Computational modeling of a spectrum of arrhythmia-associated CACNA1C mutations that prolong cardiac action potential duration

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Results

I. Functional analysis of mutant channel: G1911R affects Cav1.2 availability and VDI

С

D

صّ ₋₁₅

-20

1.0

0.8

ă 0.6

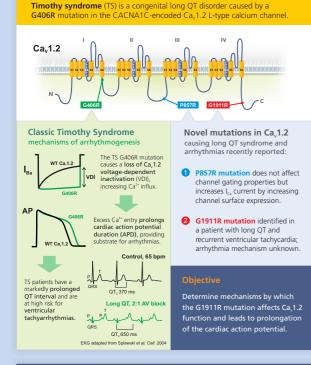
0.2

0.0

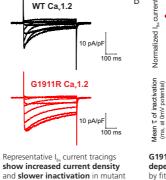
-80 -60

5 0.4

SSI protocol



A



compared to wild-type channels.

B

Wild-type G1911R G1911R decreases voltagedependent inactivation, measured by fitting a single exponential (at 0mV potential) and comparing the tau of inactivation (*p < 0.05).

100 m

250

200

150

100

50

Current-voltage relationship shows an increase in current density in the G1911R mutant compared to WT Ca,1.2.

Wild Type

- G1911R

-1.0

0.8

0.6

0 2

0.0

20

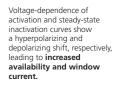
0.4 0

SSA protocol

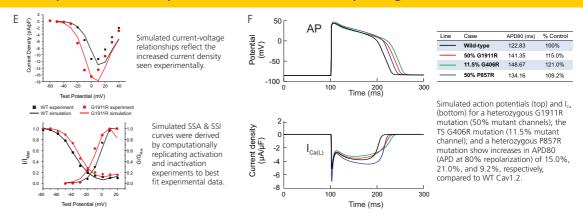
-40 -20 0

Test Potential

50 mV 500 ms



II. Computational action potential simulation: G1911R prolongs APD

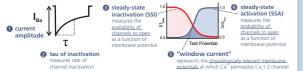


Methods

I. Whole-cell patch clamp: characterize effect of mutation on channel gating.

- WT or G1911R Ca,1.2 channels transfected into HEK293T cells
- Ba²⁺ as charge carrier to eliminate Ca²⁺-dependent inactivation; isolate VDI





II. Computational modeling: predict effect of mutations on APD.

Luo-Rudy myocyte model (Circ. Res. 1994) . modified Ca,1.2 SSI (f_.) and SSA (d_.) gating equations in Luo-Rudy myocyte model to match experimental WT/mutant recordings

simulated action potentials and l_{ca}currents for G1911R, G406R, and P857R mutations

Conclusions

- The combined effects of the G1911R mutation to diminish VDI of $l_{\rm ca}$ and increase window current are predicted to prolong the cardiac APD to a degree nearly as severe as the "classic" Timothy syndrome mutation.
- A mutation that affects current density only, such as the P857R mutation, can also prolong the APD, but to a lesser degree.
- This study broadens our understanding of the mechanisms by which a gain of function of ${\rm l}_{\rm cs}$ can contribute to QT prolongation.

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