T-tubule localization of the inward-rectifier K^+ channel in mouse ventricular myocytes: a role in K^+ accumulation

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- 1. The properties of the slow inward 'tail currents' (I_{tail}) that followed depolarizing steps in voltage-clamped, isolated mouse ventricular myocytes were examined. Depolarizing steps that produced large outward K⁺ currents in these myocytes were followed by a slowly decaying inward I_{tail} on repolarization to the holding potential. These currents were produced only by depolarizations: inwardly rectifying K⁺ currents, I_{K1} , produced by steps to potentials negative to the holding potential, were not followed by I_{tail} .
- 2. For depolarizations of equal duration, the magnitude of I_{tail} increased as the magnitude of outward current at the end of the depolarizing step increased. The apparent reversal potential of I_{tail} was dependent upon the duration of the depolarizing step, and the reversal potential shifted to more depolarized potentials as the duration of the depolarization was increased.
- 3. Removal of external Na⁺ and Ca²⁺ had no significant effect on the magnitude or time course of I_{tail} . BaCl₂ (0.25 mM), which had no effect on the magnitude of outward currents, abolished I_{tail} and I_{K1} simultaneously.
- 4. Accordingly, I_{tail} in mouse ventricular myocytes probably results from K⁺ accumulation in a restricted extracellular space such as the transverse tubule system (t-tubules). The efflux of K⁺ into the t-tubules during outward currents produced by depolarization shifts the K⁺ Nernst potential (E_{K}) from its 'resting' value (close to -80 mV) to more depolarized potentials. This suggests that I_{tail} is produced by I_{K1} in the t-tubules and is inward because of the transiently elevated K⁺ concentration and depolarized value of E_{K} in the t-tubules.
- 5. Additional evidence for the localization of I_{K1} channels in the t-tubules was provided by confocal microscopy using a specific antibody against Kir2.1 in mouse ventricular myocytes.

The complex structure of cardiac muscle has made interpretation of the results of early electrophysiological studies of the membrane currents in multicellular tissue difficult. One of the most important problems in multicellular tissue is the accumulation of K^+ in the intercellular clefts, which are only 20-30 nm wide. These changes in the extracellular K⁺ concentration can distort the magnitude and time course of K⁺ currents (Almers, 1972; Attwell & Cohen, 1977; Attwell et al. 1979; Cohen & Kline, 1982). Many of these problems have been greatly reduced by the use of enzymatically isolated single cardiac cells. However, single myocytes have numerous invaginations of the sarcolemmal membrane that could act as restricted spaces in which ions might accumulate during activation of transmembrane currents. For example, adult mouse ventricular myocytes have an extensive transverse tubule (t-tubule) system whose surface area makes up approximately 50% of the total sarcolemmal membrane of the cells (Forbes *et al.* 1984). These tubules are small, about 50-100 nm in diameter (Forbes *et al.* 1984), and hence the tubule lumen could readily accumulate ions during current flow across the t-tubule membrane.

During the course of a voltage-clamp study of the K⁺ currents in single adult mouse ventricular myocytes (Fiset *et al.* 1997), it was noted that outward currents produced by depolarizing voltage-clamp steps were followed by a transient inward 'tail current' (I_{tail}) when the ventricular cell was repolarized to a holding potential near the cell's resting membrane potential (-80 mV). In contrast, no inward I_{tail} were observed following

depolarizing steps in adult mouse a trial myocytes. The present study was undertaken to investigate the properties and the origin of I_{tail} in adult mouse ventricular myocytes. The results of this study show that I_{tail} is due to K⁺ accumulation within the t-tubule system of mouse ventricular myocytes and subsequent activation of inwardly rectifying K⁺ (I_{K1}) channels.

METHODS

Isolation of adult mouse ventricular myocytes and electrophysiological methods

Ventricular myocytes were isolated from adult male Swiss-Webster or CD-1 mice (6-9 weeks of age) by enzymatic dissociation, as described previously (Fiset et al. 1997). The animals were heparinized, anaesthetized by inhalation of methoxyflurane and then killed by cervical dislocation. After perfusion of the heart with enzyme solutions, the ventricular tissue was triturated gently with a Pasteur pipette for 10-15 min. Rod-shaped single myocytes were then collected and stored in Kraft-Brühe (KB) solution at 4°C until use 2-6 h later. During cell isolation solutions were maintained at 37 ± 1 °C and were equilibrated with 100% O₂. Isolated cells were placed into a recording chamber (volume $\sim 200 \ \mu$) on the stage of an inverted microscope (Diaphot; Nikon, Tokyo, Japan) and superfused with a Hepes-buffered Tyrode solution at $1-2 \text{ ml min}^{-1}$. All experiments were carried out at a room temperature (RT) of 20-22 °C. Whole-cell voltage-clamp recordings were made with a patch-clamp amplifier (EPC-7; List-Medical Electronics, Darmstadt, Germany). Patch pipettes had resistances in the range $1.5-4 \text{ M}\Omega$ when filled with pipette solution. Series resistance in the whole-cell mode was in the range $4-8 \text{ M}\Omega$; 80-90% series resistance compensation was always used. Voltage-clamp currents were lowpass filtered at 1-3 kHz (four-pole Bessel), digitized at 4-10 kHz with a 12 bit analog-to-digital converter (DT2821; Data Translation, Marlborough, MA, USA), and stored in a microcomputer using Cellsoft software (D. Bergman, University of Calgary). The same software controlled the programmable stimulator that generated the voltage-clamp protocols.

Membrane currents were normalized to cell capacitance (pA pF⁻¹) when required. Capacitive currents were produced by a 5 mV depolarization from a holding potential of -80 mV and integrated to determine the cell capacitance.

Immunofluorescence analysis and confocal microscopy

Ventricular myocytes were freshly isolated from 6- to 9-week-old male mice and plated on laminin-coated coverslips (15 μ g ml⁻¹) for 1 h at 37°C, 5% CO₂. Cells were fixed (2% paraformaldehyde, pH 7.4, 10 min, 4°C) and subsequently permeabilized (0.1% Triton X-100, 10 min, RT). Three washes with phosphate-buffered saline (PBS) followed each step. To prevent non-specific binding, cells were blocked with 10% normal donkey serum (NDS) for 1 h at RT. After overnight incubation with a rabbit polyclonal primary anti-Kir2.1 antibody (1:200; Alomone Laboratories, Jerusalem, Israel), cells were washed with PBS and incubated for 1 h at RT with a tetramethylrhodamine-isothiocyanate-conjugated donkey anti-rabbit antibody (1:500; Jackson ImmunoResearch, West Grove, PA, USA). In control experiments, the primary antibody was omitted. Cells were then washed three times with PBS and incubated for 30 min at RT with wheat germ agglutinin (WGA) conjugated to Oregon Green 488 (1:10; Molecular Probes) (Sedarat et al. 2000; Takeuchi et al. 2000). After washing the cells with PBS, coverslips were mounted on slides using 1 mg ml^{-1} p-phenylenediamine diluted with 75% glycerol and examined using a Zeiss Axiovert 100M microscope coupled to a Zeiss LSM 510 laser scanning confocal system.

Solutions and drugs

The Hepes-buffered Tyrode solution contained (mM): 140 NaCl, 5.4 KCl, 1 CaCl₂, 1 MgCl₂, 10 Hepes, 5.5 glucose (pH adjusted to 7.4 at RT with NaOH). BaCl₂ was added to the solution at the concentrations indicated. Where indicated, NaCl was replaced by an isomolar amount of *N*-methyl-D-glucamine, and CaCl₂ was replaced by MgCl₂; the pH of this 'Na⁺-/Ca²⁺-free' solution was adjusted to 7.4 with HCl. The patch pipette solution contained (mM): 110 potassium aspartate, 20 KCl, 8 NaCl, 1 MgCl₂, 1 CaCl₂, 10 EGTA, 4 K₂ATP, 10 Hepes (pH adjusted to 7.2 with KOH). Solutions were continuously gassed with 100 % O₂.

Statisticl analysis

All results are presented as means \pm S.E.M. Statistical significance was determined by Student's t test for paired or unpaired observations, or ANOVA, where appropriate. Statistical analyses were performed using Origin 5.0 software. A probability of P < 0.05was considered significant.

RESULTS

Slow I_{tail} follow depolarization of voltage-clamped mouse ventricular myocytes

Figure 1A illustrates a representative example of a family of membrane currents recorded from a voltage-clamped mouse ventricular myocyte, produced by 750 ms voltage steps over the range of membrane potentials from -120to +50 mV, from a holding potential of -80 mV. Several features of these currents are of note.

(1) The outward membrane currents in adult mouse ventricular myocytes, which are carried primarily by K^+ (Fiset *et al.* 1997), are very large. For the cell shown in Fig. 1, the peak current produced by the step to +50 mV corresponded to a current density of about 90 pA pF⁻¹.

(2) Outward currents declined in two distinct phases during the depolarizing steps. It has previously been shown that there are at least two different transient components of the outward K^+ current in mouse ventricular myocytes (Fiset *et al.* 1997; Nerbonne, 2000; Trépanier-Boulay *et al.* 2001).

(3) Outward currents were followed by slow, transient I_{tail} on repolarization to the holding potential. This is illustrated more clearly in Fig. 1*B*, which shows the family of currents at the end of the depolarizing steps and after repolarization to the holding potential, on expanded time and current scales. Note that I_{tail} followed only the depolarizing voltage steps; no 'tail currents' were associated with the inward currents produced by hyperpolarizing steps.

Figure 1*C* shows the current–voltage relationships for the inward and outward currents, and for I_{tail} . As shown in Fig. 1*A* and *B*, I_{tail} was not produced by steps to potentials that generated inward currents; this was true even at potentials where the magnitude of the inward and outward currents was equal. For example, the currents produced by steps to -120 and +10 mV were of approximately equal magnitude at the end of the steps, but I_{tail} was associated only with the +10 mV step.

Figure 1*C* shows that the magnitude of I_{tail} increased as the magnitude of the outward current at the end of the depolarizing step increased.

Figure 2 summarizes the main properties of I_{tail} , averaged from nine different ventricular myocytes. Figure 2Ashows the relationship between the current at the end of the depolarizing step (I_{end}) and I_{tail} . The ratio I_{tail}/I_{end} varied considerably from cell to cell; for example, for 1 s steps to +30 mV, this ratio ranged from 0.2 to 1.07. The average value of this ratio was independent of membrane potential over the range -20 to +40 mV. Figure 2Bsummarizes the kinetic properties of I_{tail} . The time course of decay of I_{tail} (at -80 mV) was described approximately by a single exponential function. The time constant varied from cell to cell over about a threefold range (e.g. from 29.7 to 77 ms) for the same nine myocytes as above (for I_{tail} following +30 mV depolarizing steps). The average time constant was not significantly dependent upon membrane potential over the range -30 to +40 mV.

The time constant was also independent of the magnitude of I_{tail} ; Fig. 2*C* shows a plot of the relative amplitude of I_{tail} (normalized to I_{tail} following a +40 mV step) as a function of membrane potential from -30 mV to +40 mV. Note that although I_{tail} increased in magnitude by about 10-fold over this range of potentials, the time constant of decay did not change significantly.

It is unlikely that channel deactivation is the mechanism that is responsible for generating I_{tail} in mouse ventricular myocytes. The major part of the outward current in these cells is carried by K⁺; hence, current tails produced by deactivation would be expected to have their reversal potential near the K⁺ Nernst potential ($E_{\rm K}$), which is about -83 mV in the recording conditions used in these experiments (see Methods). Consequently, large inward $I_{\rm tail}$ would not be expected to occur at a holding potential of -80 mV (i.e. near $E_{\rm K}$). The data shown in Fig. 3 provide further evidence that channel deactivation is not responsible for $I_{\rm tail}$. In these experiments, ventricular cells



Figure 1. Slow, inward 'tail currents', I_{tail} , follow the depolarization of voltage-clamped, adult mouse ventricular myocytes

A, family of currents from a voltage-clamped, isolated mouse ventricular myocyte. The voltage-clamp protocol is shown in the inset; the holding potential was -80 mV; 0.75 s steps to potentials between -120 and 50 mV (-120, -100, -30, -10, 10, 30 and 50 mV) were applied at 0.1 Hz. Slowly decaying, inward currents followed the depolarizing (\bullet), but not the hyperpolarizing (\Box), voltage-clamp steps. The dashed line indicates zero current in this, and all subsequent figures. *B*, currents at the end of the depolarizing steps and after repolarization to -80 mV are shown on expanded time and current scales. Note that the magnitude of the inward I_{tail} increased as the magnitude of the outward current at the end of the step increased (\bullet). No I_{tail} followed hyperpolarizing steps (\Box). *C*, current–voltage relationship for currents obtained at the end of the voltage-clamp steps (\bullet), and peak I_{tail} (\bigcirc).



Figure 2. Summary of the properties of I_{tail}

A, plot of the ratio of magnitude of I_{tail} to current at the end of the depolarizing step (I_{end}) as a function of the step potential. Data (means \pm S.E.M.) were averaged from nine different ventricular myocytes. Slow I_{tail} were measured at a holding potential of -80 mV. B, plot of the time constant of decay of I_{tail} as a function of depolarizing step potential. I_{tail} was fitted to single exponential functions. Data (means \pm S.E.M.) were averaged from nine cells (same as in A). C, normalized magnitude of I_{tail} , as a function of depolarizing step potential. Current magnitudes were normalized to I_{tail} at +40 mV. Data (means \pm S.E.M.) were averaged from nine cells (same as above). depolarizing step decreased continually as the duration of the step increased. This is shown more clearly in the plots in Fig. 3*B*, in which the magnitudes of the outward current at the end of the step and $I_{\rm tail}$ are compared for each pulse duration. $I_{\rm tail}$ was small during depolarizing steps of brief duration (e.g. < 50 ms) where the outward current was largest. The magnitude of $I_{\rm tail}$ increased in parallel with step duration up to a limit of 100–150 ms, and then declined slightly for longer steps. This relationship between outward current and $I_{\rm tail}$ magnitudes cannot be explained by a simple channel deactivation mechanism.

Na^+-Ca^{2+} exchange does not generate I_{tail}

Slow inward I_{tail} can be generated in ventricular myocytes by the electrogenic Na^+ -Ca²⁺ exchanger (Egan *et al.* 1989; Bouchard *et al.* 1993). The rise in $[Ca^{2+}]_i$ produced during depolarization, due to the influx of Ca²⁺ through L-type Ca²⁺ channels and/or Ca²⁺-induced release from sarcoplasmic reticulum stores, activates the Na⁺-Ca²⁺ antiporter. The voltage dependence of Na⁺-Ca²⁺ exchange results in an inward current when the membrane is repolarized near -80 mV (Bouchard et al. 1993). However, this is not likely to underlie I_{tail} . The intracellular solution contains a high concentration of EGTA (10 mM) as a Ca²⁺ buffer, so it is unlikely that intracellular free Ca²⁺ concentration increases during depolarizing steps. Figure 4 shows data that make it clear that Na^+-Ca^{2+} exchange does not generate I_{tail} . In these experiments, currents produced by a depolarizing step (20 mV, 1 s) were recorded in control conditions and then in Na⁺ and Ca²⁺-free Hepes-buffered Tyrode solution (see Methods). The depolarizing step in both control and Na⁺ and Ca^{2+} -free solutions produced I_{tail} of equal magnitude and duration upon repolarization to the holding potential of -80 mV. Since Na⁺-Ca²⁺exchange is completely blocked in Na⁺ and Ca²⁺-free solution (Bers *et al.* 1990), the lack of effect of Na⁺ and Ca²⁺-free solution on I_{tail} makes it improbable that Na⁺-Ca²⁺ exchange was responsible for generating the slow I_{tail} .

Blocking I_{K1} abolishes I_{tail}

Barium, at concentrations less than 1 mm, is a specific blocker of the inwardly rectifying K^+ current, I_{K1} , in mouse ventricular myocytes. Figure 5 shows that 0.25 mM $BaCl_2$ blocks both I_{K1} and I_{tail} in mouse ventricular myocytes. In the experiment shown, a depolarizing step (+20 mV, 50-200 ms duration) that produced a large outward current, and I_{tail} on repolarization, was followed by a hyperpolarizing step (-120 mV, 100 ms) that activated inward I_{K1} . Application of BaCl₂ completely blocked both $I_{\rm K1}$ and $I_{\rm tail}$, but had no effect on the outward currents. In addition, the time courses of block of $I_{\rm K1}$ and I_{tail} as the BaCl₂ solution was washed into the recording chamber were identical (data not shown). These data provide strong evidence that I_{tail} is generated by K⁺ flux through $I_{\rm K1}$ channels. As the holding potential (-80 mV) was close to the nominal $E_{\rm K}$ under these recording conditions (see Methods), only very small I_{K1} would be



Figure 3. Effect of duration of depolarization on the magnitude of I_{tail}

A, family of currents produced by a series of voltage-clamp steps (20 mV) of increasing duration (10–285 ms; inset). Note that I_{tail} increased in magnitude with increasing step duration, while the outward current at the end of each step decreased. B, plots of outward current at the end of the depolarizing step (upper graph) and I_{tail} magnitude (lower graph), as a function of depolarizing step duration.

expected to occur. The large, inward I_{tail} implies that the K⁺ gradient is transiently reduced by the depolarizing voltage-clamp steps, resulting in a depolarizing shift of E_{K} .

The 'reversal potential' of I_{tail} is time dependent

Figure 6A illustrates experiments designed to measure the reversal potential of I_{tail} . A depolarizing step (200 ms,

+10 mV) was followed by 1 s steps to membrane potentials between -30 and -90 mV. The I_{tail} appeared to reverse at potentials between approximately -50 and -70 mV. Figure 6A shows currents recorded from the same myocyte before and after the addition of 0.25 mM BaCl₂. The main effect of this concentration of BaCl₂ was to block inward I_{tail} : this is particularly evident for the

Figure 4. I_{tail} is not generated by Na⁺-Ca²⁺ exchange

A, current produced by a 20 mV, 1 s voltage-clamp step in control Hepes-buffered solution. Right panel: current is shown on expanded time and current scales. B, current in the same cell, after removing external Na⁺ and Ca²⁺ (see Methods). Note that I_{tail} was unaffected by the removal of Na⁺ and Ca²⁺.



steps to -80 and -90 mV. In contrast, outward tail currents, especially those produced by the most positive steps in this protocol (i.e. -30 and -40 mV), were not changed significantly by $BaCl_2$. Figure 6B shows the BaCl₂-sensitive difference currents that were obtained by subtracting currents before and after the addition of BaCl₂. These difference currents, measured 'immediately' (2 ms) following the depolarizing steps, showed very pronounced inward rectification; their reversal potential was about -60 mV. The outward currents produced by steps positive to -60 mV were very small compared with the inward currents produced by the steps negative to -60 mV. This rectification is clearly apparent from an inspection of the difference currents plotted on an expanded time scale in Fig. 6B. It is also apparent from the records in Fig. 6B that the reversal potential of the difference currents changed with time; the current at -70 mV reversed direction from inward to outward about 50 ms after the end of the depolarizing step, while the current at -80 mV reversed about 100 ms after the depolarizing step. Figure 6C compares peak currentvoltage relationships for difference I_{tail} , measured 'immediately' (2 ms) after the end of the depolarizing step and 1 s after the end of the step. The current-voltage relationship measured 2 ms after the end of the depolarizing step reversed at -62.9 mV and had strong inward rectification, while the currents measured at 1 s reversed at -86.0 mV, and had a negative slope conductance at membrane potentials positive to about -70 mV. Figure 6C also shows the current-voltage relationship measured at 75 ms after the depolarizing step; the reversal potential was -74.5 mV. In experiments with seven different myocytes (including that in Fig. 6), the mean reversal potential of $BaCl_2$ -sensitive I_{tail} measured within 2 ms after the depolarizing step was -60.3 ± 1 mV, while the reversal potential measured at 1 s was -84.5 ± 0.8 mV. The latter value is very close to $E_{\rm K}$ under the recording conditions used in these experiments. We repeated a similar experiment measuring the reversal potential of I_{tail} using a higher extracellular K^+ concentration. Figure 6D shows a representative current-voltage relationship measured at 2 ms, 75 ms and 1 s after the depolarizing step in the presence of 20 mm external K⁺. For all three measurements, the current reversed at values very close to the nominal $E_{\rm K}$ (e.g. -49.5 mV with the present recording conditions). Data averaged from four different ventricular myocytes indicate that the currents reversed at -47.3 ± 2.5 ,



Figure 5. Ba^{2+} blocks I_{tail} and the inwardly rectifying K⁺ current, I_{K1} , in mouse ventricular myocytes

A, voltage-clamp currents recorded in control solution. The voltage-clamp protocol (inset) consisted of a depolarizing step to 20 mV (35, 160, 285 ms), and a hyperpolarizing step to -120 mV (100 ms). The inward current produced by the hyperpolarizing step is $I_{\rm K1}$. B, currents after the addition of 0.25 mM BaCl₂. Note that both $I_{\rm tail}$ and $I_{\rm K1}$ were abolished, but there was no significant effect on the outward currents.



Figure 6. Reversal potential of $BaCl_2$ -sensitive I_{tail}

A, voltage-clamp currents produced by a two-step protocol (right, inset), consisting of a 200 ms depolarizing step to +10 mV, followed by a series of 1 s steps to -30, -40, -60, -80 and -90 mV. In control conditions (left panel), the tail currents reversed near -60 mV. Large, inward currents were produced by steps to -80 and -90 mV. After the addition of 0.25 mM BaCl₂ (right panel), the large inward I_{tail} were almost completely abolished, but there was relatively little effect on the outward I_{tail} . *B*, BaCl₂-sensitive I_{tail} , obtained by subtracting currents in BaCl₂ from control currents. Voltage steps were from -30 to -90 mV, in 10 mV increments (-50 and -70 mV steps are not shown in *A*). The lower panel shows the initial parts of I_{tail} on an expanded time scale. Note that the currents for -70 and -80 mV are initially inward, but become net outward within about 100 ms after the end of the depolarizing step. *C*, current–voltage relationship for BaCl₂-sensitive I_{tail} measured 2 ms (O), 75 ms (**1**) and 1 s (**0**) after the end of the depolarizing step. *D*, current–voltage relationship for I_{tail} solution containing 20 mM K⁺. Smooth curves through each set of points are fifth-order polynomial regressions: reversal potentials for each set of points were obtained from these polynomials by interpolation (see text for values).

 -50.0 ± 1.3 and -50.9 ± 1.1 mV, when measured at 2 ms, 75 ms and 1 s, respectively. These data show clearly that in the higher external K⁺, the change in extracellular K⁺ concentration, and the time-dependent change in the apparent reversal potential of $I_{\rm tail}$ due to K⁺ efflux were minor.

Decay time course of $I_{ m tail}$ depends upon presence of $I_{ m K1}$

The data in Figs 5 and 6 suggest strongly that I_{tail} is generated by I_{K1} . These data can be explained by assuming that K⁺ accumulates in an extracellular space in which there is relatively restricted diffusion between this space and the 'bulk' external solution, and I_{tail} is produced by current flow through I_{K1} channels in the membranes of the restricted space. The decay of I_{tail} reflects the 'clearing' of this space of excess K⁺ by both the processes of diffusion, and transport of K⁺ into the cell cytosol through I_{K1} channels. This implies that if either of these major transport processes were impeded, the decline of K⁺ concentration in the restricted space would be slowed. Figure 7 shows the results of an experiment that tests this idea by 'blocking' I_{K1} by repolarizing to a membrane potential where I_{tail} is always very small. In this experiment, the potential was -60 mV; as shown in Fig. 6, at this membrane potential I_{tail} was always a small, outward current. The amount of excess K⁺ remaining at selected times after the end of a depolarizing step was measured using a brief (10 ms) 'test' step from -60 to -80 mV; this step activated a measurable I_{tail} at each time (Fig. 7A). The currents produced by this series of test steps were compared with I_{tail} produced by repolarization to the holding potential of -80 mV. Figure 7*B* compares the time course of I_{tail} at -80 mV with the time course of 10 ms test-step current amplitudes, applied between 0 and 480 ms after the end of a 300 ms depolarizing step of +10 mV. The decay of I_{tail} , and therefore the decline of accumulated K^+ , was very much slower when I_{K1} was inhibited by repolarizing to -60 mV. The best-fit, singleexponential function fitted to I_{tail} at -80 mV had a time constant of about 46 ms; the best-fit, single-exponential for the test-step currents had a time constant of 161 ms.



Figure 7. Time course of decay of I_{tail} depends upon current flow

A, upper panel, voltage-clamp currents produced by a 300 ms, +10 mV step, from a holding potential of -80 mV (inset). A large inward I_{tail} was produced on repolarization to -80 mV. Lower panel, series of superimposed currents produced by the same depolarizing step, but followed at different times by a 10 ms 'test' step to -80 mV, from a potential of -60 mV. Currents shown were produced by test steps that were applied between 0 and 400 ms after the end of the depolarizing step, in 50 ms increments (see inset). Vertical arrows indicate currents from test steps at 300, 350 and 400 ms. Time and current scales are the same for both panels. *B*, comparison of the time course of I_{tail} recorded at a membrane potential of -80 mV (continuous line), and the magnitude of the test step currents as a function of time after the end of the depolarizing step (\bullet). The dotted line through the test currents is a best-fit, double-exponential function, with time constants of 112.4 and 329.3 ms. The amplitudes of the fast and slow components were -1.03 and -0.38 nA, respectively. The horizontal arrow indicates the level of I_{tail} (at -80 mV) at 1 s after the repolarizing step; the double-exponential function was constrained to fit to this level.

A two-exponential function gave a better fit to the test current time course, with fast and slow time constants of 112.4 and 329.3 ms, respectively (Fig. 7*B*). Single exponential functions were fitted to both I_{tail} at -80 mV and test step currents from a potential of -60 mV. For each cell, the time constant of I_{tail} at -80 mV (50.7 ± 4.1 ms) was considerably smaller than of the test step (75.2 ± 3.2 ms P < 0.001; Table 1).

To confirm the importance of the inward K⁺ flux, we repeated the protocol presented in Fig. 7, comparing two different test potentials (e.g. -80 and -120 mV). The decline of the I_{tail} proceeded faster at the more hyperpolarized potential (37.9 ± 3.5 ms at -120 mV compared to 50.7 ± 4.1 ms at -80 mV, P = 0.03). According to the idea that the slow I_{tail} results from the slow clearing of accumulated K⁺ by simple diffusion and inward transport of K⁺ through I_{K1} channels, the decay of the currents produced by this series of test steps was faster for the test pulse that produced the larger inward current (62.9 ± 4.7 ms at -120 mV compared to $75.2 \pm 3.2 \text{ ms at} -80 \text{ mV}, P = 0.03$), hence carrying away more K⁺ per test pulse.

Table 1. Comparison of the time course of current decay at test pulses of -80 and -120 mV

	-80 mV		-120 mV	
	au (ms)	n	au (ms)	n
$I_{ m tail}$	50.7 ± 4.1	13	$37.9 \pm 3.5 *$	9
Train of pulses	$75.2 \pm 3.2 \ddagger \ddagger$	15	$62.9 \pm 4.7 * \ddagger$	14
Single pulse at				
various times	$120.5 \pm 11.4 \ddagger$	15	87.7 ± 7.7 *‡	9

Time constants obtained from a single exponential function for the decline of the tail current (I_{tail}) and of the currents obtained with the train of pulses or with a single pulse applied at various times. Data are given as means \pm s.e.m. *P = 0.03 vs. corresponding parameter at -80 mV; $\dagger P < 0.001$ vs. I_{tail} ; $\ddagger P < 0.01$ vs. pulse.

We then evaluated the contribution of a simple diffusion by using the following protocol. After a depolarizing pulse, the cells were held at -60 mV for various lengths of time, and we then applied one hyperpolarizing test pulse (at -80 mV or -120 mV) to estimate the amplitude of I_{tail} . Single exponential functions fitted to the currents produced with this protocol were compared to those



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obtained with the current elicited by the train of pulses (Table 1). For the hyperpolarizing steps of -80 mV, the time constant of the current decay obtained with the train of pulses $(75.2 \pm 3.2 \text{ ms})$ was considerably smaller than that of the current decay obtained with the single pulse applied at various times $(120.5 \pm 11.4 \text{ ms})$ P = 0.004). Similarly, at -120 mV, the time constant of current decline was also faster with the train of pulses $(62.9 \pm 4.7 \text{ ms})$ compared to that obtained with the single pulse applied at various times $(87.7 \pm 7.7 \text{ ms}, P = 0.004)$. The single-pulse protocol shows that clearing of K⁺ from the restricted space is slowed when inward flux is reduced, leaving simple diffusion as the major clearing mechanism. In addition, the decay of the test-pulse current amplitude obtained with the single pulse applied at various times was significantly faster at -120 mV(Table 1) compared to -80 mV. Again, this confirms the importance of the inward flux for the clearing of the K⁺ in the restricted space.

Immunolocalization of Kir2.1 in mouse ventricular myocytes

There is evidence that I_{K1} consists of different members of the Kir2 family. For instance, Kir2.1, Kir2.2 and Kir2.3

have all been found in guinea-pig cardiomyocytes (Liu et al. 2001) and Kir2.1 and Kir2.2 are both present in mouse and rat heart (Kubo et al. 1993; Ishihara & Hiraoka, 1994; Takahashi et al. 1994; Fiset et al. 1998; Nakamura et al. 1998; Leonoudakis et al. 2001; Zaritsky et al. 2001). However, in mammalian ventricles where I_{K1} is present, Kir2.1 has the highest level of transcript when compared to Kir2.2 or Kir2.3 (Brahmajothi et al. 1996; Liu et al. 2001). A recent study by Zaritsky et al. (2001) has provided clear evidence that the Kir2.1 gene is essential for I_{K1} in mouse ventricular myocytes. They demonstrated that ventricular myocytes isolated from the Kir2.1 knockout mouse lack a detectable I_{K1} , indicating that Kir2.1 is the major component of I_{K1} in mouse. In addition, Nakamura *et al.* (1998) have reported that antisense oligonucleotides against Kir2.1 partially suppress I_{K1} in rat ventricular myocytes, also demonstrating that Kir2.1 has an essential role in $I_{\rm K1}$ in rat ventricle.

The data presented so far suggest strongly that I_{tail} is generated by I_{K1} , and is probably due to K⁺ accumulation within the t-tubule system of isolated mouse ventricular myocytes. To provide additional evidence in support of this hypothesis, we investigated the localization of



Figure 9. No significant t-tubule localization of Kir2.1 in mouse a trial myocytes, consistent with the absence of $I_{\rm tail}$ in these cells

A, phase-contrast image of a mouse atrial cell. B-C, same cell as in A showing the immunofluorescence detection of Kir2.1 (B) and WGA (C). D, superimposition of the images presented in B and C, revealing little or no overlap (yellow) between Kir2.1 and WGA labelling in the t-tubules. E, phase-contrast image of the control atrial cell presented in F, showing that no immunofluorescence staining was observed when the primary antibody was omitted. G, family of currents from a voltage clamped, isolated mouse atrial myocyte. The voltage-clamp protocol is shown in the inset; the holding potential was -90 mV; 500 ms steps to potentials between -110 and +50 mV in 10 mV increments applied at 0.1 Hz.

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Kir2.1, which is the major component of I_{K1} in mouse heart (Kubo et al. 1993; Ishihara & Hiraoka, 1994; Fiset et al. 1998; Zaritsky et al. 2001). Immunofluorescent localization of Kir2.1 established clearly that $I_{\rm K1}$ is preferentially localized in the t-tubule system of the mouse ventricular myocytes. Figure 8A shows that Kir2.1 immunoreactivity was predominantly in a regularly spaced arrangement of transversely oriented bands along the entire length of the cell. Double labelling was performed with anti-Kir2.1 and WGA to confirm that Kir2.1 labelling was localized to the plasma membrane. WGA was used to visualize sarcolemmal and t-tubule membranes since it binds selectively to N-acetylglucosamine and sialic acid residues of the glycocalyx (Peters et al. 1979; Wright, 1984; Takeuchi et al. 2000; Sedarat *et al.* 2000). Figure 8B shows that WGA clearly stained the peripheral sarcolemma, the t-tubules and the intercalated disks. Superimposition of Kir2.1 and WGA staining showed important co-localization (as visualized by the vellow coloration; Fig. 8C). Pixels with a vellow coloration were observed on the t-tubule membranes, confirming the presence of Kir2.1 in these membrane structures.

Absence of I_{tail} in mouse atrial myocytes

Labelling of single atrial myocytes with a Kir2.1-specific antibody revealed immunofluorescence staining at the peripheral plasma membrane (Fig. 9A) without organized fluorescence in the cell interior. The WGA labelling in mouse atrial myocytes showed no well defined t-tubule system in these cells. In fact, small invaginations from the surface corresponding to the t-tubules were observed with WGA (Fig. 9B). Those striations were not significantly labelled with the anti-Kir2.1 antibody. The localization of Kir2.1 in atrial cells was confirmed by the pattern of distribution of co-localized Kir2.1 and WGA staining, which revealed significant co-localization only at the cell periphery (Fig. 9C). Consistent with these observations were the electrophysiological data that showed that no slowly decaying, inward currents followed the depolarizing voltage-clamp steps in isolated mouse atrial myocytes (Fig. 9G).

DISCUSSION

This study provides an explanation for the slow inward I_{tail} that follows membrane depolarization of voltageclamped mouse ventricular myocytes. The properties of I_{tail} suggest strongly that these currents result from the accumulation of K⁺ in a restricted extracellular space associated with the ventricular myocytes. K⁺ accumulation occurs during outward current flow through I_{K} channels in the restricted compartment; on repolarization to more negative potentials (-80 mV), current through the K⁺ channels is augmented because of the elevated extracellular concentration of K⁺ and the shift of E_{K} to depolarized values. The decay of I_{tail} reflects the decline of the elevated K⁺ back to its resting value and results from a combination of the passive diffusion of K^+ out of the restricted space plus the influx of K^+ into the cytosol through $I_{\rm K}$ channels.

All of the properties of I_{tail} can be accounted for by the following model:

(1) The increase in I_{tail} magnitude with increases in outward current measured at the end of the depolarizing step (Figs 1 and 2). Increased I_{tail} magnitude reflects an increased K⁺ accumulation, which would result from a larger outward current.

(2) The non-monotonic relationship between depolarizing step duration and I_{tail} magnitude (Fig. 3). The K⁺ concentration at the end of the depolarizing step depends on a balance between influx of K⁺ into the restricted space through delayed rectifier channels and K⁺ efflux via passive diffusion. Because of the time-dependent decline in outward K⁺ current during a depolarizing step, due to inactivation, accumulated K⁺ would initially increase, then decline or possibly reach a steady state as the duration of the depolarizing voltage-clamp step is increased.

(3) Lack of effect of external Na⁺ and Ca²⁺ on I_{tail} (Fig. 4). Neither of these ions have significant effects on I_{K1} .

(4) Simultaneous block of I_{tail} and I_{K1} by BaCl₂ (Fig. 5). Since I_{tail} is postulated to be generated by K⁺ influx through I_{K1} channels in the restricted space membranes, both of these currents must be blocked identically.

(5) The time-dependent change in the apparent reversal potential of I_{tail} (Fig. 6). This arises because the concentration of accumulated K⁺ in the restricted extracellular space, and hence E_{K} , declines with time after the end of the depolarizing voltage-clamp step.

(6) Slowed time course of decline of K⁺ concentrations when I_{K1} is removed (Fig. 7). Clearance of the restricted space occurs mainly via two processes: passive diffusion out of the space and inward current flow through I_{K1} channels within the space. Voltage clamping to a membrane potential where I_{K1} is small removes this contribution due to current flow, and only passive diffusion remains.

The accumulation phenomenon for K^+ occurs in part because the outward currents in mouse myocytes are very large. However, a similar phenomenon may occur in other rodent myocytes (e.g. rat). For example, there are slow, inward I_{tail} apparent in current records obtained in rat ventricular myocytes (Shimoni *et al.* 1995; Ward *et al.* 1995). It is likely that 'accumulation tail currents' would be small in myocytes from the hearts of bigger mammals, where the major outward current is due to the rapid component of the delayed rectifier (I_{Kr}), rather than large Ca²⁺-independent transient outward K⁺ current (I_{to}) and delayed rectifier currents, as in mouse and rat. The most likely structures in adult mouse ventricular myocytes that could act as restricted extracellular spaces for K⁺ accumulation are the sarcolemmal caveolae and the t-tubule system. Adult mouse ventricular myocytes have many surface caveolae and a particularly extensive and complex t-tubule system (Forbes & Sperelakis, 1973; Forbes *et al.* 1984). The t-tubule system of the adult mouse ventricle comprises about 3.2% of the volume of the myocytes. The surface density of the t-tubule system membrane of adult mouse ventricular myocytes is about $0.5 \ \mu\text{m}^2 \ \mu\text{m}^{-3}$: this implies that roughly 50% of the total membrane area of adult mouse ventricular myocytes is made up of t-tubule membranes.

Using a specific antibody against Kir2.1, we have provided molecular evidence that $I_{\rm K1}$ is predominantly localized in the t-tubule system of mouse ventricular myocytes. Moreover, in adult mouse atrial cells where the t-tubule membrane system is very limited, I_{tail} was not observed on repolarization to the holding potential. Results of immunofluorescence microscopy performed on mouse atrial myocytes, where the immunolocalization of Kir2.1 showed mainly staining to the cell periphery, are consistent with the electrophysiological data obtained in these cells. In combination, these observations provide evidence that I_{tail} is likely to result from K⁺ accumulation in the t-tubule system in mouse ventricular myocytes. This is the first report of localization of I_{K1} (Kir2.1) in the t-tubules of mouse ventricular myocytes. Taken together, the results of this study suggest strongly that I_{tail} in mouse ventricular myocytes is produced by the activation of inward I_{K1} in the t-tubules, resulting from K⁺ accumulation in these membrane structures.

Reports by other groups strongly support our findings. Recently, a study by Leonoudakis et al. (2001) showed that Kir2.2 is also localized in striated bands corresponding to t-tubules in rat ventricular myocytes. This finding is interesting considering that Zaritsky *et al.* (2001) have demonstrated that by removing the gene encoding Kir2.2, there was a 50% reduction in $I_{\rm K1}$, suggesting that Kir2.2 can also contribute to I_{K1} . Previous studies have shown that changes in I_{K1} magnitude closely correlate with the density of t-tubules in cardiomyocytes. Primary cultures of ventricular myocytes have been associated with a reduction of I_{K1} (Wahler, 1992; Schackow et al. 1995; Christé, 1999) and a parallel reduction in the density of t-tubules (Delcarpio *et al.* 1989). Christé has reported recently that cultured rabbit ventricular myocytes lose a large portion of I_{K1} and ATPsensitive K^+ current (I_{K-ATP}) conductance in parallel with a decay in cell membrane capacitance (Christé, 1999). These data suggest that a major part of I_{K1} and I_{K-ATP} are located in the t-tubules of rabbit ventricular myocytes, since a reduction in membrane capacitance has been attributed mainly to the disappearance of the t-tubules (Lipp et al. 1996; Mitcheson et al. 1996). In addition, using immunofluorescence and immunoelectron microscopy, Takeuchi et al. (2000) have demonstrated that Kv4.2,

which generates much $I_{\rm to}$ in cardiomyocytes, is expressed mainly in the t-tubules in rat ventricular myocytes. Other studies have shown a preferential localization of ion channels to the t-tubule membrane. For instance, during the postnatal development of rabbit ventricular myocytes, it has been reported that there is a close temporal correlation between formation of the t-tubules and expression of dihydropyridine receptors (Sedarat et al. 2000). T-tubule membrane systems are well developed in mammalian ventricular myocytes and several previous studies have indicated that ion accumulation can occur within the t-tubular system in the isolated ventricular myocytes of other species. Yasui et al. (1993) recorded inward I_{tail} in single guinea-pig ventricular myocytes after treatment of the cells with nicorandil, an activator of $I_{\text{K-ATP}}$, which induced large outward currents in these myocytes. Yasui *et al.* interpreted their results using a similar model to that presented above; outward $I_{\text{K-ATP}}$ during a depolarizing step resulted in K⁺ accumulation in t-tubules, and the inward I_{tail} produced on repolarization due to I_{K1} in the t-tubule membranes. Guinea-pig ventricular myocytes have an extensive t-tubule system, with a volume fraction of 2.5-3.2% and a surface area density of about 0.4 $\mu m^2 \mu m^{-3}$ (Forbes & Van Niel, 1988), similar to the mouse ventricular myocytes. Shepherd & McDonough (1998) also demonstrated slow diffusion in the t-tubules of guinea-pig ventricular myocytes. They rapidly increased [Ca²⁺] around guinea-pig ventricular myocytes while measuring the time course of the increase in the amplitude of voltage-dependent Ca^{2+} currents (I_{Ca}). $I_{\rm Ca}$ increased in two phases: 30–40% of the current change occurred with a time constant of about 20 ms, while the remainder of the change occurred 10-fold more slowly. These authors suggested that the slow component reflected the slow equilibration of [Ca²⁺] between the bulk solution and the lumen of the t-tubules. Yao et al. (1997) performed similar rapid solution-change experiments using adult rabbit and rat ventricular myocytes. Following a rapid (4 ms) removal of external Ca^{2+} , I_{Ca} declined slowly in both rabbit and rat ventricular myocytes, although the rate of decline of the current in rat myocytes ($t_{90} = 910$ ms, the time required for 90% decline) was considerably slower than that in rabbit ventricular myocytes ($t_{90} = 241$ ms). Changes in K⁺ concentration appeared to equilibrate much more rapidly with both types of cell. The time to reach a stable membrane potential after a rapid change in external K⁺ occurred with a t_{90} of 64 ms for rabbit cells and 137 ms in rat myocytes. Yao et al. (1997) suggested that sarcolemmal structures like t-tubules, surface caveolae and negatively charged proteins all contribute to the slowed diffusion of cations in myocardial cells.

In conclusion, our data, along with data from other investigators, suggest that K^+ channels are localized preferentially in the t-tubules. Considering that cardiac K^+ channels play very important roles in regulating both action potential duration and excitation-contraction coupling and since the t-tubule is the major site of excitation–contraction coupling in ventricular cells, this t-tubule localization is not totally unexpected. Additional studies exploring the physiological consequence of the t-tubule localization of K^+ channels are required and may provide important insights into the functional role of cardiac K^+ channels under physiological and pathological conditions.

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