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The intriguing effect of ethanol and nicotine on acetylcholine-sensitive potassium current *I*_{KAch}: Insight from a quantitative model

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Abstract

Recent experimental work has revealed unusual features of the effect of certain drugs on cardiac inwardly rectifying potassium currents, including the constitutively active and acetylcholine-induced components of acetylcholine-sensitive current (IKAch). These unusual features have included alternating susceptibility of the current components to activation and inhibition induced by ethanol or nicotine applied at various concentrations, and significant correlation between the drug effect and the current magnitude measured under drug-free conditions. To explain these complex drug effects, we have developed a new type of quantitative model to offer a possible interpretation of the effect of ethanol and nicotine on the I_{KAch} channels. The model is based on a description of I_{KAch} as a sum of particular currents related to the populations of channels formed by identical assemblies of different α -subunits. Assuming two different channel populations in agreement with the two reported functional IKAch-channels (GIRK1/4 and GIRK4), the model was able to simulate all the above-mentioned characteristic features of drug-channel interactions and also the dispersion of the current measured in different cells. The formulation of our model equations allows the model to be incorporated easily into the existing integrative models of electrical activity of cardiac cells involving quantitative description of IKAch. We suppose that the model could also help make sense of certain observations related to the channels that do not show inward rectification. This new ionic channel model, based on a concept we call population type, may allow for the interpretation of complex interactions of drugs with ionic channels of various types, which cannot be done using the ionic channel models available so far.

Introduction

Our recent experimental work has revealed unusual features of the effect of ethanol and nicotine on cardiac ionic membrane channels responsible for inwardly rectifying potassium currents ($I_{\rm Kir}$); both drugs exhibited inhibition or activation depending on drug concentration. Furthermore, the drug effect correlated with the current magnitude measured under drug-free conditions [1–4]. A similar dualistic concentration-dependent characteristic was previously **Competing interests:** The authors have declared that no competing interests exist.

Abbreviations: c, Drug concentration; *f*_i, Generally: f^{th} -fraction of identical channel populations; f_1 , f_2 , 1st and 2nd fraction of identical channel populations for I_{KAch} ; Δf , Random shifts between fractions (f_1 $+\Delta f$, $f_2 -\Delta f$, F, Relative current (I/I_0); g(U), Approximation of the voltage-dependent conductivity of IKAch; G1.t, G2.t, Conductivity of the 1^{st} and 2^{nd} channel population; G_1 , G_2 , Steadystate conductivity of the 1st and 2nd channel population; G_{k_1} , G_{k_2} , Partial steady-state conductivities of the 1st and 2nd channel population related to the drug occupation of binding sites (k =0, 1 or 2); GIRK, G-protein modulated inwardly rectifying potassium channels (GIRK4, GIRK 1/4); h_{k} 1, h_{k} 2, Dimensionless parameters (G_{k} 1 = h_{k} 1 g(U), $G_{k_2} = h_{k_2} g(U)$; I, I_{Kir} , Inward rectifier potassium current (generally); *I*_i, *I*_{Kir i}, Constituents of $I(I_{Kir})$ related to individual populations (*j*) of identical channels; I_{K1} , Inward rectifier potassium current Kir2.x; IKAch, Acetylcholine-sensitive inward rectifier potassium current (Kir3.x); I_{KAch,0}, Acetylcholine-sensitive inward rectifier potassium current under drug free conditions; IKAch CONST, Constitutively active component of IKAch; IKAch ACH, Acetylcholine-induced component of IKAch; I, I, Components; j, Subscript related to individual populations of identical channels (generally *j* = 1, ..., n); k, Subscript related to occupation of binding sites; K_1 , K_2 , K_3 , K_4 , Dissociation constants; n, Number of populations of identical channels (n = 2 for I_{KAch}); PIP₂, Phospholipid phosphatidylinositol-4,5-bisphosphate; τ_1 τ_1 , Time constants; U, Membrane voltage; U_K, Equilibrium voltage for potassium ions; x_{k_j} , Probability of channels to be found in state k pertaining to j^{th} channel population.

demonstrated in the effect of halothane on GIRK channels expressed in *Xenopus laevis* oocytes [5–6].

In an attempt to explain these drug effects, we designed a quantitative model considering the known structure of the inwardly rectifying potassium channels, which are composed of various homomeric or heteromeric assemblies of channel subunits (Kir2.x, Kir3.x, and Kir6.x/ SURx) with a considerable diversity in regional expression of the underlying subunits [7] (for reviews, see [8–9]). Different combinations of the expressed subunits result in specific functional properties of the measured currents, including drug-channel interactions.

To our knowledge, the existing integral mathematical models of the inwardly rectifying potassium currents ignore the fact that these currents can be seen as the sum of separate currents. In our model, the Kir currents are described as a total of *n* particular independent currents $I_{\text{Kir,x,j}}$ via individual populations (*j*) of identical channels, i.e. channels formed by the same combination of subunits. The number *n* of constituents (each created by identical channels) is unknown. To describe the effect of ethanol on I_{K1} in ventricular and atrial cardiomyocytes [1–2], we tentatively introduced a minimum number (*n* = 3) needed to simulate the available observations [10]. Three α -subunits (Kir2.1, Kir2.2, and Kir2.3) are known to participate in the formation of I_{K1} channels (for review, see [9]).

In this study, we apply the conception of different channel populations to analyse the responses of acetylcholine-sensitive inward rectifying potassium current (I_{KAch}) to ethanol [3] and nicotine [4]. These responses showed some similar features to those revealed previously in the ethanol- I_{K1} interactions [1–2], as mentioned above. In the current version of the model, we considered two different populations of the I_{KAch} channels (n = 2), because only two combinations of α -subunit assemblies, namely heterotetramers Kir3.1/Kir3.4 (GIRK1/4) and homotetramers Kir3.4 (GIRK4), have been identified in cardiac atrial cells [11–18] (for review, see [9]). Homotetramers GIRK1 appeared to be nonconductive [14, 19]. The proposed quantitative model offers a possible mechanism for the effect of ethanol and nicotine on two components of I_{KAch} , namely the constitutively active ($I_{\text{KAch}_{-\text{CONST}}$) and the acetylcholine-induced ($I_{\text{KAch}_{-\text{ACH}}$).

Results

Kir channel populations—Model description

In the model described in our previous study [10], Kir currents (often recalculated to current densities in pA/pF) were generally expressed as:

$$I = \sum_{j=1}^{n} f_j I_j = \sum_{j=1}^{n} f_j G_j (U - U_K), \qquad (\sum_{j=1}^{n} f_j = 1).$$
(1)

In this equation, *n* denotes a number of independent components I_j of the current *I* (which are related to individual channel populations formed by different identical assemblies of α -subunits), f_j denotes fraction of the j^{th} identical channel population, and *U*, U_K , and G_j denote membrane voltage, equilibrium voltage for potassium ions, and fractional conductivities, respectively. This concept was a useful tool for describing complex interactions of I_{K1} (Kir2.x) with ethanol [10]. As mentioned above, a minimum number n = 3 was selected to enable reproduction of available experimental results and to keep the model as simple as possible.

In the case of I_{KAch} , we confined the number of channel populations to n = 2 because GIRK4 homotetramers and GIRK1/4 heterotetramers are the only functional assemblies underlying I_{KAch} in cardiac cells (for review, see [18]). Significant differences occur between GIRK4 and GIRK1/4 channels (e.g. different sensitivity to $[\text{Na}^+]_i$), although both tetramers share ~44% sequence identity [16]. The two components of I_{KAch} ($I_{\text{KAch}_\text{CONST}}$ and $I_{\text{KAch}_\text{ACH}}$) were described by the same equations, but with different numeric values for the parameters.

Previous studies have shown that a hydrophobic *pocket* in the cytoplasmic domain is a putative activating binding site for alcohols and other substances [20–22]. The pocket may be occupied by one or more ethanol molecules, indicating a sequential bond. Alternate sites were shown to be responsible for inhibition [20]. As little is known about the underlying mechanisms, we used the simple description of inhibition to display experimental results.

Both populations (f_1 , f_2) may include activation and inhibition binding sites, one of which is likely to dominate. However, as the simplest way to comply with the described concept, we introduced the putative pocket related to activation to the first population (j = 1) of identical channels. Two additional binding sites related to inhibition were assigned to the second population (j = 2), as illustrated in Fig 1.

The symbols x_{0_1} and x_{0_2} in Fig 1 denote probabilities of the respective channel population to be found drug-free. The probabilities that channels belonging to population j = 1 are occupied by one or two drug molecules are designated x_{1_1} , x_{2_1} . Probabilities that the first or the second binding site in channels of population j = 2 are occupied by a drug molecule are designated x_{1_2} and x_{2_2} , respectively. If all channels of the given population were found in one of the three states shown in the schemes above, the conductivity of this population at a given membrane voltage U would take one of the corresponding values G_{0_1} , G_{1_1} , G_{2_1} , or G_{0_2} , G_{1_2} , G_{2_2} .

Transient changes of Kir currents evoked by application of ethanol (and some other drugs) appeared to be slow, lasting up to 10^2 seconds [1–2, 23]. If we assume (in line with Šimurda *et al.* [10]) that the drug binding velocity is much higher than the subsequent (allosteric) conformational changes governing channel conductivities, then the probabilities x_{k_j} (*k* refers to occupation of binding sites, *j* to populations of identical channels) may be assumed to keep their steady state values depending only on the drug concentration *c* and the dissociation constants K_1 , K_2 , K_3 and K_4 . In this case, x_{k_j} can be expressed as solutions of linear algebraic equations related to the above schemes describing drug-binding in the channel population 1 and 2.

$$x_{0_1} = \frac{1}{1 + \frac{c}{K_1} + \frac{c^2}{K_1 K_2}}, \quad x_{1_1} = x_{0_1} \frac{c}{K_1}, \quad x_{2_1} = x_{0_1} \frac{c^2}{K_1 K_2}, \quad (2A)$$

$$x_{0-2} = \frac{1}{1 + \frac{c}{K_3} + \frac{c}{K_4}}, \quad x_{1-2} = x_{0-2} \frac{c}{K_3}, \quad x_{2-2} = x_{0-2} \frac{c}{K_4}.$$
 (2B)

Evidently,

$$x_{0_{-1}} + x_{1_{-1}} + x_{2_{-1}} = 1, \quad x_{0_{-2}} + x_{1_{-2}} + x_{2_{-2}} = 1.$$

Conductivities of the first and the second population (G_1 and G_2 , respectively) under steady state conditions may then be expressed as

$$G_{1} = x_{0_{-1}}G_{0_{-1}} + x_{1_{-1}}G_{1_{-1}} + x_{2_{-1}}G_{2_{-1}}, \quad G_{2} = x_{0_{-2}}G_{0_{-2}} + x_{1_{-2}}G_{1_{-2}} + x_{2_{-2}}G_{2_{-2}}, \tag{3}$$

and in agreement with (1)

$$I = (f_1 G_1 + f_2 G_2)(U - U_K).$$
(4)



Fig 1. A schematic representation of the drug-channel interactions. One or two drug molecules can be bound in the pocket related to activation (left). One drug molecule can be bound to one of two binding sites related to inhibition (right). $K_1, K_2, K_3, K4$ —dissociation constants; xk_j —probabilities of channels to be found in state k (k = 0, 1, or 2) pertaining to j^{th} channel population (j = 1 or 2); $G_{k_{-1}}, G_{k_{-2}}$ —partial steady-state conductivities of the 1st and 2nd channel population related to the drug occupation of binding sites.

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Under drug-free conditions (c = 0),

$$I_0 = (f_1 G_{0,1} + f_2 G_{0,2})(U - U_K).$$
(5)

The conductivities are generally voltage dependent. If, however, the drug does not interfere with the structures responsible for inward rectification, the drug effect alone is voltage independent, and all conductivities may be regarded as equally dependent on the membrane voltage. The ratio

$$F = \frac{f_1 G_1 + f_2 G_2}{f_1 G_{0-1} + f_2 G_{0-2}} = \frac{I}{I_0}$$
(6)

may then be interpreted as a voltage-independent indicator of the drug effect. In this case, we can express all the above-defined conductivities as products of voltage-independent dimensionless parameters h_{k_j} and a common voltage-dependent conductivity g(U) describing inward rectification:

$$G_{k_{-j}} = h_{k_{-j}}g(U), \tag{7A}$$

$$g(U) = 0.8 \left(\frac{0.325}{1 + e^{\frac{U+50}{5}}} + \frac{3}{1 + e^{\frac{U+150}{52}}} \right).$$
(7B)

As mentioned above, the subscripts k = 0, 1 or 2 refer to states differing by a drug occupation of binding sites while j = 1 or 2 refer to populations of the identical channels. The function g (U) was obtained as a formal mathematical expression (in μ S) from the current-voltage relationship, as recorded under control conditions [3].

In this study, the conductivities under the effect of the drug were regarded as equilibrated. The experimental data related to I_{K1} [1–2] also included transient changes of the current in response to the onset of ethanol. Corresponding transients in conductivities were described in ref. [10] by differential equations. Unfortunately, similar transients of I_{KAch} following drug application were not available. For possible future use, the corresponding equations are given in the Discussion.

Effect of ethanol on I_{KAch}

The concentration dependences of the effect of ethanol on $I_{\text{KAch}_\text{CONST}}$ and $I_{\text{KAch}_\text{ACH}}$ are illustrated in Fig 2. The experimental results as obtained on rat atrial cardiomyocytes [3] are simulated by the solution of the model equations with the numeric values of the model parameters summarized in Table 1. The model parameters were set by repeated attempts to assure good fit with available experimental data. The number of different populations were confined to n = 2 in agreement with the current knowledge of the structure of the I_{KAch} channels. The values of parameters (Table 1) remained unaltered in all simulations relating to the effect of ethanol, except for simulations of those experimental results that were interpreted as resulting from dispersion in fractions of the individual channel populations. Table 1 shows the mean values of the fraction f_1 (while $f_2 = 1 - f_1$). The results of simulations related to the dispersion of the channel populations were obtained by replacing f_1 with $f_1 + \Delta f$. The Δf values are given in the figure legends.

Fig 2A shows the steady-state concentration dependence of the effect of ethanol on I_{KAch_-} CONST (left) and I_{KAch_ACH} (right) in a relative scale. The quantity $F = I/I_0$ (6) is plotted against ethanol concentration; simulated results (black lines) are compared with experimental data from ref. [3] (circles ± SE). The blue and violet lines show how the model allows for dispersion of the measured values. Individual cells are assumed to exhibit dispersion due to various representations of the I_{KAch} -channel populations, which is expressed in the model as redistribution between the fractions f_1 and f_2 . Changes in the concentration dependence induced by 10% of the channels shifting between f_1 (channels showing activation) and f_2 (channels showing inhibition) are plotted. Blue lines correspond to $f_1 = 0.68 + 0.1$ and $f_2 = 0.32-0.1$; violet lines to $f_1 = 0.68-0.1$ and $f_2 = 0.32 + 0.1$.

The model enables us to transform currents from the relative to the absolute scale (Fig 2B) by multiplying the relative current values by the corresponding current densities at zero ethanol concentration ($I = F I_0$ according to Eq (6)). Any redistribution of the channels between f_1 and f_2 is accompanied by a change in the control current density (approximately the values at the lowest concentration in Fig 2B), as a consequence of differences in the conductivities $G_{0,1}$ and $G_{0,2}$ of the individual channel populations. The marked dispersion of the current *I* at low drug concentrations in absolute values (Fig 2B) is reflected in an increased dispersion at higher concentrations after normalization to I_0 (Fig 2A).

In Fig 2C, the mean ethanol concentration dependences of both components of I_{KAch} (black lines in Fig 2B) are expressed as total of their presumptive constituents related to the individual channel populations (red and green lines).

Another peculiarity of the effect of ethanol on both components of I_{KAch} was the correlation between the drug effect (inhibition or activation) and the current density in the absence of ethanol [3]. A similar property was observed and simulated in the interaction of I_{K1} with ethanol in ventricular and atrial cardiomyocytes [1–2, 10]. The solution of the model equations appeared to reproduce the drug effect as a nonlinear function of the current density in the absence of ethanol (Fig 3, full lines). The results of the model are compared in Fig 3 with experimental data from different cells (filled symbols) and with the results of linear regression (dashed lines). The simulated variations of the current in the control ($I_{\text{KAch}_{CONST}_{contr}}$ and $I_{\text{KAch}_{ACH}_{contr}}$) resulted from continuous redistribution between fractions f_1 and f_2 . The model offers a prospective explanation of the correlation as a consequence of different conductivities of the channel populations and their different sensitivities to ethanol.

In the current form of the model, the drug effect itself is regarded as voltage-independent so that all the parameters in Table 1 are constants. Under this assumption, the model is restricted to a description of the effects of drugs which do not considerably interfere with the



Fig 2. Analysis of the steady-state concentration dependence of the effect of ethanol on the constitutively active (I_{KAch_CONST}) and acetylcholine-induced (I_{KAch_ACH}) currents. Values of the parameters as summarized in Table 1 were used for calculations except for variations of f_j needed to simulate dispersion of the measured currents. A: Relative changes of I_{KAch_CONST} and I_{KAch_ACH} as functions of ethanol concentration; black lines-model; circles-experimental data (± SE according to [3]; 4–10 cells in the tested concentrations for I_{KAch_CONST} and 6–15 cells in the tested concentrations for I_{KAch_ACH}). The data are related to the current densities at zero ethanol concentration. Simulations of the data dispersion result from small variations $\Delta f = 0.1$ in fractions of individual populations of the I_{KAch} channels ($f_1+\Delta f$ -blue lines, $f_1-\Delta f$ -violet lines). B: Simulated absolute values of I_{KAch} resulting from multiplication of the relative values (in part A) by the current densities at zero ethanol concentration (I_0 , Eq (5)). C: Contributions of the presumptive two current constituents (red and green lines) to the resulting ethanol concentration dependence of I_{KAch} at the basal setting of parameters according to Eq (4); $I_1 = G_1 (U-U_K)$, $I_2 = G_2 (U-U_K)$.

Current	f1 -	$K_1 = K_2$ [mM]	<i>K</i> ₃ [mM]	<i>K</i> ₄ [mM]	h _{0_1}	h _{1_1}	h _{2_1}	h _{0_2}	h _{1_2}	h _{2_2}
I _{KAch_CONST}	0.68	6	0.5	150	0.013	0.2	0.035	0.09	0.09	0.09
I _{KAch_ACH}	0.68	36	0.4	150	0.13	0.85	0.09	0.9	0.7	0.05

Table 1. Parameters of the model for simulations of the ethanol effect on $I_{\rm KAch}$.

 f_1 -mean value of the first fraction of identical channel populations ($f_2 = 1 - f_1$); K_1 , K_2 , K_3 , K_4 -drug dissociation constants; h_{k_j} -dimensionless parameters related to the steady-state conductivities of the j^{th} channel population; k refers to occupation of binding sites.

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structures responsible for the non-linear current-voltage relationship (inward rectification). Otherwise, the parameters are to be regarded as voltage dependent.

The experimental current-voltage relations of $I_{\text{KAch}_\text{CONST}}$ and $I_{\text{KAch}_\text{ACH}}$ in the absence and presence of ethanol (see Fig 4A and 4B in ref. [3]) were simulated by evaluation of the expressions of *I* and I_0 from Eqs (4) and (5). In Fig 4 the experimental and the model results are compared. The differences between the cells showing activation and those showing inhibition of $I_{\text{KAch}_\text{ACH}}$ (Fig 4B) are interpreted and simulated as a consequence of differences between the fractions f_1 and $f_2 = 1 - f_1$ of channel populations in individual cells.

Effect of nicotine on I_{KAch}

The effect of nicotine on both components of I_{KAch} in rat atrial myocytes (Fig 5) displayed main features similar to those displayed by the effect of ethanol (Fig 2). However, the voltage dependence of the effect was not negligible in the case of I_{KAch_ACH} . Thus, one of the model parameters was considered as dependent on the membrane voltage. The effect of nicotine on I_{KAch} was analysed preferentially at -50 mV (i.e. in the range of positive currents) in contrast to the effect of ethanol which was measured mostly at -110 mV (negative currents). To



Fig 3. Correlation between the relative effect of ethanol (eth) and the current density of I_{KAch} components under ethanol-free conditions (contr). Experimental results from individual cells at an ethanol concentration of 20 mM (circles) were adopted from the original set of data from Bébarová *et al.* [3] (9 cells for I_{KAch_CONST} and 14 cells for I_{KAch_ACH}). The full lines result from the model simulations with values of parameters as stated in Table 1. Dashed lines result from the linear regression analysis of the experimental data; the correlation coefficient was significant (P < 0.05) in I_{KAch_ACH} . The variations of the current in control were simulated by continuous redistribution between channel populations (fractions f_1 and $f_2 = 1 - f_1$).



Fig 4. Effect of ethanol (20 mM) on the current-voltage relation of I_{KAch} **components.** Experimental (left panels; 4 cells for $I_{\text{KAch}_\text{CONST}}$; 14 cells for $I_{\text{KAch}_\text{ACH}}$ —7 cells for the activation and 7 cells for the inhibition) and simulated (right panels) data. Activation and inhibition

were simulated by redistribution of the channels between fractions f_1 and f_2 : $(f_1 \rightarrow f_1 + \Delta f, f_2 \rightarrow f_2 - \Delta f)$. A: Simulation (right panel; $\Delta f = 0.19$) of a representative record of $I_{\text{KAch}_\text{CONST}}$ (left panel). B: Simulations (right panels) of representative records of $I_{\text{KAch}_\text{ACH}}$ (left panels) showing the ethanol-induced activation (upper panels, $\Delta f = 0.15$) and the inhibition (lower panels, $\Delta f = -0.6$). To better simulate the illustrated representative experimental records, the conductivity g(U) was multiplied by 0.5.

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reproduce experimental results measured at -50 mV in ref. [4], the values of the parameters summarized in Table 2 were inserted into the model. They were set the same way as in the case of ethanol, i.e. by repeated attempts to assure good fit with experimental data. The value of f_1 related to drug-free conditions was left the same as in the above-described simulations of the ethanol effect (fixed to $f_1 = 0.68$). In contrast, the values of parameters h_{0_1} and h_{0_2} had to be changed in simulations of the nicotine effect due to differences in the experimentally obtained current-voltage relations of the two separate sets of cells.

The simulated concentration dependences of nicotine effect (Fig 5) are presented in the same way as those of ethanol (Fig 2). Relative changes in the experimental and simulated components of I_{KAch} were plotted against the nicotine concentration in Fig 5A. The steady-state effect of nicotine in $I_{\text{KAch}_\text{CONST}}$ (predominantly activation) was significantly stronger than the effect in $I_{\text{KAch}_\text{ACH}}$. In the latter case, we even observed inhibition at 400 nM. After recalculation to absolute scale (Fig 5B), the currents have a positive sign at -50 mV. Fig 5C shows the resolution of $I_{\text{KAch}_\text{CONST}}$ and $I_{\text{KAch}_\text{ACH}}$ into constituents as predicted by the model.

As in the case of ethanol, the effect of nicotine correlated with the magnitude of I_{KAch} density in the absence of nicotine (at 4, 40, and 400 nM–Fig 6). The results of linear regression applied to the experimental data are plotted for all concentrations explored (dashed lines). The model simulations revealed nonlinear dependences (full lines).

The voltage dependence of the effect of nicotine was available only for $I_{\text{KAch}_\text{CONST}}$ measured at a concentration of 40 nM (Fig 1E in ref. [4]). In Fig 7 model results are compared with experimental data. The nicotine effect was calculated as a result of an increase of the fraction of the channel f_1 by $\Delta f = 0.19$ on account of the same decrease in f_2 .

As mentioned above, the effect of nicotine was not strictly voltage independent. The concentration dependence of the relative drug effect on $I_{\text{KAch}_A\text{CH}}$ measured at -110 mV was shifted towards inhibition in comparison with the results at -50 mV (Fig 3C in ref. [4]). The present model reproduced this shift (Fig 8) as a result of a decrease of the single parameter h_{2_1} related to activation.

Discussion

The potassium inward rectifying channels are known to be composed of several different assemblies of α -subunits. These structural differences are manifested by various functional properties of the channels. However, to our knowledge, Kir currents carried by different populations of identical channels (i.e. sharing the same composition and ordering of α -subunits) have not been specified in the available integral models. While the channels responsible for I_{K1} in the ventricular and atrial cells are composed of three subunits (Kir2.1, Kir2.2 and Kir2.3), the I_{KAch} channels are homo- or heterotetramers comprised of two subunits (Kir3.1 and Kir3.4) (for review, see [9]). It has been well established that activity of the channels exhibiting inward rectification is modulated by a large variety of intracellular factors (e.g. PIP₂, Na⁺, Mg² ⁺, polyamines, pH and cAMP-dependent PK [9, 24–25]) and by various drugs [26–34], including anaesthetics and alcohols [6, 20, 23, 35–36].

Our recently published experimental studies focusing on the effects of ethanol and nicotine on Kir currents in cardiac cells [1-4] have shown that the interactions of these drugs with the



Fig 5. Analysis of the steady-state concentration dependence of the effect of nicotine on the constitutively active (I_{KAch_CONST}) and acetylcholine-induced (I_{KAch_ACH}) currents. Values of the parameters as summarized in Table 2 were used for calculations except for variations of f_j needed to simulate dispersion of the measured currents. A: Relative changes of I_{KAch} components as functions of the nicotine concentration; black lines-model; circles-experimental data (± SE according to ref. [4]; 5–7 cells in the tested concentrations for I_{KAch_CONST} and 5–12 cells in the tested concentrations for I_{KAch_ACH}). The data are related to the current densities at zero nicotine concentration. Simulations of the data dispersion result from small variations $\Delta f = 0.07$ in fractions of individual populations of the I_{KAch} channels ($f_1+\Delta f$ blue lines, $f_1-\Delta f$ -violet lines). B: Simulated I_{KAch} in absolute scale resulting from multiplication of the relative values (in part A) by the current densities at zero ethanol concentration (I_0 , Eq (5)). C: Contributions of the presumptive two current constituents (red and green lines) to the resulting nicotine concentration dependence of I_{KAch} at the basal setting of parameters according to Eq (4); $I_1 = G_1 (U-U_K)$, $I_2 = G_2 (U-U_K)$.

Current	f_1	$K_1 = K_2$	<i>K</i> ₃	K_4	h_{0_1}	h_{1_1}	h _{2_1}	h_{0_2}	h_{1_2}	h _{2_2}
	-	[mM]	[mM]	[mM]	-	-	-	-	-	-
I _{KAch_CONST}	0.68	50	1	∞	0.008	0.107	0.036	0.06	0.049	0
I _{KAch_ACH}	0.68	0.2	110	520	0.065	0.085	0.085	0.45	0.41	0.025

Table 2. Parameters of the model for simulations of the nicotine effect on $I_{\rm KAch}$.

 f_1 -mean value of the first fraction of identical channel populations ($f_2 = 1 - f_1$); K_1 , K_2 , K_3 , K4 -drug dissociation constants; hk_j -dimensionless parameters related to the steady-state conductivities of the jth channel population; k refers to occupation of binding sites.

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Kir channels exhibited several characteristic features differing from the simple inhibition observed in the effect of ethanol on most other currents [37].

To quantitatively describe these unusual observations, we tried to develop a model that would reproduce available experimental results and suggest possible mechanisms underlying the drug effects. One version of our model respecting the molecular structure of the channels was tested on the observed ethanol- I_{K1} interaction in rat ventricular and atrial myocytes [10]. In the present study, the model is modified to reproduce the effects of ethanol and nicotine on the components of I_{KAch} (constitutively active I_{KAch_CONST} and acetylcholine induced I_{KA-ch_ACH}) [3–4].

Dual drug effect

A common feature of the effect of ethanol and nicotine on I_{KAch} (similarly to the effect of ethanol on I_{K1}) is the unusual action of these drugs manifested by activation or inhibition at different drug concentrations (or in some cases at the same drug concentration in various cells) [1, 3–4]. $I_{\text{KAch}_\text{CONST}}$ was only activated by ethanol and nicotine throughout the whole range of the investigated concentrations (Figs 2 and 5). However, $I_{\text{KAch}_\text{ACH}}$ exhibited a slight inhibition at the lower ethanol concentrations, activation reaching a maximum at the medium concentrations, and a subsequent decrease of the activation effect (Fig 2). In the case of nicotine



Fig 6. Correlation between the relative effect of nicotine (nic) on I_{KAch} components and the current density under nicotine-free conditions (contr). Relative I_{KAch_CONST} (left); relative I_{KAch_ACH} (right). Nicotine concentrations of 4, 40, and 400 nM (black, red, and blue lines and symbols, respectively; 5, 6 and 5 cells for I_{KAch_CONST} at 4, 40 and 400 nM nicotine, respectively; 5, 11 and 7 cells for I_{KAch_ACH} at 4, 40 and 400 nM nicotine, respectively). Symbols: experimental results from individual cells according to Bébarová *et al.* [4]. Dashed lines: linear regression analysis of the experimental data for each concentration separately. Full lines: simulated results using values of parameters as stated in Table 2. The variations of the current in control were simulated by continuous redistribution between the channel populations (fractions f_1 and $f_2 = 1 - f_1$).



Fig 7. Effect of nicotine (40 nM) on the current-voltage relation of $I_{\text{KAch}_\text{CONST}}$. Experimental record (left panel; 3 cells) and simulated data (right panel). Activation of the current was simulated by redistribution of the channels between fractions: $(f_1 \rightarrow f_1 + \Delta f, f_2 \rightarrow f_2 - \Delta f, \Delta f = 0.19)$.

(Fig 5), the activation even reversed into an inhibition at the higher concentrations. Thus, it seems that the net effect of ethanol/nicotine modulation of Kir channels is determined by an alternant susceptibility to activation or inhibition.

These complex drug effects are interpreted in our model as resulting from two or more simpler processes related to the diversity of structures of the individual Kir channels formed by combination of the different α -subunits. Individual populations of identical channels are assumed to show different conductivities, and the total recorded whole cell current is described as a sum of *n* independent particular constituents exerting the drug-induced activation or



Fig 8. Steady-state concentration dependence of the effect of nicotine on I_{KAch_ACH} at -50 mV (above) and -110 mV (below) on a relative scale (currents under nicotine related to the respective control values; 5–12 cells in individual concentrations). The shift of the concentration dependence at -110 mV towards inhibition was simulated by a decrease of the activation related parameter h_{2_1} (from 0.085 to 0.037). All other parameters as summarized in Table 2 remained unaltered.

inhibition. To reproduce experimental results, n = 3 was sufficient to simulate the ethanol- I_{K1} interaction. However, in the present study, we reduced the number of different channel population to n = 2, in agreement with the prevailing notion that the I_{KAch} channels are composed of two different α -subunits forming assemblies of homo- and heterotetramers GIRK4 and GIRK1/4. For simplicity, we have assigned the activation effect to one of the two populations and the inhibition effect to the other.

In experiments on expressed GIRK channels, Arial *et al.* [20] demonstrated that shortchain alcohols may be bound to hydrophobic pockets (formed by two adjacent subunits of the tetramer). These pockets, which are able to bind one or two ethanol molecules, are assumed to be responsible for ethanol activation of the GIRK channels. If the binding of a single molecule causes activation more effectively than the binding of two molecules, the interaction of ethanol with the pocket results in the biphasic concentration dependence (Fig 2C).

The alcohol-binding pocket is not involved in the alcohol-dependent inhibition because mutations in the pocket had no effect on the alcohol-dependent inhibition, suggesting that the inhibition was related to an alternate site. Surprisingly, a mutation in the GIRK4 pore-helix (GIRK4^{S143T}) converted 1-butanol from an inhibitor to an activator (S3 Fig in ref. [20]).

It is worth mentioning that a dual effect similar to that described in this study was also demonstrated in the effect of the small molecule of halothane (volatile anaesthetic) on the GIRK channels expressed in *Xenopus oocytes* [5–6]. In line with our results, the constitutive active basal current was activated by halothane throughout the examined concentration range, while the acetylcholine-induced current was inhibited at lower concentrations of the drug, but activated at higher concentrations.

Interestingly, the dual effect of ethanol was also observed in the pentameric ligand-gated ion channels, consistent with a two-site model of the ethanol-induced activation and inhibition. Moreover, short-chain alcohols interacted with the excitatory site, while long-chain alcohols interacted with the inhibitory site [38–40].

Similarities and differences in the effects of drugs on Kir currents

The relative changes induced by both ethanol (Fig 2) and nicotine (Fig 5) were considerably greater in $I_{\text{KAch}_{\text{CONST}}}$ than in $I_{\text{KAch}_{\text{ACH}}}$. Furthermore, while $I_{\text{KAch}_{\text{CONST}}}$ was always activated, $I_{\text{KAch}_{\text{ACH}}}$ was activated at moderate concentrations of both drugs and slightly inhibited at a low concentration of ethanol and high concentrations of nicotine.

The peculiar behaviour of the Kir3.x (GIRK) channels under the effect of ethanol has been explained by different affinities of these channels for phosphatidylinositol 4,5-bisphosphate (PIP₂) in the absence and presence of acetylcholine [20,41]. All Kir channels are directly activated by phospholipid PIP₂ with various affinities [25, 42–45]. It has been demonstrated that the channels with high affinity for PIP₂ (such as Kir2.x channels that are essentially constitutively active) show high currents, but low sensitivity to activation by ethanol and other drugs [46]. The high affinity for PIP₂ is also accompanied by a tendency toward drug-induced inhibition [20].

The Kir3.x (GIRK) channels in the absence of acetylcholine exhibit a low affinity for PIP₂, and are characterised by a low basal current with a high tendency to be activated by ethanol [20]. This corresponds to a small basal current ($I_{\text{KAch}_{\text{CONST}}}$) and marked drug-induced current activation observed in rat atrial cells (Figs 2 and 5). Acetylcholine markedly increased the I_{KAch} (very likely due to the increased channel affinity for PIP₂) and suppressed sensitivity to drug-induced activation. Concurrently, the inhibitory effects of ethanol at low concentrations (Fig 2) and of nicotine at high concentrations (Fig 5) were revealed. There is an analogy between the ethanol effect on I_{KAch} ACH and I_{K1} [1]. In both cases the channels bind PIP₂

strongly, and thus the basal current is high and shows a tendency to be partially inhibited at low concentrations of ethanol. Our model does not describe the role of PIP₂ explicitly; however the impact of PIP₂ is implicit in values of the parameters $h_{\rm k}$ j.

Inverse correlation between I_{KAch} in control and the drug effect

An inverse correlation between the amplitude of I_{KAch} in the control conditions and the relative effect of drugs was another peculiar observation reproduced by the model simulations (Figs 3 and 6). In accord with the considerations in the previous section, the drug effect decreased with the absolute value of the control current. The low $I_{\text{KAch}_\text{CONST}}$ currents were always activated, while the $I_{\text{KAch}_\text{ACH}}$ currents (which were one order higher) were activated or inhibited at different drug concentrations. The model parameters were set primarily to reproduce the concentration dependences of the drug effect, including dispersion. The inverse correlation between the drug effect and I_{KAch} in control conditions was then also reproduced without readjusting the parameters, but only by varying the ratio of the fractions f_1 / f_2 .

Voltage dependence of the drug effects on I_{KAch}

The presented model was formulated under the assumption that the drug effects were voltage independent. Consequently, the parameters in Tables 1 and 2 were regarded as voltage independent. In other words, the interaction of drugs with the mechanism responsible for inward rectification was assumed to be negligible. This condition was not fully satisfied in the case of nicotine. Compared with the relative effect of nicotine on $I_{\text{KAch}_\text{ACH}}$ at -50 mV, the effect assessed at -110 mV was shifted towards inhibition at all applied concentrations (Fig 3C in [4]). A similar shift (between the effect at -110 and -40 mV) was observed under the effect of relatively large molecules of local anaesthetic bupivacaine (pure inhibitor) on GIRK1/4 channels [41]. In our model, simulation of the shift of relative concentration dependence at -110 mV against that at -50 mV (Fig 8) required a reduction of the single parameter (h_{2_1}) responsible for the activation.

Limitations of the model

The inwardly rectifying potassium currents are newly understood to be sums of individual constituents related to the families of identical channels characterized by the same structure. Although the model is based on the current knowledge of channel structure and function, the need for simple quantitative description did not allow for the capturing of detailed processes at the molecular level. In particular, the processes responsible for the voltage dependence (inward rectification) are not included, and the current-voltage relations are described by a formal mathematical expression. The model parameters are voltage-independent constants in order to comply with the observation that the ethanol effect itself appeared to be approximately voltage-independent. Nevertheless, one parameter had to be regarded voltage-dependent to simulate the impact of the voltage on concentration dependence of the nicotine effect (Fig 8).

The number of populations of identical channels is generally unknown. The value of n = 2 was chosen in agreement with the current view that the I_{KAch} channels in the atrial cells are restricted to two functional assemblies of α -subunits GIRK1/4 and GIRK4. Their specific assignment to the two channel populations in the model remains undetermined. It is unclear to what extent GIRK4 homo-tetramers versus GIRK1/4 hetero-tetramers dominate in cardiac cells [16]. The option of $f_1 = 0.68$, i.e. $f_1/f_2 = 2.125$ appeared to be the optimal mean value to display all the observed phenomena as a result of random variations of a single parameter f_1/f_2 . Simultaneously, the concentration dependence of the drug-effects, the observed scatter of the measured I_{KAch} values, and the dependence of the drug-effects on the baseline currents

must be displayed. The observed scatter of the measured I_{KAch} currents is attributed to random variations of the f_1/f_2 ratio according to the model; however, the proportion of the experimental error may not be negligible.

As mentioned above, the experimental data related to I_{K1} included transient changes of the current in response to the onset of ethanol. Corresponding transients in conductivities were described by Šimurda *et al.* in ref. [10] by differential equations. Drug-induced changes of I_{KAch} during transition to their steady-states have not yet been reported, and so such data is missing in this study. For possible future simulations, the model could be supplemented by the differential Eq (8). The conductivities G_1 and G_2 in Eq (3) refer to their steady-state values. Their transient changes at a given voltage in response to sudden drug application (designated here $G_{1,t}$ and $G_{2,t}$) are assumed to obey first-order differential equations

$$\frac{G_{1,t}}{dt} = \frac{1}{\tau_1} (G_{1,t} - G_1), \quad \frac{G_{2,t}}{dt} = \frac{1}{\tau_2} (G_{2,t} - G_2), \tag{8}$$

where τ_1 and τ_2 are time constants. The corresponding time courses of the current in absolute and relative scale (I_t and F_t) can be obtained if the quantities G_1 and G_2 in Eqs (4) and (6) are substituted by $G_{1,t}$ and $G_{2,t}$.

Localization of the binding sites for inhibition has not yet been fully elucidated. Assignment of activation to the first population (f_1) and inhibition to the second population (f_2) is more or less arbitrary. It is possible that each population includes both mechanisms with activation predominating in one and inhibition in the other. Their detailed incorporation into the model would unduly complicate mathematical description without bringing new insights into the problem. The number of parameters that should be fitted to the given experimental data would increase several times. Our main goal was to show that the mere fact that Kir channels are composed of populations with different functional properties is able to explain the complex drug-effects, not only in Kir2.x but also in Kir3.x channels.

Conclusions

We conclude that the proposed model may serve as a useful tool for analysis and simulation of complicated manifestations of the interaction of some substances with the Kir channels. This approach is flexible and can be easily modified if new essential experimental findings become available. An analogical approach could be also used to describe other types of ionic channels—channels that do not show inward rectification but that may exhibit several possible channel populations with specific properties. Moreover, the model can be readily built into the existing integral models of electrical activity of cardiac cells in order to upgrade the description of Kir currents and their interaction with drugs.

Materials and methods

Experimental data

The experimental data were obtained from our previously published studies [3–4], which give detailed information on the materials and methods. The constitutively active I_{KAch} (I_{KAch} -_{CONST}) was evaluated as the current sensitive to its specific inhibitor tertiapin-Q (300 nM) and I_{KAch} active the steady-state application of 3 μ M acetylcholine (Fig 9). Both currents were measured at the end of a 500-ms rectangular pulse, from -85 mV to -110 or to -50 mV. To measure voltage dependence of the drug effects, 3-s ramp pulses between -110 and -10 mV were applied. In all experiments, the holding voltage was set



Fig 9. Representative time courses of the membrane current density during application of specific inhibitor of the I_{KAch} channels tertiapin (Ter; 300 nM) to reveal the constitutively active component of I_{KAch} (I_{KAch_CONST} ; left panels), or during application of acetylcholine (Ach; 3 μ M) to activate the acetylcholine-activated component of I_{KAch} (I_{KAch_ACH} ; right panels), both in the absence and in the presence of ethanol (A) or nicotine (B) at selected concentrations.

to -85 mV and the stimulation frequency to 0.2 Hz. When necessary, the previously published data were recalculated to obtain the intended graphs (Fig 3 and Fig 6 left).

Computer simulations

The Eqs (2)–(7), describing the effect of ethanol and nicotine on the rat atrial $I_{\text{KAch}_\text{CONST}}$ and $I_{\text{KAch}_\text{ACH}}$ [3–4] with values of parameters as defined in Tables 1 and 2, were solved numerically using the computational software MATLAB v. 7.2 (MathWorks, Inc.). The MATLAB files for modelling of ethanol and nicotine effects on I_{KAch} are available from authors upon request.

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References

- Bébarová M, Matejovič P, Pásek M, Šimurdová M, Šimurda J. Dual effect of ethanol on inward rectifier potassium current *I*K1 in rat ventricular myocytes. J Physiol Pharmacol. 2014; 65: 497–502. PMID: 25179082
- Hořáková Z, Matejovič P, Pásek M, Hošek J, Šimurdová M, Šimurda J. Effect of ethanol and acetaldehyde at clinically relevant concentrations on atrial inward rectifier potassium current *IK*1: separate and combine effect. J Physiol Pharmacol. 2016; 67: 339–351. PMID: 27511995
- Bébarová M, Matejovič P, Pásek M, Hořáková Z, Hošek J, Šimurdová M, et al. Effect of ethanol at clinically relevant concentrations on atrial inward rectifier potassium current sensitive to acetylcholine. Naunyn-Schmiedeberg s Arch Pharmacol. 2016; 389: 1049–1058.
- Bébarová M, Matejovič P, Švecová O, Kula R, Šimurdová M, Šimurda J. Nicotine at clinically relevant concentrations affects atrial inward rectifier potassium current sensitive to acetylcholine. Naunyn-Schmiedeberg s Arch Pharmacol. 2017; 390: 471–481.
- Weigl LG, Schreibmayer WG. Protein-gated inwardly rectifying potassium channels are targets for volatile anesthetics. Mol Pharmacol. 2001; 60: 282–289. https://doi.org/10.1124/mol.60.2.282 PMID: 11455015
- Milovic S, Steinecker-Frohnwieser B, Schreibmayer W, Weigl LG. The sensitivity of G protein-activated K+ channels toward halothane is essentially determined by the C terminus. J Biol Chem. 2004; 279: 34240–34249. https://doi.org/10.1074/jbc.M403448200 PMID: 15175324
- Dhamoon AS, Pandit SV, Sarmast F, Parisian KR, Guha P, Li Y, et al. Unique Kir2.x properties determine regional and species differences in the cardiac inward rectifier K+ current. Circ Res. 2004; 94: 1332–1339. https://doi.org/10.1161/01.RES.0000128408.66946.67 PMID: 15087421
- Ehrlich JR. Inward rectifier potassium currents as a target for atrial fibrillation therapy. J Cardiovasc Pharmacol. 2008; 52: 129–135. https://doi.org/10.1097/FJC.0b013e31816c4325 PMID: 18670367
- Hibino H, Inanobe A, Furutani K, Murakami S, Findlay I, Kurachi Y. Inwardly rectifying potassium channels: their structure, function, and physiological roles. Physiol Rev. 2010; 90: 291–366. <u>https://doi.org/ 10.1152/physrev.00021.2009 PMID: 20086079</u>
- Śimurda J, Śimurdová M, Bébarová M. Inward rectifying potassium currents resolved into components: modeling of complex actions. Pflugers Arch–Eur J Physiol. 2018; 470: 315–325.
- Krapivinsky G, Gordon EA, Wickman K, Velimirovic B, Krapivinsky L, Clapham DE. The G-proteingated atrial K+ channel /KACh is a heteromultimer of two inwardly rectifying K+-channel proteins. Nature. 1995; 374: 135–141. https://doi.org/10.1038/374135a0 PMID: 7877685
- Corey S, Krapivinsky G, Krapivinsky L, Clapham DE. Number and stoichiometry of subunits in the native atrial G-protein-gated K+ channel, *I*KACh. J Biol Chem. 1998; 273: 5271–5278. https://doi.org/ 10.1074/jbc.273.9.5271 PMID: 9478984
- Corey S, Clapham DE. Identification of native atrial G-protein-regulated inwardly rectifying K+ (GIRK4) channel homomultimers. J Biol Chem. 1998; 273: 27499–27504. PMID: 9765280
- Kennedy ME, Nemec J, Corey S, Wickman K, Clapham DE. GIRK4 confers appropriate processing and cell surface localization to G-protein-gated potassium channels. J Biol Chem. 1999; 274: 2571–2582. https://doi.org/10.1074/jbc.274.4.2571 PMID: 9891030
- Bender K, Wellner-Kienitz MC, Inanobe A, Meyer T, Kurachi Y, Pott L. Overexpression of monomeric and multimeric GIRK4 subunits in rat atrial myocytes removes fast desensitization and reduces inward rectification of muscarinic K+current (IK(ACh)). Evidence for functional homomeric GIRK4 channels. J Biol Chem. 2001; 276: 28873–28880. https://doi.org/10.1074/jbc.M102328200 PMID: 11384974
- Touhara KK, Wang W, MacKinnon R (2016) The GIRK1 subunit potentiates G protein activation of cardiac GIRK1/4 hetero-tetramers. eLife; 5:e15750. https://doi.org/10.7554/eLife.15750 PMID: 27074664
- Mirshahi T, Logothetis DE. Molecular determinants responsible for differential cellular distribution of G protein-gated inwardly rectifying K+ channels. J Biol Chem. 2004; 279: 11890–11897. PMID: <u>14703518</u>
- Inanobe A, Kurachi Y. Membrane channels as integrators of G-protein-mediated signaling. Biochim Biophys Acta. 2014; 1838: 521–531. https://doi.org/10.1016/j.bbamem.2013.08.018 PMID: 24028827

- Lewohl JM, Wilson WR, Mayfield RD, Brozowski SJ, Morrisett RA, Harris RA. Gprotein coupled inwardly rectifying potassium channels are targets of alcohol action. Nat Neurosci. 1999; 2: 1084–1090. https://doi.org/10.1038/16012 PMID: 10570485
- Aryal P, Dvir H, Choe S, Slesinger PA. A discrete alcohol pocket involved in GIRK channel activation. Nat Neurosci. 2009; 12: 988–995. https://doi.org/10.1038/nn.2358 PMID: 19561601
- Mahajan R, Ha J, Zhang M, Kawano T, Kozasa T, Logothetis DE. Computational model predicts that Gβγ acts at a cleft between channel subunits to activate GIRK1 channels. Sci Signal. 2013; 6:ra69. https://doi.org/10.1126/scisignal.2004075 PMID: 23943609
- Toyama Y, Kano H, Mase Y, Yokogawa M, Osawa M, Shimada I. Structural basis for the ethanol action on G-protein–activated inwardly rectifying potassium hannel 1 revealed by NMR spectroscopy. PNAS. 2018; 115: 3858–3863. https://doi.org/10.1073/pnas.1722257115 PMID: 29581303
- Bodhinathan K, Slesinger PA. Molecular mechanism underlying ethanol activation of G-protein-gated inwardly rectifying potassium channels. Proc Natl Acad Sci USA. 2013; 110: 18309–18314. https://doi. org/10.1073/pnas.1311406110 PMID: 24145411
- Treiber F, Rosker C, Keren-Raifman T, Steinecker B, Gorischek A, Dascal N, et al. Molecular basis of the facilitation of the heterooligomeric GIRK1/GIRK4 complex by cAMP dependent protein kinase. Biochim Biophys Acta. 2013; 1828: 1214–1221. <u>https://doi.org/10.1016/j.bbamem.2012.12.016</u> PMID: 23305758
- Li J, Lü S, Liu Y, Pang C, Chen Y, S Zhang, et al. Identification of the conformational transition pathway in PIP2 opening Kir channels. Sci Rep. 2015; 5: 11289. <u>https://doi.org/10.1038/srep11289</u> PMID: 26063437
- Christé G, Tebbakh H, Šimurdová M, Forrat R, Šimurda J. Propafenone blocks ATP-sensitive K+ channels in rabbit atrial and ventricular cardiomyocytes. Eur J Pharmacol. 1999; 373: 223–232. https://doi. org/10.1016/s0014-2999(99)00217-4 PMID: 10414443
- Kobayashi T, Washiyama K, Ikeda K. Inhibition of G protein-activated inwardly rectifying K+ channels by various antidepressant drugs. Neuropsychopharmacology. 2004; 29: 1841–1851. <u>https://doi.org/10. 1038/sj.npp.1300484</u> PMID: 15150531
- Kobayashi T, Washiyama K, Ikeda K. Inhibition of G protein-activated inwardly rectifying K+ channels by ifenprodil. Neuropsychopharmacology. 2006; 31: 516–524. <u>https://doi.org/10.1038/sj.npp.1300844</u> PMID: 16123769
- Caballero R, Dolz-Gaitón P, Gómez R, Amorós I, Barana A, González de la Fuente M, et al. Flecainide increases Kir2.1 currents by interacting with cysteine 311, decreasing the polyamine-induced rectification. PNAS. 2010; 107: 15631–15636. https://doi.org/10.1073/pnas.1004021107 PMID: 20713726
- Walsh KB. A real-time screening assay for GIRK1/4 channel blockers. J Biomol Screen. 2010; 15: 1229–1237. https://doi.org/10.1177/1087057110381384 PMID: 20938046
- Ferrer T, Ponce-Balbuena D, López-Izquierdo A, Aréchiga-Figueroa IA, de Boer TP, van der Heyden MAG, et al. Carvedilol inhibits Kir2.3 channels by interference with PIP2-channel interaction. Eur J Pharmacol. 2011; 668: 72–77. https://doi.org/10.1016/j.ejphar.2011.05.067 PMID: 21663737
- Liu QH, Li XL, Xu YW, Lin YY, Cao JM, Wu BW. A novel discovery of *I*K1 channel agonist: zacopride selectively enhances *I*K1 current and suppresses triggered arrhythmias in the rat. J Cardiovasc Pharmacol. 2012; 59: 37–48. https://doi.org/10.1097/FJC.0b013e3182350bcc PMID: 21921806
- Vanheiden S, Pott L, Kienitz MC. Voltage-dependent open-channel block of G protein-gated inward-rectifying K+ (GIRK) current in rat atrial myocytes by tamoxifen. Naunyn-Schmiedeberg's Arch Pharmacol. 2012; 385: 1149–1160.
- Gómez R, Caballero R, Barana A, Amorós I, DePalm SH, Matamoros M, et al. Structural basis of drugs that increase cardiac inward rectifier Kir2.1 currents. Cardiovasc Res. 2014; 104: 337–346. <u>https://doi.org/10.1093/cvr/cvu203 PMID: 25205296</u>
- Kobayashi T, Ikeda K, Kojima H, Niki H, Yano R, Yoshioka T, et al. Ethanol opens G-protein-activated inwardly rectifying K+ channels. Nat Neurosci. 1999; 2: 1091–1097. <u>https://doi.org/10.1038/16019</u> PMID: 10570486
- Yamakura T, Lewohl JM, Harris RA. Differential effects of general anesthetics on G protein–coupled inwardly rectifying and other potassium channels. Anesthesiology. 2001; 95: 144–153. <u>https://doi.org/ 10.1097/00000542-200107000-00025</u> PMID: <u>11465552</u>
- Bébarová M, Matejovič P, Pásek M, Ohlídalová D, Jansová D, Šimurdová M, et al. Effect of ethanol on action potential and ionic membrane currents in rat ventricular myocytes. Acta Physiol (Oxf). 2010; 200: 301–314.
- Zuo Y, Aistrup GL, Marszalec W, Gillespie A, Chavez-Noriega LE, Yeh JZ, et al. Dual action of n-alcohols on neuronal nicotinic acetylcholine receptors. J Mol Pharmacol. 2001; 6: 700–711.

- Borghese CM, Henderson LA, Bleck V, Trudell JR, Harris RA. Sites of excitatory and inhibitory actions of alcohols on neuronal α2 β4 nicotinic acetylcholine receptors. J Pharmacol Exper Ther. 2003; 307: 42–52.
- Murail S, Howard RJ, Broemstrup T, Bertaccini EJ, Harris RA, Trudell JR, et al. Molecular mechanism for the dual alcohol modulation of cys-loop receptors. PloS Comput Biol. 2012; 8: e1002710. https://doi. org/10.1371/journal.pcbi.1002710 PMID: 23055913
- Zhou W, Arrabit C, Choe S, Slesinger PA. Mechanism underlying bupivacaine inhibition of G proteingated inwardly rectifying K+ channels. PNAS. 2001; 98:6482–6487. https://doi.org/10.1073/pnas. 111447798 PMID: 11353868
- 42. Huang CL, Feng S, Hilgemann DW. Direct activation of inward rectifier potassium channels by PIP2 and its stabilization by Gβγ. Nature. 1998; 391: 803–806. <u>https://doi.org/10.1038/35882</u> PMID: 9486652
- Zhang H, He C, Yan X, Mirshahi T, Logothetis DE. Activation of inwardly rectifying K+ channels by distinct PtdIns(4,5)P2 interactions. Nat Cell Biol. 1999; 1: 183–188. <u>https://doi.org/10.1038/11103 PMID: 10559906</u>
- 44. Xie LH, John SA, Ribalet B, Weiss JN. Activation of inwardly rectifying potassium (Kir) channels by phosphatidylinosital-4,5-bisphosphate (PIP2): interaction with other regulatory ligands. Prog Biophys Mol Biol. 2007; 94: 320–335. https://doi.org/10.1016/j.pbiomolbio.2006.04.001 PMID: 16837026
- 45. Lacin E, Aryal P Glaaser IW, Bodhinathan K, Tsai E, Marsh N, et al. Dynamic role of the tether helix in PIP2-dependent gating of a G protein–gated potassium channel. J Gen Physiol. 2017 Jul 18. pii: jgp.201711801. https://doi.org/10.1085/jgp.201711801 PMID: 28720589
- Logothetis DE, Jin T, Lupyan D, Rosenhouse-Dantsker A. Phosphoinositidemediated gating of inwardly rectifying K⁺ channels. Pflugers Arch—Eur J Physiol. 2007; 455: 83–95.