Interaction between Tetraethylammonium and Permeant Cations at the Inactivation Gate of the HERG Potassium Channel

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Abstract: The fast inactivation of the human ether-à-go-go related gene product (HERG) channel is a form of C-type inactivation and is decelerated by external tetraethylammonium (TEA) and potassium. From the time constant of inactivation, the dissociation constants of TEA ($K_{TEA}^{\text{K}}$) and potassium ($K_{K}^{\text{K}}$) to the inactivation-impeding site were evaluated. $K_{TEA}^{\text{K}}$ was found to exhibit unexpected voltage dependence: $K_{TEA}^{\text{K}}$ decreased with depolarization. This was opposite the voltage dependence of $K_{K}^{\text{K}}$ on inactivation, in which permeating potassium impeded closure of the inactivation gate upon binding to a site in the pore (a “foot-in-the-door” mechanism). Further experiments on inactivation revealed anomalous mole fraction effects between permeating alkali cations and TEA, while no anomalous effects were seen between permeating ion species ($K^+$, $Rb^+$, $Cs^+$). The results indicate that TEA and permeating ions impede inactivation through binding to different but closely interacting sites. $K_{TEA}^{\text{K}}$ was influenced by permeating ions through their bindings in the pore. As the size of the occupied ion was increased the dissociation constant of TEA to the ion-occupied pore decreased. Thus, we conclude that an ion bound to the inactivation-impeding site in the selectivity filter is located in close proximity to TEA bound at the entrance of the filter. The order of affinity of alkali cations for the inactivation-impeding site, $Rb^+ > Cs^+ > K^+$, indicated that the selectivity of the site differed significantly from permeation selectivity, $K^+ > Rb^+ > Cs^+$. [Japanese Journal of Physiology, 53, 25–34, 2003]

Key words: anomalous mole fraction effect, ion selectivity, kinetics, selectivity filter, Xenopus oocytes.

Potassium channels exhibit strict ion selectivity uncompromised by a high permeation rate, the mechanism of which has been attributed to transitions and interactions of ions in multiple sites of the selectivity filter [1]. Recently, static pictures of the distribution of ions in the pore have been revealed by X-ray crystallography of the KcsA channel [2, 3]. Studies of molecular dynamics have highlighted events occurring in the time scale of picoseconds for transitions from one site to the others [4–10]. Dynamic characteristics of the ion-binding sites of the selectivity filter will enable the gap between the microscopic static picture and macroscopic ionic flow to be bridged and provide novel insights on the mechanisms of ion permeation and selectivity. Here, we extracted the individual characteristics of permeation sites from electrophysiological measurements of ion flow. We utilized fast inactivation of the human ether-à-go-go related gene product (HERG) channel and its sensitivity to tetraethylammonium (TEA) and permeating ions to examine the properties of the external mouth of the pore.

Fast inactivation of the HERG channel occurs by a form of C-type inactivation in which permeating ions impede the closure of the inactivation gate by a foot-
in-the-door mechanism; the inactivation gate closes upon emptying of the external filter [11–19]. Consequently, the fast inactivation gate continuously monitors the microscopic vacant status of the outer selectivity filter and actuates the gate to close, which can be evaluated by the time constant of inactivation from macroscopic measurements. TEA has been a versatile tool to explore the characteristics of potassium channels. Extracellular TEA binds to potassium channels with a high dissociation constant and blocks the potassium current [20]. Electrophysiological experiments have shown that external and internal TEA bind to the opposite ends of the selectivity filter and interact with each other through ions in the filter [21]. Studies of molecular dynamics have revealed that the stability of TEA binding depends strongly on ion distributions in the selectivity filter [22, 23]. For potassium channels exhibiting C-type inactivation, TEA slows the inactivation rate [11, 24]. The location of the binding site for TEA is the outer entrance of the selectivity filter, where TEA decelerates the inactivation. Thus, the dissociation constant of TEA evaluated by the deceleration effects on inactivation will reveal interactions between TEA at the entrance and ions in the permeating sites. Here, the following strategies were employed to characterize a site in the selectivity filter: The fast inactivation gate of the HERG channel was used as a sensor to detect the vacancy of a site in the selectivity filter. TEA acting from the extracellular side was used as a reporter of ion distribution in the selectivity filter. A strategy combining these approaches has allowed us to elucidate the characteristics of the inactivation-impeding site of the selectivity filter, which exhibited a marked difference in the binding selectivity from the permeation selectivity and contributes to TEA affinity through the occupying ion species.

MATERIALS AND METHODS

Oocyte isolation and cRNA injection. Female *Xenopus laevis* were anesthetized in tricaine solution (1.5 g/l) for 30 min at 4°C. Ovarian lobes were removed through a small abdominal incision. The follicular layer was removed enzymatically for 1.5 h with collagenase (Sigma, St. Louis, MO, USA) in Ca²⁺-free Barth’s solution (88 mM NaCl, 1 mM KCl, 2.4 mM NaHCO₃, 0.82 mM MgSO₄, and 5 mM HEPES, pH 7.6). HERG cRNA was prepared from cDNA (a gift provided by G. A. Robertson, University of Wisconsin, USA) [25] with a commercial kit (mMessage mMachTwo-electrode voltages) to a heterologous expression system (*Xenopus* oocytes). Oocyte isolation and cRNA injection. Two days after injection using a Dagan CA1 amplifier (Dagan Corporation, Minneapolis, IL, USA) at 23°C. The resistance of glass pipettes (borosilicate) filled with 3 M KCl was 0.5–1.0 MΩ. The cut-off frequency of the filter was 1 kHz and the sampling rate was 3 kHz. The bath solution contained 100 mM XCl (X: K, Rb, Cs, Na, NMDG [N-methyl-D-glucamine]); 1.8 mM CaCl₂, 1 mM MgCl₂, and 5 mM HEPES, pH 7.5. Except for solutions used for mole fraction experiments ([X]+ [TEA]=100 mM), NMDG was added as an inert cation. Hereafter, square brackets mean concentration. Raw two-electrode voltage clamp data were not leakage or capacitance subtracted. The current traces were fitted by an exponential function, \( I(t) = A \exp\left(-t/t_{\text{inact}}\right) + C \), where \( A \) is an amplitude and \( C \) is a constant, using pClamp (Axon Instruments, Inc., Union City, CA, USA). The reversal potential was obtained when oocytes were bathed in a 100 mM XCl (X: K, Rb, Cs, Na, and NMDG) solution. The permeability ratio \( (P_X/P_K) \) was calculated using the Goldman–Hodgkin–Katz equation [1], in which intracellular potassium concentration was estimated from the reversal potential in the pure potassium solution. Data were analyzed using Origin (MicroCal Software, Inc., Northampton, MA, USA) and Mathematica (Wolfram Research, Inc., Champaign, IL, USA). All data are shown as mean±SE (\( n=5–9 \)). For the fitted parameters, error propagations were taken into account [27].

RESULTS

Strange voltage dependence of the dissociation constant of TEA

The fast inactivation of the HERG channel was investigated by applying electrophysiological methods to a heterologous expression system (*Xenopus* oocytes). A double-pulse protocol enlightened the inactivation process (Fig. 1A): From the holding voltage of −80 mV, a long depolarizing pulse (3.0 s) at +20 mV activated the channel, which subsequently underwent a very fast inactivation process (C→O→I), where C, O, I represent the closed-, open-, and inactivated-state, respectively. Then, a short interpulse (30–50 ms) at −100 mV recovered the channel from inactivation (I→O: deinactivation). Finally, the second depolarizing pulse, at various voltages, manifested the pure inactivation process (O→I). The current traces for the
fast inactivation are shown for the membrane potentials from −50 to +50 mV recorded in the absence of external potassium (Fig. 1B). As the membrane potential was depolarized, the current inactivated more rapidly and more completely.

The effect of TEA on inactivation at +50 mV is shown in Fig. 1C. As the TEA concentration increased, the time course of inactivation was significantly prolonged. The slower inactivation has been described for open channel blockers, where the blocked channel does not gate (inactivate) unless it is unblocked [28], which is represented by the following scheme.

\[
B \xrightarrow{k_{\text{block}}} \text{O} \xrightarrow{k_{\text{DInact}}} \text{I} \xrightarrow{k_{\text{unblock}}} B
\]

where B represents the blocked-state; \(k_{\text{inact}}\) and \(k_{\text{DInact}}\) are the inactivation and deinactivation rate constants; and \(k_{\text{block}}\) and \(k_{\text{unblock}}\) are the blocking and unblocking rate constants. The dissociation constant of TEA \(K_{\text{TEA}}\) is \(k_{\text{unblock}}/k_{\text{block}}\).

From Scheme 1 the time constant of inactivation \(\tau_{\text{TEA}}\) can be formulated as a function of TEA concentration:

\[
\tau_{\text{TEA}} = \frac{1}{k_{\text{inact}}} \frac{1+\text{[TEA]}/K_{\text{TEA}}}{1+K_I(1+\text{[TEA]}/K_{\text{TEA}})}
\]

where \(K_I\) is the equilibrium constant for inactivation (\(=k_{\text{DInact}}/k_{\text{inact}}\), all equations are listed in APPENDIX).

The \(K_{\text{TEA}}\) value was evaluated from the data shown in Fig. 1D inset, where normalized \(\alpha_{\text{TEA}}\) (\(=\text{[TEA]}/[\text{I}]\)) was plotted as a function of TEA concentration and fitted by Eq. A1. Fig. 1F: The voltage dependency of the \(K_{\text{pot}}\) values. \(\alpha_{\text{TEA}}\) was plotted as a function of potassium concentration and was fitted by Eq. A2.

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**Fig. 1. Inhibitory actions of TEA and potassium on inactivation gating.** A: Voltage-clamp experiments of the HERG channel expressed in Xenopus oocytes. A command pulse protocol and representative current traces in the absence of external potassium (100 mM NMDG) are shown. A double-pulse protocol was applied to investigate the inactivation process. The current traces demarcated by a broken line are expanded and shown in B. B: Inactivating current traces from −50 to 50 mV. The broken line indicates the zero current level. C, E: Current traces in the presence of external TEA (C) and potassium (E) measured at +50 mV. The current amplitudes were normalized. D: The voltage dependency of the \(K_{\text{TEA}}\) values. Inset: The concentration dependency of the \(\tau_{\text{TEA}}/\tau_0\) values. \(\tau_{\text{TEA}}/\tau_0\) was plotted as a function of TEA concentration and fitted by Eq. A1. F: The voltage dependency of the \(K_{\text{pot}}\) values. \(\tau_{\text{pot}}/\tau_0\) was plotted as a function of potassium concentration and was fitted by Eq. A2.
where \([K^-]_o\) represents the extracellular potassium concentration and \(O_{occ}\), \(O_{vct}\), and \(I\) indicate the ion-occupied open state, ion-vacant open state, and inactivated state, respectively. In this scheme, the ion-vacant state does not refer to a channel state without any ions in the pore. Rather, one of the ion binding sites, upon occupied inactivation is impeded (an inactivation-impeding site), is vacant, leaving the rest of the sites unconsidered. Here, the simplest case for the transition between two permeating states of the pore is shown. This model represents the shift model deduced from ion distribution in the KcsA potassium channel [3, 5]; \(k_{bind}\) and \(k_{unbind}\) are the ion-binding and ion-unbinding (from outer site) rate constants, respectively, \(k_{esc}\) and \(k_{eq}\) are the ion-escaping and ion-filling (from inner sites) rate constants. The steady-state probabilities of the open states are expressed using these four rate constants [29]. \(K^K\) is the dissociation constant of potassium and is given by \((k_{unbind}+k_{esc})/k_{bind}\). \(r^K\) is expressed as follows:

\[
\tau^K = \frac{1}{k_{inact}^0} \frac{1 + [K]/K^K}{1 + K_i (1 + [K]/K^K)}
\]

where the apparent inactivation rate, \(k_{inact}^0 \left( = \frac{p^{inact}}{p^{vct}} \times k_{inact}\right)\), is the rate in the absence of external potassium.

The \(K^K\) value was evaluated from concentration dependencies of \(\tau^K\) (Eq. 2). The normalized \(\tau^K\) values \((\tau^K/\tau^0)\) were fitted by Eq. A2 (see APPENDIX), in which non-linearity was parameterized by the non-zero value of \(K_i\). The \(K^K\) values for different membrane potentials are shown in Fig. 1F. In contrast to TEA action, the \(K^K\) value increased significantly as the membrane potential was depolarized. Thus, the ion binding site impeding the inactivation locates in the pore under the influence of the electric field (i.e., a mechanism known as “foot-in-the-door”).

**Interactive action of TEA and K⁺ on HERG inactivation**

The effects of external potassium on \(K^{TEA}\) were examined. In Fig. 2 and inset, \(K^{TEA}\) evaluated from \(\tau_{inact}\) was shown, in which the extracellular potassium concentration was fixed. In the inset, \(\tau_{inact}\) was plotted as a function of the TEA concentration. Two sets of data for different potassium concentrations (0 and 20 mM at ±50 mV) are shown. The data points were fitted by straight lines. From Eq. 1, different slopes indicate different \(K^{TEA}\) values. Therefore, \(K^{TEA}\) was affected by external K⁺. This is good evidence that TEA and potassium interact with each other in the inactivation process. It is noted that the extrapolated lines intersected at the third quadrant, which occurs between uncompetitive inhibitors (see the following section on the mole-fraction effect) [30].

Voltage dependencies of \(K^{TEA}\) for four different potassium concentrations are shown in Fig. 2. \(K^{TEA}\) at 2 mM K⁺ exhibited similar voltage dependency with a sigmoidal shape to those observed at a potassium concentration of zero. On the other hand, at 20 mM K⁺, the voltage dependency of \(K^{TEA}\) was almost abolished. This sigmoidal shape and its abolition by external potassium were intriguing features of the interaction between potassium and TEA. Comparing \(K^{TEA}\) (Fig. 1D) with \(K^K\) (Fig. 1F) as a function of voltage, it is noted that a mirror-image relationship exists between the two dissociation constants: As the membrane potential depolarizes, the \(K^{TEA}\) value decreases, which coincides with the increase in the \(K^K\) value. At potentials with higher \(K^K\) values, the occupancy of potassium on the inactivation-impeding site should be lower and those channels have lower dissociation constant to TEA. Thus, it is likely that the dissociation constant of TEA is controlled by the existence of a
potassium ion on a specific site of the selectivity filter. A plausible model of these interactions, in which TEA binds to both vacant and occupied pores, was introduced (Scheme 3).

![Scheme 3](image)

where \( B \) and \( O \) represent the ion-vacant blocked state and ion-occupied blocked state, respectively, and \( K_{\text{TEA}}^{\text{vct}} \) and \( K_{\text{TEA}}^{\text{occ}} \) are the dissociation constants of TEA to the vacant and occupied pores. \( \tau_{\text{inact}} \) is expressed by Eq. A3 (see APPENDIX).

**Anomalous mole-fraction effect between permeating ions and blocking TEA**

To elucidate the mechanism of interaction between two inhibitors, mole fraction experiments were performed. In the Fig. 3A inset, current traces recorded at \(+50\) mV in the pure (K+ or TEA) and mixed (K+ and TEA) external solutions are shown. The time course of inactivation in the 1:1 mixed solution was not between those in two pure solutions. In the figure, the concentration dependency of \( \tau_{\text{inact}} \) in the pure potassium solution (the open triangle symbol) is shown as a reference, as well as the pure TEA effect plotted in the opposite direction of the abscissa (the open square symbol). In mole fraction solutions, the \( \tau_{\text{inact}} \) values deviated upward significantly from the linear relationship that is roughly a sum of the pure effects, and showed the maximum value in the mixed solution (Fig. 3A). This anomalous mole-fraction effect indicates that TEA and potassium bind to different sites and impede inactivation.

The anomalous mole-fraction effect was analyzed quantitatively by fitting the data to Eq. A3, where the parameters obtained hitherto were used for new parameter \( (K_{\text{TEA}}^{\text{TEA}}) \) fitting. From the fitted parameters, hypothetical lines were drawn for competitive and uncompetitive actions. In the figure, the straight broken line indicates a competitive action (TEA does not bind to the ion-occupied pore; \( K_{\text{vct}}^{\text{occ}} = \infty \)) and the curved broken line indicates an uncompetitive action (TEA can bind to the channel with the same dissociation constant whether a site in the pore is occupied by an ion or not; \( K_{\text{TEA}}^{\text{TEA}} = K_{\text{TEA}}^{\text{vct}} \)). The data points fall between the competitive and uncompetitive lines, indicating that \( K_{\text{TEA}}^{\text{TEA}} \) is not equal to but larger than \( K_{\text{vct}}^{\text{TEA}} \). A similar result is shown in Fig. 2 and its inset: \( K_{\text{TEA}}^{\text{TEA}} \) increased as the fixed concentration of potassium was raised. This happens only when \( K_{\text{vct}}^{\text{TEA}} > K_{\text{TEA}}^{\text{TEA}} \) (see Eqs. A4 and A5). Both experiments indicate that TEA binding to the ion-occupied pore is weaker than that to the vacant pore.

We also used other alkali cations. Outward currents at positive potentials were little affected by Rb+ and Cs+, and inactivating currents were recorded at +50 mV in the pure (K+ or TEA) and mixed (K+ and TEA) external solutions. The time course of inactivation in the 1:1 mixed solution was not between those in two pure solutions. In the figure, the concentration dependency of \( \tau_{\text{inact}} \) in the pure potassium solution (the open triangle symbol) is shown as a reference, as well as the pure TEA effect plotted in the opposite direction of the abscissa (the open square symbol). In mole fraction solutions, the \( \tau_{\text{inact}} \) values deviated upward significantly from the linear relationship that is roughly a sum of the pure effects, and showed the maximum value in the mixed solution (Fig. 3A). This anomalous mole-fraction effect indicates that TEA and potassium bind to different sites and impede inactivation.

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mV. As shown in Fig. 3B and C, inactivation was slowest for the mixed solution of an alkali cation and TEA. Prominent anomalous mole-fraction effects on \( \tau_{\text{inact}} \) were seen for Cs\(^+\), where data points approached the uncompetitive curve. Rb\(^+\) and TEA also exhibited an anomalous effect. In this case, the data points located at the middle of the competitive and uncompetitive lines. In contrast to the interaction between ions and TEA, mole fraction experiments for two species of alkali cations (K\(^+\) vs. Rb\(^+\) and K\(^+\) vs. Cs\(^+\)) showed no anomalous behavior (Fig. 3D). The results indicate that permeating ions compete for a site in the pore. These mole fraction experiments revealed that interaction between TEA and ions depends largely on ion species and that the inactivation-impeding site for permeating ions is localized rather than spread over the selectivity filter.

Anomalous mole-fraction effects for different ion species were analyzed quantitatively. We focused on the dissociation constant of TEA to the ion-occupied pore, \( K_{\text{TEA}} \), since it reflects the pattern of the data points and is sensitive to ion species. The \( K_{\text{TEA}} \) value was the largest for the K\(^+\)-occupied pore and decreased as the ionic radius of the occupied ion increased (\( K_{\text{TEA}} \) for K\(^+\) (1.33 Å)=20.7±3.97 mM; for Rb\(^+\) (1.48 Å)=14.9±2.55 mM; for Cs\(^+\) (1.69 Å)=8.8±1.15 mM (n=5)). How do such small differences in the ionic radii reflect on the sizable differences in the dissociation constant of TEA? Larger ions do not become an obstacle for TEA binding, but rather increases the affinity. Possible mechanisms will be discussed.

**Ion specificity of the ion-binding site impeding inactivation**

Binding of the alkali ions alone to the inactivation-impeding site in the selectivity filter was evaluated from the concentration-dependence of the \( \tau_{\text{inact}} \) values at +50 mV (Fig. 3A–C). The \( K_{\text{ion}} \) values for alkali cations are shown in Table 1. The selectivity of the inactivation-impeding site was Eisenman’s sequence II (Rb\(^+\) > Cs\(^+\) > K\(^+\)) [1]. As a reference, the permeability coefficients were obtained from the reversal potentials under bi-ionic conditions (see METHODS) and the permeability ratio is shown in Table 1. The selectivity sequence was: K\(^+\) > Rb\(^+\) > Cs\(^+\). A discrepancy appeared between the affinity to the inactivation-impeding site and permeation selectivity. Thus, a local property of the inactivation-impeding site was extracted from the multiple ion-binding sites in the selectivity filter.

**Table 1. Ion selectivity for the inactivation-impeding site and throughput permeation.**

<table>
<thead>
<tr>
<th>Ion</th>
<th>K(^+)</th>
<th>Rb(^+)</th>
<th>Cs(^+)</th>
</tr>
</thead>
<tbody>
<tr>
<td>( K_{\text{ion}} ) (mM)</td>
<td>18.6±1.86</td>
<td>5.49±0.80</td>
<td>8.22±1.18</td>
</tr>
<tr>
<td>( P_{\text{fl}}/P_{\text{k}} )</td>
<td>1</td>
<td>0.95±0.02</td>
<td>0.24±0.01</td>
</tr>
</tbody>
</table>

The dissociation constants of alkali cations to the inactivation-impeding site were evaluated from the data shown in Fig. 3 (open symbols) (n=5). The permeability ratios (\( P_{\text{fl}}/P_{\text{k}} \)) were calculated from the reversal potentials (n=10).

**DISCUSSION**

In this paper, we have investigated the action of TEA and permeant ions interacting with the inactivation gate and obtained a detailed picture of the inactivation-impeding site of the HERG channel. TEA and an alkali cation bind to different sites and both impede inactivation. The binding site of TEA should locate at the entrance of the pore, since the voltage-dependence of \( K_{\text{TEA}} \) is eliminated as the potassium concentration is raised (Fig. 2). On the other hand, the binding site of the permeating ions is located in the pore, which is indicated by voltage-dependent \( K_{\text{ion}} \). The mole fraction experiments between alkali cations revealed that the inactivation-impeding action of permeating ions is competitive. In the selectivity filter, inactivation is impeded by an ion bound to a localized site.

Recent progress on the permeation mechanism in the KcsA channel assumes that permeating ions move in the selectivity filter in a concerted manner: The predominant ion distribution in the selectivity filter has been proposed to be W–I–W–I or I–W–I–W for four binding sites (from the extracellular side) where I represents a permeating ion and W represents a water molecule. Ion permeation operates preferentially by transition between these two distributions [3, 5]. Here, two permeation states in our permeation-coupled inactivation scheme (Scheme 3) are allocated to the ion distributions in the shift model (Fig. 4). In our experiments, the results plotted in Fig. 1F show that raising external concentration decelerates inactivation. In the shift model, increased external potassium boosts the shift in the distribution from W–I–W–I to I–W–I–W. Thus, our results, based on the shift model, reveal that inactivation proceeds from W–I–W–I distribution and W–I–W–I corresponds to the “vacant” state (Fig. 4).

TEA binds more tightly to the vacant pore than to the occupied pore, thus, the apparent dissociation constant of TEA reflects the vacant probability of the pore. Lower \( K_{\text{TEA}} \) values at depolarized potentials with their sigmoidal voltage dependency (Fig. 2) indi-
cate that vacant probability increased with depolariza-
tion. This was true even in the absence of external 
potassium. The site should be filled by potassium 
from the inner sites, and hyperpolarization facili-
tates the filling. This seemingly opposite voltage depen-
dency is explained in the shift model (Fig. 4) if the 
intracellular potassium association rate (k_{oc}) over-
whelms the transition rate from I–W–I–W to W–I– 
I–W (k_{oc}). The shift model also supports the high 
voltage dependence of K_{ic}, since permeating potassium 
ions reach the inactivation-impeding site through 
concerted movements with larger effective charge. 
The shift model simplified the permeation issues and aided 
in the interpretation of our experimental data.

The inactivation-impeding site in the “Shaker” 
potassium channel has been proposed to be located in 
the outer selectivity filter [16]. By assuming an exclu-
sive shift model for the permeation mechanism, we 
concluded that the W–I–W–I distribution is the “va-
cant” state. If an ion occupies either the first site or 
the third site of the selectivity filter, conformational 
change for inactivation cannot be initiated. We found 
that the binding of TEA is governed by the ion species 
occupied in the pore, with the dissociation constant 
being decreased as the size of the occupied ion is 
increased. Such ion-size sensitivity and decremental 
size-dependency strongly suggest that the inactiva-
tion-impeding site should locate close to bound TEA 
(i.e., the outermost site of the selectivity filter). If the 
inactivation-impeding site is the third position, the 
difference in ion size is smeared at the TEA binding 
site. In the case of close proximity between bound ion 
and TEA, how is the electrical repulsion force be-
tween closely located TEA and an ion overcome? In a 
molecular dynamics study, TEA and potassium were 
located adjacent at the entrance of the pore without in-
serting a water molecule between them [5]. The inter-
action free energy was not unrealistically high.

Here, we propose a hypothesis that bound TEA and 
an ion interact electrostatically. Permeating ions 
bound in the selectivity filter are solvated by eight car-
bonyl oxygen atoms [2]. Bound ion in the outermost 
site of the selectivity filter exposes only a part of its 
surface to the pore entrance. Thus, TEA can bind to 
the pore entrance being subjected to the attenuated 
electrostatic repulsion force from the buried ion in the 
outermost site. If they are in close contact, the dis-

cance between TEA and the ion becomes r_{TEA}+r_{ion}. In 
Fig. 5, the free energy of the TEA binding calculated 
from the K_{TEA} values (Fig. 3 legends) is plotted as a 
function of the distance. Here, the slope represents the 
inverse of the dielectric constant (e_{oc}=6.5), assuming 
that the difference of the free energy is attributed ex-
clusively to electrostatic interaction. This value repre-
ts the environment surrounding an alkali cation-
TEA pair around the entrance of the selectivity filter.

The vacant site is filled by alkali cations from out-

Fig. 4. A model for interaction among permeating 
ions, TEA, and the inactivation gate. This model imple-
ments the permeation mechanism of the shift model into 
the vacancy-coupled inactivation scheme (Scheme 3). Per-
meation occurs through transition between two ion distribu-
the distribution from W–I–W–I to I–W–I–W with the rate of 
[K^{+}]_{Woc}, where k_{oc} is the binding (second-order) rate con-
stant from the extracellular side. The opposite shift occurs 
by intracellular potassium with the rate of [K^{+}]_{Ioc}, where k_{oi} 
is the binding rate constant from the inside. Transition 
between two ion distributions also takes places without the 
contribution of ions outside the selectivity filter. The first-
order rate constants for the transition are k_{oi} and k_{oc}.

Fig. 5. TEA binding energy and ionic radius occupied 
in the pore. The free energy (∆G) of the TEA binding to the 
occupied pore as a function of an inverse of the distance 
between an ion and TEA (r_{ion}+r_{TEA}; ionic radius of 
1.33 Å for K^{+}, 1.48 Å for Rb^{+}, and 1.69 Å for Cs^{+} and the 
effective radius of 4.5 Å for TEA) [1, 22, 23]. The ∆G values 
were calculated from the K_{TEA} values at +50 mV. The data 
points were fitted by an electrostatic equation [30, 31], 
\[ \Delta G = \frac{1}{r} \left( \frac{1}{r_{TEA}} + r_{ion} \right) \]

\[ \text{K} \]

\[ \Delta G_{TEA} \text{(kJ/mol)} \]

\[ 1/(r_{ion}+r_{TEA}) \text{ (1/\text{nm})} \]
side with the selectivity of Rb\(^+\) > Cs\(^+\) > K\(^+\). Thus, the outermost site is neither just a dehydration site nor a site with comparable selectivity for the throughout permeation. By focusing on a single permeation site, a discrepancy between local and overall properties was elucidated. Once bound, an ion contributes a part of the TEA binding site, thus controlling TEA affinity. The characteristics of the outermost site of the selectivity filter elucidated here, based on dynamic measurements, contribute invaluably to further understanding the mechanism of ion permeation and selectivity.

**Note added in proof:** Since this paper was submitted, a paper has appeared by L. Guidoni and P. Carioni (J Recept Signal Transduct Res 22: 315–331, 2002). They calculated TEA binding for different ion distributions in the selectivity filter of the KcsA potassium channel and exhibited that a quasi-planar form of TEA bound to the K–W–K–W configuration with only 0.7 kcal/mol less stable than to the W–K–W–K configuration.

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**REFERENCES**

distribution for permeating ions in the selectivity
the inactivation-impeding site being occupied (Oocc)
divided into two ensembles of states with respect to
when the outer selectivity
pressed as follows:

\[
\frac{D_{\text{Inact}}}{p_{\text{Inact}}} = \frac{1}{K_{\text{DInact}}} \left( \frac{1 + [K]/K^K}{1 + K_1(1 + [K]/K^K)} \right)
\]  

The five-state model
The inhibitory action of TEA and permeating ions
on inactivation was expressed as a five-state model
(Scheme 3). A formula was derived under the
assumptions that (1) the channel immediately reaches
the steady-state among the open states (empty and occu-
perations between open states with different ion distrib-
ions are much faster than those between open and in-
activated states. The steady-state probability of the oc-
occupied-open and vacant-open states can be expressed
by the ratio of polynomials of ion concentrations. The
order of the polynomial depends on the number of the
ion-binding sites for multi-ion permeation. Here, the
simplest case, which represents not only one-ion pore
but also the shift model deduced from ion distribution
in the KcsA potassium channel [3, 5], is shown. The
inactivation-impeding site is filled by a permeating
ion from either an outer or inner site. Thus, the transi-
ption from vacant pore to occupied pore was expressed
by four rate constants with concentration-dependent
(the second-order) and independent (the first-order)
facets. The vacant probability becomes

\[
p_{\text{vct}} = p_{\text{vct}}^0 \frac{1}{1 + K(1 + [K]/K^K)}
\]

The apparent inactivation rate, \( k_{\text{Inact}}^p = p_{\text{vct}}^0 \times k_{\text{Inact}} \),
is the rate in the absence of external potassium. The
normalized time constant is expressed as follows:

\[
\tau^o = 1 + \frac{[K]/K^K}{1 + K_1(1 + [K]/K^K)}
\]  

\[
\tau_o = 1 + \frac{[K]/K^K}{1 + K_1(1 + [K]/K^K)}
\]  

where \( K_{\text{TEA}} \) is the dissociation constant of TEA,
and \( K_1 \) is the equilibrium constant for inactivation
\( (= k_{\text{DInact}}/k_{\text{Inact}}) \). When \( K_1 \) is negligible, the equation
becomes linear. The normalized time constant is ex-
pressed as follows:

\[
\tau^p \tau^o = 1 + \frac{[\text{TEA}]/K_{\text{TEA}}}{1 + K_1(1 + [\text{TEA}]/K_{\text{TEA}})}
\]  

Permeating potassium impedes inactivation by the
foot-in-the-door mechanism. Inactivation proceeds
when the outer selectivity filter is vacant [16]. Ion dis-
tribution for permeating ions in the selectivity filter is
divided into two ensembles of states with respect to
the inactivation-impeding site being occupied (Oocc) or
vacant (Ovct). The steady-state approximation for
the probability of open states was applied since transi-
tions between open states with different ion distrib-
utions are much faster than those between open and in-
activated states. The steady-state probability of the oc-
occupied-open and vacant-open states can be expressed
by the ratio of polynomials of ion concentrations. The
order of the polynomial depends on the number of the
ion-binding sites for multi-ion permeation. Here, the
simplest case, which represents not only one-ion pore
but also the shift model deduced from ion distribution
in the KcsA potassium channel [3, 5], is shown. The
inactivation-impeding site is filled by a permeating
ion from either an outer or inner site. Thus, the transi-
ption from vacant pore to occupied pore was expressed
by four rate constants with concentration-dependent
(the second-order) and independent (the first-order)
facets. The vacant probability becomes

\[
p_{\text{vct}} = p_{\text{vct}}^0 \frac{1}{1 + K(1 + [K]/K^K)}
\]

The apparent inactivation rate, \( k_{\text{Inact}}^p = p_{\text{vct}}^0 \times k_{\text{Inact}} \),
is the rate in the absence of external potassium. The
normalized time constant is expressed as follows:

\[
\tau_o = 1 + \frac{[K]/K^K}{1 + K_1(1 + [K]/K^K)}
\]  

\[
\tau_o = 1 + \frac{[K]/K^K}{1 + K_1(1 + [K]/K^K)}
\]  

where \( [X] \) represents the concentration of an alkali-
cation and the dissociation constant of the X ion is
\( K_{\text{Ion}} \). The dissociation constant of TEA to the vacant
pore is \( K_{\text{TEA}} \) and that to the occupied pore is \( K_{\text{TEA}} \).
The apparent dissociation constant of TEA (\(K_{\text{TEA}}\)) is determined by a combination of these dissociation constants (\((1-p_{\text{emp}}^0)/K_{\text{occ}}^\text{TEA} + p_{\text{emp}}^0/K_{\text{vct}}^\text{TEA}\)).

From this equation, qualitative features of interaction between inhibitors can be predicted. For example, potassium dependency of the \(K_{\text{TEA}}\) value can be expressed as:

\[
K_{\text{TEA}} = \frac{1 + \frac{p_{\text{emp}}^0}{K_{\text{lon}}} [X]}{1 - \frac{p_{\text{emp}}^0}{K_{\text{TEA}}^\text{occ}} + \frac{p_{\text{emp}}^0}{K_{\text{TEA}}^\text{vct}} + \frac{p_{\text{emp}}^0}{K_{\text{TEA}}^\text{occ} K_{\text{lon}}} [X]}
\]

(A4)

Differentiating the equation in terms of the concentration leads to a simple relationship: The increase in \(K_{\text{TEA}}\) with increased permeant concentration happens only with \(K_{\text{occ}}^\text{TEA} > K_{\text{TEA}}\).

When the deinactivation rate is negligible (\(K_{\text{i}} \equiv 0\) or \(p_{\text{emp}}^0 k_{\text{inact}} \gg k_{\text{Dinact}}\), which is the case at positive potentials), Eq. A3 becomes a simplified form:

\[
\tau_{\text{inact}} = \frac{1}{K_{\text{inact}}^0} \left\{ 1 + \frac{[X]}{K_{\text{lon}}^0} + \left( \frac{1}{K_{\text{TEA}}} + \frac{1}{K_{\text{occ}}^\text{TEA} K_{\text{lon}}^0} [X] \right) [\text{TEA}] \right\}
\]

(A5)

For fixed concentrations of an ion, \([\text{TEA}]\) dependency of \(\tau_{\text{inact}}\) becomes linear. Extrapolating each line for different concentrations of an ion to the negative region discriminates a mode for the inhibitory action between the ion and TEA. If an ion and TEA are competitive, TEA cannot bind to the occupied pore \((K_{\text{occ}}^\text{TEA} = \infty)\), thus, the slope is fixed and the lines are parallel. If they are uncompetitive (TEA can bind to vacant and occupied pores equally \((K_{\text{occ}}^\text{TEA} = K_{\text{vct}}^\text{TEA} = K_{\text{TEA}})\)), all lines cross on the abscissa axis at \(-K_{\text{TEA}}\). If they are neither competitive nor uncompetitive, lines still intersect in the negative region at the coordinate of

\[
\{x, y\} = \left\{ -K_{\text{occ}}^\text{TEA}, \frac{1}{p_{\text{emp}}^0 k_{\text{inact}}} \left( 1 - \frac{K_{\text{occ}}^\text{TEA}}{K_{\text{TEA}}} \right) \right\}
\]

The crossing point locates at the third quadrant, if \(K_{\text{occ}}^\text{TEA} > K_{\text{TEA}}\) (\(K_{\text{occ}}^\text{TEA} > K_{\text{vct}}^\text{TEA}\) from Eq. A4).