Simulated Blocking Potassium Channels Medication on Variant Mutant SCN5A Sodium Channels

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Abstract

Two gene-mutant sodium channels, 1795insD and I1768V, were considered to be the possible molecular markers in the initiation of long QT (LQT) syndrome. The 1795insD, which decreases the channel availability and enhances the inactivation, and I1768V, which increases the channel availability and enhances the rate of recovery from inactivation, have the ability to induce LQT regardless of their heterogeneous physical characteristics. However, their arrhythmic susceptibility with the use of certain antiarrhythmic medications has not yet been examined closely. In this study, the two mutant SCN5A channels were explored to elucidate the interactions among various potassium channels, IK₁, IKr, and IKs with simulated antiarrhythmic medications by computer modeling. The two mutant SCN5A Markov models, adapted to fit into Rudy's ventricular cell model, performed numerical calculation by using cvode, an ODE solver, with C code in a 4-node PC cluster. In this study, our previously developed S1-S2 protocol was used to investigate the cell excitability in simulated blocking potassium channel medication. The results are as follows: (1) by blocking IK₁ from 10% to 80%, the needed injection charges to initiate an action potential for 1795insD were smaller than I1768V's; (2) by blocking 70% IKr with current stimuli at the rate of 0.5Hz, the action potential of I1768V at the 4th beat began to display premature repolarization; and (3) by blocking 40%-50% IKs with current stimuli at the rates of 0.5 Hz and 1 Hz respectively, the action potential of I1768V at the 3rd beat and the 7th beat began to display premature repolarization. Accordingly, the blockage of IK₁ could demonstrate both positive and negative effects on the two mutant SCN5A channels, as it may enhance or reduce the channel availability while increasing or decreasing the charge threshold. In addition, the blockage of IKs for I1768V might cause serious premature repolarization than the blockage of IKr.

2. Methods

2.1. The model

Two gene-mutant sodium channels, 1795insD and I1768V based on Markov modeling, were incorporated into Rudy phase II cell model. The cell model was used to simulate the membrane excitability and ionic current activity under whole cell voltage clamp mode and current clamp mode. The relationships between membrane voltage and ionic currents are shown in Equation (1):

\[
\frac{\partial V}{\partial t} = \frac{1}{C} (I_{ion} + I_{st})
\]  

Where \(V\) represents the membrane voltage, \(C\) represents the membrane capacitance, \(I_{ion}\) represents the summation of total ionic currents, and \(I_{st}\) represents the...
2.2. I-V relationship

To better understand the physical properties of the two mutant sodium channels, the current-voltage relationship simulation was performed on voltage clamp mode. In the same manner, the activation and inactivation curves of two mutant sodium channels were described by model simulation.

2.3. Cell excitability

To understand the interactions between potassium blockage and cell excitability under mutant sodium channels, the S1-S2 stimulus protocol was applied on the cell model to observe the possible changes of action potentials. Where S1 represents the conditional current stimulus to invoke action potential, S2 represents the test current stimulus.

2.4. The vulnerability of arrhythmias

To test the vulnerability of mutant sodium channel under various potassium blockers, the initiation of early after-depolarization was observed. In Figure 1, the initiation of early after-depolarization can prolong the action potential duration. Therefore, long QT interval can display on E.C.G [3].

The system differential equations of ionic channels described by Rudy’s phase II model and equation (1) were sent to a partial differential equation solver, CVode, running on a 4-node Linux cluster at the lab of system physiology and biology in Chung Hua University.

3. Results

3.1. Electrophysiological prosperities of sodium channel

To understand the electrophysiological characteristics of three sodium channels including wild type, 1795insD and I1768V, the current –membrane voltage (I-V) relationship were measured by their activation and inactivation process. In Figure 2A, the three sodium channels had similar I-V relationship during channel activation.

![Figure 2A. Activation curves of three sodium channels including wild type, 1795insD and I1768V.](image)

In the inactivation curve test, the three sodium channels showed large differences in electrophysiology. The inactivation curve of 1795insD was shifted to the left of wild type’s; however, the inactivation curve of I1768V was shifted to the right. The result in Figure 2B indicated that the 1795insD enhances the inactivation process.

2.5. Numerical computation
Figure 2B. The inactivation curves of three sodium channels: wild type, 1795insD, and I1768V.

As shown in Figure 3, the recovery rates from inactivation of three sodium channels were tested based on Rudy’s protocol[1], the I1768V on the left curve showed faster recovery rate than the wild type and 1795insD.

Figure 3. The curves of recovery rates from inactivation.

3.2. **Cell excitability under potassium blockers**

To better understand the interactions between cell excitability and potassium channel blockers, an S1-S2 current stimulus protocol was used to measure relative ratio of sodium currents by testing stimulus (S2) to the conditional stimulus (S1). In Figure 4A, the cell excitability was decreased with increased IK1 blockage in mutant and non-mutant sodium channels. However, I1798 showed higher excitability than 1795insD and the wild type sodium channel in IK1 blockage. As shown in Figure 4B, I1768V and 1795insD had the same degree of cell excitability despite of increased blockage of IKs from 40% to 70%. In Figure 4C, the cell excitability of three sodium channels remained the same despite of increased IKr blockage. However, the I1768V had highest excitability among the three sodium channels under the same blockage of IKr.

Figure 4 A,B,C. Cell excitability vs. various potassium channel blocker. The S1-S2 interval was 1500 ms.
3.2. Vulnerability vs Potassium Blockers

The initiation of early afterdepolarization of action potentials was used as the vulnerability in this study. In Figure 5A, the stimulus interval was 1500 ms and Iks was 60% blockage, which resulted in early afterdepolarization and degenerated into arrhythmias on the incorporated I1768V cell model. In Figure 5B, the blockage of Iks was decreased to 50% and the stimulus interval was reduced to 1000 ms, and the early afterdepolarization was initiated. In Figure 5C, non-sustained early afterdepolarization was observed on the 80% of blockage of Ikr at 1500 ms stimulus interval.

4. Discussion

Based on our simulations, the needed injection charges to initiate an action potential for 1795insD were smaller than I1768V’s by blocking IK1 from 10% to 80%. Accordingly, the blockage of IK1 could demonstrate both positive and negative effects on the two mutant SCN5A channels, as it may enhance or reduce the channel availability shown in Figure 4 while increasing or decreasing the charge threshold. Based on the simulations shown in Figure 5, it was found that the blockage of Iks for I1768V might cause more serious premature afterdepolarization than the blockage of Ikr.

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References