

# QT/TDP Risk Screen: A Web-based Tool for the Early Identification of Drug-Induced Proarrhythmic Risk

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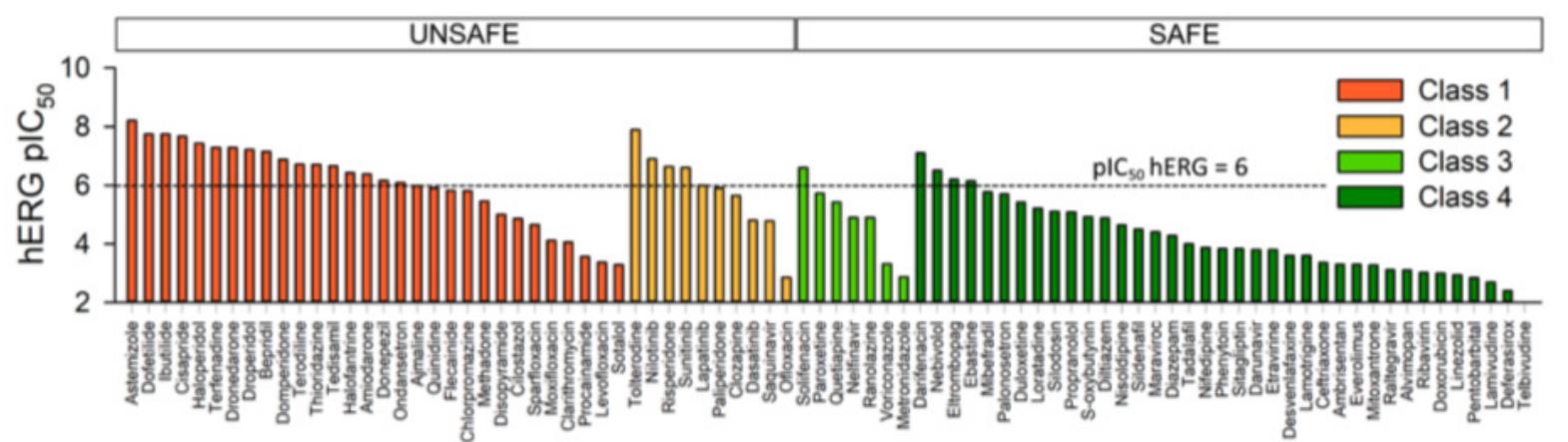
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## PURPOSE

It is well recognized that **early identification of drug-induced proarrhythmic safety risks** is crucial to drug development for **ethical, animal sparing and costs reduction** considerations<sup>1</sup>. The availability, however, of **easily accessible, user-friendly tools for real time assessments of the proarrhythmic potential** of compounds has been lacking. The novel **Tx index**<sup>2</sup>, implemented in the presented web-based tool, was applied to a dataset of **84 drugs compounds**.

## METHODS

QT/TdP Risk Screen is based on **206,766 cellular simulations** of compound-induced effects on Action Potential Duration (APD) in isolated **Endo-, Mid-, and Epi-cardial cells** and on **7,072 tissue simulations** on QT prolongation in a **1D-virtual tissue**<sup>3</sup>. Simulations were performed by blocking the **slow and fast components of the delayed rectifier current (IK<sub>s</sub> and IK<sub>r</sub>, respectively)** and the **L-type calcium current (ICa<sub>L</sub>)** at different concentration levels. Based on these simulations, **four Tx indices** were defined as the ratio of **drug concentration** leading to a **10% prolongation** of the **APD<sub>Endo</sub>, APD<sub>Mid</sub>, APD<sub>Epi</sub>** or **QT interval** over the maximum Effective Free Therapeutic Plasma Concentration (EFTPC), respectively. A dataset of **44 non-torsadogenic** and **40 torsadogenic** drug compounds was used to validate the performance of the tool. **hERG test** (positive response: hERG pIC<sub>50</sub> > 6) was applied to the **84 compounds** to compare performances<sup>2</sup> (Figure 1). QT/TdP Risk Screen was built and integrated on the **cloud-based InSilicoTrials platform**, in compliance with the highest standards of **security** and **privacy**.



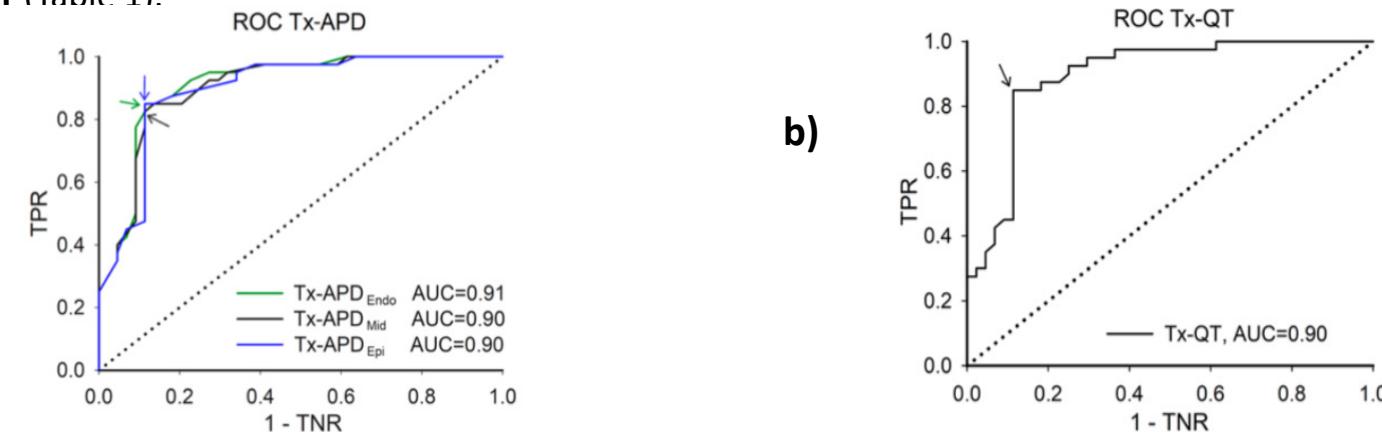
**Figure 1. Torsadogenic risk classification of the 84 compounds using pIC<sub>50</sub> hERG > 6.** CredibleMeds categories: **Class 1**, compounds with risk of TdP; **Class 2**, compounds with possible risk of TdP; **Class 3**, compounds with conditional risk of TdP; and **Class 4**, drugs that should be avoided by patients with congenital Long QT Syndrome.

## References

- [1] Chi KR *Nat. Rev. Drug Disc.* (2013) 12, 565-567; [2] Romero L et al. *J. Chem. Inf. Model.* (2018) 58, 867-878. [3] O'Hara T et al. *PLoS Comput. Biol.* (2011) 7(5): e1002061 doi: 10.1371/journal.pcbi.1002061. [4] Sager, PT et al. *Am. Heart J.* (2014) 167, 292-300. [5] Cantilena L et al. *Clin. Pharmacol. Ther.* (2006) 79, P29.

## RESULTS

Receiver operating characteristic (ROC) curves were constructed on the **four estimated Tx indices** for each compound in the dataset to enable the identification of **torsadogenic potential cut-off values**<sup>2</sup>. These were identified as **8, 8, and 6.4** for **Tx-APD<sub>Endo</sub>, Tx-APD<sub>Mid</sub>, Tx-APD<sub>Epi</sub>** and as **9.2** for **Tx-QT**, respectively. The classification of the **84 compounds** resulted in an accuracy ranging between **87%** and **88%** for the **four Tx indices Tx-APD<sub>Endo</sub>, Tx-APD<sub>Mid</sub>, Tx-APD<sub>Epi</sub> and Tx-QT** (Figure 2). The classification of the **84 compounds** resulted in an accuracy ranging between **87%** and **88%** for the **four Tx indices Tx-APD<sub>Endo</sub>, Tx-APD<sub>Mid</sub>, Tx-APD<sub>Epi</sub> and Tx-QT** (Table 1).



**Figure 2. In silico isolated cellular Tx-APD and Tx-QT assays.** a) ROC curves for the Tx-APD and optimal cut-off points where sensitivity and specificity are maximal (arrows) for the Tx-APD<sub>Endo</sub> (green), Tx-APD<sub>Mid</sub> (black), and Tx-APD<sub>Epi</sub> (blue) (8, 8, and 6.4, respectively). b) ROC curve for the Tx-QT and optimal cut-off point where sensitivity and specificity are maximal for the Tx-QT (arrow, 9.2). The area under the curve (AUC) values are also indicated.

Endo			Mid			Epi			QT		
Tx	TdP+	TdP-									
+	34	5	+	34	6	+	34	5	+	34	5
-	6	39	-	6	38	-	6	39	-	6	39
A	TPR	TNR									
0.87	0.85	0.89	0.86	0.85	0.86	0.87	0.85	0.89	0.87	0.85	0.89

**Table 1. Confusion matrices for the Tx-APDs and Tx-QT.** The values of the true positive rate (TPR), true negative rate (TNR), and accuracy (A) are also indicated.

## CONCLUSIONS

hERG block alone exhibits poor predictive performance<sup>4</sup>. When applying the **in vitro hERG test** to the **84 compounds**, it exhibited a **TPR** of **55%**, a **TNR** of **89%**, and an **A** of **73%**, in close agreement with previous studies<sup>5</sup>. In comparison, the **in silico Tx tests** described in this study yield **TPRs** of **85%**, **TNRs** of **86%-89%** and **As** of **86%-87%**. This method does not include drug effects on Na<sup>+</sup> channels, which is related to the misclassification of 3 compounds (quetiapine, ranolazine, and lamotrigine - significant Na<sup>+</sup> channels blockers at EFTPC). Future work will include this channel. The presented web-based tool is a **highly innovative method** for an **accurate torsadogenic risk assessment**. Each assessment required only a **few seconds** of computational time.