

RESEARCH PAPER

High potency inhibition of hERG potassium channels by the sodium-calcium exchange inhibitor KB-R7943

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BACKGROUND AND PURPOSE

KB-R7943 is an isothiourea derivative that is used widely as a pharmacological inhibitor of sodium–calcium exchange (NCX) in experiments on cardiac and other tissue types. This study investigated KB-R7943 inhibition of hERG (human ether-à-qo-qo-related gene) K⁺ channels that underpin the cardiac rapid delayed rectifier potassium current, I_{Kr}.

EXPERIMENTAL APPROACH

Whole-cell patch-clamp measurements were made of hERG current (I_{hERG}) carried by wild-type or mutant hERG channels and of native rabbit ventricular I_{Kr}. Docking simulations utilized a hERG homology model built on a MthK-based template.

KEY RESULTS

KB-R7943 inhibited both I_{hERG} and native I_{Kr} rapidly on membrane depolarization with IC_{50} values of ~89 and ~120 nM, respectively, for current tails at -40 mV following depolarizing voltage commands to +20 mV. Marked I_{hERG} inhibition also occurred under ventricular action potential voltage clamp. I_{hERG} inhibition by KB-R7943 exhibited both time- and voltage-dependence but showed no preference for inactivated over activated channels. Results of alanine mutagenesis and docking simulations indicate that KB-R7943 can bind to a pocket formed of the side chains of aromatic residues Y652 and F656, with the compound's nitrobenzyl group orientated towards the cytoplasmic side of the channel pore. The structurally related NCX inhibitor SN-6 also inhibited I_{hERG} , but with a markedly reduced potency.

CONCLUSIONS AND IMPLICATIONS

KB-R7943 inhibits I_{hERG}/I_{Kr} with a potency that exceeds that reported previously for acute cardiac NCX inhibition. Our results also support the feasibility of benzyloxyphenyl-containing NCX inhibitors with reduced potential, in comparison with KB-R7943, to inhibit hERG.

Abbreviations

AP, action potential; AV, atrioventricular; DMEM, Dulbecco's minimum essential medium; $n_{\rm H}$, Hill coefficient; hERG, human Ether-à-go-go-Related Gene; I_{hERG}, hERG potassium channel ionic current; I_{Kr}, rapid delayed rectifier K⁺ current; k, slope factor for voltage-dependent activation or inactivation relations; KB-R7943, 2-[2-[4-(4-nitrobenzyloxy)phenyl] ethyl]isothiourea methane sulphonate; NCX, sodium calcium exchange; SEA0400, 2-[4-[(2,5-difluorophenyl)methoxy] phenoxy]-5-ethoxyaniline; SN-6, 2-[[4-[(4-nitrophenyl)methoxy]phenyl]methyl]-4-thiazolidinecarboxylic acid ethyl ester; TRPC, canonical transient receptor potential channel; $V_{\rm m}$, membrane potential; $V_{0.5}$, half maximal activation or inactivation voltage



Introduction

Sodium–calcium exchange (NCX) proteins are expressed in many tissue types and are recognized to be important for cellular Ca²⁺ ion homeostasis (Dipolo and Beauge, 2006). In the heart, the cardiac NCX isoform (NCX1) contributes both to Ca²⁺ homeostasis and to electrogenesis due to its stoichiometry, and, in principle, partial NCX inhibition may be beneficial in some cardiac pathologies (Philipson and Nicoll, 2000; Shigekawa and Iwamoto, 2001; Watanabe *et al.*, 2006; Toth *et al.*, 2009; Zhang and Hancox, 2009). At present, however, there are no high-affinity selective NCX inhibitors available for clinical use.

KB-R7943 (2-[2-[4-(4-nitrobenzyloxy)phenyl]ethyl] isothiourea methane sulphonate) was developed as the first NCX-specific inhibitor (Iwamoto et al., 1996; Watanabe et al., 2006). It was initially reported to inhibit preferentially outward over inward cardiac NCX current (I_{NCX}) when the NCX was activated unidirectionally (Watano et al., 1996), but with bi-directional NCX activation, the compound was found to inhibit both NCX modes similarly, with IC50 of ~1 μM (Kimura et al., 1999). KB-R7943 was reported also to affect cardiac Na+, Ca2+ and inwardly rectifying K+ currents with higher IC₅₀ values (Watano et al., 1996; Tanaka et al., 2002). Although KB-R7943 transpires not to be entirely selective for the NCX, its ease of use and effectiveness as an NCX blocker means that it has been employed widely as a pharmacological tool for the study of the cardiac NCX in physiological and pathophysiological conditions (Amran et al., 2003; Toth et al., 2009).

The rapid delayed rectifier K⁺ current (I_{Kr}) plays an important role in cardiac action potential (AP) repolarization (Sanguinetti and Mitcheson, 2005; Sanguinetti and Tristani-Firouzi, 2006). KCNH2-encoded hERG (human Ether-à-go-go-Related Gene) protein forms the pore-forming subunit of I_{Kr} channels (Sanguinetti et al., 1995; Trudeau et al., 1995; Alexander et al., 2011). Due to structural features of the channel, hERG is able to interact with chemically and therapeutically diverse drugs that are associated with the acquired Long QT Syndrome (Sanguinetti and Mitcheson, 2005; Sanguinetti and Tristani-Firouzi, 2006). Consequently, it is customary for all promising drug candidates now to be tested for hERG channel inhibitory activity (ICH, 2005; Shah, 2005; Hancox et al., 2008). Despite the long use of KB-R7943 in experiments utilizing cardiac cell, tissue and intact heart preparations and the hERG channel's high susceptibility to pharmacological blockade, there is currently no information available regarding effects of the compound on hERG channel current (I_{hERG}). One study has reported partial inhibition of guinea-pig ventricular outward delayed rectifier K⁺ current (I_K) by a single concentration (10 µM) of KB-R7943 (Tanaka et al., 2002). However, guinea-pig I_K is a composite current comprised of both I_{Kr} and the slow delayed rectifier current, I_{Ks} (Sanguinetti and Jurkiewicz, 1990), and the investigation by Tanaka et al. (2002) neither determined concentration-dependence of the compound's effect on composite I_K, nor identified any specific inhibitory effect of KB-R7943 on I_{Kr} as opposed to I_{Ks}. In a very recent study of the role of NCX in spontaneous activity of cells from the rabbit atrioventricular (AV) node, KB-R7943 led to membrane potential depolarization at a concentration (5 µM) that inhibited completely inward NCX current

(Cheng *et al.*, 2011). This seemingly paradoxical effect might, in principle, be accounted for by an inhibitory action on an AV nodal I_{Kr} , as this is a major repolarizing current in these cells, which lack I_{Ks} (Mitcheson and Hancox, 1999; Sato *et al.*, 2000). Given the lack of direct evidence either for or against KB-R7943 actions on hERG/ I_{Kr} , the present investigation was conducted to determine whether or not KB-R7943 is able to inhibit hERG and native cardiac I_{Kr} and if so to characterize the underlying mechanism. The results: (i) establish KB-R7943 as a potent I_{hERG}/I_{Kr} inhibitor; (ii) provide insight into the nature of interactions between the KB-R7943 molecule and the hERG channel pore; and (iii) demonstrate that the structurally related NCX inhibitor SN-6 inhibits I_{hERG} much less potently than does KB-R7943.

Methods

Maintenance of cell lines stably expressing hERG

 I_{hERG} measurements were made from HEK 293 cells stably expressing wild-type (WT) hERG [donated by Professor Craig January; (Zhou et al., 1998)] or the hERG S6 mutations Y652A and F656A (Milnes et al., 2003a), or transiently expressing the hERG pore helix mutation S624A (transfections were conducted using lipofectamine LTX using CD8 as a transfection marker; see Zhang et al., 2010). Cells were passaged using enzyme-free cell dissociation solution (Millipore, Watford, UK) and were plated onto glass coverslip shards in 40 mm Petri dishes (Zhang et al., 2010). Dulbecco's minimum essential medium with Glutamax-1 (DMEM; Gibco, Paisley, UK) was supplemented with 10% fetal bovine serum, together with 50 μg·mL⁻¹ gentamycin and 400 μg·mL⁻¹ geneticin (G418; Gibco) (Zhang et al., 2010). Cells were incubated at 37°C (5% CO₂) for a minimum of 1 day before electrophysiological recording.

Isolation of rabbit ventricular myocytes

Male New Zealand White rabbits (\sim 2–3 kg) were killed in accordance with the UK Home Office Animals (Scientific Procedures) Act, 1986; hearts were rapidly removed, and ventricular myocytes were isolated from the right ventricle using previously described methods (Hancox *et al.*, 1993; Howarth *et al.*, 1996). Before use, cells were kept at 4°C in Kraft–Brühe medium (Isenberg and Klockner, 1982). For the native I_{Kr} data presented in Figure 7, each drug concentration was tested on cells from two or more hearts.

Electrophysiological recording

Cells were superfused in an experimental chamber mounted on the stage of an inverted microscope (Eclipse TE2000-U, Nikon) with a Tyrode's solution containing (mM) 140 NaCl, 4 KCl, 2 CaCl₂, 1 MgCl₂, 10 glucose, 5 HEPES (titrated to pH 7.45 with NaOH). For experiments in which a high [K⁺]_e (94 mM) was used, KCl was substituted for NaCl in this solution to attain the total desired [K⁺]. Experimental solutions were applied using a home-built, warmed solution delivery system capable of changing the bathing solution surrounding a cell in <1 s (Levi *et al.*, 1996). Patch pipettes (A-M Systems, Sequim, WA, USA) were pulled using a Narishige vertical

electrode puller (Narishige PP-83) and heat-polished to a final resistance of 2–3 $M\Omega$ (Narishige MF-83). The pipette solution contained (mM) 130 KCl, 1 MgCl₂, 5 EGTA, 5 MgATP, 10 HEPES (titrated to pH of 7.2 with KOH). The pipette and external solutions were identical for I_{hERG} and I_{Kr} measurements. Series resistance values typically lay between 4 and 7 $M\Omega$ and were compensated by ~70% or more. Measurements were made at 35–37°C. Action potential voltage clamp (AP clamp) experiments employed a ventricular AP waveform identical to that used in prior experiments from our laboratory (e.g. Zhang $et\ al.$, 2010).

KB-R7943 and SN-6

KB-R7943 and SN-6 (Tocris, Bristol, UK) were dissolved in dimethyl sulphoxide (DMSO) to produce stock solutions of 1 μ M-100 mM, which were kept at -20°C. The stock solutions were diluted, as necessary, with Tyrode's solution to give a final DMSO concentration in experimental superfusate of 1/1000.

Docking simulations

KB-R7943 was docked into open state hERG pore homology models constructed on a MthK-based template (Witchel *et al.*, 2004). Docking was performed using the Flexidock module of Sybyl. Docking runs were set up with at least 40 different configurations of the drug within the hERG pore binding pocket comprising the amino acid residues T623, S624, V625, G648, Y652, F656, S660 and all atoms within 5 Å. Rotation of side chain bonds within the residues listed was allowed during the docking. 60 000 generations of the genetic algorithm were calculated in each docking run.

Analysis

Data are presented as mean \pm SEM. Statistical analyses were performed using Microsoft Excel (Microsoft Corporation) and Prism (GraphPad Software Inc.), whilst fits to particular data

sets were performed either using Prism or Clampfit of pClamp 10.0 (Axon instruments, Molecular Devices). Comparisons were made using one-way ANOVA, paired t-test or unpaired t-test as appropriate; P < 0.05 was taken as significant.

Concentration–response data were fitted with a Hill equation of the form:

Fractional block =
$$1/(1+10 \wedge ((\text{LogIC}_{50}-X)*n_{\text{H}}))$$
 (1)

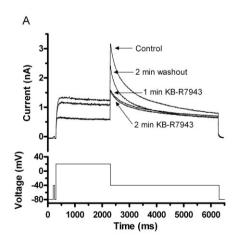
where Fractional block refers to the degree of inhibition of I_{hERG} or I_{Kr} by a given concentration of KB-R7943 (X, the logarithm of concentration); IC_{50} is [KB-R7943] producing half-maximal inhibition of I_{hERG} or I_{Kr} , and n_{H} is the Hill coefficient for the fit.

For voltage-dependent activation and inactivation, the half-maximal activation or inactivation voltage $(V_{0.5})$ and slope factor (k) values were derived from fits to the relevant data with standard Boltzmann functions. Time courses of current activation and deactivation were determined respectively by fitting data with single or bi-exponential functions.

Results

Concentration-dependent inhibition of I_{hERG} by KB-R7943

A standard voltage protocol (lower trace of Figure 1A), as used in previous studies of I_{hERG} pharmacology from our laboratory (e.g. McPate *et al.*, 2008; Zhang *et al.*, 2010), was applied from a holding potential of -80 mV in initial experiments to investigate I_{hERG} inhibition by KB-R7943. The protocol incorporated a brief (50 ms) pre-pulse from -80 to -40 mV before the +20 mV test command, in order to quantify the instantaneous current at -40 mV. Comparison between this instantaneous current and the peak outward I_{hERG} tail amplitude on repolarization to -40 mV enabled the accurate measurement



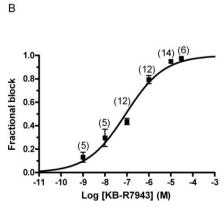


Figure 1

Concentration-dependent inhibition of I_{hERG} by KB-R7943. (A) Upper panel shows representative I_{hERG} traces in normal Tyrode's solution, after 1 and 2 min exposure to 100 nM KB-R7943 and after 2 min of washout. Lower panel shows voltage protocol (start-to-start interval of 12 s). (B) Concentration–response relation for the action of KB-R7943 on I_{hERG} tails. Tail current amplitude was measured as the difference between peak tail current and the current level attained during the brief pre-pulse to -40 mV. Numbers of replicates at each concentration are given in parentheses. Data were fitted with Equation 1 to give IC_{50} and Hill coefficient (n_H) values mentioned in the Results text.



of I_{hERG} tail amplitude (McPate et al., 2008; Zhang et al., 2010). Figure 1A shows representative traces illustrating the effect of 100 nM KB-R7943. The compound produced a rapid reduction in both end-pulse current and tail current amplitudes, with inhibition complete within ~1-2 min (current amplitudes in KB-R7943 were similar at 1 and 2 min of drug exposure). The extent of inhibition of I_{hERG} tails recorded at -40 mV was quantified for a range of concentrations between 1 nM and 30 µM in order to construct a concentrationresponse relation (Figure 1B). A fit to the resulting data with Equation 1 yielded an estimated IC₅₀ value of 88.6 \pm 25.7 nM and an $n_{\rm H}$ value for the fit of 0.50 \pm 0.06. However, the IC₅₀ derived from the line of best fit to the data may slightly overestimate potency, as the experimentally derived extent of I_{hERG} tail inhibition by 100 nM KB-R7943 with this protocol was 43.6 \pm 2.6% (n = 12), suggestive of an IC₅₀ closer to 100 nM. The inhibitory effect of KB-R7943 was partially reversible, with recovery to $91.8 \pm 4.1\%$ of control amplitude following exposure to 1 nM KB-R7943 (n = 5) and to 40.5 \pm 6.5% of control amplitude following exposure to 30 µM (n = 6) of the compound.

Voltage-dependence of I_{hERG} inhibition by KB-R7943

The protocol used to investigate voltage-dependence of inhibition of I_{hERG} by KB-R7943 is shown in the lower traces of Figure 2A (control) and 2B (in 100 nM KB-R7943). From a holding potential of -80 mV, 2 s depolarizing voltage commands were applied to a range of test potentials between -40 and $+40\,\text{mV}$, and tail currents were then recorded during a 4 s repolarization step to -40 mV (McPate et al., 2008; Zhang et al., 2010); 100 nM KB-R7943 appeared to produce a dual effect on IhERG. At potentials of approximately -10 mV and more positive to this, the compound produced a marked inhibition of I_{hERG} both during and following the depolarizing voltage command (Figure 2A and B); however, at more negative potentials, this was not evident, and at -30 and -40 mV, an increase in elicited I_{hERG} was seen (Figure 2A and B). Figure 2Ci and Di shows mean I–V relations for end-pulse (Figure 2Ci) and tail (Figure 2Di) currents. The data in Figure 2Di were used to derive halfmaximal activation parameters: for the pooled mean data plot shown, the derived $V_{0.5}$ and k values were, respectively, in control -15.0 and 6.9 mV, and in KB-R7943 these were -24.6 and 6.6 mV. When fits were made to data from individual cells, the mean $V_{0.5}$ and k values then obtained were: control -14.1 ± 2.3 and 6.7 ± 0.5 mV (n = 6), and following KB-R7943, these were $-24.9 \pm 2.6 \text{ mV}$ (P < 0.01 vs. control) and 6.2 \pm 0.5 mV (P > 0.05 vs. control). Figure 2Cii and Dii shows, respectively, plots of mean fractional block of endpulse current (Figure 2Cii) and tail current (Figure 2Dii). Both plots indicate marked voltage-dependence of the observed effect (P < 0.01 for each; one-way ANOVA across the potential range from -40 to +40 mV). In Figure 2Dii, activation curves for I_{hERG} are also plotted. The range of steepest change in fractional inhibition coincides with the steep part of the activation curves. The leftward shift in activation with KB-R7943 is likely to account for the increase in current seen at negative voltages in the membrane potential range examined.

Effects of KB-R7943 on time-dependent activation and deactivation of I_{herg}

An 'envelope of tails' protocol was used to investigate the development of inhibition of I_{berg} by KB-R7943 with time following membrane depolarization (e.g. Milnes et al., 2003a; Zhang et al. 2010). The protocol is shown as the lower set of traces in each of Figure 3A and B, with representative sets of current traces shown above the voltage protocol. As is characteristic of IhERG with this protocol (e.g. Milnes et al., 2003a; Zhang et al., 2010), in both control and drug-containing solutions, I_{hERG} tail amplitude increased progressively as the duration of the depolarizing command to +20 mV increased, with current amplitude in 100 nM KB-R7943 smaller than that for the corresponding command pulse in control solution. Notably, even with relatively brief depolarizations, current was suppressed by KB-R7943. Figure 3C shows a plot of mean time course of development of the currents in control and KB-R7943, with monoexponential fitting to derive activation time constant (τ) values. Fits to the pooled mean data yielded activation τ values of ~59 and ~90 ms in control and KB-R7943, whilst the mean values derived from fits to data from each experiment yielded mean τ values of 60.7 \pm 7.7 ms (control) and 99.7 \pm 15.7 ms (KB-R7943; n = 8; P < 0.01 vs. control). Figure 3D shows a plot of fractional inhibition of I_{bERG} against corresponding test pulse durations, focusing on the first ~110 ms of the protocol. There was little difference in inhibition at the different time points [ANOVA analysis across the full range of test pulse durations (up to 810 ms) showed no significant differences; P > 0.05; n = 8]. Thus, I_{hERG} block by KB-R7943 was evident rapidly on membrane depolarization. Effects of the agent on current deactivation were quantified by assessing deactivation time-constant (τ) values for tail currents elicited in control and 100 nM KB-7943, by the protocol shown in Figure 1A: the derived τ values were τ_{fast} of 294 \pm 24 ms and τ_{slow} of 1969 \pm 147 ms in control and τ_{fast} of 350 \pm 26 ms and τ_{slow} of 2332 \pm 122 in KB-R7943 (n = 6; P < 0.05 for both τ_{fast} and τ_{slow}). In summary, KB-R7943 slowed the deactivation time course of IhERG; it also produced a modest slowing of I_{hERG} activation at a test voltage (+20 mV) at which full I_{hERG} activation could be attained in both control and drug conditions (Figure 2D).

KB-R7943 and I_{hERG} inactivation

Effects of KB-R7943 (100 nM) on the voltage-dependence of I_{hERG} inactivation were assessed using the 'availability' protocol shown in Figure 4A. This was comprised of an initial depolarizing step to +40 mV to activate and inactivate I_{hERG}, followed by 2 ms repolarization steps to potentials ranging from +50 to -140 mV, to relieve channel inactivation to differing extents; the membrane potential was then stepped back to +40 mV for 100 ms, and the amplitude of resulting current transients was used to assess IhERG availability (cf. Milnes et al., 2003a; McPate et al., 2005). Figure 4B shows mean availability plots, from which inactivation $V_{0.5}$ values were obtained for control solution and following exposure to KB-R7943. The mean $V_{0.5}$ values derived from fits to data from each of eight cells were -48.4 ± 3.7 mV in control and -57.0 \pm 6.1 mV in KB-R7943 (P > 0.05), whereas the k values were 26.9 ± 2.5 and 24.1 ± 2.3 mV, respectively (P > 0.05). Thus, there was no statistically significant effect of KB-R7943 on the

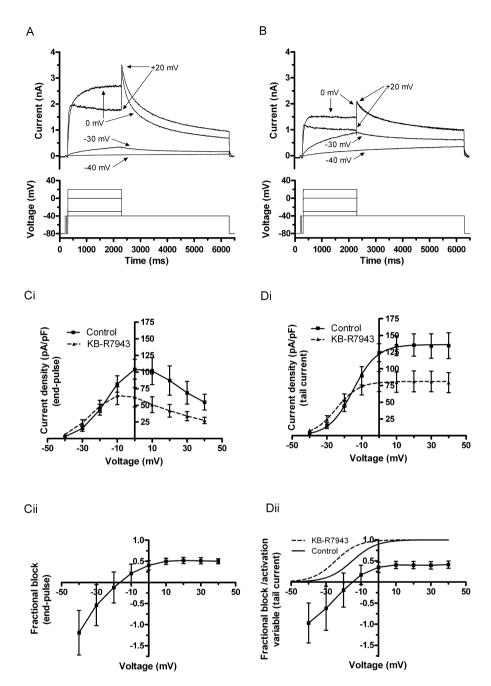


Figure 2

Voltage-dependence of I_{hERG} inhibition by KB-R7943. Upper traces show representative I_{hERG} records in control (A) and in the presence of 100 nM KB-R7943 (B). Lower traces show corresponding voltage steps in the experimental protocol. Currents were evoked by a series of 10 mV increments of step depolarizations between -40 and +40 mV from a holding potential of -80 mV. However, for clarity of display, only selected steps are shown. (Ci) Mean I–V relation for end-pulse currents in control and in the presence of 100 nM KB-R7943 (n = 6 cells). (Cii) Mean fractional block of end-pulse currents against voltage. (Di) Mean I–V relation for I_{hERG} tails in control and in the presence of KB-R7943 (n = 6 cells). Data were fitted with a Boltzmann equation to give $V_{0.5}$ and k values in the Results text. (Dii) Corresponding plot of mean fractional block of tail currents. Superimposed on this plot are continuous plots describing voltage-dependent activation of I_{hERG} in control and KB-R7943.

voltage-dependence of inactivation. The effect of promoting increased I_{hERG} inactivation on the inhibitory action of KB-R7943 was assessed using the protocol shown in Figure 4C (lower trace; cf. Ridley *et al.*, 2003). From a holding potential of –80 mV, membrane potential was stepped first to 0 mV for

3 s, followed by 4 s of depolarization to +80 mV and then a return to 0 mV. In both control and 100 nM KB-R7943, depolarization to 0 mV elicited sizeable I_{hERG} , which was reduced when membrane potential was further stepped to +80 mV (due to more channels becoming inactivated) and which



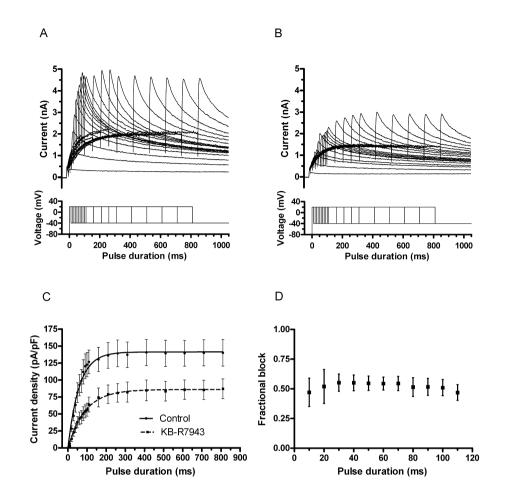


Figure 3

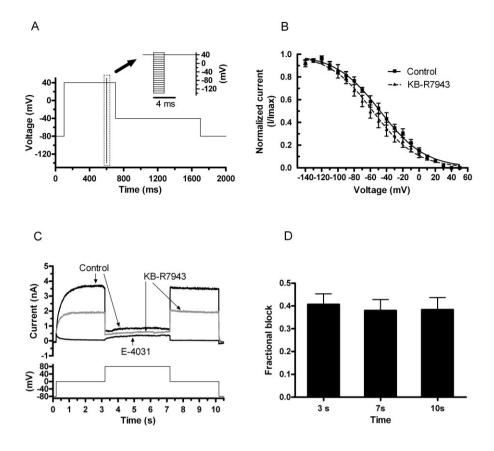
Time course of I_{hERG} activation and KB-R7943. (A, B) Upper traces show representative currents elicited by envelope of tails protocol shown as lower traces in each panel: (A) currents in control and (B) during exposure to KB-R7943. (C) Mean plots of current amplitude against depolarizing pulse duration (n = 8), fitted with a single-exponential equation to yield τ values given in the Results text. (D) Plot of mean level of fractional block, focusing on the first 110 ms of the protocol. There was no statistically significant time-dependent difference in this (one-way ANOVA).

then increased on the step back to 0 mV. Currents were sampled just before the end of the first step to 0 mV, the step to +80 mV and of the return to 0 mV. Measurements were made in control solution, during a single application of the protocol after a 2 min equilibration period in KB-R7943 and after application of a high concentration (5 μ M) of the selective hERG/I_{Kr} inhibitor E-4031 to allow subtraction of residual leak current. Figure 4D compares mean levels of fractional block of I_{hERG} at the three time points (n=7), showing there to be no significant difference between these (P>0.05; ANOVA; cf. Ridley *et al.*, 2003).

Effect of mutation of S6 aromatic amino acid residues on the action of KB-R7943

For a number of high-affinity hERG channel inhibitors, I_{hERG} blockade involves drug binding within the inner cavity at a site involving aromatic residues (Y652 and F656) that are rendered accessible on channel gating (e.g. Mitcheson *et al.*, 2000; Sanguinetti and Mitcheson, 2005; Sanguinetti and Tristani-Firouzi, 2006). Alanine mutants of these residues (Y652A and F656A) were therefore employed in order to explore further I_{hERG} inhibition by KB-R7943 (cf. Milnes *et al.*,

2003a; Ridley et al., 2004). Figure 5A shows data for Y652AhERG. The same voltage protocol was employed as for WT I_{hERG} in Figure 1. As indicated in Figure 5Ai and Aii, 100 nM KB-R7943 produced markedly less inhibition of Y652A-hERG in comparison with WT-hERG. Three KB-R7943 concentrations (100 nM, 1 µM and 10 µM) were tested on Y652AhERG, and mean data from these experiments are shown in Figure 5B (with the concentration-response relation for WT I_{hERG} shown overlaid). A fit to the data with Equation 1 gave an estimated IC50 for Y652A-hERG of 1.13 \pm 0.14 μM , with an $n_{\rm H}$ of 1.08 \pm 0.16. Thus, the potency of $I_{\rm hERG}$ inhibition by KB-R7943 was ~13-fold lower for Y652A-hERG than for WT-hERG. The F656A-hERG mutant is comparatively poorly expressing (Mitcheson et al., 2000; Milnes et al., 2003a), and so, as in previous studies from our laboratory (e.g. Milnes et al., 2003a; Ridley et al., 2004), this mutant was studied using a high [K⁺]_e (94 mM), employing the voltage protocol shown in Figure 5Ciii. Additional experiments were performed to assess inhibition of WT IhERG under these conditions (Figure 5Ci and D). The estimated IC₅₀ for WT I_{hERG} was $1.20 \pm 0.02 \,\mu\text{M}$, with an n_{H} of 0.75 ± 0.01 , indicating reduced potency of the compound against WT I_{hERG} with raised [K⁺]_e



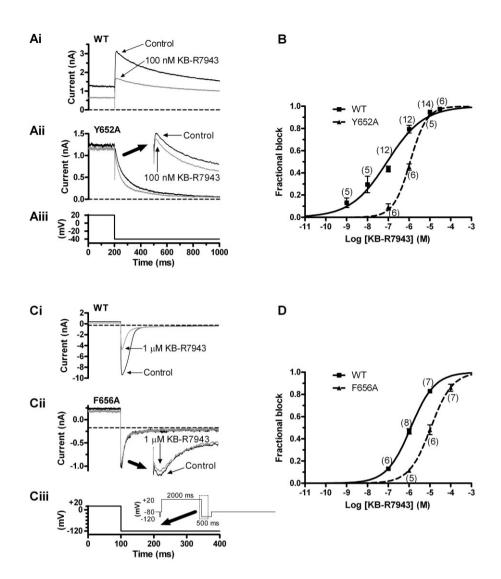
 I_{hERG} inactivation and KB-R7943. (A) Protocol used to investigate I_{hERG} 'availability'. From -80 mV, the membrane potential was stepped to +40 mV for 500 ms; this was then followed by 2 ms repolarization steps to potentials ranging from -140 to +50 mV. Membrane potential was then stepped back to +40 mV for 100 ms (inset shows portion of protocol encompassing the transition to and from repolarizing steps). The amplitude of current transients elicited by the second step to +40 mV was used to assess I_{hERG} availability. (B) Plots against voltage of normalized current transient amplitude in control and in 100 nM KB-R7943 (n = 8 cells); a Boltzmann equation was used to derive the $V_{0.5}$ and k values shown in the Results text. (C) Three-step protocol used to assess the effect of promoting I_{hERG} inactivation on the action of KB-R7943. Protocol is shown as lower trace; upper traces show representative currents in control, in the presence of 100 nM KB-R7943 and following exposure to 5 μM E-4031. (D) Mean level of fractional block at three time points during the protocol (n = 7). The three time points are given with reference to the start of the acquisition period (200 ms before the initial depolarization to 0 mV).

(cf. Wang et al., 1997; Ridley et al., 2004; Barrows et al., 2009); 1 μM KB-R7943 produced comparatively little inhibition of F656A I_{hERG} under these conditions (compare Figure 5Ci and Cii); the estimated IC50 for inhibition of F656A-hERG was 11.09 \pm 0.23 μ M (with an $n_{\rm H}$ of 0.84 \pm 0.01), ~9-fold higher than its WT control (Figure 5D). In order to investigate whether the compound may also be able to interact with the base of the hERG channel pore helix (cf. Mitcheson et al., 2000), additional experiments were performed using the S624A mutation, using the same recording conditions and protocol as used to study outward WT I_{hERG} (Figure 6A and B). Figure 6A shows effects of 100 nM KB-R7943 on S624A I_{hERG}; the compound reduced the tail current by ~30%. Figure 6B shows concentration-response data for the S624A mutant: the derived IC₅₀ was 216.7 \pm 31 nM ($n_{\rm H}$ = 1.13 \pm 0.17), \sim 2.4-fold that for WT I_{hERG.} Considered together, these observations implicate inner cavity binding interactions in the I_{hERG} inhibitory effect of KB-R7943.

In order to pursue further the nature of the interactions between KB-R7943 and hERG, ligand docking simulations

were conducted in which interactions between the drug and channel were investigated in an open-channel configuration homology model (see Methods). Low energy binding configurations for KB-R7943 were dominated by extended conformations in which the drug lay approximately parallel to the pore axis with the positively charged thiourea group oriented towards the selectivity filter and the nitrobenzyl group oriented towards the cytoplasmic side of the pore (Figure 6C). The drug bound within a pocket comprising the side chains of Y652 and F656 enabling stacking interactions between aromatic rings on the drugs and the aromatic side chains of Y652 and F656; this is consistent with a significant contribution to binding with these residues and is in good agreement with the Y652A and F656A data shown in Figure 5. In low-energy configurations, the positively-charged thiourea group of KB-R7943 was also seen either to extend close to the ring of hydroxyl side chains of S624 or to locate near the C-terminus of a pore helix and to make a hydrogen bond with the backbone carbonyl group of G648. An additional thiourea hydrogen bond with the S624 side chain of





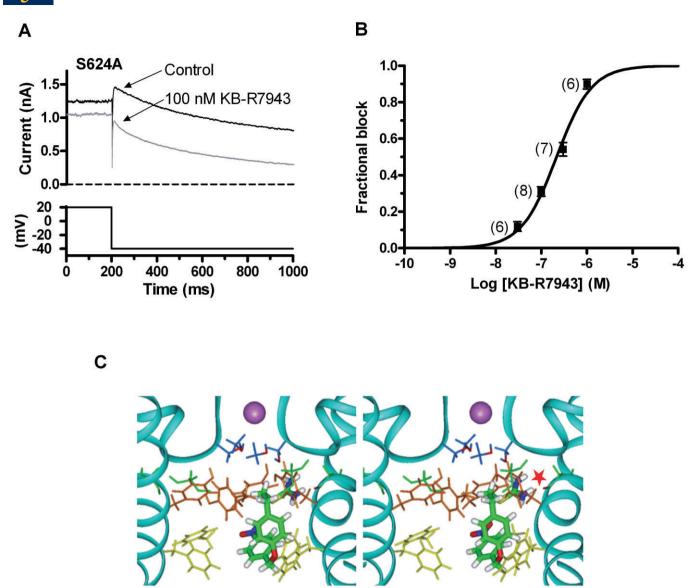
Effects of the Y652A and F656A mutations on the action of KB-R7943. (A) The traces in panels Ai and Aii show, respectively, WT I_{hERG} and Y652A I_{hERG} elicited in the absence and presence of 100 nM KB-R7943. The corresponding voltage protocol is shown in panel Aiii. (B) Mean data indicating effects of three concentrations of KB-R7943 on Y652A I_{hERG} , fitted by Equation 1 to give the IC_{50} and n_H values given in the text. The concentration–response relation for WT I_{hERG} (identical to that shown in Figure 1) is shown overlain, for comparative purposes. (C) The traces in panels Ci and Cii show, respectively, WT I_{hERG} and F656A I_{hERG} elicited in the absence and presence of 1 μ M KB-R7943. The corresponding voltage protocol is shown in panel Ciii and its inset. Experiments performed with high (94 mM) [K†]_e. (D) Mean data indicating effects of three concentrations of KB-R7943 on each of WT and F656A I_{hERG} tails, measured at -120 mV, fitted by Equation 1 to give the IC_{50} and n_H values given in the text. For A and C, the horizontal dashed lines are drawn at the level of the current at the pre-pulse of -40 or -120 mV, against which peak tail current amplitudes were measured. For B and D, numbers in parentheses indicate numbers of replicates at each drug concentration tested on WT and mutant channels.

the same subunit is possible in these poses, but would be anticipated to make only a small contribution to the binding energy, given the modest effect of the S624A mutation on I_{hERG} block.

Effects of KB-R7943 under AP clamp and on native I_{Kr}

In order to evaluate the effect of KB-R7943 on I_{hERG} elicited by a dynamic physiological waveform, action potential (AP)

voltage-clamp experiments were performed. Current measurements were made first in control solution and then in the presence of 100 nM KB-R7943. I_{hERG} throughout the repolarizing phase of the AP, was reduced in the presence of KB-R7943 (Figure 7A), with peak current reduced by 43.5 \pm 1.9% (n = 7). The voltage at which maximal I_{hERG} occurred was shifted by approximately -9.8 ± 1.2 mV in the presence of KB-R7943, which is similar to the extent of shift of I_{hERG} activation $V_{0.5}$ seen in Figure 2. In a subsequent set of experiments, inhibition by KB-R7943 of native ventricular I_{KF} was

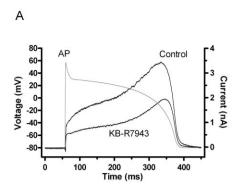


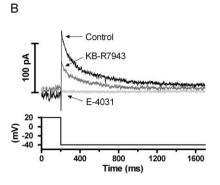
Effects of the S624A mutant on the action of KB-R7943 and simulated docking to the open hERG channel. (A) Representative records of S624A I_{hERG} tails (upper traces) elicited on repolarization to -40 mV following a 2 s depolarization to +20 mV from -80 mV (lower trace shows corresponding portion of the voltage protocol) in control solution (standard Tyrode's solution) and following exposure to 100 nM KB-R7943. (B) Mean \pm SEM fractional inhibition by SN-6 concentrations between 30 nM and 1 μ M, fitted by Equation 1 to give the IC_{50} and n_H values presented in the Results text. Numbers in parentheses indicate numbers of replicates at each drug concentration. (C) Stereoview of representative low energy score configuration for KB-R7943 docked into the open hERG channel pore homology model based on the MthK crystal structure. The protein backbone for two of the four subunits of the channel pore tetramer is indicated with a ribbon and a K^+ ion (shown in mauve) occupies the S3 site in the selectivity filter. Gly648 (green) and the side chains of S624 (blue), Y652 (orange) and F656 (yellow) are shown as sticks. The red star marks a potential hydrogen bond between the thiourea group of KB-R7943 and the backbone carbonyl of G648 (green with red carbonyl oxygen).

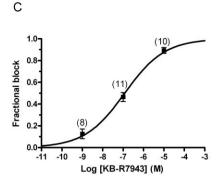
assessed. From a holding potential of -80 mV, membrane potential was briefly stepped to -40 mV (for 100 ms; both to inactivate fast Na $^+$ channels and provide a reference current level for tail current measurement), then to +20 mV (for 500 ms), then back to -40 mV for 1.5 s to elicit I_{Kr} tails. Figure 7B shows representative tail current traces in control solution, following 100 nM KB-R7943 and after subsequent application of a high concentration (10 \mu M) of E-4031. The I_{Kr}

tail was markedly inhibited by KB-R7943, and the current remaining was abolished by E-4031 (thereby confirming identity of the tail current as I_{Kr}); 100 nM KB-R7943 inhibited native I_{Kr} by 46.5 \pm 4.1% (n = 11), which is similar to the extent of WT I_{hERG} inhibition produced by this concentration (P > 0.05; unpaired t-test). Figure 7C summarizes the effects of three concentrations of KB-R7943 on native I_{Kr} (1 nM, 100 nM and 10 μ M). The data were fitted with Equation 1 to









Effects of KB-R7943 on I_{hERG} under AP clamp and on native I_{Kr} . (A) Representative records of I_{hERG} under AP clamp conditions in control and in 100 nM KB-R7943, shown with the AP command waveform overlaid. AP commands were applied every 3 s, and online leak subtraction was performed using a P/4 protocol (Hancox *et al.*, 1998). (B) Inhibition of native I_{Kr} by 100 nM KB-R7943. Membrane potential was stepped to -40 mV for 100 ms, +20 mV for 500 ms, then back to -40 mV for 1500 ms and finally back to the holding potential. The start-to-start interval for the protocol was 10 s. The figure focuses on the repolarization step, which elicited I_{Kr} tails. Representative traces are shown in control, following exposure to 100 nM KB-R7943 and following 10 μ M E-4031. (C) Concentration—response relation for the action of KB-R7943 on I_{Kr} tails. Tail current amplitude was measured as the difference between peak tail current and the current level attained during the 100 ms to -40 mV. Numbers of replicates at each concentration are given in parentheses. Data were fitted with Equation 1 to give IC₅₀ and n_H values mentioned in the Results text.

estimate an IC₅₀ value for I_{Kr} inhibition: the derived value was 120.3 \pm 27.5 nM, with an $n_{\rm H}$ value for the fit of 0.44 \pm 0.04. As observed for I_{hERG}, deactivation of I_{Kr} was slowed by KB-R7943: with $\tau_{\rm fast}$ of 81 \pm 7 ms and $\tau_{\rm slow}$ of 499 \pm 65 ms in control and $\tau_{\rm fast}$ of 144 \pm 22 ms and $\tau_{\rm slow}$ of 754 \pm 192 ms in 100 nM KB-R7943 (P<0.05 for each time constant; n=6).

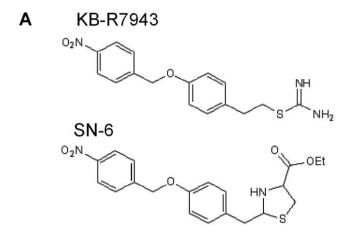
hERG inhibition by the structurally related NCX inhibitor SN-6

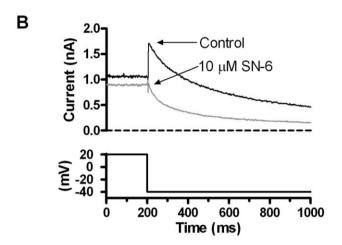
SN-6 (2-[4-(4-nitrobenzyloxy) benzyl] thiazolidine-4carboxylic acid ethyl ester) is an NCX inhibitor that shares structural similarity to KB-R7943 (Figure 8A) and inhibits NCX1 (Iwamoto et al., 2004) and native cardiac I_{NCX}, apparently with improved selectivity (Niu et al., 2007). Therefore, in a final series of experiments, we investigated the propensity of this compound to inhibit I_{hERG}. Figure 8B shows the effect of 10 μM SN-6 on the amplitude of I_{hERG} tails on repolarization to -40 mV from +20 mV, whilst Figure 8C shows mean data across a range of concentrations from 1 nM to 100 µM. SN-6 produced a concentration-dependent inhibition of I_{hERG} but was considerably less potent than KB-R7943 in this regard, with an estimated IC₅₀ of 10.4 \pm 3.3 μ M and $n_{\rm H}$ of 0.25 ± 0.03 .

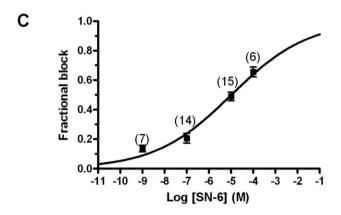
Discussion and conclusions

Although KB-R7943 is known not to be entirely selective for NCX (Amran *et al.*, 2003; Watanabe *et al.*, 2006), until now there has been no direct evidence for $I_{\text{hERG}}/I_{\text{Kr}}$ inhibition by this compound. Examination of previously published data by Tanaka *et al.* (2002) on the effect of 10 μ M KB-R7943 on guinea-pig composite I_{K} (both in respect of raw currents and 'tail' current–voltage relations in Figure 4 of that study) indicates that I_{Ks} is likely to have predominated in their record-

ings. Accordingly, the KB-R7943-sensitive current in that study is likely to have been comprised predominantly of I_{Ks}, although experimental data under selective recording conditions are not shown (Tanaka et al., 2002). Our data demonstrate unequivocally that KB-R7943 inhibits both hERG and native I_{Kr} at nanomolar to micromolar concentrations, with IC_{50} values of ~100 nM. The majority of I_{hERG} inhibitors exhibit IC₅₀ values in the µM range (Shah, 2005), and in this context, KB-R7943 can fairly be considered to be a potent I_{bERG} inhibitor. We observed that, in addition to inhibiting I_{hERG}, KB-R7943 also modified voltage-dependent activation properties of the underlying channels. A similar dual effect has been observed for both low and high potency inhibitors that inhibit hERG in a gating-dependent fashion. For example, both desethylamiodarone and 4-aminopyridine inhibit IhERG at the majority of tested voltages, whilst producing an apparent augmentation of current at negative voltages - associated with a leftward voltage shift in current activation, as seen here for KB-R7943 (Ridley et al., 2003; Zhang et al., 2010). This feature of KB-R7943 inhibition is concordant with a gating-dependent channel blocking mechanism; however, our data from the envelope of tails protocol (Figure 3) indicate that the drug produces substantial hERG channel inhibition rapidly on membrane depolarization, and, on that basis, a possible contribution of closed channel block cannot entirely be excluded. The data in Figure 4 suggest that KB-R7943 exhibits little preference for inactivated over activated channels (cf. Ridley et al., 2003; Zhang et al., 2010), whilst the positive (as opposed to inverse) voltagedependence and neutral time-dependence of inhibition seen here suggest that a preferential closed channel blocking mechanism is unlikely (cf. Milnes et al., 2003b). In addition, the fact that altering the fraction of inactivated WT IhERG channels did not significantly change the extent of observed I_{hERG} inhibition (Figure 4) is concordant with the notion that decreased WT I_{hERG} block with high [K⁺]_e and inward K⁺ flux







Effects of SN-6 on I_{hERG} . (A) Structural formulae of KB-R7943 and SN-6 (structures from http://www.tocris.com). (B) Representative records of I_{hERG} tails (upper traces) elicited on repolarization to -40 mV following a 2 s depolarization to +20 mV from -80 mV (lower trace shows corresponding portion of the voltage protocol) in control solution and following exposure to $10~\mu$ M SN-6. (C) Mean \pm SEM fractional inhibition by SN-6 concentrations between 1 nM and $100~\mu$ M, fitted by Equation 1 to give the IC₅₀ and $n_{\rm H}$ values presented in the Results text. Numbers in parentheses indicate the numbers of replicates at each drug concentration.

(Figure 5) may be accounted for by a direct interaction (electrostatic repulsion or 'knock off') between the permeant ion and KB-R7943 (Wang *et al.*, 1997; Ridley *et al.*, 2004; Barrows *et al.*, 2009). It is notable that the Hill coefficient ($n_{\rm H}$) values for the concentration-dependence of KB-R7943 inhibition of WT I_{hERG} (Figure 1) and native I_{Kr} (Figure 7) lay close to 0.5. This raises the possibility that KB-R7943 block of WT I_{hERG}/I_{Kr} might involve negative co-operativity (cf. McPate *et al.*, 2006). However, whilst this was seen for outward I_{hERG} with normal [K⁺]_e, when [K⁺]_e was elevated, the $n_{\rm H}$ was closer to 1 (0.75) for inward I_{hERG} measurement in high [K⁺]_e.

The impact of alanine substitution at Y652 and F656 seen here for KB-R7943 (Figure 5) is significant, though less marked than observed previously for some other high-affinity blockers (e.g. the Y652A mutation increased the IC50 for MK-499 block of I_{hERG} by ~94-fold, whilst F656A increased the IC₅₀ by ~650-fold, with both mutations also profoundly affecting the potency of cisapride and terfenadine; Mitcheson et al., 2000). Pore helix mutations, including \$624A, also influence significantly the potency of I_{hERG} inhibition by high-affinity inhibitors including methanesulphonanilides as well as that of cisapride and terfenadine (Mitcheson et al., 2000; Kamiya et al., 2006; Kamiya et al., 2008). Here, the S624A mutation exerted a relatively modest effect on blocking potency (Figure 6), suggesting that this residue may not be a key constituent of the drug binding site. It is notable, however, that although it was studied with an identical protocol and measurement conditions to WT IhERG, S624A hERG exhibited an $n_{\rm H}$ value close to 1 for concentration-dependent I_{hERG} inhibition, concordant with some influence of the residue on the manner of drug-channel interaction. Y652AhERG also exhibited an $n_{\rm H}$ for concentration-dependent inhibition close to 1. Thus, in electrophysiological experiments manipulation of the direction/magnitude of K+ flux and mutagenesis of S624 and Y652 residues influenced the steepness of observed inhibitory concentration-dependence, whilst Y652 and F656 were implicated as significant components of the KB-R7943 binding site. This was borne out by our docking simulations (Figure 6), which, additionally, demonstrate a likely orientation of the KB-R7943 molecule within the open channel pore. Stacking interactions between aromatic rings on the drugs and the aromatic side chains of Y652 and F656, together with potential interactions involving the positively charged thiourea group of KB-R7943 (electrostatic interactions with the pore helix dipole charge and hydrogen bond interactions with G648 carbonyl), may contribute to the relatively high affinity of this compound. Additional docking simulations (not shown) showed that SN-6 is unable to make equivalent hydrogen bond interactions to those seen for KB-R7943, and so these may contribute to the observed difference in inhibitory potency between KB-R7943 (Figures 1 and 7) and SN-6 (Figure 8).

In the first study of KB-R7943, the compound was found to inhibit Na⁺-dependent Ca²⁺ uptake into NCX1-transfected CC39 cells, rat aortic smooth muscle cells and cardiac myocytes, with IC₅₀ values ranging between 1.6 and 2.4 μ M (Iwamoto *et al.*, 1996), whilst Na⁺-dependent Ca²⁺ influx into Na⁺-loaded cardiac sarcolemmal vesicles was inhibited with an IC₅₀ of 5.4 μ M, and Na⁺o-dependent Ca²⁺ efflux was inhibited with an IC₅₀ of 11–13 μ M (Iwamoto *et al.*, 1996). When I_{NCX} from guinea-pig ventricular myocytes was elicited in a



unidirectional manner, outward and inward I_{NCX} were inhibited, respectively, with an IC $_{50}$ of ~320 nM and 17 μM (Watano et al., 1996). For bi-directional guinea-pig ventricular I_{NCX} , the reported IC_{50} was ~1 μ M (Kimura *et al.*, 1999), whilst KB-R7943 was able to inhibit outward canine ventricular I_{NCX} more extensively than inward $I_{\text{NCX}}\text{,}$ but with EC_{50} values, respectively, of ~4.7 and 3.4 μM (Birinyi et al., 2005). As highlighted in the Introduction, the compound can also inhibit cardiac Na⁺ and Ca²⁺ and inwardly rectifying K⁺ currents at micromolar concentrations (Watano et al., 1996; Tanaka et al., 2002). There is evidence for additional off-target effects of the compound from experiments on non-cardiac preparations. For example, KB-R7943 inhibits recombinant Ca_v1.2 channels with an IC₅₀ of ~7.3 μM (Ouardouz et al., 2005) and canonical transient receptor potential channels with IC50 values of 460 nM (TRPC3), 710 nM (TRPC6) and 1.38 μM (TRPC5) (Kraft, 2007). At 10–20 μM, the compound can significantly inhibit store-operated Ca²⁺ entry into cultured neurons and astrocytes (Arakawa et al., 2000). In cultured hippocampal neurons, it inhibits NMDA receptormediated increases in cytosolic calcium with an IC50 of 13.4 µM (Brustovetsky et al., 2011), and it can inhibit mitochondrial Ca²⁺ uptake [with an IC₅₀ of 5.5 μM in permeabilized HeLa cells (Santo-Domingo et al., 2007)] and mitochondrial complex I (Brustovetsky et al., 2011). Another study, focusing on acutely isolated CA1 hippocampal neurons, has reported inhibition of two populations of NMDA channels with high-affinity (IC₅₀ of 0.8 μM) and lowaffinity (IC₅₀ of ~11 μM) block by KB-R7943 (Sobolevsky and Khodorov, 1999). KB-R7943 has also been reported to inhibit a non-selective cation channel implicated in chemosensory transduction (with an IC₅₀ of ~11.7 µM; Pezier et al., 2009) and to activate vascular large conductance Ca²⁺-activated K⁺ channels (EC₅₀ of ~6.8 µM; Liang et al., 2008). Significantly, when the results of the present study are considered against the background of the compound's overall profile of effects, it is clear that (i) KB-R7943 inhibits hERG/I_{Kr} at least as potently has been reported previously for cardiac I_{NCX}, and (ii) none of the above-mentioned off-target effects is as potent as that seen in the present study for $hERG/I_{Kr}$. Thus, $hERG/I_{Kr}$ channel inhibition appears to be the most potent off-target pharmacological action thus far identified for KB-R7943.

Given the results of the present study, it seems reasonable to conclude that in experiments on cardiac cells, tissues and intact hearts significant I_{hERG}/I_{Kr} inhibition can be anticipated to occur at NCX-blocking concentrations of KB-R7943. It follows, therefore, that the propensity of the agent to inhibit hERG/I_{Kr} channels needs to be taken into account when interpreting experimental data with the compound obtained from cardiac preparations from species in which I_{Kr} participates in AP repolarization [these include human, dog, rabbit and guinea-pig (Sanguinetti and Jurkiewicz, 1990; Tamargo et al., 2004)] or pacemaking (Zaza et al., 1997; Mitcheson and Hancox, 1999; Sato et al., 2000). For example, it is noteworthy that a low concentration of KB-R7943 (i.v. injection to plasma levels of ~31 nM) has been reported to prolong atrial effective refractory period (AERP) of anaesthetized dogs (Miyata et al., 2002), an effect that was attributed to action(s) of KB-R7943 other than NCX inhibition (Miyata et al., 2002), and that is concordant with hERG/I_{Kr} inhibition [I_{Kr} participates in atrial AP repolarization and pharmacological I_{Kr} blockade is known to exert an atrial class III anti-arrhythmic action (Torp-Pedersen et al., 2000; Tamargo et al., 2004; Doggrell and Hancox, 2005)]. The hERG/I_{kr}-inhibition by KB-R7943 demonstrated here also explains recently reported membrane potential depolarization of rabbit AV node cells at compound concentrations sufficient to inhibit inward I_{NCX} (Cheng et al., 2011). This highlights the importance of using alternative/complementary interventions to KB-R7943 to alter NCX function in investigations of the role of the NCX in cardiac pacemaking (e.g. Sanders et al., 2006; Cheng et al., 2011). It is worth noting that in some experimental settings, at some drug concentrations, the ability of KB-R7943 to exert multiple actions may mean that overt AP or QT interval prolongation due to diminished I_{Kr} may not be evident, and hence, the consequences of I_{Kr} blockade may feasibly be overlooked, despite contributing to the overall actions of the compound. Indeed, if the overall experimental objectives of a cardiac study employing KB-R7943 do not require that I_{Kr} be present, it may be prudent to select a species in which I_{Kr} is not important for ventricular repolarization or to employ an alternative means of NCX inhibition. Considered in isolation, our KB-R7943 data might be interpreted to suggest that benzyloxyphenyl-based pharmacological strategies for NCX inhibition would be hampered by concurrent potent hERG inhibition. However, our findings with SN-6 (Figure 8) indicate that this need not necessarily occur to the extent that is evident here for KB-R7943. SN-6 has been reported to inhibit ⁴⁵Ca²⁺ uptake in NCX1-transfected fibroblasts with an IC₅₀ of 2.9 µM (Iwamoto et al., 2004) and to inhibit outward and inward components of bi-directional I_{NCX} from guinea-pig ventricular myocytes with IC_{50} values of 2.3 and 1.9 μM (Niu et al., 2007). Thus, the compound is comparable with KB-R7943 in its NCX-inhibitory potency, but the present study shows it to be ~100-fold less potent as a hERG inhibitor; it also affects other currents less potently than KB-R7943 (Niu et al., 2007). It remains to be established whether benzyloxyphenyl-based NCX inhibitors entirely devoid of hERG activity can be produced. However, SN-6 may offer a useful complementary or alternative NCX inhibitor to KB-R7943 for use in experiments on cardiac preparations that contain I_{Kr}/hERG channels. An alternative to either compound is SEA0400, which inhibits I_{NCX} with greater potency than does KB-R7943 (Tanaka et al., 2002; Birinyi et al., 2005) and for which the available data suggest exerts little effect on cardiac composite delayed rectifier K+ current (Tanaka et al., 2002). Finally, it is worth noting that hERG channel expression is not restricted to the heart (Sanguinetti and Tristani-Firouzi, 2006; Larsen, 2010); consequently, the hERGchannel blocking action of KB-R7943 may also need to be taken into account when the compound is used in experimental studies of hERG-expressing non-cardiac tissue types.

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Conflicts of interest

None.

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KB-R7943 inhibition of hERG K+ channels



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