A model of the ventricular cardiac action potential. Depolarization, repolarization, and their interaction

CH Luo and Y Rudy

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A Model of the Ventricular Cardiac Action Potential
Depolarization, Repolarization, and Their Interaction
Ching-hsing Luo and Yoram Rudy

A mathematical model of the membrane action potential of the mammalian ventricular cell is introduced. The model is based, whenever possible, on recent single-cell and single-channel data and incorporates the possibility of changing extracellular potassium concentration [K]o. The fast sodium current, \( I_{Na} \), is characterized by fast upstroke velocity \( (V_{\text{max}}=400 \text{ V/sec}) \) and slow recovery from inactivation. The time-independent potassium current, \( I_{K1} \), includes a negative-slope phase and displays significant crossover phenomenon as \( [K]_o \) is varied. The time-dependent potassium current, \( I_K \), shows only a minimal degree of crossover. A novel potassium current that activates at plateau potentials is included in the model. The simulated action potential duplicates the experimentally observed effects of changes in \( [K]_o \) on action potential duration and rest potential. Physiological simulations focus on the interaction between depolarization and repolarization (i.e., premature stimulation). Results demonstrate the importance of the slow recovery of \( I_{Na} \) in determining the response of the cell. Simulated responses to periodic stimulation include monotonic Wenckebach patterns and alternans at normal \( [K]_o \), whereas at low \( [K]_o \), nonmonotonic Wenckebach periodicities, aperiodic patterns, and enhanced supernormal excitability that results in unstable responses (“chaotic activity”) are observed. The results are consistent with recent experimental observations, and the model simulations relate these phenomena to the underlying ionic channel kinetics. (Circulation Research 1991;68:1501–1526)

In the late 1970s, two models of the electrical activity of cardiac cells were formulated based on the formalism introduced by Hodgkin and Huxley.1 McAllister et al2 developed a model of the cardiac Purkinje fiber action potential. Subsequently, Beeler and Reuter3 published a model of the electrical activity of the mammalian ventricular myocyte (referred to as the B-R model in the present paper). The B-R model was based on experimental data that were available at the time from voltage-clamp studies. These data were subject to limitations in available voltage-clamp techniques and their application to multicellular preparations of cardiac muscle.4 In addition, the concentrations of ions in the extracellular clefts of the multicellular preparations were unknown.

With the development of single-cell and single-channel recording techniques in the 1980s, the limitations of voltage-clamp measurements were overcome and the intracellular and extracellular ionic environments could be controlled. The data from single-channel recordings provide the basis for a quantitative description of channel kinetics and membrane ionic currents. In 1985, DiFrancesco and Noble5 developed a model of the Purkinje fiber action potential based on available single-cell and single-channel data. Our goal is to incorporate, whenever possible, recent experimental information that have accumulated since the formation of the B-R model into the formulation of a modified model (referred to as the L-R model in the text) of the mammalian ventricular action potential. The work presented here constitutes the first phase of this effort. In this paper we formulate the fast inward sodium current and the outward potassium currents. We use the model to investigate phenomena that are dominated by these currents and are only minimally influenced by the slow inward current. Therefore, we retain the B-R formulation of the slow inward current to support the plateau of the action potential.

This paper focuses on the depolarization and repolarization phases of the action potential and on phenomena that involve interaction between these processes. These include supernormal excitability,
Wenckebach periodicity, and aperiodic response of the cell to periodic stimulation. These phenomena were observed recently in single ventricular cells and in Purkinje fibers.5,7 In the B-R model, the extracellular concentrations of ions are fixed. Because changes in extracellular potassium concentration exert a strong effect on the time course of repolarization,8,9 we introduce in the model the dependence of the potassium currents on potassium concentration. We also incorporate a negative-slope characteristic of the time-independent potassium current10,11 and a novel potassium channel that activates at plateau potentials.12 The fast sodium current is characterized by fast upstroke velocity \((V_{\text{max}})^{13}\) and slow recovery from inactivation,14 a property that strongly influences the response of the cell to premature stimulation.

In addition to the study of mechanisms that determine the behavior of the single cell, an accurate model of the action potential is important to simulation studies of propagation of excitation in cardiac tissue. Our studies of propagation15,16 were limited by the inability to simulate important situations of physiological and clinical significance such as the effects of elevated extracellular potassium concentration, an important aspect of ischemia. In addition, our model simulations of reentry17 demonstrated the importance of the interaction between depolarization and repolarization in the induction and maintenance of reentrant arrhythmias. The need for an accurate representation of this interaction to further elucidate mechanisms underlying abnormal propagation and arrhythmogenesis provided yet another motivation for the development of the action potential model presented here.

Methods

The general approach is based on a numerical reconstruction of the ventricular action potential by using Hodgkin-Huxley-type formalism.1 The rate of change of membrane potential \(V\) is given by

\[
dV/dt = -(1/C) (I + I_{\text{Na}}) \tag{1}
\]

where \(C\) is the membrane capacitance, \(I_{\text{Na}}\) is a stimulus current, and \(I\) is the sum of six ionic currents: 
- \(I_{\text{Na}}\), a fast sodium current;
- \(I_{\text{K1}}\), a time-dependent potassium current;
- \(I_{\text{Kt}}\), a time-independent potassium current;
- \(I_{\text{Kpp}}\), a plateau potassium current; and
- \(I_{\text{L}}\), a time-independent background current. The ionic currents are determined by ionic gates, whose gating variables are obtained as a solution to a coupled system of eight nonlinear ordinary differential equations. The ionic currents, in turn, change \(V\), which subsequently affects the ionic gates and currents. The differential equations are of the form

\[
dy/dt = (y_s - y)/\tau_y \tag{2}
\]

where

\[
\tau_y = 1/(\alpha_y + \beta_y)
\]

and

\[y_s = \alpha_y/(\alpha_y + \beta_y)
\]

\(y_s\) represents any gating variable, \(\tau_y\) is its time constant, and \(y_s\) is the steady-state value of \(y\). \(\alpha_y\) and \(\beta_y\) are voltage-dependent rate constants. In addition, \(\alpha_{K1}\) and \(\beta_{K1}\) of the \(I_{K1}\) channel depend on extracellular potassium concentration.

The integration algorithm used to solve the differential equations is based on the hybrid methods introduced by Rushton Larsen18 and Victorri et al.19 Briefly, the algorithm uses an adaptive time step that is always smaller than \(\Delta t_{\text{max}} = 1\) msec. For time intervals of relatively slow changes in \(V\) \((\Delta V \leq \Delta V_{\text{min}} = 0.2\) mV), \(\Delta t\) is set equal to \(\Delta V_{\text{max}}/V\), where \(\Delta V_{\text{max}} = 0.8\) mV. For time intervals of fast changes in \(V\) \((\Delta V > \Delta V_{\text{max}})\), \(\Delta t\) is set equal to \(\Delta V_{\text{min}}/V\). If this \(\Delta t\) results in \(\Delta V > \Delta V_{\text{max}}\), \(\Delta t\) is reduced until the condition \(\Delta V < \Delta V_{\text{max}}\) is met. During the stimulus, a fixed time step (0.05 or 0.01 msec) is used to minimize variability in the stimulus duration caused by the time discretization procedure.20

Rate constants of ionic gates were obtained by parameter estimation with an adaptive nonlinear least-squares algorithm developed by Dennis et al.21 All computer programs were coded in FORTAN 77 (Microsoft, Seattle), and all simulations were implemented (double precision) on a Macintosh IICx computer.

Formulation of Equations for Ionic Currents

All ionic currents are computed for 1 cm² of membrane. Membrane capacity is set at 1 \(\mu\)F/cm².22 The formulation is based on experimental data adjusted to 37°C by using a \(Q_{10}\) adjustment factor. Ionic concentrations for standard preparations are \([K]\) = 5.4 mM, \([K]\) = 145 mM, \([Na]\) = 18 mM,13,23 \([Na]\) = 140 mM, and \([Ca]\) = 1.8 mM. \([Ca]\) varies during the action potential; we set \([Ca]\) = 2 \times 10⁻⁷ mM as an initial value under standard conditions. We assume that a short-term stimulation does not appreciably affect the ionic environment of the cell under normal conditions and, therefore, the ionic concentrations (except \([Ca]\)) do not change dynamically in our simulations. The complete set of equations for all ionic currents is provided in Table 1.

\(I_{\text{Na}}\): Fast sodium current. The model of the fast sodium channel incorporates both a slow process of recovery from inactivation and adequate maximum conductance that results in a realistic rate of membrane depolarization. We adopt the activation \((m)\) and inactivation \((h)\) parameters of Ebihara and Johnson4 (we refer to the Ebihara-Johnson model of \(I_{\text{Na}}\) as the E-J model). The formulation of these parameters is based on data from cardiac cells (chicken embryo) and results in a realistic rate of depolarization \((V_{\text{max}} = 300\) V/sec). However, it does not include the property of slow recovery. Following the methods of Beeler and Reuter,3 we incorporate a slow inactivation gate \((j)\) to represent this slow process. As suggested by Haas et al.24 the steady-
TABLE 1. Formulations of Ionic Currents

**Inward currents**

Fast sodium current

\[ I_{Na} = 23 \cdot m^3 \cdot h \cdot j \cdot (V - E_{Na}) \]

For \( V \leq -40 \text{ mV} \)

\[ a_0 = a_0 = 0, \quad b_0 = 1/(0.13[1 + \text{exp}(V + 10.66) - 11.1]) \]

\[ b_0 = 0.3 \cdot \text{exp}(-2.535 \cdot 10^{-7} V)/(1 + \text{exp}(-0.1(V + 32))] \]

For \( V < -40 \text{ mV} \)

\[ a_0 = 0.135 \cdot \text{exp}(80 + V)/\text{exp}(80) = 6.8, \quad b_0 = 3.56 \cdot \text{exp}(0.079V) + 3.1 \cdot 10^{4} \cdot \text{exp}(0.35V) \]

\[ a_0 = -1.2714 \cdot 10^{-3} \cdot \text{exp}(0.244V) - 3.474 \cdot 10^{-3} \cdot \text{exp}(-0.04391V) \cdot (V + 37.78)/(1 + \text{exp}[0.311 \cdot (V + 79.23)]) \]

\[ b_0 = 0.1212 \cdot \text{exp}(-0.01052V)/(1 + \text{exp}(-0.1378(V + 40.14))] \]

For all range of \( V \)

\[ a_0 = 0.32(V + 47.13)/(1 - \text{exp}(-0.1(V + 47.13))), \quad b_0 = 0.08 \cdot \text{exp}(V/11) \]

Slow inward current

\[ I_s = 0.09 \cdot d \cdot f \cdot (V - E_{Na}), \quad E_s = 7.7 - 13.0287 \cdot \ln([Ca]) \]

\[ a_0 = 0.095 \cdot \text{exp}[-0.01(V - 5)]/(1 + \text{exp}[-0.072(V - 5)]) \]

\[ b_0 = 0.07 \cdot \text{exp}[-0.017(V + 44)]/(1 + \text{exp}[0.05(V + 44)]) \]

\[ a_0 = 0.012 \cdot \text{exp}[-0.008(V + 28)]/(1 + \text{exp}[0.15(V + 28)]) \]

\[ b_0 = 0.0065 \cdot \text{exp}[-0.02(V + 30)]/(1 + \text{exp}[-0.2(V + 30)]) \]

Calcium uptake: \( d([Ca]) \cdot \text{d}t = -10^{-4} \cdot I_s + 0.07(10^{-4} - [Ca]) \)

**Outward currents**

Time-dependent potassium current

\[ I_{K} = G_{K} \cdot X \cdot X_{i} \cdot (V - E_{K}), \quad G_{K} = 0.282 \cdot \sqrt{[K]/5.4} \]

\[ X_{i} = \text{exp}(0.04(V + 77))/1 \cdot ((V + 77) \cdot \text{exp}(0.04(V + 35)) \]

\[ \alpha_{X} = 0.0005 \cdot \text{exp}(0.083(V + 50))/(1 + \text{exp}(0.057(V + 50))) \]

\[ \beta_{X} = 0.0013 \cdot \text{exp}[-0.06(V + 20)]/(1 + \text{exp}[-0.04(V + 20)]) \]

Time-independent potassium current

\[ I_{K1} = G_{K1} \cdot X_{1} \cdot (V - E_{K1}), \quad G_{K1} = 0.6047 \cdot \sqrt{[K]/5.4} \]

\[ \alpha_{K1} = 1.02/[1 + \text{exp}(0.2385 \cdot (V - E_{K1} - 59.215))] \]

\[ \beta_{X1} = 0.49124 \cdot \text{exp}(0.08032 \cdot (V - E_{K1} + 5.476) + \text{exp}(0.06175 \cdot (V - E_{K1} - 59.431))] \]

Plateau potassium current

\[ I_{Kp} = 0.0183 \cdot K_p \cdot (V - E_{Kp}), \quad E_{Kp} = E_{K1} \]

\[ K_p = 1/[1 + \text{exp}(7.488 - V)/5.98)] \]

Background current

\[ I_b = 0.03921 \cdot (V + 59.87) \]

Total time-independent potassium current

\[ I_{K1,T} = I_{K1} + I_{Kp} + I_b \]

state values of \( j \) (\( j \)) are obtained by setting \( j = h_{\text{sag}} \), where \( h_{\text{sag}} \) is from the E-J model. The time constant of \( j \) (\( \tau \)) is set equal to the \( \tau \) of the B-R model. The rate constants \( \alpha \) and \( \beta \) are obtained by using the parameter estimation procedure mentioned above.\( ^{21} \) The sodium current is

\[ I_{Na} = G_{Na} \cdot m^3 \cdot h \cdot j \cdot (V - E_{Na}) \quad (3) \]

where \( G_{Na} \) is the maximum conductance of the sodium channel (23 mS/cm\(^2\))\(^{4,25} \); \( E_{Na} \) is the reversal potential of sodium \[ E_{Na} = (RT/F) \cdot \ln ([Na_{i,j}]/[Na]) \]; and \( m, h, j \), and \( \alpha \) are obtained as solutions to Equation 2 with the appropriate rate constants. Note that \( E_{Na} \) computed with the E-J model is 29 mV since \([Na_{i,j}] = 40 \text{ mM} \) in chicken embryo heart cells.\(^{25} \) In our model we set \( E_{Na} = 54.4 \text{ mV} \) based on \([Na_{i,j}] = 18 \text{ mM} \) in mammalian ventricular cells.\(^{13,23} \)

\( I_{Na} \): Slow inward current. Representation of \( I_{Na} \) is the same as in the B-R model. The formulation is provided in Table 1.
\[ I_K = G_K \cdot X \cdot X_i \cdot (V - E_K) \] (4)

and introduce the \([K]\), dependence through

\[ \bar{G}_K = 0.282 \cdot \sqrt{[K]/5.4} \]

and

\[ E_K = (RT/F) \cdot \ln \left( \frac{[K]_0 + PR_{NaK} \cdot [Na]_0}{[K]_0 + PR_{NaK} \cdot [Na]_0} \right) \]

where \( PR_{NaK} = 0.01833 \) is the \( Na/K \) permeability ratio.\(^27,28\) For this value of \( PR_{NaK} \) and \([K]_0 = 5.4 \) mM, the computed \( E_K \) is \(-77 \) mV, a result that is consistent with the measurements of Beeler and Reuter.\(^3\) Also, note that \( G_K = 0.282 \) mS/cm\(^2\) for \([K]_0 = 5.4 \) mM. This value is obtained from the fully activated current \( I_K = I_0 / X \) (B-R model,\(^3\) Figure 1) for \( V = -100 \) mV, a potential at which \( X_i = 1 \).

\( X_i \) introduces the inward rectification property of \( I_K \). It is obtained from the B-R expression for \( I_K \) by factoring out \( G_K \cdot (V - E_K) \) at \([K]_0 = 5.4 \) mM. The formulation is provided in Table 1. To verify that this formulation correctly introduces the \([K]\), dependence of \( I_K \) we plotted the computed fully activated current \( I_K \) for \([K]_0 = 5.4, 50, \) and 150 mM (Figure 1). The behavior is consistent with the experimental observations of McDonald and Trautwein\(^29\), Matsura et al.,\(^27\) and Shibasaki.\(^20\) Note the strong inward rectification and the minimal crossover between curves of different \([K]\).

**\( I_{K1} \): Time-independent potassium current.** Sakmann and Trube\(^10,11\) using patch-clamp techniques, demonstrated two important properties of \( I_{K1} \): 1) square

![Figure 1](image1.png)

**Figure 1.** Current-voltage (I-V) curves of the fully activated time-dependent potassium current \( I_K \) (current denoted \( I_K \) in the figure) for \([K]_0 = 5.4, 50, \) and 150 mM. Note minimal crossover between curves of different \([K]\), and strong inward rectification.

root dependence of single-channel conductance on \([K]\), 2) high selectivity for potassium (i.e., \( E_K \sim \text{Nernst potential of potassium} \)). Kurachi\(^30\) identified an inactivation gate \( (K1) \) of the \( I_{K1} \) channel. This gate, in addition to its dependence on membrane potential, depends on \( E_{K1} \) and therefore on \([K]_0 \). K1 closes at high potentials and therefore \( I_{K1} \) has no contribution at this range. In addition, the time constant of K1 is small (\( \tau_{K1} = 0.7 \) msec at \( V = -50 \) mV for \([K]_0 = 5.4 \) mM), so it can be approximated by K1, (its steady-state value).

Based on these findings, we formulate the \( I_{K1} \) current as follows:

\[ I_{K1} = \bar{G}_{K1} \cdot K1 \cdot (V - E_{K1}) \] (5)

where

\[ E_{K1} = (RT/F) \cdot \ln \left( \frac{[K]_0}{[K]_1} \right) \]

\[ K1 \cdot = \alpha_{K1} / (\alpha_{K1} + \beta_{K1}) \]

\[ \bar{G}_{K1} = 0.6047 \cdot \sqrt{[K]_0/5.4} \]

To verify that this formulation correctly introduces the \([K]\), dependence of \( I_{K1} \) we plotted the computed current-voltage curve for \([K]_0 = 10, 20, 40, 75, \) and 150 mM (Figure 2). The results resemble the single-channel measurements of Kurachi (see Figure 12B of Reference 30). Note the strong inward rectification, the crossover between curves of different \([K]\), the zero contribution at high potentials, and the negative slope over a certain potential range. These characteristics reflect the voltage and \([K]\), dependence of the K1 gate.

![Figure 2](image2.png)

**Figure 2.** Current-voltage (I-V) curves of the time-independent potassium current \( I_{K1} \) for different \([K]\), (indicated in the figure in mM). Note the large degree of crossover between curves of different \([K]\), strong inward rectification, negative slope of the I-V curves, and zero-current contribution at high potentials.
It should be emphasized that our definition of $I_{K1}$ differs from $I_{K1}$ of the B-R model. The B-R $I_{K1}$ includes contributions from other currents at plateau potentials. Recently, it was recognized that the current at plateau potentials results from a time-independent, $[K]_o$-insensitive channel ($I_{Kp}$)\(^2\) and a background current ($I_b$).\(^{32-34}\) We formulate $I_{Kp}$ to simulate the behavior of the $I_{Kp}$ current measured by Yue and Marban\(^{12}:\)

$$I_{Kp} = G_{Kp} \cdot Kp \cdot (V-E_{Kp})$$  \hspace{1cm} (6)

where $G_{Kp} = 0.0183$, $E_{Kp} = E_{K1}$, and $Kp = 1/[1+exp((7.488-V)/5.98)].$ In addition, $I_b$ is a background current\(^{32-34}\) that can be formulated as

$$I_b = G_b \cdot (V-E_b)$$  \hspace{1cm} (7)

where $G_b = 0.03921$ and $E_b = -59.87$ mV. We define the total time-independent potassium current, $I_{K1(T)}$, as

$$I_{K1(T)} = I_{K1} + I_{Kp} + I_b$$  \hspace{1cm} (8)

The parameters (conductances, gates, and reversal potentials) of $I_{K1}$, $I_{Kp}$, and $I_b$ are obtained by using a parameter estimation technique\(^{21}\) to fit the whole-cell $I_{K1(T)}$ measured by Sakmann and Trube (Figure 4A of Reference 11) for different values of $[K]_o$. A comparison of the measured and simulated $I_{K1(T)}$ is shown in Figure 3 for different values of $[K]_o$. Note that the negative slope (which is not a property of the B-R $I_{K1}$) and crossover phenomena are well duplicated by the model. Also, all curves for different values of $[K]_o$ converge at plateau potentials. This is because $I_{K1}$, which depends on $[K]_o$, has zero contribution at this potential range, while $I_{Kp}$ and $I_b$ are $[K]_o$ insensitive.

**Results**

**Ionic Currents and the Action Potential**

The goal of this paper is to investigate, at the ionic channel level, the mechanisms of various electrophysiological phenomena related to depolarization, repolarization, and their interaction. We limit the simulations to phenomena that are dominated by the fast inward sodium current and the outward potassium currents. Figure 4A shows simulated action potentials for different extracellular concentrations of potassium ($[K]_o$ = 3, 4, 5.4, and 7 mM). For the typical physiological concentration of potassium ($[K]_{o,normal} = 5.4$ mM), the following characteristics are observed: the threshold potential is at $-60$ mV, the membrane rate of depolarization ($V_{\text{max}}$) is $400$ V/sec, the maximum plateau potential is $17.7$ mV, action potential duration (APD) at 90% repolarization is $366$ msec, resting potential is $-84$ mV, and the overshoot potential is $41.7$ mV. Note that $V_{\text{max}}$ is more than three times that of the B-R model ($V_{\text{max,B-R}} = 115$ V/sec). The fast $V_{\text{max,L-R}}$ reflects the higher sodium channel conductance in our model and is in good agreement with experimental data from mammalian ventricular cells.\(^{13}\) Also, $V_{\text{max,L-R}}$ is faster than that of the E-J model ($V_{\text{max,E-J}} = 300$ V/sec). This difference results from different reversal potentials for the sodium current ($E_{Na,L-R} = 54.4$ mV), whereas

$E_{Na,E-J} = 29.0$ mV) and, as a result, greater driving force in the L-R model. The difference in the reversal potential reflects different intracellular concentrations of sodium in the mammalian ventricular cell ($[Na]_i = 18$ mM) and the chicken embryo ($[Na]_i = 40$ mM). The overshoot is in good agreement with experimental measurements in guinea pig ventricular cells.\(^{31,35}\)

The major simulated effects of changes in extracellular potassium concentration are on APD and rest potential ($V_{\text{rest}}$), whereas effects on plateau potentials are minimal. These results are in agreement with experimental observations.\(^{8,9}\) As $[K]_o$ increases, $V_{\text{rest}}$ becomes less negative and APD decreases. For $[K]_o = 3$ mM, $V_{\text{rest}}$ is $-95.5$ mV and APD is $485$ msec; for $[K]_o = 7$ mM, $V_{\text{rest}}$ is $-78.2$ mV and APD is $322.6$ msec. For the typical value $[K]_o = 5.4$ mM, APD is $366$ msec. This value is in the range of $305$ msec with a standard deviation of $55$ msec measured by Isenberg and Klöckner\(^{31}\) in the guinea pig single ventricular...
simulate various physiological phenomena and elucidate their mechanism.

Fast Depolarization and Slow Recovery of $I_{Na}$

Recovery of excitability. As discussed above, $I_{Na}$ in the L-R model is characterized by a larger conductance and a faster rate of depolarization (large $V_{max}$) than in the B-R model. Also, the L-R model incorporates a fast process of inactivation and a much slower process of recovery from inactivation. In this section we demonstrate the importance of the fast (realistic) rate of depolarization caused by the kinetics of $I_{Na}$ in determining membrane behavior. Recovery of excitability can be described in terms of $t_{U}$ and $t_{th}$ (defined below) using the protocol of Figure 5. Test pulses ($S_2$) 1.77 $\mu$A in amplitude and 100 msec in duration are applied at various ($S_1$, $S_2$) intervals from a brief suprathreshold stimulus ($S_0$), which excites a conditioning action potential. Because $S_1$, $S_2$ $>$ $APD_U$, $S_2$ is always applied during phase 4 of the action potential. The time from the onset of $S_2$ to threshold is $t_{th}$ ($V=V_{th} \sim -60$ mV). The time from the onset of $S_2$ to the action potential upstroke as determined from $V_{max}$ is $t_U$. As the $S_1$, $S_2$ interval decreases, $t_U$ and $t_{th}$ increase, providing a measure of the excitability of the membrane; the longer $t_U$ and $t_{th}$, the less excitable the membrane.

Figure 6 shows the relation between $t_U$ and $t_{th}$ obtained from the protocol of Figure 5. Panel A duplicates the experimental results of Delmar et al (Figure 5B in their paper). Panel B shows the simulated results of the same protocol using the B-R membrane model but with the sodium channel replaced by the L-R formulation. Panels C and D depict simulated results with the original E-J and B-R sodium channel representations, respectively.

The results demonstrate that the experimental behavior (panel A) is duplicated accurately by the L-R and E-J models (panels B and C). However, B-R kinetics of the sodium channel (panel D) causes
significant deviations from the actual experimental behavior. These include a strong deviation from the linear relation for larger $t_{th}$ (arrow) and a larger slope (≈2.5 times, note different $t_U$ scale in panel D) of the linear portion for smaller $t_{th}$. The slope is close to 1.0 for the L-R and E-J models, duplicating the experimental behavior well and indicating that the increase in $t_U$ is determined by the increase in $t_{th}$ (subthreshold response). This is because the latency from threshold potential to $V_{max}$ is very small (about 2 msec) when the realistic conduction and kinetics of the cardiac sodium channel are incorporated in the model ($G_{Na,L-R}=4$ mS, $G_{Na,B-R}=23$ mS). In contrast, the large slope of the B-R model indicates a significant latency from threshold potential to $V_{max}$ (greater than 25.5 msec), resulting from the unrealistic conduction and kinetics of the sodium channel in that model.

**Time course of reactivation.** The simulations in this section demonstrate the importance of the slow recovery property of $I_{Na}$ in determining the membrane reactivation. As mentioned in previous sections, the E-J model does not include the process of slow recovery from inactivation represented by the j gate. Panel A of Figure 7 duplicates the experimental reactivation data of Ebihara et al (Figure 3 of Reference 36). Panel B shows the results of a simulation with the L-R model. The protocol used both in the experiment and in the model simulations is also illustrated in panel B. Panels C and D are simulations with the B-R and E-J models, respectively. The protocol is as follows: two depolarizing voltage pulses with 45-mV amplitude are applied after preclamping the membrane at −65 mV for 300 msec. The first pulse of duration $T_1$ inactivates the sodium channel. $T_2$, the interval between the depolarizing pulses, is varied to provide different reactivation times before the second test pulse of a constant duration (65 msec) is applied. The membrane response to the second pulse in terms of the peak inward current is plotted as a function of $T_2$.

The experimental behavior (panel A) is well duplicated by the L-R model (panel B). The individual curves in panel A are obtained for $T_1$ in the range 1–200 msec, while in panel B the range is 1–50 msec. This quantitative difference probably reflects the different species (chicken embryo and guinea pig). The B-R model (panel C) deviates from the experimentally observed behavior, especially for $T_1=1$ msec. As explained below, this is caused by the unrealistic $I_{Na}$ kinetics of the B-R model. The E-J model (panel D) completely fails to duplicate the dependence on $T_2$, and the curves obtained for different $T_1$ values overlap.

The E-J model (panel D) fails to separate the curves for different $T_1$ values because slow recovery (j gate) is not part of this model, so that inactivation and reactivation of the sodium channel are controlled by the fast inactivation h gate. Because in this model h is almost completely inactivated in 1 msec when the membrane potential is depolarized to −20 mV ($\tau_h=0.33$ msec at −20 mV), all curves for $T_1\geq 1$ msec overlap. In contrast, the L-R model (panel B) generates different reactivation curves for different values of $T_1$, as observed experimentally. This is because the model incorporates a slow inactivation j gate ($\tau_j=4.34$ msec at −20 mV). This implies that for $T_1=1$ and 5 msec the j gate at the end of $T_1$ is inactivated to a different degree, resulting in different reactivation curves. For $T_1\geq 20$ msec, the j gate is completely inactivated at a membrane potential of −20 mV, and all the curves overlap. In addition, since for $T_1>1$ msec the recovery process is controlled by the j gate, all reactivation curves can be fit by a single exponential. In particular, for $T_1\geq 20$ msec all curves are represented by the same time constant and
FIGURE 7. Reactivation of the sodium channel. The protocol is illustrated in panel B. Peak inward current obtained during the second pulse is normalized by the maximum current measured during the first pulse and is plotted as a function of the interval between the depolarizing pulses ($T_2$). Panel A: Experimental results of Ebihara et al. for $T_1 = 1, 100,$ and 200 msec. Panel B: Present (L-R) model. Panel C: Beeler-Reuter (B-R) model. Panel D: Ebihara-Johnson (E-J) model. In each panel, the different curves are obtained for different values of $T_1 (1, 5, 20,$ and 50 msec). Note that all curves overlap in the E-J model simulation because of unrealistically fast recovery (slow inactivation $j$ gate is not included in this model).

FIGURE 8. Membrane responsiveness and sodium channel kinetics. Maximum upstroke velocity ($V_{max}$) values (normalized by its maximum value) are indicated by discrete points. Curves for the fast ($h$) and slow ($j$) inactivation gates (or only $h$ for the Ebihara-Johnson [E-J] model) are plotted as solid lines. Both are shown as a function of membrane potential ($V$). The protocols are illustrated in panel A and are described in the text. Panel A: Experimental data of Gettes and Reuter (Figure 7 of Reference 14). Panel B: Present (L-R) model. Panel C: Beeler-Reuter (B-R) model. Panel D: E-J model.
(τ = 70 msec), which is very close to the τ value at V = -65 mV (τ = 77.4 msec), as expected.

Membrane responsiveness. This simulation demonstrates the importance of the slow inactivation j gate in determining the membrane response during the repolarization phase of an action potential. Results are shown in Figure 8. Panel A is the experimental data of Gettes and Reuter (Figure 7 of Reference 14). Panels B, C, and D are the L-R, B-R, and E-J models, respectively. The protocols are illustrated in panel A. Curve 1 is obtained from protocol 1 by applying a suprathreshold stimulus that scans the repolarization phase of an action potential. Curve 2 is obtained from protocol 2 by stimulating the membrane after 500 msec of preclamping at different potential levels. For both curves Vmax is shown (discrete points) as a function of the membrane potential at the onset of the stimulus. In panels B–D the solid curves are h ∙ j (or only h for the E-J model) computed for the same protocols.

By using protocol 1, the range of membrane potential over which Vmax drops from its maximum value to 0.2 of the maximum is well simulated by the L-R and B-R models. This normalized value (0.2) is obtained experimentally (panel A, curve 1) for a membrane potential of -76 mV. The corresponding simulated values are -78 mV for both models. In contrast, curve 1 in the E-J model (panel D) is shifted to less negative potentials. This is because the E-J model does not incorporate a slow recovery process of the sodium channel. An equivalent statement is that in the E-J model j is always 1. However, the L-R and B-R models incorporate a slow recovery j gate that is not fully recovered during the repolarization phase, shifting the curve to a more negative range of membrane potentials.

Curve 2 reflects steady-state behavior of the membrane since the stimulus was applied after a long (500-msec) preconditioning period. The L-R model correctly simulates the behavior of Vmax for these conditions (compare curve 2 in panels B and A). In contrast, a shift of curve 2 to more negative potentials is observed in the B-R model (panel C). This difference reflects the different characteristics of h and j, in the two models.

Note that the Vmax behavior is closely approximated by the h ∙ j curves (deviations are caused by currents other than Im). This implies that the membrane responsiveness is controlled by the h and j gates (because of its fast time constant, m obtains its maximum value of 1 within 0.2 msec and its effect is negligible). Because curve 2 is obtained under steady-state conditions, it is closely approximated by h ∙ j. The difference between curves 1 and 2 results from the fact that during repolarization (curve 1) h and j do not reach their steady-state values. The E-J model does not incorporate a j gate. Therefore, the difference between curves 1 and 2 in panel D implies that the rate of repolarization is too fast for h to reach steady state. Of course, the same is true for the much slower j gate. Since during repolarization j is almost completely inactivated and since τj > τm, it follows that the membrane responsiveness is mostly determined by the j gate.

Supernormal Excitability

Supernormality, which can be defined as greater than normal excitability during or immediately after the action potential repolarization phase, is a known property of cardiac preparations at low extracellular potassium concentrations.737-39 The strength–interval curve, used to investigate supernormality, is obtained from the protocol shown in Figure 5 by applying a test pulse, S2, after an action potential excited by a stimulus, S1. By varying the S1S2 interval and the current amplitude of S2, the threshold current (Ith) is obtained as a function of the S1S2 interval. Based on such excitability measurements, Spear and Moore found that 1) supernormality disappears when [K]0 is elevated above 5 mM and 2) the supernormal period at fixed [K]0 is independent of the APD and is determined by the characteristics of the membrane potential during repolarization. Ith and Is of the L-R model can change their reversal potentials and conductances in response to variations in [K]0. We used the L-R model to simulate and investigate supernormality. Figure 9 shows the simulated strength–interval curves for [K]0 = 7, 4.6, 4, and 3 mM, covering late phase 3 and early phase 4 of the action potential. The simulated protocol followed the experimental protocol described above (also see Figure 5). Ith is defined as the critical current amplitude for which the peak sodium current is greater than 4 μA (1% of the maximum current for a fully recovered membrane and suprathreshold stimulus). Note that test stimuli of very short duration (T = 0.5 msec) were used to investigate the instantaneous membrane excitability at that S1S2 interval. The abscissa is normalized by setting the 70% repolarization of the action potential as time zero. Clearly, supernormality is observed for [K]0 ≤ 4.6 mM (but not for [K]0 = 7 or 5.4 mM) as a notch (local minimum followed by a local maximum) in the strength–interval curve. With decreasing [K]0, the peak-to-peak amplitude of the notch increases and so does the width of the “supernormal window.” This time window is defined as the interval during which the threshold is lower than the local maximum that follows the notch. The supernormal window is indicated in the figures by two broken vertical lines.

Plotting Ith versus the potential difference (∆V = Vth – V) for the membrane to reach the threshold potential (Vth), the linear relation of Figure 10 is obtained for all values of [K]0. This linear relation implies that the membrane response during repolarization can be characterized in terms of the membrane potential and the threshold potential for all values of [K]0. It also implies that the threshold current does not depend on APD, as observed experimentally by Spear and Moore.39

To explain the underlying mechanism of supernormality in terms of membrane channel kinetics, the
following simulation was carried out for [K]₀ = 7 mM. The j gate of the sodium channel was clamped at the value of 1 for all times and the strength–interval curve was computed (Figure 11A). The unclamped strength–interval curve is also plotted for comparison. With j=1, sodium channel recovery from inactivation is controlled by the h gate and its much faster kinetics (τₜ << τₗ). Under these conditions, supernormality is observed in Figure 11A even for [K]₀ = 7 mM. In contrast, when j is free to vary according to its normal kinetics, no supernormality is observed for this concentration of Kₜ, in agreement with the experimental behavior.²⁵ In Figure 11B, V and V₄₃ are plotted as a function of time during repolarization. The curve for V is the same for both j=1 and j-unclamped conditions (because the sodium channel does not contribute to V during repolarization). On the other hand, the V₄₃ curve is influenced by the j gate kinetics. For the clamped condition (j=1) V₄₃ decreases earlier in time, reflecting faster recovery of membrane excitability. Also, the value of V₄₃ just after the absolute refractory period (arrows) is larger, followed by a steep decrease (approximately 3.5 V/sec) over a period of 6 msec. During the same interval, the membrane potential drops at a rate of about 1.33 V/sec. This implies that ΔV (defined as V₄₃−V) is decreasing rapidly during this interval, bringing about a proportional decrease in I₃₆ (Figure 10) that constitutes the descending portion of the notch in the strength–interval curve (Figure 11A, DES). Beyond this interval, the membrane potential drops faster than V₄₃, increasing ΔV and, therefore, I₃₆ (ascending portion of the notch, ASC in Figure 11A). This nonmonotonic behavior of ΔV is the basis for the notch in the strength–interval curve and, hence, for supernormality. For larger values of t, both V₄₃ and V approach their steady-state values with ΔV (and hence I₃₆) approaching a constant. When the j gate is free to vary according to its normal kinetics (j-unclamped in Figure 11B), the V₄₃ curve is shifted to the right toward later times. As a result, the drop in the V₄₃ curve occurs over an interval during which V decreases slowly and, in particular, always slower than V₄₃ (0.5 and 1.4 V/sec, respectively). Consequently, ΔV (and I₃₆) is monotonically decreasing (j-unclamped in Figure 11A), and no supernormality is observed. It is clear that the nonmonotonic, supernormal behavior is determined by the shape of the V₄₃ curve, which in turn is determined by the recovery of the sodium channel. As will be explained below, the sodium channel recovers faster at low [K]₀.
creating the possibility of a supernormal phase when \([K_o]_{\text{s}}\leq 4.6\ \text{mM}\).

A clear demonstration of the dominant effect of the sodium recovery process on the \(V_{th}\) curve is provided in Figure 11B. When both the h gate and j gate are clamped at the value of 1, \(V_{th}\) is constant, except for a small decrease (for \(t<30\ \text{msec}\)) caused by \(I_d\) (this decrease in \(V_{th}\) disappears when we set \(I_d=0\) during the stimulation). The threshold potential is also affected by \(I_{K1(T)}\) and \(I_K\); however, the contribution is very small.

Figure 12 shows \(V\) and \(V_{th}\) as a function of time for different \([K_o]\), in the range of 3–7 mM (j is free to vary according to its normal kinetics). As \([K_o]\) decreases from 7 to 3 mM, the resting membrane potential changes from -78.2 to -95.5 mV (Figure 12A), resulting in a decrease of \(\tau_h\) from 33 to 7 msec during late phase 3 and phase 4 of the action potential (arrows in Figure 12A). Therefore, the sodium channel recovers earlier for lower \([K_o]\) (Figure 12B) during a phase of fast decrease of \(V\). As a result, a phase is created during which \(V\) decreases faster than \(V_{th}\). In addition, the membrane potential reaches a phase of very slow decrease (i.e., end of ascending portion of the supernormal notch) later in time for lower \([K_o]\) (arrows in Figure 12A). The result of these two effects is the appearance of a supernormal window for \([K_o]\leq 4.6\ \text{mM}\) and a widening of this window as \([K_o]\) is further decreased (Figure 9). The peak-to-peak amplitude of the supernormal notch also increases when \([K_o]\) is decreased. This is because the local minimum (see Figure 9) decreases, reflecting a smaller \(\Delta V\) caused by the faster decrease of \(V_{th}\) for lower \([K_o]\) (Figure 12B). Also, the local maximum following the notch increases with decreas-
ing \([K]_0\) reflecting a larger \(\Delta V\) caused by more negative rest potentials.

It should be emphasized that the E-J representation of the sodium channel does not include a slow inactivation \(j\) gate. Therefore, the E-J strength–interval curves are similar to the strength–interval curves obtained by the L-R model with \(j=1\). As shown above (Figure 11A), under these conditions, supernormality is a property of the model even for high extracellular concentrations \((|K|_0|=7 \text{ mM})\) in contradiction to experimental observations. This, of course, is a nonphysiological behavior that results from a nonphysiological fast recovery of the sodium channel.

The simulations described above were performed with a test stimulus, \(S_2\), of 0.5-msec duration. Experimentally, Chialvo et al.7 used a stimulus of 20 msec. Our choice of a short stimulus was dictated by the resolution needed for scanning the supernormal window whose minimum duration is 12 msec. Also, we were interested in determining instantaneous membrane excitability without the influence of a long stimulus duration. To investigate the effects of the stimulus duration \((T)\) on membrane excitability, we redefine the strength–interval curve in terms of threshold charge introduced by the stimulus \((Q_{th}=I_{th} \cdot T\), Figure 13). In Figure 13A, \(Q_{th}\) is plotted for an interval of 80 msec that includes the supernormal phase. Stimuli of two durations \((T=0.5 \text{ and } 20 \text{ msec})\) are compared. For \(T=20\) msec, \(Q_{th}\) is larger (note different scales), the amplitude of the supernormal notch is twice as large, and the notch is shifted toward earlier time. Figure 13B covers a later time interval that includes late phase 4 of the action potential. During this time interval, the behavior of \(Q_{th}\) for \(T=0.5\) and 20 msec is very different. For \(T=0.5\) msec, \(Q_{th}\) displays a very small and slow increase, while for \(T=20\) msec, \(Q_{th}\) displays a relatively large decrease. It should be commented that similar results were obtained for all \([K]_0\) values, except no supernormality was observed for \([K]_0>4.6\) mM \((|K|_0=4 \text{ mM} \text{ is shown here as an example})\).

Figure 14 provides an explanation for these differences between strength–interval curves constructed with stimuli of different durations. Figure 14A shows the total charge \((Q_{\text{total}})\) carried by all ionic channels during critical subthreshold stimuli for a short \((T=0.5 \text{ msec})\) and a long \((T=20 \text{ msec})\) stimulus duration (note the different scales). For both durations, \(Q_{\text{total}}\) is positive, indicating a net positive charge leaving the cell. For \(T=0.5\) msec, \(Q_{\text{total}}\) is negligible compared with the stimulus charge \(Q_{th}\) (Figure 13). However, for \(T=20\) msec, \(Q_{\text{total}}\) is about half the magnitude of \(Q_{th}\) (Figure 13, note the different scales). This significant amount of charge that is lost to the extracellular medium during a long stimulus is the basis for the difference between the strength–interval curves of Figure 13. The \(Q_{th}\) supplied by a very short stimulus can be regarded as the amount of charge needed to bring the membrane to threshold with minimal charge losses to the extracellular medium. In contrast, the \(Q_{th}\) supplied by a long stimulus can be divided into two components: \(Q_{th}\) supplied by a very short stimulus and the total charge \((Q_{\text{total}})\) lost to the extracellular medium during the duration of the stimulus. The fact that \(Q_{th}\) includes \(Q_{\text{total}}\) explains the large values of \(Q_{th}\) including the large amplitude of the maximum that follows the supernormal notch (Figure 13A) when a long stimulus duration is used. Figure 14B shows the contributions of the different ionic channels to \(Q_{\text{total}}\) for a stimulus of 20 msec applied at different times, \(t\), during phase 2 to phase 4 of an action potential. For large \(t\), all contributions except \(Q_{K(T)}\) approach very small values, identifying \(I_{K(T)}\) as the reason for the large rheobase needed when long stimuli are used (Figure 13B). During phase 3 \((t<20 \text{ msec})\), a large decrease in \(Q_{\text{total}}\) from
its peak value is observed as the 20-msec stimulus is applied earlier in time (Figure 14A, arrow). This is caused by an increase in the magnitude of the (negative) \( Q_0 \) and a decrease in \( Q_{KI(T)} \) as \( t \) decreases (Figure 14B). \( Q_0 \) increases since, for a small \( t \), the stimulus is applied during a phase when \( I_{K1} \) is not completely inactivated. \( Q_{KI(T)} \) decreases because the stimulus is applied during the negative slope phase of \( I_{K1} \). This additional contribution of 2.3 nanocoulombs to the decrease in \( Q_{0} \) is a substantial fraction of the amplitude of the supernormal notch (6 nC). The result is a supernormal notch of about twice the amplitude as compared with that for a 0.5-msec stimulus (Figure 13A). The relatively fast decrease of \( Q_{0} \) during phase 4 (\( t=30−450 \) msec in Figure 13B, \( T=20 \) msec), when long stimuli are used, reflects a similar behavior of \( Q_{K} \) (Figure 14B). Note that contributions from all other currents are constant during this phase; in particular, \( Q_{KI(T)} \) is constant since \( I_{KI(T)} \) is time independent and the membrane potential during this phase is almost constant. Therefore, for long stimuli, the membrane excitability during phase 4 is determined by the kinetics of \( I_{K} \) (the time-dependent X gate). In contrast, \( Q_{0} \) is almost constant over the entire phase 4 when short stimuli are used (Figure 13B, \( T=0.5 \) msec). This is because for short stimuli \( Q_{0} \) is determined by \( V_{Na} \) (negligible contribution from \( Q_{total} \), which do not change significantly during phase 4. As will become clear in the next section, the behavior of the strength–interval curve during phase 4 strongly influences the response of the cell to repetitive stimulation. For short stimulus duration one expects minimal dependence of the response on the timing of stimulation during phase 4 (constant excitability, Figure 13B, \( T=0.5 \) msec). In contrast, for long stimuli, the response is strongly influenced by the timing (Figure 13B, \( T=20 \) msec). This implies that to investigate rate-dependent response patterns (e.g., Wenckebach periodicity), stimuli of long duration should be used.

**Wenckebach Periodicity and Channel Behavior**

Wenckebach periodicity is defined as a periodic, rate-dependent activation failure in cardiac tissue. Recently, this phenomenon was observed in isolated guinea pig or rabbit ventricular myocytes and in sheep cardiac Purkinje fibers.7 In this section we study the Wenckebach phenomenon by simulating periodic pacing of single myocytes with the use of the L-R membrane model. The response of the membrane to periodic stimulation can be analyzed in terms of the strength–interval curve and the APD.
restoration curve. During phases 3 and 4 of an action potential, the strength–interval curve describes the recovery of membrane excitability, whereas the restitution curve describes the recovery of APD of premature responses.\textsuperscript{41,42} The simulated strength–interval and APD restitution curves for $[K]_o=5.4$ mM are shown in Figure 15. Both curves display a monotonic behavior that is determined by the time course of deactivation of the $I_K$ $X$ gate. In the following simulations, a stimulus strength of 3.06 $\mu$A that corresponds to threshold stimulation at $S_1S_2=882$ msec was used (indicated by the lower arrow in Figure 15). Stimulus duration of 20 msec was used for reasons described above (conclusion of previous section) and to be consistent with the experimental protocols of Chialvo et al\textsuperscript{7} and Delmar et al.\textsuperscript{6} Stimulation was applied at a basic cycle length (BCL) ranging from 160 to 2,000 msec. Simulated activation patterns for different BCLs are shown in Figure 16A. A staircase plot of response-to-stimulus (R:S) ratios versus BCL is shown in Figure 16B (numbers in the figure indicate stimulus-to-response [S:R] ratios). The staircase pattern demonstrates phase-locking at S:R ratios of integer values (1:1, 2:1, 3:1, etc.) and sharp transitions between these values through many noninteger S:R ratios. This pattern is in good agreement with the experimental observations of Chialvo et al\textsuperscript{7} To elucidate the underlying mechanism of the Wenckebach phenomenon observed in single cells the following simulations were performed.

As we discussed in the previous section, the post-refractory period (phase 4 of an action potential) is dominated by the $X$ gate of $I_K$.\textsuperscript{6,28} Therefore, the Wenckebach phenomenon is likely to be related to the value of $X$ just before the stimulation is applied ($X_3$). To test this assumption, $X_3$ of a given beat was clamped to the value of $X_3$ of the preceding beat. Independent of the particular beat where clamping was applied the following behavior was observed: 1) S:R pattern always progressed to S+1:R+1 pattern and 2) the superimposed action potentials of the two beats with the same $X_3$ overlapped because of equal time latency from stimulus to response. An example of this procedure is shown in Figure 17 for an initial

\begin{figure}
\centering
\includegraphics[width=\textwidth]{figure16.png}
\caption{Periodic rate-dependent patterns for normal $[K]_o=5.4$ mM. Panel A: Examples of activation patterns (calibration bar, 52.5 mV). Numbers on the left indicate basic cycle length (BCL), whereas ratios on the right are the resulting stimulus-to-response (S:R) patterns. The train of stimuli is indicated below each pattern. Panel B: The staircase plot of response-to-stimulus (R:S) ratios as a function of BCL. The numbers inside the panel indicate the S:R ratios.}
\end{figure}

\begin{figure}
\centering
\includegraphics[width=\textwidth]{figure17.png}
\caption{Rate-dependent block is studied by resetting the activation ($X$) gate of the time-dependent potassium current value at onset of stimulation ($X_3$). Panel A: Stimulus-to-response (S:R) ratio of 6:5 changes to 7:6 by setting $X_3=X_4$. The train of stimuli is shown below the pattern. Panel B: The action potentials for the pattern in panel A are redrawn with the onset of stimulation for all beats set at time zero. The superimposed third and fourth action potentials (indicated by * ) clearly indicate that beat 4 is exactly the same as beat 3 because of their equal values of $X_3$. Calibration bars for action potentials are 35 mV in both panels.}
\end{figure}
example. Beat 1 in the figure refers to the first beat after activation failure during the previous period. The APDU of the second beat (APDU2) is smaller than the APDU of the first beat (APDU1). This is due to a larger decrease (7.7 msec) in APD than the increase (6 msec) of tu from the first beat to the second beat (note APDU=APD+tu). The relatively large decrease of APD from beat 1 to beat 2 reflects the twice-longer recovery time before beat 1 that results in a long APDU (arrow b in Figure 15) as compared with APD2 (arrow a in Figure 15). Also, a corresponding large increase in XS is observed. Starting from beat 2, APDU and tu increase monotonically, whereas APD remains almost constant and XS increases very slowly. The monotonic increase in APDU implies a progressively shorter recovery time between beats starting from beat 2. As a result, X is progressively less deactivated at the time of the stimulation (XS increases monotonically), until failure occurs. When we set XS for all beats equal to XS of the first beat, activation failure does not occur and the pattern is always 1:1. This mechanism is consistent with the explanation of Delmar et al.28

A deviation from this behavior is observed for the first two beats (Figure 18). APDU decreases from beat 1 to beat 2, implying a longer recovery time before beat 3 than before beat 2. However, in spite of the longer recovery time before beat 3, XS3 is greater than XS2. This seemingly anomalous behavior results from the very different initial conditions at the onset of beats 1 and 2. Beat 1 occurs at an interval of 2·BCL from the last successful beat of the previous period, implying a much longer (about twice) recovery time than that for all other beats. As a result, X is almost completely deactivated at the onset of beat 1 (XS1=0.007), reaching a maximum value of 0.4215 at plateau potentials. At the onset of beat 2, because of much shorter recovery time, X is much less deactivated (XS2=0.0527), reaching a maximum value of 0.4262 during this beat. The larger maximum value of X at plateau potentials more than compensates for the increase in recovery time, resulting in XS3>XS2 (XS3=0.0529). Hence, in spite of an initial decrease in APDU, XS increases monotonically from beat to beat until failure occurs. It should be mentioned that the “anomalous” decrease in APDU from beat 1 to beat 2 is not always observed. When stimulating, for the same S:R pattern, at longer BCLs (and therefore with lower stimulus strength) we observe a monotonic increase in APDU for all beats, similar to the observations of Delmar et al.28 This is because as BCL becomes progressively longer (relative to \( \tau_X \)), the change in XS from beat 1 to beat 2 becomes progressively smaller, resulting in a smaller decrease of APD until APD approaches a constant value. In contrast, for a given S:R pattern, the increase of tu from beat 1 to beat 2 remains in the range of 5.5–7 msec for all BCLs (obviously, to keep the same S:R pattern the stimulus strength was adjusted for different BCLs). Because APDU=APD+tu, APDU in-

\[ \text{S:R}=6:5 \] (see Figure 16A). The XS of the fourth beat was clamped to the XS of the third beat. The pattern (Figure 17A) has changed to S:R=7:6 because of resetting of the fourth beat to the third beat. When the action potentials are superimposed (Figure 17B) by setting time zero at the onset of each stimulus, the action potential of the clamped beat overlaps with the previous action potential (i.e., tu and APD [defined in Figure 5] have the same values for the two beats). Also, when XS for all beats is clamped to XS of the first beat (not shown) S:R is always 1:1. Clearly, XS plays a dominant role in determining the pattern of Wenckebach periodicity.

To further elucidate the mechanism through which the IK X gate controls the activation pattern, we plotted APD, APDU, tu (see Figure 5 for definitions), and XS as a function of beat number (Figure 18). The periodic 6:5 pattern discussed above was used as an

![Figure 18. APDU, APD, XS, and tu for the 6:5 pattern are plotted as a function of beat number. Panel A: Nonmonotonic behavior of APD and APDU. Panel B: Monotonic increase in XS and tu. APDU, action potential duration from onset of stimulation to 90% repolarization; APD, action potential duration from maximum upstroke velocity to 90% repolarization; XS, activation gate of the time-dependent potassium current at onset of stimulation; tu, latency from onset of stimulation to maximum upstroke velocity](image-url)
The simulations described above identify the ionic channel kinetics responsible for the alternans phenomenon. It remains to be clarified why these alternans appear only for short BCLs (BCL<560 msec in our simulations). Under these conditions (the example we show is for BCL=448 msec) we observe a relatively large decrease (56.2 msec) in APDl from beat 1 to beat 2 that reflects a large decrease (61.8 msec) in APD. The large change in APD is possible because short BCL corresponds to the steep portion of the APD restitution curve (Figure 15). Mechanistically, the large APD change results from a relatively large increase in X gate and a decrease in f gate (Figure 19). This decrease in APDl is followed by a long recovery time of 130.4 msec to the third beat. This recovery time is long enough so that the X gate at the onset of beat 3 is relatively more deactivated in spite of its large initial value at the onset of beat 2, bringing about an increase in APDl of beat 3. The following recovery time is therefore decreased, resulting in a decrease in APDl of beat 4. This alternating pattern continues until a stable steady state is achieved with a constant APDl=327.66 msec. In contrast, for BCL>700 msec, the changes in APDl are smaller (<23 msec) since this range corresponds to the flat portion of the APD restitution curve (Figure 15). Therefore, the decrease in APDl from beat 1 to beat 2 results in a relatively small increase in the recovery time to beat 3. This recovery time is too small to compensate for the large initial value of X at the onset of beat 2. As a result, Xs increases monotonically from beat to beat and no alternans are observed.

It should be emphasized that the large changes in APDl at short BCL reflect large changes in APD. This is because at short BCL the stimuli are applied during a fast-recovery phase of the APD restitution curve (Figure 15). In contrast, for long BCL, the stimuli are applied during a slow phase of the restitution curve (Figure 15), resulting in small changes in APD. Consequently, for long BCL, APDl changes are determined by changes in tu (except for beat 1 to beat 2, where APD also changes significantly as described above). Starting from beat 2 changes in APDl are determined by tu alone (since APD approaches a constant value). As shown in Figure 18, the increase in tu starting from beat 2 results in a monotonic increase in APDl and a progressively shorter recovery time between beats until activation failure occurs. The result of this progressive decrease in membrane excitability is the appearance of noninteger S:R patterns (such as 8:7, 7:6, 6:5, etc.) between 1:1 and 2:1 and all other integer S:R ratios. These noninteger S:R patterns are observed over a BCL interval of 2 msec. In contrast, for short BCL, noninteger S:R patterns are not observed in our simulations and there is a direct transition from 1:1 to 2:1 (and between all other integer S:R ratios). This is because for short BCL APDl changes are dominated by the large changes in APD (and not by tu). Under these conditions, the recovery time does not...
not decrease progressively from beat to beat since the shortest recovery time is between beat 1 and beat 2. This is because APD_U is determined by APD, and APD of beat 1 is the longest since it occurs after a recovery time of 2 · BCL. Therefore, if activation is successful at beat 2, all subsequent stimuli result in successful responses and a 1:1 pattern is established. As described above, this 1:1 pattern is characterized by alternans in APD. However, if activation failure occurs, it always occurs at beat 2, where recovery time is the shortest. The result is a direct transition from a 1:1 to 2:1 pattern without the possibility for noninteger S:R patterns.

Supernormality and Rate-Dependent Activation

The Wenckebach activation patterns discussed above were obtained in the setting of normal extracellular potassium concentration ([K]_o=4.6 mM), for which supernormal excitability does not exist. For low extracellular potassium concentration, supernormal excitability appears during or immediately after the action potential repolarization phase (see “Supernormal Excitability”). Since the presence of supernormality modifies the strength–interval curve by introducing nonmonotonic recovery of excitability, we expect rate-dependent activation patterns under conditions of low [K]_o to be different from the regular rhythms for normal [K]_o discussed in the previous section. Recently, complex response patterns were observed experimentally by Chialvo et al. in Purkinje fibers at low extracellular potassium concentrations ([K]_o=4 mM). In this section, we will use the L-R model to simulate and investigate this phenomenon.

The strength–interval curve computed for [K]_o=4 mM and a test stimulus of 20-msec duration is plotted in Figure 20A. The horizontal dotted line indicates the current amplitude of the stimuli during repetitive stimulation. For this level of stimulation (3.13 μA), four different regions of S1S2 interval exist, as characterized by their responses. Regions 1 and 3 are excitable, whereas regions 2 and 4 are inexcitable. Note that excitable and inexcitable regions alternate, so that excitable regions are separated by an inexcitable region. Figure 20B depicts the APD_U restitution curve for the same conditions. The same four regions are identified. Note the discontinuity of this curve in region 2, corresponding to the inexcitable region in Figure 20A. Also, APD_U is almost constant in region 1, while in region 3 a fast rate of change is observed. This fast change of APD_U is caused by the relatively fast decrease of the X gate during this phase (see “Wenckebach Periodicity and Channel Behavior”).

The results of a simulated repetitive stimulation experiment with the stimulus strength indicated in Figure 20A (current amplitude of 3.13 μA) are shown as the staircase plot of the R:S ratio versus BCL of stimulation (Figure 21). Numbers in the figure indicate the S:R ratios. This format is the same as that for Figure 16B, which was obtained for [K]_o=5.4 mM, in the absence of supernormality. A comparison of these figures clearly demonstrates that the activation patterns are different under these two different condi-

**Figure 20.** Strength–interval curve (panel A) and APD_U restitution curve (panel B) for [K]_o=4 mM. In panel A horizontal dotted line indicates the stimulus amplitude used in this simulation. Excitable regions (1 and 3) are shaded to distinguish them from the inexcitable regions (2 and 4). The restitution curve in panel B is discontinuous because of the inexcitable region 2. I_a, threshold current; APD_U, action potential duration from onset of stimulation to 90% repolarization. The letters in panel B are defined in the text.

**Figure 21.** Periodic rate-dependent block for low [K]_o=4 mM. Note the nonmonotonic behavior caused by the supernormal phase (arrows). R:S, response-to-stimulus ratio; BCL, basic cycle length. The numbers in the figure indicate the stimulus-to-response ratio, S:R.
patterns are from 465 A are panel BCL. Here by showing the range of enlarged to-stimulus ratio; 22.

FIGURE 22. The first nonmonotonic range of Figure 21 is enlarged here by showing the range of basic cycle length (BCL) from 465 to 410 msec (panel A). Panel B: Related activation patterns are plotted (calibration bar, 35 mV). R:S, response-to-stimulus ratio; S:R, stimulus-to-response ratio. Numbers in panel A are S:R ratios. Arrows in panel A indicate a specific BCL.

tions. For normal [K]o (no supernormality) a monotonic progression of the activation ratio is observed (Figure 16B). In contrast, nonmonotonic changes in the activation ratio are observed for low [K]o (supernormality present) (Figure 21). Under these conditions, a stable 1:1 pattern is followed by a region of nonmonotonic behavior as BCL is decreased below 458 msec (right arrow). This nonmonotonic region covers a range of BCLs from 458 to 415 msec. Below 415 msec another regular region of a 2:1 pattern is obtained, followed by another region of nonmono-

tonic behavior (BCLs from 205 to 190 msec, left arrow). Note that the same patterns repeat for different regions of BCL. For example, a 1:1 pattern is

FIGURE 23. The second nonmonotonic range of Figure 21 is enlarged here by showing the range of basic cycle length (BCL) from 205 to 180 msec (panel A). Panel B: Highly aperiodic patterns that correspond to BCL = 200.4 msec (arrow in panel A) are plotted (calibration bar, 35 mV). In the second row of panel B, * indicates complex patterns of consecutive action potentials; F and S indicate activation failure or success, respectively. R:S, response-to-stimulus ratio. Numbers in panel A are stimulus-to-response (S:R) ratios. For action potentials with rapid sodium-dependent upstroke we define success as peak INa greater than 1% of the peak INa under fully recovered conditions. For slow response action potentials (upstroke depends on INa) we define success as an action potential of duration greater than 50 msec (not including the stimulus duration).
observed for a BCL of 500, 438, and 428 msec. These regions are separated by regions of 2:1 patterns.

The nonmonotonic regions of Figure 21 are amplified in Figures 22A and 23A. In Figure 22A, as BCL is decreased from 465 msec the activation pattern changes from 1:1 to 2:1 at a BCL of 458 msec and then returns to the 1:1 pattern at a BCL of 439 msec. This nonmonotonic transition from 2:1 to 1:1 is possible because of the nonmonotonic strength–interval curve (supernormality) of Figure 20A. The 2:1 pattern in Figure 22A occurs over a range of BCLs (between 458 and 440 msec) that corresponds to region 2 of the strength–interval curve (Figure 20A).

A stimulus at this region fails to elicit a response so that the next stimulus occurs at 2 • BCL in region 1 of the strength–interval curve, resulting in a successful response. A 2:1 pattern is thus established (see the first row in Figure 22B). The 1:1 pattern of activation for BCLs between 439 and 443 msec is also shown in Figure 22B (second row). The first action potential is of maximum duration because it is stimulated from a fully recovered membrane. The second stimulus is applied during the supernormal phase (region 3 in Figure 20A) and displays a shorter duration. This is because the stimulus is timed during the fast decreasing phase of the restitution curve (point a in Figure 20B). As a result, a longer recovery time follows this action potential and the third stimulus is applied in region 1 of the strength–interval curve (Figure 20A).

However, the duration of the third action potential (point a1 in Figure 20B) is still shorter than that of the first action potential, obtained under fully recovered conditions. Consequently, the recovery time is longer and all subsequent stimuli are applied in region 1 of the strength–interval curve, resulting in a 1:1 pattern. For a further decrease in BCL, the pattern returns to 2:1 (BCL of 432 msec), and then back to 1:1 (BCL of 428 msec). These two patterns are shown in Figure 22B (third and fourth rows, respectively). For a BCL of 432 msec, the second action potential is of short duration, corresponding to point b on the restitution curve of Figure 20B (note that the APD at b is smaller than that at point a). The following stimulus elicits an action potential of duration (b1) on the restitution curve (b1 > a1). As a result, the next stimulus is timed during region 2 of the strength–interval curve and does not elicit a successful response. A 2:1 pattern is thus established. For a BCL of 428 msec (see Figure 22B) the second response is of a very short duration, corresponding to point c in Figure 20B. The following stimulus is applied after a long recovery time, resulting in an action potential of long duration (point c1 in Figure 20B). As a result, the next stimulus is applied in region 2 of the strength–interval curve and fails to elicit an action potential. This event is followed by a long recovery time and an action potential of long duration so that the next stimulus is timed during the supernormal phase (region 3) of the strength–interval curve, resulting in a successful response of short duration. After these transitional events the pattern

is similar to that obtained for a BCL of 436 msec (see Figure 22B), and a ratio of 1:1 is established. A similar interplay between the excitability and APD at the different regions defined in Figures 20A and 20B determines the various patterns as BCL is further decreased.

In Figure 23A, as the BCL is reduced below 205 msec, irregular activation patterns are observed. An example is shown in Figure 23B for a BCL of 200.4 msec (arrow in panel A). For this particular BCL, unstable patterns are observed and no regular periodicity is established for more than 250 consecutive stimuli. This behavior is consistent with the experimental findings of Chialvo et al., who observed highly aperiodic activity (termed "chaotic activity") at a BCL of 200 msec with no repeated patterns for recording periods that encompassed 100 or more stimuli. In our simulations, after a transient chaotic pattern that lasted more than 250 consecutive stimuli a very complex pattern with very long periodicity of about 13 seconds (67 stimuli) was established.

The highly aperiodic activity occurs at a BCL below 205 msec. For this range of BCL, the stimuli are frequently applied during region 3 of the strength–interval curve (supernormal phase). In this region, the APD depends very strongly on the time of stimulation. In fact, APD, changes by 281 msec over an S1,S2 interval of 15 msec (Figure 20B). As a result, a small change in the time of stimulation during the supernormal phase can cause a very large change in the duration of the elicited action potential and in the timing of the next stimulus. Consequently, small variations in the time of stimulation in region 3 can shift the timing of the next stimulus between different
regions of the strength–interval curve, eliciting a very different response. This high sensitivity to the timing of stimulation ("initial conditions") is the basis for the unstable patterns ("chaotic activity"). It should be noted that at a BCL between 410 and 465 msec nonmonotonic behavior of the activation patterns is also observed. However, for this range of BCL there are no unstable patterns or aperiodic activity. This is because for the longer BCL, the stimuli are infrequently applied during region 3 of the strength–interval curve (supernormal phase). Moreover, a stimulus during region 3 is never followed by another stimulus in the same region. The next stimulus always falls either in region 2 (activation failure) or in region 1 (the flat portion of the restitution curve, Figure 20B). On the flat portion of the restitution curve, small changes in the time of the stimulation can cause only very small changes in APD. This has a stabilizing effect on the response so that irregular activity and unstable patterns do not occur.

To further elucidate the ionic current behavior during the highly aperiodic activity, in Figure 24 we plotted the magnitude of $I_{\text{Na}}$ and $I_T$ that corresponds to the pattern shown in Figure 23B, second row from the top. An important observation is that most of the responses during the complex patterns do not involve the fast sodium channel and depolarization is caused by $I_T$. In Figure 23B, F identifies an activation failure during the plateau of an action potential that was initiated in region 1 of the strength–interval curve. A successful activation during the plateau of an action potential that was initiated in region 3 of the strength–interval curve is identified by S in the figure. The corresponding events in the $I_T$ curve are also identified by F and S, respectively (Figure 24). The difference between the peak magnitudes of $I_T$ for these two events is very small (0.2 $\mu$A). However, the total outward current ($I_{K(T)}+I_K$) at F is 3.93 $\mu$A as compared with only 2.35 $\mu$A at S. This decrease is mostly due to reduced $I_{K(T)}$ at the lower membrane potential of S and results in a successful response, bringing about a very complex pattern of activation. It should be mentioned that this behavior involves the negative slope characteristic of the $I_{K(T)}$ curve (Figure 3). When we eliminate the negative slope in the $I_{K(T)}$ curve (not shown), unstable patterns do not appear.

**Discussion**

The goal of this study is to construct a model of the membrane action potential of the mammalian ventricular cell based, whenever possible, on recent experimental findings in single-cell and single-channel studies. In this report we reformulate three ionic currents: $I_{\text{Na}}$, $I_K$, and $I_{K(T)}$. $I_{\text{Na}}$ is inactivated by two processes—fast and slow. The slow process of inactivation also implies slow recovery of excitability. In addition, $I_{\text{Na}}$ is characterized by a large channel conductance rising in fast upstream velocity ($V_{\text{max}}=400$ V/sec) of the action potential. A newly discovered potassium channel that activates at high potentials is implemented as a component of $I_{K(T)}$ in the model. $[K]_o$ dependence of $I_K$ and $I_{K(T)}$ is a major feature of the model. This enables us to simulate various phenomena that involve changes in $[K]_o$. In this paper we focus on the rising phase, the late repolarization phase, and the postrepolarization phase of the action potential. We use the model to investigate phenomena that involve $I_{\text{Na}}$, $I_K$, and $I_{K(T)}$ during these phases such as supernormal excitability, membrane responsiveness, and Wenckebach periodicity. The results are consistent with the experimental observations, and the phenomena are well described in terms of ionic channel behavior. It should be emphasized that the present model does not include ionic pumps and exchange mechanisms. It therefore cannot simulate dynamic changes in ionic concentrations. The slow inward current in the model is adopted from the B-R model and is not consistent with recent findings related to calcium currents through the membrane such as the existence of two types of calcium channels, L and T, and the role played by $[Ca]$ in inactivation. Also, while an uptake mechanism of intracellular calcium is represented in the model, the model cannot simulate intracellular calcium transients that result from the interaction between calcium currents through the sarcolemma and calcium release and uptake by the sarcoplasmic reticulum. A second phase of the ventricular model presented here is currently being developed in our laboratory to include all of these processes based on recent experimental data. Such processes were first incorporated in a model of the Purkinje action potential by DiFrancesco and Noble in 1985.

We use the Hodgkin-Huxley formalism to describe the macroscopic (ensemble) behavior of ionic channels. This choice is consistent with (and dictated by) our goal, namely, the reconstruction of the membrane action potential that results from ensemble currents through many individual channels. However, one should practice caution when extrapolating from the ensemble behavior to the single-channel behavior, especially when the cardiac sodium channel kinetics are considered. While many single sodium channel properties correspond to similar characteristics of the ensemble current and are adequately described by the Hodgkin-Huxley kinetics, some properties are not. For example, at potentials near threshold for sodium channel activation, both channel activation and inactivation determine the kinetics of the macroscopic sodium current decay (i.e., current inactivation). The microscopic behavior of ionic channels other than the sodium channels can be adequately described by the Hodgkin-Huxley kinetics.

For the sake of clarity, the discussion below is divided into modeling aspects and physiological simulation aspects of the study.

**Modeling Considerations**

$I_{\text{Na}}$: Fast sodium current. The fast upstream velocity of the action potential has been measured in single cells. The $V_{\text{max}}$ measured by Brown et al is in
the range of 200–300 V/sec at 20–22°C. By applying a minimum \(Q_{Li} \) of 1.23,\(^{48}\) \(V_{max} \) is at least 272–410 V/sec at 37°C. \(V_{max} \) of the E-J model is 300 V/sec, while it is 400 V/sec if computed by the L-R model. The faster uptake in the L-R model results from adjusting the reversal potential from 29 mV (E-J model) to 54.4 mV to reflect the different intracellular sodium concentration in mammals ([Na]\(_i\)=18 mM)\(^{13,23}\) and chicken embryos ([Na]\(_i\)=40 mM).\(^{25}\) This adjustment gives \(V_{max} \)=400 V/sec in the range of measured values. It also results in an overshoot (41.7 mV) of the action potential that is in good agreement with the experimental findings in guinea pig ventricular cells.\(^{31,35}\)

A tetrodotoxin (TTX)-sensitive slow sodium current that affects the action potential duration has been identified in single-channel recordings.\(^{49–52}\) In guinea pig ventricular cells, a low dose of TTX (≤30 \(\mu\)M) does not affect the APD significantly.\(^{33,53}\) However, Kiyosue and Arita,\(^{52}\) using a high dose of TTX (60 \(\mu\)M), observed ~10% shortening of the APD. This slow current is different from the window current caused by the overlap of the \(m_{Na} \) and \(h \) curves at high potentials,\(^{51,52}\) as demonstrated by the fact that TTX can affect the APD with little (or no) change in \(V_{max} \).\(^{54}\) The kinetics of this slow current are still not completely elucidated, and it is not included in the L-R model. We will have to include this channel in future studies (as more data become available), especially if effects of drugs are to be studied (some local anesthetics can affect this current).

Two time constants (fast and slow) better describe the inactivation of the sodium channel.\(^{13,49,50,55,56}\) This cannot be described by the Hodgkin-Huxley kinetics with one inactivation gate. Similar to Beeler and Reuter we incorporated two Hodgkin-Huxley-type inactivation gates as suggested by Haas et al.\(^{24}\) Haas et al\(^{24}\) proposed that two inactivation gates (\(h \) and \(j \) in the L-R model) with different time constants can describe the macroscopic behavior of the sodium current inactivation. Following this strategy, the L-R model correctly simulated the behavior of the sodium channel observed experimentally (Figures 7 and 8). Without slow recovery from inactivation (a property not included in the E-J model or in the DiFrancesco-Noble model) \(I_{Na} \) recovery from inactivation follows the same time course following different periods of inactivation (Figure 7D). This is inconsistent with the experimental observations (Figure 7A) by Ebihara et al.\(^{36}\) Also, without slow recovery from inactivation, the fast recovery of \(I_{Na} \) results in an unrealistically fast recovery of excitability at a high level of membrane potential during the repolarization phase of an action potential (Figure 8D). Note that a conservative value of \(\tau \) was used in our simulations. We used a maximum \(\tau_{j} \) of ~80 msec at \(V=-65 \) mV. Values as high as 200 msec were reported in the literature.\(^{14}\)

To account for this process of slow recovery from inactivation, alternative models to that of Hodgkin and Huxley were proposed to describe an inactivation gate with two time constants (fast and slow). Chiu\(^{57}\) proposed a modified Hodgkin-Huxley formalism (second-order kinetics) to describe two processes of inactivation. However, Brown et al.\(^{13}\) found that the \(h \) curve can be described by a single exponential, a result that is also supported by the recent observations for single-channel recordings.\(^{44}\) This deviates from the prediction of Chiu’s model\(^{57}\) that \(h \) has to be described by two exponentials as a function of voltage. Recently, Markov chain models have been proposed to describe the sodium channel kinetics in cardiac cells.\(^{58}\) Based on single-channel recordings and a five-state Markov model, Scanley et al.\(^{45}\) provided a kinetic analysis of the sodium channel. The basic five-state model reproduced observed open-duration data and reopening behavior but produced only a single exponential decay of current, implying that more complex models are required.

Single-channel recordings demonstrate that near-threshold potentials for channel activation sodium channels tend to open more than once during a depolarization (reopening phenomenon).\(^{44,45}\) Also, a significant fraction of sodium channels do not open by the time of peak inward current. These microscopic phenomena determine the time course of the macroscopic current decay and provide the basis for the slow macroscopic current inactivation that is introduced in our model by the \(j \) gate.

\(I_{K} \): Time-dependent potassium current. The formulation of \(I_{K} \) in our model is based on the form suggested by Shibasaki\(^{26}\) to fit his data in nodal cells, but its parameters are obtained from the B-R model of the ventricular action potential. Note that the property of [K]\(_0\)-dependent activation is introduced as a square root dependence of the maximum conductance. In contrast, the DiFrancesco-Noble formulation\(^5\) results in a linear dependence of channel conductance on [K]\(_0\) in the physiological range of [K]\(_0\), from 2 to 10 mM. The simulated results (Figure 1) are consistent with the experimental findings by McDonald and Trautwein\(^{29}\) and Shibasaki\(^{26}\) and include minimal crossover between current–voltage curves of different [K]\(_0\), and strong inward rectification. An alternative formulation could be constructed from the data measured by Matsuura et al\(^{27}\) in guinea pig single ventricular cells. They measured the fully activated \(I_{K} \) curve [\(I_{K}(V) \)] for [K]\(_0\)=5.4 mM and proposed that \(I_{K}=|I_{K}| \cdot X^2 \) (\(I_{K} \) activation is proportional to the second power of the activation gate, \(X \)). We introduced [K]\(_0\), dependence in their formulation and investigated the behavior of \(I_{K} \) for different values of [K]\(_0\). The results displayed significant crossover between \(I_{K} \) curves of different [K]\(_0\). This is inconsistent with the experimental observations.\(^26,29\)

We then introduced this alternative formulation of \(I_{K} \) in the L-R model of the action potential. Since the surface membrane area was not given by Matsuura et al.,\(^{27}\) we adjusted the maximum conductance (\(G_{K} \)) based on the mean capacitive membrane area measured by Isenberg and Klöckner (2x10\(^{-4}\) cm\(^2\)/cell)\(^{31}\) and Kamp et al (8.52x10\(^{-5}\) cm\(^2\)/cell).\(^{60}\) This resulted
in $G_K=0.143 \text{ mS/cm}^2$ and $G_K=0.335 \text{ mS/cm}^2$, respectively, in comparison with the L-R $G_K=0.282 \text{ mS/cm}^2$. Note that specific capacitance of 1 $\mu \text{F/cm}^2$ is assumed. The (adjusted) alternative $I_k$ failed to repolarize the membrane when $[K]_o \leq 4.5 \text{ mM}$ (for $G_K=0.143 \text{ mS/cm}^2$) and $[K]_o \leq 2.2 \text{ mM}$ (for $G_K=0.335 \text{ mS/cm}^2$), and the membrane potential stayed at about $-30$ to $-40 \text{ mV}$. This is inconsistent with the experimental observations at low [K]_o. Tseng et al. observed repolarization at [K]_o = 2 mM but not at [K]_o = 1 mM. This behavior is simulated correctly by the L-R $I_k$ in spite of its smaller conductance ($G_K<0.335 \text{ mS/cm}^2$). This inability of the Matsuura et al. to repolarize the membrane in spite of the large conductance results from the X^2 dependence of $I_k$. For these reasons we formulated the L-R $I_k$ to depend on X (rather than $X^2$), as suggested by Shibasaki. It should be emphasized, however, that the behavior of $I_k$ was demonstrated in a model that retains the slow inward current formulation of the B-R model. The Matsuura $I_k$ will have to be reexamined once an accurate model of $I_{Na}$ is developed.

$I_{K(T)}$: Total time-independent potassium current. This current is equivalent to $I_{K1}$ in the B-R model and is renamed here $I_{K(T)}$ based on the following recent findings. In single-channel recordings, Kurachi et al. found that 1) an inactivation gate (K1) of $I_{K1}$ is a function of membrane potential and of the potassium Nernst potential and 2) this gate completely inactivates at plateau potentials, resulting in zero $I_{K1}$ current at these potentials. The zero contribution of $I_{K1}$ at plateau potentials was also observed by Isenberg and Shah et al. Therefore, the total (non-zero) time-independent potassium current ($I_{K(T)}$ at high (plateau) potentials must result from current through other channels. Kakei et al. and Taira et al. found that the shortening of the APD caused by potassium channel agonists (nicarandil and pinacidil) is a result of increasing the current only at high potentials. This observation supports the Yue and Marban discovery of a plateau potassium channel ($I_{Kp}$) that activates only at high potentials. This channel is characterized by a much smaller time constant than $I_K$, no detectable inactivation for more than 600 msec, and high selectivity to potassium ions. Also, the maximum conductance ($G_{Kp}$) is largely insensitive to changes in $[K]_o$. This insensitivity is consistent with the experimental observations by Isenberg and Klöckner (Figure 8 of Reference 8), Sakmann and Trube (Figure 4A of Reference 11), and Tseng et al. (Figure 2A of Reference 9) in whole-cell recordings. They observed that the current--voltage curves of $I_{K(T)}$ for different values of $[K]_o$ (1--22 mM) overlap at high potentials. This overlap implies that the plateau currents are insensitive to changes in $[K]_o$. According to the observations described above, $I_{K1}$ contributes no current at plateau potentials and $I_{Kp}$ can be treated as a component of $I_{K(T)}$ that is activated only at plateau potentials. Also, a background current ($I_b$) exists at negative potentials after blocking $I_{Na}$, $I_{K1}$, and $I_{Kp}$. We represent the $I_{K(T)}$ as the sum of these three currents to fit the data measured by Sakmann and Trube (Figure 4A of Reference 11) and obtain an excellent fit for all values of $[K]_o$ (Figure 3). It is of interest to note that the negative slope of the simulated $I_{K(T)}$ is smaller than that measured by Kurachi et al. in single-channel recordings. However, the L-R result is consistent with the experimental observations in multicellular12,63 and single-cell9,34 recordings. Note that the property of $[K]_o$-dependent activation is introduced as a square root dependence of the maximum conductance. This behavior is different from the linear dependence on $[K]_o$ over the physiological range in the DiFrancesco-Noble formulation ($[K]_o<<[K]_m$ in their Equation 13).

Physiological Simulations

Membrane response to premature stimulation. Premature stimulation can be used to investigate the recovery of membrane excitability during the repolarization phase of an action potential. The simulations show that the premature membrane response, indicated by $V_{\text{max}}$ measured in mammalian ventricular cells, is well represented by the L-R model (Figure 8). We define the absolute refractory period as the range of membrane potential for which stimulation will result in $V_{\text{max}}$ that is smaller than 20% of the fully recovered $V_{\text{max}}$. The result ($V>-78 \text{ mV}$) of the L-R model (Figure 8B) that incorporates the process of slow recovery from inactivation is very close to the experimental data ($>-76 \text{ mV}$, Figure 8A). In the absence of slow recovery from inactivation (E-J model, Figure 8D), the simulation result ($V>-69 \text{ mV}$) deviates significantly from the experimental data. Similar behavior is expected from all models of the sodium channel that do not incorporate the process of slow recovery from inactivation (e.g., Drouhard and Roberge). This comparison demonstrates the importance of the process of slow recovery from inactivation in determining the time course of recovery of membrane excitability during repolarization. Since during repolarization $j$ is almost completely inactivated and since $\tau_i \gg \tau_{in}$, it follows that the membrane responsiveness is controlled by the slow inactivation $j$ gate. Our simulations also demonstrate the importance of incorporating the correct steady-state characteristics of $I_{Na}$, inactivation [h(V)] and $j_e(V)$ in the model. The steady-state behavior predicted by the B-R model deviates from the experimental behavior of cardiac fibers (compare curve 2 in Figures 8C and 8A), whereas the L-R model (based on the $I_{Na}$ data from mammalian ventricular cells) simulates the experimental behavior correctly. It is also important to incorporate the correct $G_{Na}$. The B-R model ($G_{Na}=4 \text{ mS/cm}^2$) predicts a $V_{\text{max}}$ that is unrealistically slow (115 V/sec), while the L-R model ($G_{Na}=23 \text{ mS/cm}^2$) predicts a $V_{\text{max}}$ within the range of measured values (400 V/sec). Note that a $G_{Na}$ increase by a factor of almost 6 results in a $V_{\text{max}}$ increase by a factor of about 3. This
is because the actual rate of activation of the cardiac sodium channel (represented in the L-R model) is slower than the rate of activation used in the B-R model (modified from squid-axon data). The unrealistic low $G_{Na}$ of the B-R model also leads to an unrealistically long latency from threshold potential to $V_{\text{max}}$ (Figure 6D) that is inconsistent with the experimental findings.28

So far, we have considered the membrane response at a normal $[K]_o$, ($[K]_o$=5.4 mM). The L-R model allows changes in extracellular potassium concentration to be incorporated. We studied the membrane response to premature stimulation for different $[K]_o$. The strength–intervals curve is determined by the critical (minimum) strength of test stimuli needed to excite the membrane during the repolarization phase of an action potential. Experimentally, the stimulus strength decreases monotonically with repolarization at normal $[K]_o$; however, supernormal excitability has been identified as an interval in which nonmonotonic behavior of the strength–intervals curve occurs at low $[K]_o$.7,39 The results (Figure 9) of the L-R model simulations are consistent with the experimental observations. For $[K]_o$ > 4.6 mM, the threshold potential decreases during repolarization at a rate that is faster than the rate of repolarization (Figure 12). As a result, membrane excitability increases monotonically during repolarization. For low $[K]_o$, ([K]o≤4.6 mM) there is a range during late repolarization at which the rate of membrane repolarization is faster than the decrease of threshold potential (Figure 12).38 This results in a decrease of excitability over this range and, therefore, in a nonmonotonic recovery of membrane excitability (Figure 9). The presence of an interval during which excitability is higher than in the adjacent (previous and following) intervals is known as “supernormality.”

At the level of channel kinetics, this behavior is a consequence of different kinetics of slow recovery from inactivation of $I_{Na}$ at different values of $[K]_o$. As $[K]_o$ decreases from 7 to 3 mM, resting potential (primarily controlled by $I_{K1}$), becomes more negative (from −78.2 to −95.5 mV) so that the time constant of sodium channel recovery, dominated by the slow inactivation j gate, becomes smaller ($\tau$ decreases from 33 to 7 msec). As a result, the sodium channel recovers earlier in the repolarization phase at a range for which repolarization is faster than the rate of decrease of threshold potential, and a supernormal window is created. The interaction between $I_{K1}$ and slow recovery from inactivation of $I_{Na}$ in response to changes in $[K]_o$ determines the amplitude and width of the window of supernormal excitability (Figure 9). Without the process of slow recovery from inactivation, the sodium channel recovers unrealistically earlier so that supernormality is observed even when $[K]_o$ is as high as 7 mM (Figure 11A). This is inconsistent with the experimental observations.7,39

Periodic stimulation at normal $[K]_o$. Wenckebach rate-dependent block was originally characterized as a propagation-related process that involves decremental conduction. Wenckebach65 described this phenomenon in the atroventricular node. Wenckebach periodicity was also demonstrated in cardiac tissue that contained a region of depressed conduction separating two regions of normal excitability.66 Recently, Wenckebach periodicity was demonstrated in a single ventricular cell under periodic stimulation,5,40 indicating that the phenomenon is an intrinsic property of the cell membrane. Using the L-R model at normal $[K]_o$ (5.4 mM), we investigated the cellular mechanism of the periodic rate-dependent block. The simulations demonstrate that this process is dominated by the kinetics of the X gate of $I_K$. This is because membrane excitability is influenced to a large extent by the loss of charge to the extracellular medium through the $I_K$ channel during stimuli of relatively long duration (20 msec) (see Figures 13 and 14).

As the BCL of periodic stimulation decreases from 2,000 msec, a monotonic decrease of the R:S ratio is observed (staircase in Figure 16B). In our simulations, noninteger patterns of S:R ratios are observed in the transition interval between integer S:R ratios. For example, noninteger Wenckebach patterns (8:7, 7:6, 6:5, etc.) are found between 1:1 and 2:1 (Figure 16B). These patterns are observed in a very narrow interval of about 2 msec. The fact that these patterns occur over such a narrow time interval might be the reason why these various Wenckebach patterns could not be sustained in single myocyte preparations28,40 since small perturbations (e.g., in temperature and pipette resistance) can cause large changes relative to the narrow transition range. In multicellular preparations7,65,66 the transition intervals are longer (up to 100 msec) and stable noninteger Wenckebach patterns are observed.

The appearance of Wenckebach block patterns is related to beat-to-beat changes in APD (measured from $V_{\text{max}}$ to 90% repolarization) and the latency from onset of stimulation to $V_{\text{max}}$. Both parameters are controlled by the degree of $I_K$ deactivation (kinetics of the X gate). For long BCL (BCL>920 msec), APD of all beats in a Wenckebach pattern remains almost constant (see Figure 15), while the latency increases progressively so that deactivation time (BCL−APD−latency) of the $X$ gate decreases monotonically, bringing about a progressive beat-to-beat decrease of membrane excitability. This process continues until activation failure occurs. This is the typical (classical) Wenckebach rate-dependent block observed in cardiac Purkinje fibers7 and in single ventricular myocytes.5,40 When BCL decreases (920 msec>BCL>700 msec), the pattern of beat-to-beat changes becomes different from the typical behavior described above. APD does not remain constant, and recovery time first increases (from second to third beat) and then decreases monotonically. However, membrane excitability still decreases monotonically from beat to beat until activation failure occurs. This is because the beat-to-beat increase in the value of
the X gate at the time of stimulation more than compensates for the initial increase in recovery time (see Figure 18). This atypical (nonmonotonic) behavior of recovery time can be observed in the data of Delmar et al. for stimulation at BCL=910 msec. A careful inspection of Figure 1B in their paper demonstrates that total APD (APD+latency) of the first beat in a Wenckebach pattern is greater than that of the second beat; also, the value of plateau potential is decreased. These experimental findings are consistent with the simulations presented here (Figures 17 and 18).

When BCL further decreases (<560 msec), the increase in the value of the X gate (i.e., degree of \( I_K \) activation) does not compensate for the initial increase in recovery time. As a result, membrane excitability increases at the time of the next stimulus, resulting in an action potential of long duration. This implies a shorter recovery time and lower membrane excitability at the time of the following stimulus, so that the following action potential is of short duration. A pattern of beat-to-beat alternans in APD is thus established. The amplitude of alternans (see Figure 19) decreases monotonically until a stable activation pattern (1:1) with constant APD (smaller than that of the fully recovered action potential) is established. Noninteger Wenckebach patterns cannot be found when alternans occur. This is consistent with the experimental observations (at BCL=430 msec) in isolated rabbit ventricular myocytes. The reason is that both phenomena (noninteger Wenckebach patterns and alternans) cannot occur simultaneously. Alternans occur at small BCL, at a range where there is a strong dependence of APD on the time of stimulation (Figure 15), reflecting a strong dependence of the X gate deactivation on the time of stimulation. At this range, the changes in APD are much larger than changes in the latency from stimulus to \( V_{\text{max}} \), so that the recovery time is determined by APD. Since the first beat is of the longest APD (a fully recovered action potential), the recovery time to the second beat is the shortest and all following stimulations will be successful. Hence, a 1:1 pattern is always established when alternans are present. In contrast, noninteger Wenckebach patterns are established at a range of BCL for which large beat-to-beat changes in the latency from stimulus to \( V_{\text{max}} \) and small changes in APD are present. At this range, recovery time is determined by the latency from stimulus to \( V_{\text{max}} \), which increases monotonically from beat to beat. The result is a monotonically decreasing recovery time leading to block. Delmar et al. used the B-R model with modified \( I_K \) to investigate the Wenckebach phenomenon for long BCL. The simulated transition interval during which noninteger Wenckebach patterns were observed was longer than 20 msec. With such a long interval, one expects that systematic measurements of Wenckebach patterns in single cells will reveal persistent noninteger ratios. This is inconsistent with experimental observations.

**Periodic stimulation at low \([K]\).** The L-R model incorporates the possibility of changing \([K]\). We studied the effects of changes in \([K]\) on the cell response to periodic stimulation. At low \([K]\), the nonmonotonic behavior of the strength–interval curve (Figure 20A) allows stimuli to excite the membrane during the supernormal phase but not at adjacent (earlier or later) intervals. This discontinuity in excitability together with the high sensitivity of APD to the time of stimulation (Figure 20B) during the supernormal phase brings about nonmonotonic behavior of the R:S curve and unstable activation patterns (Figure 21). This complex behavior was observed experimentally and described in a recent paper by Chialvo et al. For long BCL (>410 msec), stimulation is seldom timed during the supernormal phase. As a result, stable but complex activation patterns are observed (such as 10:6, 12:8, 6:4, and 20:11). Also, the same activation patterns are repeated at different ranges of BCL (nonmonotonic staircase behavior, see Figure 22). For short BCL (<205 msec), stimulation is frequently timed during the supernormal phase. Consequently, highly aperiodic patterns that do not stabilize for more than 250 consecutive stimuli are observed (Figure 23). This is consistent with the experimental findings of Chialvo et al. A very small change in BCL (from 200.4 to 200.7 msec) results in very different patterns, indicating very high sensitivity to initial conditions. The unstable patterns are frequently characterized by successful activations during the early repolarization phase of an action potential. This action potential was initiated by a stimulus applied during the supernormal phase. It is characterized by low plateau potentials, in the range where \( I_{K_{\text{r}}(T)} \) exhibits a negative slope. The decrease of \( I_{K_{\text{r}}(T)} \) with depolarization at this range results in higher membrane excitability and a possibility of successful activation during the early repolarization phase of an action potential. It should be noted that when we eliminate the negative slope in the current–voltage curve of \( I_{K_{\text{r}}(T)} \), the unstable aperiodic patterns disappear. Also, the successful activation at this range is a “slow response” caused by activation of \( I_{\text{s}} \) and not of \( I_{K_{\text{r}}} \). Similar unstable aperiodic patterns were predicted by an analytic (difference-differential) model formulated by Chialvo et al. Their model is based on the relations between excitability, latency, and APD. They show that the appearance of aperiodic patterns is related to the presence of supernormality and to the shape of the APD restitution curve at short diastolic intervals. These observations are consistent with the behavior of our membrane model and, as elucidated by our simulations, reflect both large changes in the state of deactivation of the \( I_K \) channel (X gate) and the negative slope characteristic of \( I_{K_{\text{r}}(T)} \).

In summary, the membrane model developed here correctly simulates various phenomena that result from the interaction of excitation and repolarization. This is an important property of the model since reentrant arrhythmias involve this type of interaction.
between head and tail of the reentrant action potential.\textsuperscript{17} Another property of the model is the ability to introduce changes in $[\text{K}]_o$. This is also an important aspect of the model since elevated $[\text{K}]_o$ is an important aspect of ischemia that influences propagation and arrhythmogenesis.

The simulations of membrane response to premature stimulation predict a variety of rate-dependent phenomena. These include Wenckebach periodicity, beat-to-beat alternans of APD, and (because of supernormality at low $[\text{K}]_o$) unstable aperiodic response patterns that display high sensitivity to initial conditions. The occurrence of Wenckebach periodicity in our simulations support other observations that this phenomenon is an intrinsic property of single cardiac cells. This observation suggests that the single-cell response under a variety of conditions might play an important role in arrhythmogenesis. The occurrence of beat-to-beat alternans in the simulations is intriguing. Beat-to-beat alternans in APD are a property of reentrant action potentials.\textsuperscript{17,68} In a broader sense, electrical alternans are detected in many circumstances that are associated with ventricular fibrillation and, as suggested by Guevara et al.,\textsuperscript{67} may reflect the first in a cascade of period-doubling bifurcations that eventually result in chaos, that is, fibrillation.\textsuperscript{69} As with Wenckebach periodicity, the observation that alternans can be generated at the level of the single cell points to the possible important role of the single-cell response in arrhythmogenesis. Finally, the appearance of highly aperiodic patterns that are extremely sensitive to initial conditions might also be related to arrhythmogenesis since it implies that short propagation delays (\textasciitilde1 msec) can initiate very different responses in neighboring cells (or multicellular fiber bundles). Note that propagation delays of this magnitude can occur across gap junctions under a very moderate degree of cell-to-cell uncoupling.\textsuperscript{15,16} We conclude with a word of caution. The behaviors described above are the responses of isolated cells. In the heart, the behavior of cells is modulated by loading conditions and electrotonic interactions.\textsuperscript{70} The response of a cell in an ensemble may, therefore, be different from its intrinsic response. Under conditions of increased cell-to-cell uncoupling the individual response of cells (or of a group of cells, i.e., fiber bundles) may become unmasked, bringing about conditions that favor arrhythmogenesis. Verification of these hypotheses awaits further experimentation in multicellular preparations and simulations in a multicellular model of cardiac tissue.

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