Mechanisms underlying probucol-induced hERG-channel deficiency

Abstract: The hERG gene encodes the pore-forming α-subunit of the rapidly activating delayed rectifier potassium channel ($I_{Kr}$), which is important for cardiac repolarization. Reduction of $I_{Kr}$ due to genetic mutations or drug interferences causes long QT syndrome, leading to life-threatening cardiac arrhythmias (torsades de pointes) or sudden death. Probucol is a cholesterol-lowering drug that could reduce hERG current by decreasing plasma membrane hERG protein expression and eventually cause long QT syndrome. Here, we investigated the mechanisms of probucol effects on $I_{Kr}$ and hERG-channel expression. Our data demonstrated that probucol reduces SGK1 expression, known as SGK isoform, in a concentration-dependent manner, resulting in downregulation of phosphorylated E3 ubiquitin ligase Nedd4-2 expression, but not the total level of Nedd4-2. As a result, the hERG protein reduces, due to the enhanced ubiquitination level. On the contrary, carbachol could enhance the phosphorylation level of Nedd4-2 as an alternative to SGK1, and thus rescue the ubiquitin-mediated degradation of hERG channels caused by probucol. These discoveries provide a novel mechanism of probucol-induced hERG-channel deficiency, and imply that carbachol or its analog may serve as potential therapeutic compounds for the handling of probucol cardiotoxicity.

Keywords: long QT, hERG potassium channels, probucol, SGK1, Nedd4-2

Introduction

The hERG gene encodes the pore-forming subunits of channels that conduct the rapid delayed rectifier potassium current.1 Mutation of the hERG gene may cause dysfunction of rapidly activating delayed rectifier potassium channel ($I_{Kr}$) and thus lead to cardiac arrhythmia, such as congenital long QT syndrome (LQTS).2 In addition, a wide variety of potentially dangerous proarrhythmic effects of antiarrhythmic or nonantiarrhythmic drugs have been recognized to inhibit hERG channels, which could induce acquired LQT2.3,4 Diverse therapeutic compounds, agents of distinct pharmacological classes have been removed from the market; several drugs have been limited in clinical applications, due to their risk of ventricular arrhythmias.3 Since most of these adverse reactions are preventable, increased knowledge of drug application associated with repolarization abnormalities and careful evaluation of risk factors is strongly recommended. At present, there are no effective therapeutic methods to restore or enhance hERG-channel function. Therefore, looking for compounds or therapies is extremely necessary right now.

Probucol is a cholesterol-lowering drug initially developed in the treatment of coronary artery disease,3 which reduces $I_{Kr}$ by decreasing membrane hERG-channel expression and thus causes LQTS.6 Although probucol was identified as the hERG inhibitor responsible for LQTS, the underlying mechanism is still not fully understood. Therefore, a better mechanistic understanding of probucol-induced hERG-channel abnormality may
help us to develop therapeutic compounds that are prone to the
disruption of hERG-channel processing and trafficking.

In this study, we investigated the mechanism underlying
probucol-induced hERG-channel deficiency. It has been
confirmed that SGK1 expression is regulated during both
discrete developmental stages and pathological conditions,
such as hypertension, drug interferences, diabetic neuropathy,
ischemia, trauma, and neurodegenerative diseases. Over
the past few years, there has been increasing evidence that over-
expression of SGK1 is able to increase the current and expres-
sion level of the mature proteins of hERG channels localized
in the membrane. SGK1 and its isoforms are expressed in
every tissue, including the heart, and play a significant role in
the regulation of hERG-channel expression. Interestingly, the
downstream targets of SGK1 include the E3 ubiquitin ligase
Nedd4-2, which is responsible for substrate recognition in the
glorification process of proteins. Also, it has been reported that
SGK inhibits Nedd4-2 activity by phosphorylating Nedd4-2
at Ser-444 residues present in the WW domains; this conse-
quently enhances cell-surface expression of the hERG chan-
nel, which is a direct substrate of Nedd4-2. Also, probucol
and As2O3 are the two main kinds of drugs to decrease hERG
expression and subsequently cause LQTS. Therefore, we
wondered whether these two drugs, which can induce
hERG-channel deficiency, were involved in the ubiquitina-
protein-degradation pathway. Based on this, we became
interested in SGK1 expression after treatment with different
concentrations of probucol or As2O3 in hERG-HEK293 cells.
As the research proceeded much further, our data provided
evidence that the SGK1–Nedd4-2 axis represents a novel
mechanism for probucol to decrease hERG-channel expression
and eventually cause LQTS. Therefore, we discovered that carbachol (CCh) impeded the reduction of
the 155 kDa hERG expression induced by long-term treatment
of probucol through enhancement of the phosphorylation level
of Nedd4-2 as an alternative to SGK1.

Materials and methods
Reagents
Probucol and CCh were purchased from Sigma-Aldrich
Co (St Louis, MO, USA). In cellular experiments, hERG-
HEK293 cells were incubated with different concentrations
of probucol for 48 hours with or without different concentra-
tions of CCh, which were diluted in cultured medium.

Cell culture
Experiments were performed on HEK293 cells stably
expressing the wild-type hERG gene. Cells were cultured
in Dulbecco’s Modified Eagle’s Medium (HyClone, Logan,
UT, USA) supplemented with 10% (v/v) fetal bovine
serum (Thermo Fisher Scientific, Waltham, MA, USA) and
400 mg/mL Geneticin (G-418; Thermo Fisher Scientific) in
an atmosphere of 95% humidified air and 5% CO2, at 37°C.
For electrophysiological studies, the cells were harvested
from the culture dishes by 0.25% trypsinase, washed twice
with standard Dulbecco’s Modified Eagle’s Medium, and
stored in this medium at room temperature for later research.
Cells were investigated within 8 hours of harvest.

Western blot analysis
Western blot experiments were used to detect the expression
levels of interest proteins. Drugs were diluted and added to
hERG-HEK293 cells for 48 hours at 37°C before analysis
by Western blot. The cells were washed using ice-cold
phosphate-buffered saline, and total protein was harvested
with radioimmunoprecipitation-assay buffer containing 1%
protease inhibitor (Sigma-Aldrich Co). Protein (100 μg)
per sample was separated using 8% sodium dodecyl sulfate
polyacrylamide gel electrophoresis, and then transferred
onto nitrocellulose membranes (Agilent Technologies, Santa
Clara, CA, USA). The membrane was blocked afterward
using 5% nonfat milk and 0.1% Tween 20 in Tris-buffered
saline and immunoblotted overnight using appropriate primary
antibodies at 4°C with gentle shaking. Goat antirabbit
or goat antimouse fluorochrome-labeled Alexa Fluor 800
(dilution 1:8,000; Li-Cor Inc, Lincoln, NE, USA) was used
as a secondary antibody. The Odyssey infrared fluorescent
scanning system (Li-Cor Inc) was used to detect signals of
membrane proteins. To quantify the Western blot data, the
intensities of proteins of interest in each gel are firstly normal-
ized to their respective actin intensities, then the normalized
intensities are compared with the intensity of control group
and expressed as relative values to their controls.

Patch-clamp recording techniques
The whole-cell patch-clamp technique was used to measure
hERG currents. Heat-polished patch pipettes had final
resistances of 2–4 MW when filled with a pipette solu-
tion containing 130 mM KCl, 1 mM MgCl2-6H2O, 10 mM
HEPES, 5 mM Mg-adenosine triphosphate, 5 mM ethylene
glycol tetraacetic acid, and 0.1 mM guanosine triphosphate,
ph 7.3 with KOH. The standard 5 mM K+ bath solution con-
tained 136 mM NaCl, 5.4 mM KCl, 5 mM HEPES, 1 mM
MgCl2-6H2O, 1 mM CaCl2, and 10 mM glucose, ph 7.4 with
NaOH. For electrophysiological recordings, submaximally
confluent cells were used after 1–3 days of culture at a seeding
density of 1–4×10^5 cells per flask (Nest Biotechnology, Wuxi, People’s Republic of China). The cells were harvested from the plate by treatment with 0.25% trypsin and 0.02% ethylenediaminetetraacetic acid, and transferred to a small cell bath mounted on the stage of an inverted microscope (IX-70; Olympus Corporation, Tokyo, Japan), where they were allowed to attach to the glass bottom for about 10 minutes. The cells were then superfused continuously at a ratio of 1.5 mL/min. The experiment set a holding potential voltage from −80 mV to voltages between −70 and 70 mV in 10 mV increments. Tail currents upon repolarization to the holding potential of −80 mV after the depolarizing pulse to 50 mV were used for the current amplitudes in control and treated groups. Patch-clamp experiments were all performed at room temperature. An Axopatch 200B patch-clamp amplifier was used to record membrane current. Computer software (Clampex 9.2; Molecular Devices LLC, Sunnyvale, CA, USA) was used to generate voltage-clamp protocols and acquire data. Capacitance and series-resistance compensation were optimized. Data were recorded on a computer via Digidata 1322A, and analyzed using Clampfit 9.2 (Molecular Devices LLC) and Prism (GraphPad Software Inc, La Jolla, CA, USA) software. Graphical fits of the data were made using previously described standard equations.14,15

Statistical analyses

Data are expressed as means ± standard error of mean. A one-way analysis of variance or Student’s t-test was used to determine the significance of differences between control and test groups. A P-value of 0.05 or less was considered significant (two-tailed). Boltzmann distribution was used to fit voltage-dependent activation and inactivation curves, and a single exponential function was used to fit the curve of recovery from inactivation (reactivation). All graphs were drawn by GraphPad Prism 5.0.

Results

Probucol significantly reduced SGK1 expression

The stress-responsive SGK belongs to a family of serine/threonine kinases that is under acute transcriptional control by several stimuli.16 It has been demonstrated that overexpression of SGK1 increased the current and expression level of the membrane-localized mature proteins of hERG channels stably expressed in hERG-HEK293 cells.8 Probucol reduces hERG-channel expression in the plasma membrane in a concentration-dependent manner.6 Therefore, we used a concentration gradient of probucol to investigate the mechanisms for probucol-induced hERG-channel deficiency. Cells were cultured in either control medium or probucol (30, 50, and 100 μM) containing medium for 48 hours; hERG-expression levels and I_{hERG} were then examined. As depicted in Figure 1A, probucol significantly reduced SGK1 expression in a concentration-dependent manner, indicating that probucol reliably downregulated SGK1 expression. In our previous research, we demonstrated that As2O3 can damage the I_{hERG} current via disturbing its trafficking to cellular membrane.17 In order to estimate whether As2O3-induced hERG-channel inhibition was involved in the alteration of SGK1 level, we substituted the probucol with 3 μM As2O3.

![Figure 1](https://www.dovepress.com/)

**Figure 1** Probucol (Pro) significantly reduced the expression level of SGK1.

**Notes:** (A) Downregulation of SGK1 expression after treatment with Pro at different concentrations of Pro (30, 50, and 100 μM) for 48 hours determined by Western blotting. (B) Western blotting showing SGK1 expression in hERG-HEK293 cells under control conditions, with 3 μM As2O3 treatment for 24 hours. *P<0.05 versus control; n=6.
After incubation of 3 μM As₂O₃ for 24 hours, the SGK1 protein level was also detected as previously. Figure 1B shows that As₂O₃ had no significant effect on SGK1 expression.

**Phosphorylation level of Nedd4-2 decreased due to reduced expression of SGK1**

Nedd4-2, also known as E3 ubiquitin ligase, is responsible for substrate recognition and ubiquitin transfer to target proteins. The phosphorylated form of Nedd4-2 (p-Nedd4-2) leads to the inactivation of its catalytic activity. It has been confirmed that Nedd4-2 can be phosphorylated by SGK1 in response to various stimuli. We have previously shown that probucol significantly reduces SGK1 expression. Since Nedd4-2 is the direct downstream target of SGK1, we suspected that reduced SGK1 levels may result in decreases of phosphorylated Nedd4-2 levels. Therefore we proposed the hypothesis that probucol treatment decreases hERG-channel expression by increasing Nedd4-2-mediated hERG ubiquitination. Interestingly, our data provide evidence that probucol treatment significantly decreased p-Nedd4-2, but not the total level of Nedd4-2 (Figure 2A and B). Meanwhile, As₂O₃ is also a major drug we do research on. To determine whether As₂O₃-induced hERG-channel abnormality results from enhanced ubiquitination, we performed our experiment by detecting the effect of As₂O₃ on Nedd4-2 and p-Nedd4-2 expression. As shown in Figure 2C and D, As₂O₃ had no significant effect on either Nedd4-2 or p-Nedd4-2. These findings indicated that As₂O₃-induced hERG-channel abnormality was probably due to other reasons, like trafficking deficiency.

**Carbachol rescued probucol-induced hERG-channel deficiency**

CCh, which is a cholinomimetic drug that binds and activates the acetylcholine receptor and increases the expression

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**Figure 2** Probucol treatment downregulated the phosphorylated Nedd4-2 (p-Nedd4-2), but not the total level of Nedd4-2. **Notes:** (A) There was no effect on the total level of Nedd4-2 in the absence and presence of probucol (30, 50, and 100 μM) for 48 hours. (B) Probucol reduced the p-Nedd4-2 level in a concentration-dependent manner. n=6, *P<0.05 and **P<0.01 versus control. (C, D) Western blotting showing Nedd4-2 and p-Nedd4-2 expression level in hERG-HEK293 cells in the presence and absence of 3 μM As₂O₃ treatment for 24 hours.
and function of hERG. CCh-mediated increase in hERG expression was accomplished by enhancement of the phosphorylation level but not the total level of Nedd4-2.20 To further demonstrate probucol-induced hERG-channel deficiency caused by Nedd4-2 involved enhancement of ubiquitination, hERG-HEK293 cells were cultured with different concentrations (25 and 50 μM) of CCh and 100 μM probucol for 48 hours; hERG-expression levels and \( I_{\text{HERG}} \) were then examined. Previous research has demonstrated that hERG-channel protein is primarily synthesized in the endoplasmic reticulum as the immature core-glycosylated form with a molecular mass of 135 kDa, and then transported to the Golgi apparatus for glycosylation to become the mature and functional form with a molecular mass of 155 kDa.21 Consistent with our previous study, the expression level of hERG, especially the mature form, declined approximately 50% compared to the control group with the incubation of probucol. However, in the probucol and CCh coinubicated groups, the inhibition of hERG channels was reversed. As shown in Figure 3A, CCh (25 and 50 μM) reversed the downregulation of hERG 155 kDa channels induced by 100 μM probucol. However, CCh (50 μM) did not reverse the \( \text{As}_3\text{O}_3 \)-induced hERG-channel deficiency (Figure 3B). Meanwhile, we also detected the hERG current after treatment with probucol and CCh. Likewise, CCh (25 and 50 μM) reversed the downregulation of \( I_{\text{HERG}} \) induced by 100 μM probucol (Figure 3C and D).

To demonstrate further whether CCh-imped probucol-induced reduction of 155 kDa hERG expression was the result of alteration of p-Nedd4-2, we detected the p-Nedd4-2 expression level under equivalent conditions. Interestingly, the increased p-Nedd4-2 was accompanied by an increased expression of the mature hERG band. Figure 3E illustrates that CCh (25 and 50 μM) treatment significantly increased p-Nedd4-2 compared with the probucol-treatment group. It has been demonstrated that blockade of the PKC pathway abolishes the CCh-induced enhancement of hERG channels.20 Based on this, we used the PKC inhibitor HA to eliminate the effect of CCh on Nedd4-2, and the Western blot results showed that after treatment with HA, CCh did not reverse the reduced p-Nedd4-2 caused by probucol (Figure 3F).

**Effects of CCh on APD prolonged by probucol on neonatal cardiac myocytes**

To test whether our analysis in heterologous expression systems might be extended to ventricular cardiomyocytes, we studied the effects of extracellular application of probucol and CCh on action potentials (APs) evoked in neonatal cardiac myocytes. Figure 4 illustrates the representative traces showing the effects of CCh on AP duration (APD) prolonged by probucol. Changes in AP waveform were described by using 50% and 90% repolarization of APD (APD\(_{50}\) and APD\(_{90}\), respectively).

Neonatal cardiac myocytes were cultured under control conditions or in the presence of 100 μM probucol for 48 hours. We found that APD was significantly prolonged on neonatal cardiac myocytes (Figure 4). To demonstrate the effect of CCh on the prolongation of cardiac APs after probucol treatment, we added CCh to myocytes incubated with 100 μM probucol. Results showed that APD\(_{50}\) was prolonged to 142.6 milliseconds by probucol, 25 μM CCh shortened APD\(_{50}\) to 102.6 milliseconds, and 50 μM CCh shortened APD\(_{90}\) to 89.7 milliseconds on neonatal cardiac myocytes (Figure 4).

**Probucol damaged hERG current but did not affect channel kinetics after long-term treatment**

To identify the damage of hERG current induced by probucol depended on the decrease of protein level but not channel functions, patch-clamp recordings were used to detect the long-term effect of probucol on hERG-channel kinetics. Figure 5A shows the effect of probucol and coinubcation of probucol with CCh on the activation curve after incubation for 48 hours. The \( V_{1/2} \) was \(-7.98±1.48 \text{ mV} \) in the control group and \(-9.42±1.93 \text{ mV} \) in the probucol group. Probucol did not shift the activation curve. Figure 5B illustrates the inhibited effect of probucol and the reversion effect of CCh (25 and 50 μM) on hERG current and its protocol, the results of which were consistent with previous studies.22 Figure 5C shows the effect of probucol on the time-constant curve. As depicted, probucol did not change the time constant or the onset of inactivation and recovery from inactivation. Based on these results, we came to the conclusion that long-term incubation of probucol damaged hERG current, but did not affect channel kinetics.

**Discussion**

A great variety of compounds interfere with hERG function and have the potential to cause LQTS by directly blocking the channel or reducing channel-membrane expression.21 Increasing awareness of drug-induced hERG channel-trafficking deficiency will help to reduce further the potentially fatal adverse cardiac events associated with acquired LQTS.

Probucol is a cholesterol-lowering drug initially developed in the treatment of coronary artery disease.3 By decreasing hERG channel-membrane density, probucol can cause LQTS and torsades de pointes arrhythmia in patients and
Figure 3 Carbachol (CCh) rescued the reduction of 155 kDa hERG expression and $I_{\text{hERG}}$ induced by probucol (Pro).

Notes: (A) CCh impeded the Pro-reduced 155 kDa of the hERG channel. (B) CCh did not reverse As$_2$O$_3$-induced hERG-channel deficiency. (C, D) Voltage-clamp protocol and representative hERG-current traces recorded from hERG-HEK293 cells. Consistent with A, CCh likewise reversed the reduced hERG current caused by Pro. (F) CCh (25 μM, 50 μM) reversed the reduced p-Nedd4-2 expression level. (G) CCh (25 μM, 50 μM) did not reverse the reduced p-Nedd4-2 expression level after exposure HA. $^*P<0.05$ versus control; $^\#P<0.05$ versus Pro; $^{**}P<0.01$ versus control.

Abbreviations: HA, HA-100 (hydrochloride); p-Nedd4-2, phosphorylated Nedd4-2.
sudden cardiac death in experimental animals.\cite{9,24} However, the mechanisms of probucol-reduced hERG-channel expression seem to be quite unspecific. Therefore, a better mechanistic understanding of probucol-induced hERG-channel deficiency may help us to develop therapeutic compounds that are prone to the disruption of hERG-channel processing and trafficking.

As we know, hERG proteins are primarily synthesized in the endoplasmic reticulum and then transported to the Golgi apparatus for glycosylation, and finally transported to the cell membrane for expression. Misfolded or misassembled proteins are retained in the endoplasmic reticulum by its quality-control system, altering forward trafficking of hERG channels to the plasma membrane.\cite{21,25} Our data show that probucol (30, 50, and 100 μM) treatment decreases the mature 155 kDa band of hERG, but does not reduce the intracellular forms of the immature hERG 135 kDa band (Figure 3A), suggesting that probucol may not block hERG forward trafficking. On the other hand, we demonstrated that probucol damaged hERG current, but did not affect channel kinetics after long-term treatment. This seems to be a result of enhanced degradation of plasma membrane hERG protein induced by probucol treatment. It has been demonstrated that Nedd4-2 plays an important role in the degradation pathway of hERG retrograde trafficking.\cite{26} The mature hERG channels are degraded by ubiquitin ligase Nedd4-2 via enhanced channel ubiquitination. In the present study, we confirmed that Nedd4-2 is involved in the process

Figure 4 The effects of carbachol (CCh) on the action potential duration (APD) prolonged by probucol (Pro) on neonatal cardiac myocytes. Notes: (A) Representative AP traces from control (black line), Pro (red line), 100 μM Pro/25 μM CCh group (blue line), and 100 μM Pro/50 μM CCh group (pink line) on neonatal cardiac myocytes. (B) APD\textsubscript{90} was prolonged by Pro and recovered by 25 μM (50 μM) CCh. **P<0.05 versus control; *P<0.05 versus probucol; n=4.
Figure 5 Probenecid (Pro) damaged the hERG current, but did not affect channel kinetics after long-term treatment.

Notes: (A) Voltage-dependent activation curves for the control and following exposure to Pro or Pro and carbachol (CCh) for 48 hours. (B) Voltage-clamp protocol and representative current tracing for steady-state inactivation. (C) The effect of Pro and Pro coincubated with CCh on inactivation curve after treatment for 48 hours. (D) Voltage-clamp protocol and representative current tracing for the onset of inactivation. (E) Voltage-clamp protocol and representative current tracing for recovery from inactivation. (F) The effect of Pro and Pro coincubated with CCh on the time constant for onset of inactivation and recovery from inactivation after treatment for 24 hours. n=10.
of probucol-disrupted hERG surface expression. Probucol treatment significantly reduced p-Nedd4-2 expression levels (Figure 2B). Interestingly, it has been reported that CCh can efficiently increase p-Nedd4-2, but not the total level of Nedd4-2, and eventually increase the expression level of the hERG channel.20 To address further whether CCh could rescue the probucol-disrupted \( I_{\text{hERG}} \) and hERG-channel expression, we coincubated probucol and CCh in hERG-HEK293 cells for 48 hours. Figure 3A illustrates that CCh was able to rescue the decreased 155 kDa hERG-channel expression induced by probucol. Likewise, hERG current was also reversed by long-term coincubation with CCh (Figure 3B–D).

Cardiac toxicities caused by administration of probucol can be controlled by complex signaling pathways or molecular mechanisms. Our present research shows that the SGK1–Nedd4-2 axis may provide a plausible explanation for probucol-induced LQTS and torsade de points, but the results did not illustrate the involvement of any other upstream protein kinases or signaling pathways that could affect the expression and function of serine/threonine kinase SGK1. Whether there are other molecules or protein kinases that contribute to the expression and function of SGK1 needs further investigation.

\( \text{As}_2\text{O}_3 \) has long been of biomedical interest, since it is widely used to treat cancers like acute myeloid leukemia and other conditions.27 However, the cardiotoxicity of LQTS restricts its clinical applications. Previous studies have shown that \( \text{As}_2\text{O}_3 \) can decrease the expression of hERG channels by disturbing its trafficking to the cell membrane. In this research, we wanted to figure out whether \( \text{As}_2\text{O}_3 \) treatment could also affect the SGK1–Nedd4-2 signaling pathway. Our data indicated that in the presence or absence of 3 \( \mu \text{M} \text{As}_2\text{O}_3 \), there was no significant difference between the two groups in the activity of the SGK1–Nedd4-2 signaling pathway and CCh did not rescue the \( \text{As}_2\text{O}_3 \)-induced hERG-channel deficiency (Figure 3B), which means CCh enhances the expression level of hERG channels mainly by the SGK1–Nedd4-2 signaling pathway.

In summary, the present study provides evidence that probucol does not block the hERG channel, but accelerates degradation of the mature hERG channel and inhibit hERG current by decreasing SGK1 expression level and eventually enhancing the ubiquitination of the hERG channel through the E3 ubiquitin ligase Nedd4-2, whereas CCh can rescue probucol-disrupted hERG surface expression by enhancing the phosphorylation level of Nedd4-2 as an alternative to SGK1. These interesting findings may help us in identifying strategies and therapeutic compound classes to prevent or treat QT-interval prolongation associated with administration of probucol.

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Disclosure
The authors report no conflicts of interest in this work.

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