



# Corrigendum: Optimization of an In silico Cardiac Cell Model for Proarrhythmia Risk Assessment

# **OPEN ACCESS**

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#### Specialty section:

This article was submitted to Computational Physiology and Medicine, a section of the journal Frontiers in Physiology

Received: 22 November 2017 Accepted: 27 November 2017 Published: 06 December 2017

#### Citation:

Dutta S, Chang KC, Beattie KA, Sheng J, Tran PN, Wu WW, Wu M, Strauss DG, Colatsky T and Li Z (2017) Corrigendum: Optimization of an In silico Cardiac Cell Model for Proarrhythmia Risk Assessment. Front. Physiol. 8:1025. doi: 10.3389/fphys.2017.01025 Sara Dutta<sup>1</sup>, Kelly C. Chang<sup>1</sup>, Kylie A. Beattie<sup>1</sup>, Jiansong Sheng<sup>1</sup>, Phu N. Tran<sup>1</sup>, Wendy W. Wu<sup>1</sup>, Min Wu<sup>1</sup>, David G. Strauss<sup>1</sup>, Thomas Colatsky<sup>2</sup> and Zhihua Li<sup>1\*</sup>

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Keywords: Torsade-de-Pointes (TdP), Comprehensive *in vitro* Proarrhythmia Assay (CiPA), rapid delayed rectifier potassium current (IKr), *in silico* cardiac cell model, drug block, proarrythmia risk, model optimization

## A corrigendum on

## Optimization of an In silico Cardiac Cell Model for Proarrhythmia Risk Assessment

by Dutta, S., Chang, K. C., Beattie, K. A., Sheng, J., Tran, P. N., Wu, W. W., et al. (2017). Front. Physiol. 8:616. doi: 10.3389/fphys.2017.00616

In the original article, there was a mistake in **Figure 6** as published. In the top left panel the qNet gray value should be 0.109 instead of 0.011. The corrected **Figure 6** appears below. The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way.

**Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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**FIGURE 6** | AP traces for mexiletine **(A)** at 1x Cmax (black solid line) and 10x Cmax (gray dashed line) without (left panel) and with 95% IKr reduction (right panel); verapamil **(B)** at 1x Cmax (black solid line) and 3x Cmax (gray dashed line) without (left panel) and with 98% IKr reduction (right panel); and ranolazine (black solid line) and cisapride (dashed gray line) **(C)** at 25x Cmax without (left panel) and with 75% IKr reduction (right panel) for a CL of 2,000 ms. Corresponding APD90 (ms) and qNet ( $\mu$ C/ $\mu$ F) values are reported in black for mexiletine 1x Cmax, verapamil 1x Cmax and ranolazine 25x Cmax and in gray for mexiletine 10x Cmax, verapamil 3x Cmax and cisapride 25x Cmax. Note the IKr reduction (simulated by scaling the IKr maximum conductance) is applied in addition to the drug block effect and is used to assess the system's robustness against EADs (see Results section).