and are shown in the figures or described in legends. The obtained charge versus voltage (QV) and conductance versus voltage (Gv) curves were fitted with the following Boltzmann equation: \( y = 1/[1 + \exp(-(V - V_{1/2})/k)] \), where \( V \) represents the applied voltage, \( V_{1/2} \) is the midpoint potential at which 50% of the total charge has moved or half of the channels have opened, and \( k \) is the slope factor. The time constants of the activating and deactivating gating currents (I_act and I_deac) were determined by fitting the decaying part of the current traces with a single or double exponential function. The kinetics of the ionic currents were determined by fitting the rise in current activation (I_rise) or decay during deactivation (I_decay) with a single or double exponential function, respectively. When a double exponential fit was used, a weighted mean time constant was calculated based on the contribution (amplitude) of each component.

Signals for simultaneous ionic currents and fluorescence recordings were not subtracted and analogically filtered at 5–10 kHz. Fluorescence traces were further digitally filtered at 1–5 kHz. The procedure for fitting the fluorescence changes induced by the pulses was similar to that described in the previous paragraph. All results are expressed as means ± SEM, with \( n \) being the number of oocytes analyzed.

RESULTS

Strong correlation between VSD movement and BC gate opening/closure in Shaker channels

To investigate the coupling between the VSD and the BC gate during channel opening and closure, we first recorded ionic currents (reporting on the BC gate status) in Shaker channels expressed in Xenopus oocytes. Next, gating currents (reporting VSD movement) were...
of $-47.5 \pm 1.1$ mV and a slope of $8.0 \pm 0.9$ mV ($n = 6$; Fig. 1 C). To obtain the voltage dependence of VSD deactivation, gating currents were generated by pulsing from a constant depolarized voltage to more negative voltages (deactivation protocol; Fig. 1 D). Integration of the $I_{\text{Off}}$ gating currents produced during the repolarizing test pulses resulted in a $QV$ curve with a $V_{1/2}$ of $-63.7 \pm 1.2$ mV and a slope of $9.9 \pm 1.0$ mV ($n = 5$; Fig. 1 C). A similar approximately $-15$-mV difference in the $V_{1/2}$ values of the $QV$ curves between the voltage dependence of activation and deactivation has been ascribed to either an intrinsic difference of voltage sensitivity of the VSD between its resting or active position or to an undetected slow gating component in $I_{\text{Off}}$ (Lacroix et al., 2011).

Figure 1. Ionic and gating currents from Shaker-H4-IR; behavior of the VSD and channel gate. (A) Activating ionic currents ($I_{\text{ac}}$) recorded from Shaker-H4-IR using the pulse protocol shown on top. (B) Gating current recordings after depletion of K+ and external TEA block. The holding potential was $-130$ mV, and oocytes were depolarized in 5-mV increments from $-120$ to 80 mV. Background leak and capacitive currents were subtracted with a $-P/4$ protocol using a $-120$-mV holding potential. The inset is a scaled-up view of the $I_{\text{Off}}$ currents highlighting the gradual slowing in $I_{\text{Off}}$ decay when prepulse depolarization voltages became stronger (red trace is $I_{\text{Off}}$ for a $-50$-mV pulse). (C) Voltage dependence of both activating and deactivating charge movement ($QV$ curve) and BC gate opening ($GV$ curve). The $GV$ curve (circles) was obtained from the ionic tail currents during the repolarization step of pulse protocols shown in A. The voltage dependence of gating charge activation ($QV$ curve displayed with open triangles; $Q_{\text{ac}}$) was obtained by integrating the repolarizing gating currents ($I_{\text{Off}}$) of the activation protocol shown in B. The voltage dependence of gating charge deactivation was obtained by integrating the repolarizing gating currents of the deactivation protocol shown in D ($QV$ curve displayed with closed triangles; $Q_{\text{deac}}$). Both $GV$ and $QV$ values were normalized, and the curves shown are for both $GV$ and $QV$ the average fit to a Boltzmann equation. (D) Superposition of scaled ionic (red traces) and gating currents (black traces) obtained from the same oocyte using the voltage protocol shown on top. Scale bars for gating and ionic currents are shown in black and red, respectively. (E, top) A scaled-up view of the overlapping deactivation ionic ($I_{\text{deac}}$) and gating ($I_{\text{Off}}$) currents from D at $-110$-mV repolarizing voltage. (bottom) The respective $I_{\text{deac}}$ and $I_{\text{Off}}$ currents were normalized and superimposed. On the left, the $I_{\text{Off}}$ is inverted to highlight the overlap of the fast component in $I_{\text{deac}}$ with the rising phase observed in $I_{\text{Off}}$. On the right, a scaled-up view of the slow component in $I_{\text{deac}}$ that matches the $I_{\text{Off}}$ decay. (F) Voltage dependency of the time constants ± SEM of $I_{\text{Off}}$ decay (open triangles; $n = 8$), $I_{\text{Off}}$ decay (closed triangles; $n = 7$), the rising phase of $I_{\text{Off}}$ (red circles; $n = 7$), $I_{\text{deac}}$ (gray circles; $n = 7$), the fast component of $I_{\text{deac}}$ (yellow inverted triangles; $n = 7$), the slow component of $I_{\text{deac}}$ (green inverted triangles; $n = 7$), and the weighted $I_{\text{deac}}$ kinetics (blue circles; $n = 7$). Note the superposition of the slow $I_{\text{deac}}$ component with $I_{\text{Off}}$ decay and the fast $I_{\text{deac}}$ component with the rising phase in $I_{\text{Off}}$. Error bars represent SEM.
The time constant of the IgON decays matched the time constant of ionic current activation Iac, that directly reflects the opening of the BC gate (Fig. 1, A, B, and F). In accordance with a previous study, the ionic current deactivation Ideac was analyzed with a double exponential function yielding a fast and a slow component (Stefani et al., 1994). To correlate IgOFF and Ideac correctly, the kinetics of both were analyzed pairwise in the same oocyte (Fig. 1, D and E). It is well established that the rate of decay of IgOFF decreases as the amplitude of the depolarizing prepulse increases and exceeds the threshold for opening the BC gate (about −50 mV for Shaker; Fig. 1 B; Hoshi et al., 1990; Bezanilla et al., 1994). Concomitantly with this slowing, IgOFF displays an unambiguous rising phase that is presumably caused by the stabilization of the conducting conformation of the channels (Batulan et al., 2010). The direct pairing analysis shows that the fast component of the Ideac kinetics matched the time course of the rising phase in IgOFF (Fig. 1, E and F). This rising phase reflects an initial rate-limiting energy barrier for the deactivation VSD transition coming from its activated state. As shown previously, the slow component in Ideac followed IgOFF decay (Fig. 1, E and F; Bezanilla et al., 1991). Apparently, the fast Ideac component had the largest contribution in the overall process of channel closure, which is reflected in the weighted average Ideac kinetics (Fig. 1 F).
Synchronized slowing in \( I_{\text{gOFF}} \) and \( I_{\text{Ideac}} \) kinetics in \textit{Shaker} upon prolonged membrane depolarizations

We previously showed that the overall kinetics of deactivating gating currents in \textit{Shaker} channels displays a biphasic slowing down upon prolonged depolarizations (Fig. 2 A; Lacroix et al., 2011). Similarly, our present analysis of \( I_{\text{gOFF}} \) with varying durations of depolarization showed that there were two noticeable slowing phases (Fig. 2, C and E). The fast developing slowing process developed with a time constant \( \tau_f \) of 8.5 ± 1.0 ms and has been linked to the stabilizing interaction imposed by pore opening (Kanevsky and Aldrich, 1999; Batulan et al., 2010). The second, slower developing slowing period had a time constant \( \tau_s \) of 1,100 ± 100 ms, which corresponds to the VSD relaxation (Lacroix et al., 2011).

As there seems to be a strong correlation between the gating and ionic current kinetics, the time constant of \( I_{\text{Ideac}} \) is expected to also increase upon prolonged depolarizations. In agreement with the slowing in VSD return, there appeared a slow decaying \( I_{\text{Ideac}} \) component that gradually increased in amplitude when the depolarization prepulse durations were prolonged (Fig. 2 B). Similar to the \( I_{\text{gOFF}} \) behavior, there were two distinct slowing processes in \( I_{\text{Ideac}} \) kinetics (Fig. 2, D and E). The first, a fast developing process, occurred within a time window of 20 ms and most likely reflects the open pore stabilization process that occurs upon channel opening. The speed of channel closure slowed down further upon prolonged depolarizations, reaching a steady-state maximal value with prepulse durations of 5 s or longer. Plotting the \( I_{\text{Ideac}} \) kinetics at −50 mV as a function of depolarization time showed that the two slowing processes, with a \( \tau_f \) of 4.8 ± 0.6 ms and a \( \tau_s \) of 670 ± 90 ms (\( n = 5 \)), were similar to the slowing processes in \( I_{\text{gOFF}} \). Moreover, upon the first and second slowing process, both \( I_{\text{gOFF}} \) and \( I_{\text{Ideac}} \) kinetics increased by a factor of four and a factor of two, respectively (Fig. 2 E). Thus, as both the time development and the magnitude of the slowing process in \( I_{\text{gOFF}} \) and \( I_{\text{Ideac}} \) were similar, both processes appeared to be well correlated.

Because differences in recording conditions for ionic and gating current measurements might have affected the observed slowing processes in \( I_{\text{gOFF}} \) and \( I_{\text{Ideac}} \) kinetics, we monitored VSD and BC gate movements simultaneously using site-directed fluorimetry (Cha and Bezanilla, 1997). To achieve this, we covalently attached the conformational reporter TMRM to a cysteine inserted at the top of S4 (M356) in \textit{Shaker} channels. This allowed us to simultaneously record the changes in fluorescence (\( \Delta F \)) that report on the VSD movements and the amplitude and kinetics of the ionic currents reflecting opening/closure of the BC gate. Fig. 3 A shows examples of \( I_{\text{Ideac}} \) and recovery in \( \Delta F \) traces during repolarizing pulses to −40 mV after prepulses of variable durations to 20 mV. The time constants of both \( I_{\text{Ideac}} \) and \( \Delta F \) were calculated and plotted as a function of the depolarization duration, and the results show an identical biphasic slowing for both \( I_{\text{Ideac}} \) and \( \Delta F \) (Fig. 3, B and C). The first slowing developed with a \( \tau_f \) of 35 ± 16 ms for \( I_{\text{Ideac}} \) and 24 ± 15 ms for \( \Delta F \), and the second one had a \( \tau_s \) of 1,450 ± 280 ms for \( I_{\text{Ideac}} \) and 1,670 ± 250 ms for \( \Delta F \) (\( n = 5 \); the slowing processes in \( I_{\text{Ideac}} \) are slower here than in WT \textit{Shaker}, presumably because of the presence of the

\[ \Delta F = 2 \mu A \times 0.8\% \]
\[ \Delta F = 2 \mu A \times 0.5\% \]
\[ \Delta F = 2 \mu A \times 0.4\% \]
Figure 4. C-type inactivation properties of Shaker and Kv1.2. (A) Ionic current recordings from oocytes expressing Shaker (black trace) or Kv1.2 channels (gray trace) evoked by depolarizing the membrane potential from a holding of −80 to 20 mV. Upon prolonged depolarization, both channels display C-type inactivation, which is characterized by a reduction in current amplitude. At the end of the pulse, Kv1.2 channels display only 25 ± 3% (n = 4) current reduction, whereas Shaker channels display 52 ± 2% current decrease (n = 4). (B) Time constants of the inactivation process in Shaker (black circles) and Kv1.2 channels (gray circles) obtained by fitting the reduction in current amplitude with a single exponential function. Note that in both channels, the kinetics are voltage independent. Error bars represent SEM.

Figure 5. Ionic and gating currents from Kv1.2: behavior of the VSD and BC gates. (A) Activating ionic currents from WT Kv1.2, a member of the Shaker family that displays a more modest C-type inactivation process. (B) Gating current recordings from Kv1.2 after depletion of K+. Current recordings were obtained using the activation pulse protocol shown on top. The holding potential was −130 mV, and oocytes were depolarized in 5-mV increments from −130 to 70 mV (for clarity only, current traces are shown every 10-mV increment). Similar to Shaker, there was a gradual slowing in I_{OFF} decay when prepulse depolarization voltages became stronger (red trace is I_{OFF} after a −40-mV prepulse). (C) Superposition of scaled deactivation ionic (red traces) and gating currents (black traces) from Kv1.2 obtained from the same oocyte using the voltage protocol shown on top. The scales for gating and ionic currents are shown in black and red, respectively. (D) Voltage dependence of BC gate opening (GV curve) and charge movement (QV curves) using an activation (Q_{ac}) or deactivation (Q_{deac}) protocol. Both GV and QV values were normalized, and for both GV and QV curves, the average fit to a Boltzmann equation is shown. (E) Mean time constant ± SEM for I_{ON} decay (open triangles), I_{OFF} decay (closed triangles), rising phase of I_{OFF} (red circles), I_{ac} (gray circles), fast component of I_{deac} (yellow inverted triangles), slow component of I_{deac} (green inverted triangles), and the weighted I_{deac} kinetics (blue circles) obtained in a similar way as described for Shaker (Fig. 1), with an n of at least six independent oocytes analyzed. Error bars represent SEM.
Fig. 4 shows that after a 5-s activation pulse to 20 mV, approximately half of the Shaker channels have entered their inactivated state, whereas only a quarter of Kv1.2 channels were inactivated. Moreover, the time course of the inactivation of the potassium conductance in these two channels differs with time constants of $\tau \approx 3.5$ s in Shaker and 5.5 s in Kv1.2, respectively. Because of these differences in C-type inactivation, we wondered whether prolonged depolarizations affect the VSD deactivation and BC gate closure of Kv1.2 channels in a similar or distinct manner compared with Shaker.

To test this, we recorded gating and ionic currents from WT conducting Kv1.2 channels (Fig. 5, A–C). The voltage dependence of charge movement during activation resulted in a QV curve with a midpoint potential of $-37.9 \pm 1.0$ mV and slope factor of $5.9 \pm 0.6$ mV ($n = 6$; Fig. 5 D). The GV curve had a midpoint of $-23.3 \pm 2.6$ mV.

Kv1.2 channels display a similar VSD and BC gate behavior with prolonged depolarizations

The aforementioned results indicated that in Shaker channels, the VSD gets stabilized in an upward conformation (relaxed state) by spending longer periods at depolarizing potentials. It is conceivable that a similar mechanism exists in other Kv channels such as Kv1.2. Although Kv1.2 is a mammalian homologue of the bigger Shaker subfamily of Kv channels and thus shares a high sequence homology, both Shaker and Kv1.2 differ in their C-type inactivation process (Stühmer et al., 1989; Hoshi et al., 1990; Ramaswami et al., 1990).

TMRM linked at M356C). This confirms that the VSD and BC gate movements are strongly coupled and that both VSD return and BC gate closure slow down upon VSD relaxation.
with a slope factor of $7.9 \pm 0.5$ mV ($n = 9$), i.e., $\sim 10$ mV displaced to more positive potentials compared with the QV curve. Furthermore, the voltage dependence of gating charge deactivation, determined from integrating $I_{\text{OFF}}$ using a deactivation protocol (Fig. 5 C), resulted in a QV curve with a midpoint of $-68.3 \pm 2.0$ mV and a slope factor of $9.8 \pm 1.0$ mV ($n = 5$), which is $\sim 30$ mV more negative than the QV curve obtained from an activation protocol (Fig. 5 D). Thus, the underlying relation between the GV and QV curves was in $K_v$1.2 quite similar to Shaker. The time constants of $I_{\text{ON}}$, $I_{\text{OFF}}$, $I_{\text{ac}}$, and $I_{\text{deac}}$ were obtained with the same procedure as described previously (see section Strong correlation between VSD movement. . .) for Shaker (Fig. 5 E). Likewise, the $I_{\text{ac}}$ time constants nicely matched the $I_{\text{ON}}$ kinetics, and the slow component of $I_{\text{deac}}$ matched the decay in $I_{\text{OFF}}$ well. In addition, the fast component of $I_{\text{deac}}$ followed the rising phase in $I_{\text{OFF}}$ and, as in Shaker, appeared to contribute most to the overall ionic current deactivation, as shown by the weighted average $I_{\text{deac}}$ kinetics (Fig. 5 E).

Investigating the $I_{\text{OFF}}$ behavior as a function of the depolarizing prepulse duration indicated that there were two noticeable slowing phases (Fig. 6). As for Shaker, the first slowing followed BC gate opening and most likely results from a similar stabilizing $S4S52$–$S6$ interaction (Batulan et al., 2010). More importantly, the second slowing starts to develop for durations that exceed steady-state BC gate opening (Fig. 6 E). Therefore, this second slowing in $I_{\text{OFF}}$ indicated that the VSD of Kv1.2 becomes stabilized in a secondary activated (relaxed) state upon prolonged depolarizations. We determined the time course for the development of this slowing down in the $I_{\text{OFF}}$ kinetics by fitting the $I_{\text{OFF}}$ traces at $-60$ mV with a double exponential function. The fit yielded a $\tau_f$ of $13.5 \pm 1.8$ ms and a $\tau_s$ of $1,120 \pm 210$ mV ($n = 5$; Fig. 6). The analysis of the kinetics of $I_{\text{deac}}$ showed that BC gate closure also slows down and displays two slowing periods with a $\tau_f$ of $16.5 \pm 4.1$ ms and a $\tau_s$ of $1,250 \pm 160$ ms ($n = 7$; Fig. 6). Consequently, the development of both slowing processes in $I_{\text{deac}}$ and $I_{\text{OFF}}$ appeared very well synchronized. Furthermore, during the second slowing process, which produces a 1.5-fold slowing down in both $I_{\text{OFF}}$ and $I_{\text{deac}}$ (Fig. 6), Kv1.2 channels display only $\sim 25\%$ of C-type inactivation.

VSD relaxation is an intrinsic property of the VSD

Negative shifts in the QV curve determined from activation or deactivation gating currents have been ubiquitously reported in voltage-sensitive proteins. It is thus tempting to raise the hypothesis that these QV shifts are apparent as a result of a VSD relaxation process similar to $K_v$ channels. Accordingly, the relaxation would be an intrinsic property of a VSD with a four-transmembrane topology, and it would not be allosterically triggered by another region of the protein. To test this hypothesis, we investigated VSD relaxation in Ci-VSP, which does not contain a pore (Fig. 7). As reported previously, the kinetics of the deactivating sensing current ($I_{\text{OFF}}$; Fig. 7 A) undergoes a relatively rapid slowing down upon strong depolarizing pulses, a phenomenon that has been attributed to the binding of the phosphatase domain to the membrane (Villalba-Galea et al., 2009). However, analyzing the effect of prolonged prepulse durations revealed a second gradual slowing in $I_{\text{OFF}}$ decay time, similar to Shaker and Kv1.2 channels (Fig. 7, B and C). A shift in the QV curve of Ci-VSP sensing currents has been reported previously (Villalba-Galea et al., 2008), and, as we show here, this shift may be mostly apparent as a result of a slowing in the deactivation kinetics of VSD return (relaxation process).
the full-length Ci-VSP. However, the time course of VSD relaxation in the truncated mutant Δ239–576 that lacks the phosphatase domain (red) attached to the VSD (blue). (right) Sensing current traces (I) of the truncation mutant Δ239–576 recorded using the indicated activation protocol. (B) QV curves for the mutant Δ239–576 (red) and WT Ci-VSP (black) obtained by integrating the activating I, from pulse protocols shown in A and Fig. 7 A, respectively. (C) A superposition of I off current traces at −20 mV after 10-ms (black) and 3-s (red) depolarizations to 80 mV is displayed. Note the slowing in I off decay and the reduction in the fast component amplitude of the red trace. The inset shows the normalized tracings to better highlight the slowing in I off decay. (D) Kinetics of activating I on (open triangles) of the mutant Δ239–576 obtained from different activation voltage pulses (similar to the protocol shown in A) and of deactivating I off of the same mutant obtained using 80-mV prepulses of the indicated durations from 3 ms to 3 s (colored symbols) and pulsing to different deactivation voltages. (E) Weighted time constant of I off decay measured at −20 mV as a function of the duration of the conditioning prepulse at 80 mV. The obtained curve was fitted to a single exponential function and yielded a time constant of τ = 76 ± 11 ms (n = 5). (F) For both the full-length Ci-VSP (left) and the truncation mutant Δ239–576 (right), the normalized amount of sensing charges moved during the conditioning prepulse to 80 mV (Q on, left axis) was superimposed on the depolarization-induced slowing in the kinetics of I off decay (τ on, right axis). Fitting the normalized charge displacement as a function of depolarization duration with a single exponential function yielded τ Q on WT = 61.7 ± 0.1 ms (n = 4) for full-length Ci-VSP and τ Q on Δ239–576 = 18.8 ± 0.1 ms (n = 5) for the truncation mutant. Error bars represent SEM.

To investigate the VSD relaxation in the absence of any load attached to the S4 helix, we created the Ci-VSP truncation mutant Δ239–576 by removing the entire phosphatase domain (Fig. 8 A). This mutant displayed a voltage dependence of sensing current activation that was slightly shifted toward more negative potentials compared with full-length Ci-VSP (Fig. 8, A and B). The maximal value of I off for the deletion mutation was obtained upon repolarizing to −20 mV instead of 10 mV for the WT Ci-VSP (Fig. 8, C and D). By comparing the maximal values of I off, we found that they decayed ~10-fold faster in the truncated Δ239–576 mutant compared with the full-length Ci-VSP (I off WT = 13 ± 1 ms vs. I off Δ239–576 = 4 ± 1 ms after 5 ms prepulses and I off WT = 24 ± 3 ms vs. I off Δ239–576 = 4 ± 1 ms after 100 ms prepulses; compare Fig. 7 B with Fig. 8 D). Despite this acceleration, the decay kinetics of I off slowed down with increased prepulse depolarization time similar to the full-length Ci-VSP. However, the time course of VSD relaxation in the truncated mutant appeared to be accelerated compared with the full-length version and developed with a time constant of 76 ± 11 ms (n = 5) at 80-mV depolarization. This is about sevenfold faster than the full-length version whereby relaxation developed with a time constant of 530 ± 80 ms (Fig. 7 C).

To confirm that the slowing in I off occurred subsequently to the charge movement during activation, we determined for both the full-length Ci-VSP and the Δ239–576 mutant the time course of the sensing charge movement during the depolarizing prepulse by integrating I on. The truncation of the phosphatase domain appeared to accelerate threefold the sensing charge displacement at 80 mV (τ Q on WT = 61.7 ± 0.1 ms compared with τ Q on Δ239–576 = 18.8 ± 0.1 ms), yet there was (as in the full-length Ci-VSP) a clear temporal delay between the displacement of the sensing charge and the slowing
shown that the fast developing slowing in IgOFF kinetics (Fig. 2 and 6). It has been established immediately after channel opening. A recent study showed that this fast developing slowing in WT Ci-VSP, Shaker, and Kv1.2. Thus, the truncation mutant displayed the characteristic relaxation that developed subsequently to sensing charge displacement. This indicates that the isolated VSD from Ci-VSP is able to undergo relaxation in the absence of any load at the C-terminal end of the S4 segment.

**DISCUSSION**

Direct pairing of VSD movement with BC gate opening/closure in Shaker and Kv1.2 channels showed that in both channels, the time course of ionic current activation (Iac) followed that of Ion decay. This indicates that BC gate opening arises late in the activation sequence and is linked to one of the final transition steps of the VSD. This is in agreement with previous proposed gating schemes for the Shaker channel (Bezanilla et al., 1994; Zagotta et al., 1994; Schoppa and Sigworth, 1998) in which BC gate opening requires that all four VSDs have moved to the preactivated/activated state. Iac displayed two components, and, in agreement with a previous study, the late one followed IgOFF decay (Bezanilla et al., 1991). The fast component in Iac, as we showed here, followed the rising phase in IgOFF, indicating that it is linked to a first rate-limiting transition in the sequential steps of VSD return (Fig. 1).

The slowing in the deactivation kinetics of the gating (IgOFF) and ionic (Iac) currents in both Shaker and Kv1.2 displayed two identifiable phases, a fast developing one and a slower one (Figs. 2 and 6). It has been shown that the fast developing slowing in IgOFF kinetics is linked to channel opening (Kanovsky and Aldrich, 1999; Batulan et al., 2010). Remarkably, however, there was a similar slowing in BC gate closure (Iac) that developed within the same time window. This indicates that the slowing in VSD return or BC gate closure. This is in agreement with our observation of a time lag between BC gate opening and the establishment of the stabilizing S4S5L–S6c interactions that results in a slowing of both IgOFF and Iac kinetics.

After this initial slowing, there appeared a subsequent second slowing down in VSD return and BC gate closure when the membrane depolarization was prolonged beyond the duration of the charge movement. Similar to the first process, this second slowing phase in IgOFF and Iac kinetics was synchronized, as further indicated by our site-directed fluorimetry data from the Shaker-M356C mutant (Fig. 3). Because the first stabilization is caused by an S4S5L–S6c interaction, one might argue that a similar pore stabilization also accounts for this second slowing process. Recently, Haddad and Blunck (2011) showed that mutations within S4S5L (I358N) and S6c (F484G) that uncouple the communication between the VSD and pore domain prevented this second slowing period in IgOFF kinetics. However, both their data and our data indicate that the coupling between the VSD and BC gate is very tight and mutations within the pore module will consequently affect the transition rate toward the channel’s relaxed state. These data cannot rule out the hypothesis that this second VSD stabilization (the relaxation process) is an intrinsic property of membrane proteins possessing an S4-based VSD. In fact, the relaxation process, being an intrinsic property of the VSD, is strengthened by our observation that Ci-VSP, which does not contain a pore domain, and even its isolated VSD displays a similar slowing in VSD return as a function of the time spent in the activated state (Figs. 7 and 8). We propose that this second slowing in the kinetics of BC gate closure (Iac) and VSD deactivation (Iac) in Shaker and Kv1.2 channels is the direct consequence of a stabilization of the VSD into its relaxed state.

Although we do not rule out that the conformational changes of the pore when entering the C-type inactivated state have an impact on the movement of the VSD, our data strongly support that the described modulation in IgOFF and Iac kinetics occurs independently from the state of the K’ selectivity filter (being conductive or inactivated). First, the process of VSD relaxation is also observed in Ci-VSP and its isolated VSD version (Figs. 7 and 8). Second, the gating currents were recorded from conducting Shaker-H4-IR by depletion of potassium, and it has been described that reducing the K’ flux speeds up C-type inactivation substantially (Baukrowitz and Yellen, 1995, 1996; Starkus et al., 1997). Consequently, the slowing in IgOFF (K’-depleted conditions) and Iac (recorded in the presence of K’) would not be synchronized if C-type inactivation was involved. Moreover, Iac reports only from noninactivated channels, whereas IgOFF reports from both inactivated and noninactivated channels. Thus, if C-type inactivation were to affect the return of the VSD, there should be a discrepancy between the time-dependent changes in the kinetics of these two types of currents, which was not the case. Third, we recently showed that Shaker-W434F, a channel mutant that displays extremely fast inactivation (Yang et al., 1997), displays IgOFF slowing with a similar time development (Lacroix et al., 2011).
Coupling with the BC gate, we propose a gating scheme for both Shaker and Kv1.2 channels that accounts for the biphasic slowing process observed in both deactivating (IgOFF) and ionic (Ideac) currents (Fig. 9 A). To adequately illustrate the stabilizing effect of relaxation on both VSD deactivation and BC gate closure, we show two parallel schemes, one reflecting the status of the VSD and the other the BC gate. Both schemes are interconnected such that a stabilizing state in one scheme will result in a stabilized state in the other. Although the represented state model follows a square pathway occupying different energetic states, the forward and backward path followed by the VSD and BC gate can be structurally the same. Therefore, we provide an illustration of a linear scheme for the structural rearrangements in Shaker and Kv1.2 channels as a function of depolarization history (Fig. 9 B). Upon membrane depolarization, the channels evolve from closed over a preactivated state to the open state. Subsequently, there is a first stabilization imposed by the BC gate, and when depolarization is maintained, the VSD rearranges, resulting in a second stabilization referred to as the open relaxed state. This illustration concurs with the biphasic slowing observed in both IgOFF and Ideac, indicating the existence of three identifiable open channel conformations.

Because depolarization-induced QV shifts have been ubiquitously observed among various voltage-dependent proteins containing an S4-based VSD (Bezanilla et al., 1982; Shirokov et al., 1992; Piper et al., 2003; Kuzmenkin et al., 2003). Fourth, Kv1.2 channels whose C-type inactivation develops markedly slower than in Shaker channels (Fig. 4) displayed a VSD relaxation similar to Shaker channels. Finally, in both Shaker and Kv1.2 channels, the rate of C-type inactivation is approximately three-to fivefold slower than the time course of the VSD relaxation process. Although the exact underlying molecular mechanism for VSD relaxation is at present unknown, it involves a stabilization of the VSD in a postactivated conformation. Because of the strong coupling with the BC gate, this VSD relaxation is also reflected in the speed of gate closure. A previous study in hKv1.5 channels (the closest human homologue of Shaker) reported a similar slowing down in Ideac kinetics when depolarization times were increased (Rich and Snyders, 1998). Furthermore, as there were good reasons to conclude that this slow component in Ideac did not arise from channels recovering from an underlying hidden inactivated state, Rich and Snyders (1998) interpreted their observation as the channel having multiple open configurations. From the point of view of the kinetics of K⁺ conductance, this is indeed the case, but we show here that the underlying cause is the VSD relaxation process, and, therefore, the structural conformation of the BC gate might be similar in both open states.

Based on our previously described kinetic model that accounts for VSD relaxation at the level of gating charge movement and the observation that there is a tight coupling with the BC gate, we propose a gating scheme for both Shaker and Kv1.2 channels that accounts for the biphasic slowing process observed in both deactivating gating (IgOFF) and ionic (Ideac) currents (Fig. 9 A). To adequately illustrate the stabilizing effect of relaxation on both VSD deactivation and BC gate closure, we show two parallel schemes, one reflecting the status of the VSD and the other the BC gate. Both schemes are interconnected such that a stabilizing state in one scheme will result in a stabilized state in the other. Although the represented state model follows a square pathway occupying different energetic states, the forward and backward path followed by the VSD and BC gate can be structurally the same. Therefore, we provide an illustration of a linear scheme for the structural rearrangements in Shaker and Kv1.2 channels as a function of depolarization history (Fig. 9 B). Upon membrane depolarization, the channels evolve from closed over a preactivated state to the open state. Subsequently, there is a first stabilization imposed by the BC gate, and when depolarization is maintained, the VSD rearranges, resulting in a second stabilization referred to as the open relaxed state. This illustration concurs with the biphasic slowing observed in both IgOFF and Ideac, indicating the existence of three identifiable open channel conformations.

Because depolarization-induced QV shifts have been ubiquitously observed among various voltage-dependent proteins containing an S4-based VSD (Bezanilla et al., 1982; Shirokov et al., 1992; Piper et al., 2003; Kuzmenkin et al., 2003).
physiological mechanism that enables slowing down of a ubiquitous property of the VSD, may constitute the Xie et al., 2010). We propose here that VSD relaxation, Brew et al., 2007; Douglas et al., 2007; Shu et al., 2007; ling repetitive firing of a large variety of neurons has the selectivity filter remains highly conductive. sure may have a physiological impact, as it occurs while slower than VSD relaxation, the slowing in BC gate clo-

Kv1.2 channels in which C-type inactivation develops thus reducing the repolarizing power, slowing down channel closure has the opposite effect. Therefore, both processes affect the excitability of cells differently, and, in the case of Shaker, in which both occur simultaneously, they may counteract each other. However, in Kv1.2 channels in which C-type inactivation develops slower than VSD relaxation, the slowing in BC gate closure may have a physiological impact, as it occurs while the selectivity filter remains highly conductive.

The crucial role played by Kv1.2 channels in controlling repetitive firing of a large variety of neurons has been well established (Glazebrook et al., 2002; Dodson et al., 2003; Shen et al., 2004; Khavandgar et al., 2005; Brew et al., 2007; Douglas et al., 2007; Shu et al., 2007; Xie et al., 2010). We propose here that VSD relaxation, a ubiquitous property of the VSD, may constitute the physiological mechanism that enables slowing down of BC gate closure in Kv1.2 channels.

This research was funded by National Institutes of Health grant GM030376 (to F. Bezanilla), the Research Foundation Flanders, grants G.0256.08 (to D.J. Snyders) and 1.5.087.11N (to A.J. Labro), and grant GOA/TOP41.3016 from the concerted action fund of the University of Antwerp.

Kenton J. Swartz served as editor.

Submitted: 12 April 2012
Accepted: 13 September 2012

REFERENCE LIST


