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Multi-ion conduction bands in a simple model of calcium ion channels

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Abstract
We report self-consistent Brownian dynamics simulations of a simple electrostatic model of the selectivity filters (SF) of calcium ion channels. They reveal regular structure in the conductance and selectivity as functions of the fixed negative charge \(Q_f\) at the SF. With increasing \(Q_f\), there are distinct regions of high conductance (conduction bands) \(M_0, M_1, M_2\) separated by regions of almost zero-conductance (stop-bands). Two of these conduction bands, \(M_1\) and \(M_2\), are related to the saturated calcium occupancies of \(P = 1\) and \(P = 2\), respectively and demonstrate self-sustained conductivity. Despite the model’s limitations, its \(M_1\) and \(M_2\) bands show high calcium selectivity and prominent anomalous mole fraction effects and can be identified with the L-type and RyR calcium channels. The non-selective band \(M_0\) can be identified with a non-selective cation channel, or with OmpF porin.

1. Introduction

Voltage-gated calcium ion channels play an important role in stimulating muscle contraction, in neurotransmitter secretion, gene regulation and transmission of action potentials, based on their high selectivity for divalent calcium ions \(\text{Ca}^{2+}\) over monovalent sodium ions \(\text{Na}^+\). They exhibit the anomalous mole fraction effect (AMFE), an effective blockade of \(\text{Na}^+\) permeation by small concentrations of \(\text{Ca}^{2+}\), combined with measurable \(\text{Ca}^{2+}\) currents in the pA range [1].

The selectivity of calcium channels is defined by a narrow selectivity filter (SF) with a strong binding site formed by negatively-charged protein residues [1, 2]. Wild-type calcium channels and their mutants differ in the composition, structure (locus) and net fixed charge \(Q_f\) of these protein residues at the SF. The most-studied L-type calcium channel possesses an EEEE locus with an estimated \(Q_f \approx 3-4e\) [2, 3], where \(e = -1.6 \times 10^{-19} \text{ C}\) is the electronic charge. The ligand-gated Ryanodine receptor (RyR) calcium channel has a DDDD locus with a DDDD scaffold with an orifice of \(4.6 \times 4.6 \text{ Å}\) defined by van der Waals surfaces [6, 8].

Mutant studies show that \(Q_f\) is crucial in determining the \(\text{Ca}\) versus \(\text{Na}\) selectivity of calcium channels. Usually, mutations that influence \(Q_f\) also destroy the channel’s selectivity and hence physiological functionality [9]. However, an appropriate point mutation of the DEKA sodium channel (\(Q_f \approx 1e\)) converts it into a calcium-selective channel with a DEEA locus and \(Q_f \approx 4e\) [10]. The essentially nonselective bacterial OmpF porin (\(Q_f \approx 1e\)) can be also turned into a Ca-selective channel by introduction of two additional glutamates in the constriction zone. The resultant mutant contains a DEEE-locus (\(Q_f \approx 4e\)) and exhibits an Na current with a strongly increased sensitivity to 1mM Ca [11].

Sodium channels have structures very similar to calcium channels, but with different SF loci (and therefore different \(Q_f\)), and different lengths and radii [5–7]. The eukaryotic sodium channel has a DEKA locus with \(Q_f \approx 1e\) [5, 7]. In the recently studied bacterial NavAb channel, the four Glutamate side chains form a \(6.5 \times 6.5 \text{ Å}\) scaffold with an orifice of \(4.6 \times 4.6 \text{ Å}\) defined by van der Waals surfaces [6, 8].

Dynamic Monte-Carlo simulations of the flexible volume exclusion model of calcium and sodium channels [3, 12] show that the charge density at the SF is the first-order determinant of selectivity [3] and the \(\text{Na}^+\) to \(\text{Ca}^{2+}\) occupancy ratio decreases
monotonically as $Q_f$ increases from 1e (DEKA locus) to 4e (DEEE locus), while the pore becomes more and more Ca$^{2+}$ selective [5].

Self-consistent Brownian dynamics (BD) simulations [2] of conductance and selectivity, based on a purely electrostatic model of the L-type channel with a rigid binding site, revealed a narrow peak in Ca$^{2+}$ conductance near $Q_f = 3.2e$ [2, 13].

Although mutant studies [9–11] and simulations [2, 3, 5] have demonstrated very clearly the dominant influence of $Q_f$ on the selectivity of the calcium channel, it has remained unclear why particular values of $Q_f$ should be optimal for selectivity or how many such values may exist.

Multi-ion knock-on conductivity is assumed to be one of the main mechanisms of permeation and selectivity in the case of the potassium channel [14–17]; a similar mechanism could resolve the paradox of high selectivity and high conductance in the calcium ion channel [18, 19]. Its selectivity and conductivity are connected with strong ion-ion repulsion and multi-ion occupancy of the SF [2, 12, 19]. The discreteness of the ionic occupancy also plays a significant role in conductance and is expected to manifest as discrete multi-ion steps/bands in the dependences of occupancy and conductance on concentration, time and other parameters [20–22]. These steps/bands are related to the barrier-less knock-on conduction mechanism that has been suggested as being the underlying mechanism responsible for the conductivity and selectivity of ion channels [17]. Simplified (‘toy’) electrostatic models of an ion channel, describing it as a water-filled charged protein hub in the cell membrane, were found to reproduce significant features related to the conductivity and, in particular, to the valence selectivity between ions of different charge [13, 22–25].

In this paper, we show that the Ca$^{2+}$ conduction and Ca$^{2+}$/Na$^+$ valence selectivity of a simple calcium channel model forms a regular structure of conduction/selectivity bands (regions) as a function of $Q_f$, separated by non-conduction bands. This pattern is related to saturated, almost barrier-less, conductivity with different numbers of Ca$^{2+}$ ions involved in the conduction. The conductance peak obtained in [2] is one part of this structure. We infer that all calcium-selective channels (both wild-type and mutants) should correspond to one of these bands.

2. Methods

2.1. A simple model of the calcium channel

We use a simple, self-consistent, purely electrostatic model of a calcium ion channel to study the effects of surface charge $Q_f$ on its conduction and selectivity. This model represents the channel’s selectivity filter (SF) as a negatively charged, axisymmetrical, water-filled protein hub in the cell membrane (figure 1) similar to that used in earlier work [2, 3, 26]. We model the SF only, following [3, 26]: we omit the charged vestibule modelled in [2] because the ion–ion and ion–fixed-charge interactions inside the SF are the main determinants of selectivity [2, 3]. The negatively charged protein residues are modelled as a single, thin, uniformly charged, centrally placed, rigid ring around the SF. Located within the wall, the ring brings a net negative charge $Q_f$ of 0–6.5e. The extracellular (left) and intracellular (right) baths are filled with ionic sodium-only, calcium-only, or mixed sodium–calcium aqueous solutions. In what follows we assume an asymmetrical ionic concentration: $C_L > 0$, $C_R = 0$.

The minimum possible radius $R$ of the SF of an L-type calcium channel has been determined as being $R = 2.8$ Å. We use the value of $R = 3.0$ Å. We choose the length $L = 16$ Å for the main simulations as being close to the value used in [3]. Moving sodium and calcium ions are described as charged particles of radius $R \approx 1$ Å (matching both ions), with diffusion coefficients of $D_{Na} = 1.17 \times 10^{-9}$ m$^2$s$^{-1}$ and $D_{Ca} = 0.79 \times 10^{-9}$ m$^2$s$^{-1}$, respectively. The domain length is $L_d = 100$ Å, the domain radius is $L_r = 100$ Å and the grid size is $h = 0.5$ Å. A voltage $V = 0$–75 mV was applied between the left and right domain boundaries.

2.2. Self-consistent electrostatics of the ion channel

The electrostatic field was derived by numerical solution of Poisson’s equation:

$$-\nabla (\varepsilon \nabla u) = \sum \frac{e z_i n_i}{\varepsilon_0},$$

(1)

where $\varepsilon_0$ is the dielectric permittivity of vacuum, $\varepsilon$ is the dielectric permittivity of the medium (water or protein), $u$ is the electric potential, $e$ is the elementary charge, $z_i$ is the charge number and $n_i$ is the number density of ions.

Self-consistent electrostatic potentials and forces are calculated numerically with respect to the ion-ion interaction, channel geometry (figure 1), self-potential barrier of the dielectric boundary force [23], value of $Q_f$ and value of applied potential $V$ at each simulation step. We used a finite volume Poisson solver specially designed for use within the ion channel geometry where there are severe jumps in dielectric permittivity. The linearity of Poisson’s equation together with the superposition principle allow us to use pre-calculated lookup tables for the field components, thereby dramatically decreasing the computation time of BD simulation steps [27].

Our reduced model of the SF takes both water and protein to be homogeneous continua with dielectric constants $\varepsilon_w = 80$ and $\varepsilon_p = 2$, respectively, together with a primitive model of ion hydration (the validity of this latter approximation is discussed below). The self-consistent electrostatics of a narrow, water-filled, channel in the protein wall differs significantly from bulk electrostatics, even when the dielectric constant of the water inside the channel is taken to be the same as in the bulk [28]. The huge gradient between $\varepsilon_w = 80$ and $\varepsilon_p = 2$ results in the quasi-1D axial behaviour of the electrostatic field, and hence in single-file movement of positive ions inside the channel [25]. This effect can be interpreted as electrostatic amplification of the electric field inside the channel [25, 29]. The electrostatics of the ion channel also prohibits the entrance of any negatively charged ions due to combined influences of the dielectric boundary force and the interaction with the fixed charge [23]. Consequently, we use a 1D dynamic model to simulate the axial, single-file, movement of cations (only) inside the SF and in its close vicinity.
This generic model of the ion channel can be used to describe the SF of both calcium and sodium channels, depending on the selected values of \( R, L \) and \( Q_f \) [3, 5]. A similar model of the Gramicidin A channel is described in [30, 31].

2.3. Brownian dynamics simulations

The BD simulations are based on numerical solution of the 1D over-damped Langevin equation:

\[
x' = -D \mu x + \sqrt{2D} \xi(t),
\]

where \( x \) stands for the ion’s position, \( D \) is its diffusion coefficient, \( \mu \) is the self-consistent potential in \( k_BT/e \) units and \( \xi(t) \) is normalized white noise. The numerical solution of (2) has been implemented with the Euler forward scheme.

We use an ion injection scheme that allows us to avoid heavy-duty simulation of ionic movements in the bulk liquid. Ions are injected randomly into the vicinity of the left channel entrance at an arrival rate that simulates the diffusive ionic flux from the undisturbed bulk. The model includes a hemisphere of radius \( R_a \) at each entrance representing the boundary between the channel vicinity and the baths (we take \( R_a = R \) here). The arrival rate \( j_{arr} \) is connected to the bulk concentration \( C \) through the Smoluchowski diffusion rate: \( j_{arr} = 2\pi DR_aC \) [32].

The motion of each injected ion is simulated in accordance with (2) until it reaches a domain boundary, where it is assumed to be absorbed. Simulations continue until the chosen simulation time has been reached. The ionic current \( J \) is calculated as the averaged difference between the numbers of similar ions passing the central cross-section of the SF per second in the forward and reverse direction [17].

A number of quantities is measured during the simulations, including the sodium \( J_{Na} \) and calcium \( J_{Ca} \) ion currents, the partial ionic occupancy profiles \( P_x(x) \) along \( x \) for different concentrations, and the partial \( P_{Na} \) and \( P_{Ca} \) occupancies, in each case as functions of the respective concentrations of calcium [\( \text{Ca} \)] or sodium [\( \text{Na} \)].

The BD simulations of ion current \( J \) and occupancy \( P \) were performed separately for \( \text{CaCl}_2 \) and \( \text{NaCl} \) solutions, and also for a mixed-salt configuration, with concentrations [\( \text{Na} \)] = 30 mM and 20 \( \mu \)M \( \leq \) [\( \text{Ca} \)] \( \leq \) 80 mM. The value of \( Q_f \) was varied within the range 0–6.5e in order to cover the known variants of sodium and calcium channels [5].

2.4. Model validity and limitations

The validity of continuum electrostatics and Langevin dynamics for a narrow water-filled pore is, of course, a highly significant question in relation to simplified models of the kind discussed [33–35]. The effective screening inside the long, narrow, SF could be very different from that assumed for \( \varepsilon_w = 80 \) and the whole notion of a dielectric constant inside a cylinder with \( R = 3.0 \) Å with \( L = 16 \) Å could become ill-defined [34].

The validity of both the electrostatics and the dynamics depends on the degree of dehydration of the ion inside the channel, so it can be defined roughly by the relationship between the channel radius \( R \) and the radius of the ion’s first hydration shell \( R_h \) [28]. Continuum electrostatics and dynamics generally fail when \( R_h > R \), but still can be applied for \( R_h \approx R \) provided that one uses effective values of \( \varepsilon_w \) and the diffusion coefficients \( D_{Na}, D_{Ca} \) that are all dependent on \( R \) [28]. We estimate \( R_h \approx 3.5 \) Å for \( \text{Na}^+ \) and \( \text{Ca}^{2+} \) ions so...
the calcium channel with $R \approx 3$ Å [1] does provide some room for Na$^+$ and Ca$^{2+}$ ions to carry water molecules. Both ions are still partially hydrated, therefore, and the continuum approximation with effective values can be used inside the SF. It is shown in [28] that the effective $\varepsilon_w$ saturates to its bulk value $\varepsilon_w = 80$ for $R \approx 3.5$ Å (roughly corresponding to $R_{H}$) and is still close to it ($\varepsilon_w \approx 70$) for $R = 3$ Å. This allows us to use bulk value for $\varepsilon_w$. The effective values of the ionic diffusion coefficients also decreased significantly with decreasing $R$ in comparison with their bulk values, and are estimated as $D \sim 0.25 D_{\text{bulk}}$ for $R = 3$ Å [34].

In this research we therefore use the bulk values of $\varepsilon_w$ and $D$ as their effective values throughout the whole computational domain, including the SF, a choice that avoids the use of additional fitting parameters. This reduced model obviously represents a considerable simplification of the actual electrostatics and dynamics of moving ions and water molecules in single-file within the narrow SF [34, 36]. Nonetheless, its applicability is supported by its reproduction of the experimentally observed AMFE kinetics for the calcium channel (see figures 4 and 5).

We have performed a parametric study of the stability and variability of the simulation results to changes of the effective model parameters: the length $L$ and radius $R$ of the SF, the length $H$ of the charged ring, and the applied voltage $V$ (see the appendix). We show that the simulation results are relatively insensitive to changes in the model parameters and that the main result—the distinct conductivity and selectivity bands—exists within a range of parameter values. This stability implicitly confirms that the reported phenomena arise from the basic electrostatics of real ion channels rather than as computational artefacts of our model related to some particular values of parameters.

We note that the recent determination of the crystal structure of the sodium NavAb channel [6] provides for well-hydrated motion of ions and has a rather asymmetric mode of coordination by four residues [8]. Although this structure refers to a bacterial channel, and not a DEKA-motif channel, it may nonetheless be relevant to the Na- and Ca-selective sites in the channels under consideration. If this turns out to be the case, the model presented here, which assumes a symmetrical arrangement of fixed charges and ions inside the channel, can with advantage be modified accordingly.

3. Results and discussion

3.1. Conduction and selectivity bands of the calcium channel

Figures 2(a) and (b) demonstrate our main result - the appearance of regular structure in the Ca$^{2+}$ ion current $J$ as a function of $Q_f$ and $[Ca]$, comprising areas of high conductance (conduction bands) M0, M1, M2 separated by almost zero-conductance stop-bands. The peak separation $\Delta Q \approx 2e$ corresponds roughly to the charge on one Ca$^{2+}$ ion. Band M1 coincides with the $J$ peak from [2] which was obtained for a negative charge of $1.3 \times 10^{-19}$ C/glutamate residue giving a total charge of $Q_f = 4 \times 1.3 \times 10^{-19}$ C $\approx 3.2e$ (the authors of [2] used four residues to model the EEEE locus of an L-type calcium channel).

The Ca$^{2+}$ occupancy $P$ exhibits step-wise growth with increasing $Q_f$ (figure 2(c)). The flat steps correspond to non-conducting saturated states with $P = 0, 1, 2 \ldots$ where the potential well at the binding site is too deep to allow escape of a Ca$^{2+}$ ion, and too shallow to allow the next Ca$^{2+}$ ion to enter the SF and push the bound ion(s) out. The conduction bands M0, M1 and M2 correspond to transitions in $P$: $0 \rightarrow 1$, $1 \rightarrow 2$, and $2 \rightarrow 3$, respectively. This picture corresponds to the ‘knock-on’ mechanism of Ca$^{2+}$ conductance and selectivity [1, 2]. It has an obvious analogue in semiconductor physics, where conduction also occurs in partially filled bands [37].

The appearance of the distinct conduction bands is caused by the discreteness of the multi-ion occupancy $P$. Their existence just for Ca$^{2+}$ in the calcium channel relates to the high $Q_f$ and double-valence of Ca$^{2+}$, enhancing the electrostatic effects of valence selectivity [13].

The effective parameter values in our model ($R, L$) can be varied within restricted ranges to fit experimental data for real channels [2, 3]. Our parametric study (see appendix) shows that the simulated bands are only weakly sensitive to variations of $R$ and $L$ in the ranges: $R = 2.5$–3.5 Å, $L = 12$–20 Å. Further decrease of $L$, or increase of $R$, leads to a weakening of the bands and to their eventual disappearance (at $L = 8$ Å for $R = 3$ Å). Their disappearance for shorter SFs may be related to the existence of a minimum $L$ needed to hold two Ca$^{2+}$ ions in the SF against their mutual Coulomb repulsion.

Mixed salt simulations (figure 3(a)) show that the M1 and M2 $J_{Ca}$ peaks are decreased and shifted to the beginning of each transition regions in $P$, as compared to the corresponding peaks for a pure Ca$^{2+}$ bath, due to attenuation by Na$^+$ ions [2]. Figure 3(b) shows that the selectivity ratio $R_I = (J_{Ca}/J_{Na})$ peaks at M1 and M2, with $R_I \approx 130$ for the M1 conduction band, and that there is no selectivity outside these bands. The $J$ versus $P$
provides the highly selective self-sustained Ca2+ channel and reveals saturation of the selectivity peaks M1 and M2 bands. (*the Ca2+ concentration is constant. The selectivity bands M1, M2 show saturated conductance at nearly constant P.*

**Figure 3.** Conduction and selectivity bands for a Na+ = 30 mM, Ca2+ = 40 mM mixed salt bath. (a) Currents J versus fixed charge Qf. Curves: 1—Ca, 2—Na, 3—Ca for a pure bath (reference curve from figure 2). (b) The selectivity ratio Rs = JCa/JNa, exhibits sharp peaks for the M1 and M2 bands. (c) JCa versus Ca2+ occupancy P. The selectivity peaks M1, M2 show saturated conductance at nearly constant P.

Plots (figure 3(c)) are each drawn as a J versus [Ca] simulation at constant Qf. They show that, at P = 1 for the M1 mode, and P = 2 for M2, P remains almost constant (saturated) while J increases very rapidly. The saturated conductance appears only for distinct values of Qf where the output barrier for the bound Ca2+ ion falls close to zero due to Coulomb repulsion between the Ca2+ ions, leading to barrier-less conduction [17, 22]. The arriving ion instantly pushes out and replaces the bound ion, so that P remains constant [17]. The saturated conduction provides the highly selective self-sustained Ca2+ flux in a mixed salt bath with a permanently Ca2+-occupied channel that is blocked for Na+ ions [2].

### 3.2. AMFE kinetics and identification of conduction bands

Because the ion channels in living cells are designed to conduct particular ions selectively with high rates, we expect wild-type channels to correspond to one of the highly conductive or highly selective bands: M0 (non-selective), M1 or M2 (calcium selective). We try to identify these bands with known wild-type channels by comparing their simulated and measured conductivity and selectivity properties (see table 1).

The conductivity band M0 at Qf \(\approx 1e\) provides non-selective conductance for both Ca2+ and Na+ ions and does not exhibit mutual blockade in AMFE simulations. Such properties could correspond to a non-selective cation channel [38] or the OmpF channel with Qf \(\approx 1e\) [11].

The highly calcium selective bands M1 and M2 correspond to different modifications of the calcium channel. We now argue that, based on their Qf values and AMFE properties, the M1 and M2 bands correspond, respectively, to the L-type and RyR calcium channels.

**Figure 4.** AMFE in a mixed salt bath for the M1 channel. (a), (b) Sodium (blue, point-down, triangles) and calcium (red, point-up, triangles) currents J and occupancies P versus Ca2+ concentration [Ca] in the highly selective M1 channel for [Na+] = 30 mM. The lines are guides to the eye. M1 shows strong blockade and AMFE at P_{Ca} = 1 with a threshold of [Ca]_{50} \approx 30 \mu M. (c) Mutual occupancy profiles for Na (left, blue, curve) and Ca (right, red, curve) ions show blockade of Na ions by the first Ca ion.

Figure 4 presents the dependencies of J and P on [Ca] for the M1 band in a mixed salt configuration. As shown in (a), (b), the M1 band with Qf \(\approx 3e\) shows a strong blockade of the current J_{Na} of Na+ ions with a blockade onset at [Ca]_{50} \approx 30 \mu M. The blockade occurs after the first Ca2+ ion has occupied the SF (figure 4(c)): P_{Ca} \rightarrow 1. The strong blockade with relatively low onset agrees qualitatively with the observed properties of the L-type channel [1]. The value of Qf and the conduction mechanism for M1 also correspond to the model [2] of the L-type channel (EEE locus).

**Figure 5.** AMFE for a mixed salt bath for the M2 channel. (a), (b) Sodium (blue, point-down, triangles) and calcium (red, point-up, triangles) currents J and occupancies P versus Ca2+ concentration [Ca] in-selective M2 channels for [Na+] = 30 mM. The lines are guides to the eye. M2 shows strong blockade and AMFE at P_{Ca} = 1 with a threshold of [Ca]_{50} \approx 200 \mu M. (c) Mutual occupancy profiles for Na (left, blue, curve) and Ca (right, red, curve) ions show blockade of Na ions by two Ca ions.
shows that the $\text{Na}^+$ ion is blocked by two $\text{Ca}^{2+}$ ions bound to the SF and separated by ion-ion repulsion. The M2 channel also shows a larger calcium current than M1. The high blockade offset and high calcium current, together with the higher value of $Q_f$, can be matched to the RyR channel (DDDD locus) [4].

The identification results are collected in table 1.

<table>
<thead>
<tr>
<th>Conduction band</th>
<th>Fixed protein charge</th>
<th>Na conductivity</th>
<th>Ca conductivity</th>
<th>Blockade of Na current by Ca ions</th>
<th>AMFE</th>
<th>Channel(s)</th>
<th>Residues locus</th>
</tr>
</thead>
<tbody>
<tr>
<td>M0</td>
<td>$\approx 1e$</td>
<td>High</td>
<td>High</td>
<td>No blockade</td>
<td>No AMFE</td>
<td>Non-selective cation channel [38], OmpF channel [11]</td>
<td>Non-selective cation locus</td>
</tr>
<tr>
<td>M1</td>
<td>$\approx 3e$</td>
<td>High</td>
<td>High</td>
<td>Yes, sodium ion is blocked by one calcium ion, low blockade offset (30 $\mu$M)</td>
<td>Yes, blockade is followed by moderate calcium current</td>
<td>L-type Calcium channel [1]</td>
<td>EEEE</td>
</tr>
<tr>
<td>M2</td>
<td>$\approx 5e$</td>
<td>High</td>
<td>High</td>
<td>Yes, sodium ion is blocked by two calcium ions, higher blockade offset (200 $\mu$M)</td>
<td>Yes, blockade is followed by strong calcium current</td>
<td>RyR Channel [4]</td>
<td>DDDD</td>
</tr>
</tbody>
</table>

4. Conclusions

In conclusion, our self-consistent BD simulations in a reduced model of a calcium channel SF have revealed distinct conduction bands M0, M1 and M2 as a function of the fixed charge $Q_f$ related respectively to integer values 0,1,2 of the occupancy $P$. The M0 band appears at $Q_f = 1e$ with $P = 1$; the M1 band appears at $Q_f = 3e$ with $P = 1$; and the M2 band appears at $Q_f = 5e$ with $P = 2$. The M0 band exhibits non-selective conductivity for both calcium and sodium ions, and can be identified with OmpF channel. The M1 and M2 bands show saturated self-sustained conductivity with high calcium selectivity ($R_s = J_{\text{Ca}}/J_{\text{Na}} \approx 130$ for the M1 band) and prominent AMFE, and can be identified with the L-type (EEEEE locus) and RyR (DDDD locus) calcium channels, respectively. The existence of the bands also provides a possible explanation for the results of mutation studies [9–11] in which a change in the fixed charge was found to destroy functionality or alter the type of selectivity.

Finally, we speculate that gating, and drug-induced blockade, may correspond to switching between a conduction band and a stop-band [20].

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Appendix. Parametric study

We have performed a parametric study of the stability and variability of the simulated band structure (see above) in the face of changes in the effective model parameters: the length $L$ and radius $R$ of the SF, the length $H$ of the charged ring, and the applied voltage $V$ (see figures A1–A4 below). In fact, the characteristic structure exists within a range of parameter values, as we now describe in more detail.

The length $L$ of the SF of calcium channels is usually estimated and modelled as being in the range $L = 5–15$ Å [2, 3] but the effective length could also depend on the geometry of the SF vestibules. We have performed BD simulations with $L$ varying from 8–20 Å to investigate the parametric dependence of the conductance bands on $L$, as shown in figure A1. An increase in $L$ leads to a sharpening of the band structure and to decreasing conductance. A decrease in $L$ leads to the flattening and eventual disappearance of the band structure (at $L < = 10$ Å for $R = 3$ Å) combined with an increase in conductance.

The effect of varying $R$, shown in figure A2 reveals a significant decrease of selectivity with increasing $R$, and vice versa. The band structure eventually disappears at $R = 4.5$ Å.
Figure A2. Calcium current $J$ versus fixed charge $Q_f$ showing how the band structure changes with the radius of the SF in the range $R = 2.5–4.5$ Å. An increase of $R$ to 3–4 Å leads to a decrease in the bands’ contrast, but the general pattern of the bands is still visible. A further increase of $R$ to 4.5 Å destroys the band structure.

Figure A3. Calcium current $J$ versus fixed charge $Q_f$ showing how the band structure varies with the length of the charged cylindrical ring in the range $H = 0.5–8$ Å. The band structure is clearly not very sensitive to $H$.

Figure A4. Calcium current $J$ versus fixed charge $Q_f$ showing how the band structure depends on the membrane voltage $V = 0–75$ mV. It is evident that there is no significant variation with $V$.

References

[33] Roux B, Allen T, Bernèche S and Im W 2004 Theoretical and computational models of biological ion channels QRB 37 15–103