

1 Appendix: Comparison of in silico predictions of action
2 potential duration in response to inhibition of I_{Kr} and
3 I_{CaL} with new human ex vivo recordings

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6 **List of Figures**

7	A	Protocols for recording of the peak $I_{CaV1.2}$ current.	4
8	B	Protocols for recording of the peak I_{hERG} current.	5
9	C	Error in the cubic surface compared with the experimental ΔAPD_{90} data	8
10	D	Impact of reporting drug concentrations as measured in the bath solution or as nominal	16
12	E	Comparison of the abilities of human ventricular AP models to reproduce the APD_{90} response to I_{Kr} and I_{CaL} inhibition observed <i>ex vivo</i> , using only nominal drug concentrations as inputs for simulations	17
15	F	Correlation between baseline APD_{90} and Verapamil-induced perturbations	18
16	G	Experimental $\% \Delta APD_{90}$ measured ex-vivo under various drug conditions in human ventricular trabeculae	18
18	H	2-D maps of simulated $\% \Delta APD_{90}$ after I_{CaL} and I_{Kr} inhibition	19
19	I	Observed versus prediction plot for the tested models, using the CiPA and Pharm datasets	20

21	J	Q_{net} computed with the ORd-CiPA model, for various combinations of I_{Kr}	
22		and/or I_{CaL} inhibition	21
23	K	Stability of AP markers observed in a representative trabecula	22

24 List of Tables

25	A	Data summary of IC_{50} values for hERG and $Ca_v1.2$ block potency of the	
26		Clozapine.	6
27	B	Data summary of IC_{50} values for hERG and $Ca_v1.2$ block potency of	
28		Dofetilide.	6
29	C	Data summary of IC_{50} values for hERG and $Ca_v1.2$ block potency of	
30		Nifedipine.	7
31	D	Data summary of IC_{50} values for hERG and $Ca_v1.2$ block potency of	
32		Quinidine.	7
33	E	Data summary of IC_{50} values for hERG and $Ca_v1.2$ block potency of	
34		Verapamil.	8

35 1 In vitro measurements of I_{Kr} and I_{CaL} inhibition

36 For a consistent comparison of the APD_{90} response to drug perturbation predicted by
 37 the AP models included in the present benchmark, in some of which the dynamic hERG
 38 binding model cannot be implemented, the inhibition of ionic currents was modelled with
 39 the Hill equation:

$$I(D) = \frac{1}{1 + \left(\frac{D}{IC_{50}}\right)^h} \times I(0), \quad (1)$$

40 with I the current with drug inhibition, D the drug concentration, h the Hill coefficient,
 41 IC_{50} the half inhibitory drug concentration, and $I(0)$ the ionic current measured at baseline
 42 without any drug exposure.

43 The AP models considered in this study do not distinguish isoforms of the ion channels,
 44 thus the drug inhibition of hERG and $Ca_v1.2$ channels was modelled as equal to the drug
 45 inhibition on I_{Kr} and I_{CaL} , respectively. hERG and $Ca_v1.2$ channels are the main ion

46 channels responsible for I_{Kr} and I_{CaL} , respectively (Agrawal *et al.*, 2022; Sanguinetti *et al.*,
47 1995; Li *et al.*, 1996).

48 The experiments were carried out internally in Roche by Evgenia Gissing and Fabian
49 Häusermann, from Dr. Liudmila Polonchuk’s lab (Roche Pharmaceutical Research and
50 Early Development, Pharmaceutical Sciences).

51 1.1 Cell culture

52 The CHO crelox hERG cell line was generated and validated at Roche (Guthrie *et al.*,
53 2005). The CHO-hCa_v1.2/ β 2/ α 2 δ cell line was purchased from ChanTest (USA, Catalog
54 #CT6004). Vials with cryopreserved cells were thawed at 37°C, washed with the pre-
55 warmed IMDM cell culture medium (Gibco Life Technologies, USA) and re-suspended in
56 the extracellular solution.

57 For the hERG assay the extracellular solution contained (in mM): NaCl 80; KCl 4;
58 CaCl₂ 1; MgCl₂ 1; NMDG 40; HEPES 10; sorbitol 40; glucose 5; pH 7.2–7.4 with NaOH,
59 osmolarity 290–330 mOsm and the internal solution contained (in mM): KCl, 10; KF, 100;
60 NaCl, 10; HEPES, 10; EGTA, 20; pH = 7.0–7.4 with KOH, osmolarity 260–300 mOsm.
61 For the L-type Ca_v1.2 assay the extracellular solution contained (in mM): NaCl 80; KCl 4;
62 CaCl₂ 1.8; MgCl₂ 1; NMDG 40; HEPES 10; sorbitol 40; glucose 5; pH 7.2–7.4 with NaOH,
63 osmolarity 290–330 mOsm and the internal solution contained (in mM): KCl, 10; KF, 100;
64 NaCl, 5; HEPES, 10; EGTA, 10; Na-ATP, 4; Na-GTP, 0.1; pH = 7.0–7.4 with KOH,
65 osmolarity 260–300 mOsm.

66 1.2 Electrophysiology recordings

67 The recording of the currents was performed using automated patch clamp system
68 SynchroPatch 384 (Nanon Technologies GmbH, Germany) at 35–37°C following the
69 experimental procedure described below. On the day of the experiment, an aliquot of
70 the cell suspension in a 2:1 mixture of the HBSS and external solution was placed in the
71 Cellhotel. The cells were subsequently added into the 384-well sealchip where the currents
72 were recorded in single cells with the patch-voltage-clamp technique in the whole-cell

73 configuration at 35–37°C using the built-in 384 channel amplifier and associated software
 74 (PatchControl 384). Currents were low-pass filtered using the analog 3 kHz Bessel filter
 75 and the digital 3 kHz Lanczos filter and were digitized at 5 kHz. Series resistance was
 76 typically 2–9 MΩ and is compensated by 80%. The reported current amplitudes represent
 77 the maximal amplitude of a peak current.

78 1.3 Voltage-clamp protocols

79 The voltage-step protocols used to measure the $I_{CaV1.2}$ inhibition by drugs are shown in
 80 Figure A. In the Roche in-house protocol (Pharm dataset), the cells were held at a resting
 81 potential of -90 mV, where the baseline current amplitude was recorded. The $I_{CaV1.2}$
 82 channels were then activated with 120 ms-wide steps at 0 mV with a frequency of 0.1 Hz,
 83 where the peak current was recorded. Once the recorded peak and baseline currents were
 84 stabilised, the amplitude and kinetics of the $I_{CaV1.2}$ were recorded for 3–5 minutes without
 85 drug. Then, after each drug addition, the activity of the cells was recorded during 3
 86 minutes.

87 In the CiPA protocol, cells were held at a resting potential of -80 mV, where the
 88 baseline current was recorded. Then a stimulus 40 ms step of voltage at 0 mV was applied
 89 to record the peak current, with a 0.1 Hz frequency. Then the voltage was held at $+30$ mV
 90 for 200 ms, followed by a ramp down of voltage to return to -80 mV in 99 ms.

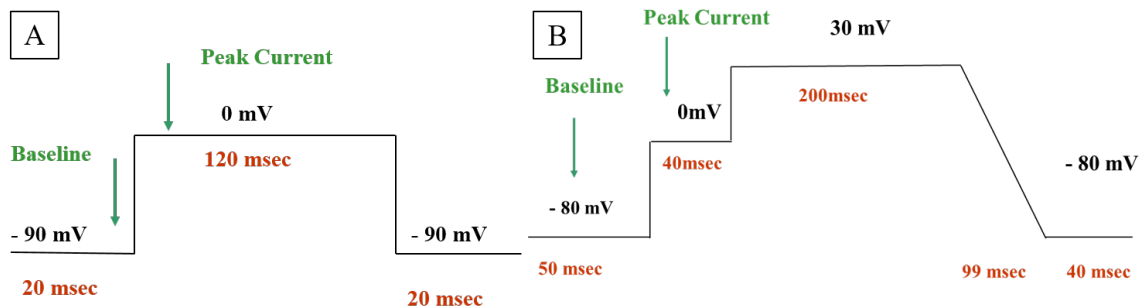


Figure A: Protocols for recording of the peak $I_{CaV1.2}$ current. **A:** Roche in-house protocol (‘Pharm’ dataset). **B:** CiPA protocol (‘CiPA’ dataset) (Li *et al.*, 2019).

91 The voltage-step protocols used to measure the hERG inhibition by drugs are shown
 92 in Figure B. In the Roche in-house protocol, the resting voltage was set to -80 mV. Then

93 the voltage was clamped to -40 mV for 100 ms, where the baseline current was recorded.
 94 The voltage was then brought to $+20$ mV for 500 ms and finally to -40 mV for 500 ms,
 95 where the peak current was recorded. Afterwards, the voltage was set back to the resting
 96 potential of -80 mV. The stimulation pattern is repeated with a frequency of 0.1 Hz.

97 In the CiPA protocol, the cells were held to the resting potential of -80 mV. A
 98 $+40$ mV voltage was then applied to them for 500 ms, followed by a -1.25 mV/ms ramp
 99 that brought the voltage down to the resting potential in 96 ms. The peak current was
 100 recorded during this ramp down. The pattern was repeated with a frequency of 0.1 Hz.

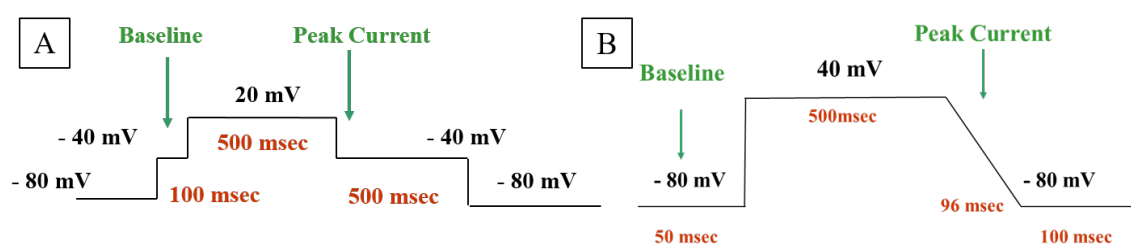


Figure B: Protocols for recording of the peak I_{hERG} current. **A:** Roche in-house ('Pharm' dataset). **B:** CiPA protocol ('CiPA' dataset).

101 1.4 Literature and new IC₅₀ data

102 The Tables A to E below summarise the IC₅₀ values that could be found in the literature
 103 for hERG and Ca_v1.2 block potency of Clozapine, Dofetilide, Nifedipine, Quinidine and
 104 Verapamil. Overall, there was good agreement between literature values and the values
 105 we present in this study. Nifedipine IC₅₀ (measured with both protocols) was however
 106 higher than literature values for Ca_v1.2. The 1.5 inter-quartile range statistical test was
 107 performed to identify outliers, and no outlier was found.

Drug	hERG block IC50 (μM)/h	Ca _v 1.2 block IC50 (μM)/h	Source
Clozapine	1.978 / 0.94	4.378 / 0.93	CiPA prot.
	2.123 / 1.05	1.676 / 0.75	Pharm prot.
	2.300 / 0.97	3.600 / 1	(Kramer <i>et al.</i> , 2013)
		5.490 / 0.94	(Li <i>et al.</i> , 2019)
	2.310 / 1	3.555 / 1	One Million Solutions in Health
	1.639 / 1	3.600 / 1	(Llopis-Lorente <i>et al.</i> , 2020)

Table A: Data summary of IC₅₀ values for hERG and Ca_v1.2 block potency of the Clozapine.

Drug	hERG block IC50 (μM)/h	Ca _v 1.2 block IC50 (μM)/h	Source
Dofetilide	0.033 / 1.17		CiPA prot.
	0.029 / 1.10	332 / 1	Pharm prot.
	0.030 / 1.2	26.7 / 1	(Kramer <i>et al.</i> , 2013)
	0.0061 / 1.08	44.5 / 3.6	CiPA GitHub
	0.005 / 1	60 / 1	(Mirams <i>et al.</i> , 2011)
	0.002 / 1		(Redfern <i>et al.</i> , 2003)
	0.002 / 1		(Crumb Jr <i>et al.</i> , 2016)
	0.037 / 2.1	182 / 1	(Okada <i>et al.</i> , 2015)
	0.025 / 1	26.7 / 1	(Llopis-Lorente <i>et al.</i> , 2020)

Table B: Data summary of IC₅₀ values for hERG and Ca_v1.2 block potency of Dofetilide.

2 Fitting of the cubic surface through the experimental ΔAPD_{90} data

Cubic surfaces were fitted through the experimental data for drug-induced ΔAPD_{90} . The cubic surface was computed following the equation:

$$z = \text{surf}(x, y, \boldsymbol{\theta}) = \theta_1 x^2 + \theta_2 y^2 + \theta_3 xy + \theta_4 x + \theta_5 y + \theta_6 x^3 + \theta_7 y^3 + \theta_8 x^2 y + \theta_9 x y^2, \quad (2)$$

with z the approximated ΔAPD_{90} , x and y the percentage of inhibition of I_{Kr} and I_{CaL}, respectively, and $\boldsymbol{\theta}$ the parameters describing the cubic surface. The location of each data point for each tested trabecula (x and y) was computed using the IC₅₀ data (Table 2 in the main text) and the measured drug concentration in the bath solution if available. The

Drug	hERG block IC50 (μM)/h	Ca _v 1.2 block IC50 (μM)/h	Source
Nifedipine	inactive	0.144 / 0.72	CiPA prot.
	inactive	0.105 / 0.85	Pharm prot.
		0.0114 / 0.67	(Li <i>et al.</i> , 2019)
	44 / 0.8	0.012 / 1.02	(Kramer <i>et al.</i> , 2013)
	275 / 1	0.060 / 1	(Mirams <i>et al.</i> , 2011)
		0.056 / 1.18	(Elkins <i>et al.</i> , 2013)
	112.25 / 1	0.060 / 1	(Llopis-Lorente <i>et al.</i> , 2020)
	0.016 / 1	Kuryshv <i>et al.</i> (2014)	

Table C: Data summary of IC₅₀ values for hERG and Ca_v1.2 block potency of Nifedipine.

Drug	hERG block IC50 (μM)/h	Ca _v 1.2 block IC50 (μM)/h	Source
Quinidine	0.820 / 1.43	6.680 / 1	CiPA prot.
	0.966 / 1.01	20.849 / 0.63	Pharm prot.
	2.371 / 1.71		(Elkins <i>et al.</i> , 2013)
	0.3 / 1	15.6 / 1	(Mirams <i>et al.</i> , 2011)
	0.72 / 1.06	6.4 / 0.68	(Kramer <i>et al.</i> , 2013)
	0.3 / 1		(Crumb Jr <i>et al.</i> , 2016)
	0.658 / 1	8.1 / 0.8	(Okada <i>et al.</i> , 2015)
	0.986 / 0.84	51.59 / 0.59	CiPA GitHub
	0.890 / 1	8.866 / 0.912	(Llopis-Lorente <i>et al.</i> , 2020)

Table D: Data summary of IC₅₀ values for hERG and Ca_v1.2 block potency of Quinidine.

116 nominal drug concentration was used otherwise.

117 The cubic surface was fitted through the experimental ΔAPD_{90} data points, by
118 minimising the cost function:

$$\mathcal{S}(\boldsymbol{\theta}) = \sum_{\text{trab}} \sum_k (\text{surf}(x_k, y_k, \boldsymbol{\theta}) - \Delta\text{APD}_{90, \text{exp}, k, \text{trab}})^2, \quad (3)$$

119 with $\Delta\text{APD}_{90, \text{exp}, k, \text{trab}}$ the experimental ΔAPD_{90} for the drug perturbation k averaged
120 over 30 consecutive APs in the trabecula trab . For each k , the associated inhibition of
121 I_{Kr} and I_{CaL} was computed using Eq. 2 and the IC50 data reported in Table 2 in the
122 main text. The minimisation of the cost function was performed using the `scipy` Python
123 package. Based on experimental observations, the following constraints were put on the
124 cubic surface during its fitting:

Drug	hERG block IC50 (μM)/h	Ca _v 1.2 block IC50 (μM)/h	Source
Verapamil	0.570 / 1.67	0.310 / 1	CiPA prot.
	0.273 / 0.98	1.381 / 0.72	Pharm prot.
	0.296 / 0.94	0.202 / 1.10	CiPA GitHub
	0.250 / 0.89	0.200 / 0.8	(Kramer <i>et al.</i> , 2013)
	0.677 / 1.43	2.685 / 0.61	(Elkins <i>et al.</i> , 2013)
	0.143 / 1	0.100 / 1	(Mirams <i>et al.</i> , 2011)
	0.7 / 1	0.1 / 1	(Crumb Jr <i>et al.</i> , 2016)
	0.212 / 1	0.347 / 1.08	(Okada <i>et al.</i> , 2015)
	0.499 / 1.1	0.201 / 1.1	(Llopis-Lorente <i>et al.</i> , 2020)

Table E: Data summary of IC₅₀ values for hERG and Ca_v1.2 block potency of Verapamil.

- 125 • $\Delta\text{APD}_{90} = 0$ ms at baseline, i.e., $\theta_6 = 0$;
- 126 • $\Delta\text{APD}_{90} \leq -50$ ms for 100% I_{CaL} block ;
- 127 • $\Delta\text{APD}_{90} \geq +320$ ms for 100% I_{Kr} block ;
- 128 • $\frac{dz}{dx} > 0$, translating that an increase in I_{Kr} inhibition prolongs the APD₉₀;
- 129 • $\frac{dz}{dy} < 0$, translating that an increase in I_{CaL} inhibition shortens the APD₉₀.

130 When the constraints were not satisfied, the cost (Eq. 3) was multiplied by 100.

131 The goodness of fit of the cubic surface to the experimental points is plotted in
 132 Figure C.

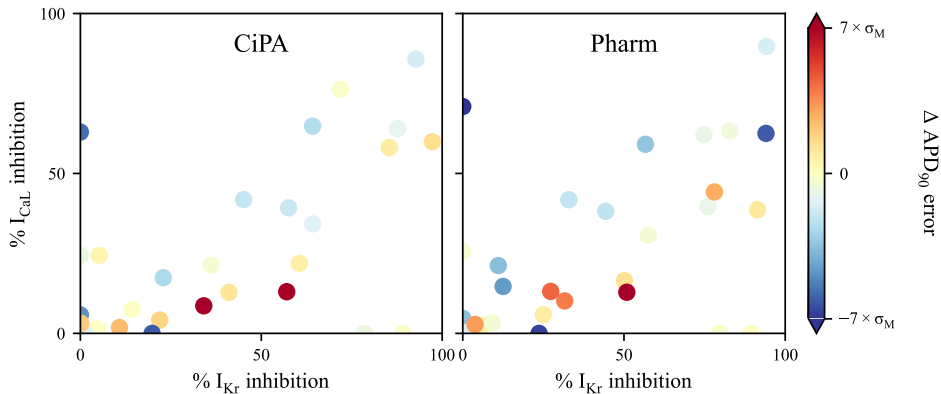


Figure C: Error (Eq. 2) in the cubic surface compared with the experimental ΔAPD_{90} data, obtained using the CiPA (**left**) and the Pharm dataset (**right**).

3 Comparison of 2D maps using nominal concentrations and using measured drug concentrations in the bath solution (when available)

As reported in Table 1, drug concentrations in the bath solution were measured for some compounds, while they were not measured for other compounds. Note that for Clozapine, the drug concentration was measured in 7 trabeculae exposed to 0.3–3 μM but not in the 4 trabeculae exposed to 0.3–30 μM . In this section we compare whether fitting the cubic surface to nominal or measured drug concentrations makes any visual change to the resulting surface, and if simulating the drug-induced ΔAPD_{90} with nominal concentrations impacts our results.

The fitting of the cubic surface was repeated three times, using differently the data for drug concentration to locate the experimental data points on the 2-D map (Eq. 2):

- using the measured drug concentration in the bath solution for each trabecula under each tested drug condition, when the drug concentration was measured. Otherwise, the nominal concentration was used. In this case, ΔAPD_{90} is reported separately for each trabecula for each drug condition ;
- averaging the measured drug concentration for all trabeculae tested with the same nominal drug concentration, when the drug concentration was measured. Otherwise, the nominal drug concentration was used. The drug-induced ΔAPD_{90} effect was averaged over the trabeculae tested with the same nominal concentration ;
- using only the nominal drug concentrations. The drug-induced ΔAPD_{90} effect was averaged over the trabeculae tested with the same nominal concentration.

The experimental data points were placed on the 2-D map following the three methods, and the corresponding cubic surfaces were fitted through these data points. The fitting of the cubic surface was repeated with the CiPA and Pharm protocols, and the results are visualised in Figure D.

159 The exact location of points was changed due to the measured concentrations in
160 the bath solution not matching exactly with the nominal concentrations. Yet, the cubic
161 surfaces fitted to the experimental points with the three methods described above were very
162 similar. Therefore, the qualitative comparison of model predictions with the experimental
163 data yields the same results when the nominal drug concentrations are used to compute
164 the drug-induced inhibition of I_{Kr} and I_{CaL} .

165 To further support that the interpretation of our results were not sensitive to the
166 discrepancy between measured drug concentrations in the bath solution and nominal drug
167 concentrations, the scores were recomputed with all nominal concentrations, similarly
168 to Figure 7. The results, plotted in Figure E, are similar to the results obtained with
169 measured drug concentrations in the bath solution (when available), and the interpretation
170 of our results therefore did not depend on the discrepancy between the measured and
171 nominal drug concentrations.

172 **4 Comparison of drug-induced ΔAPD_{90} with relative** 173 **APD_{90} change from baseline as a percentage**

174 To investigate whether drug-induced changes in APD_{90} should be reported in absolute or
175 relative values, the correlation between baseline APD_{90} and response to drug perturbation
176 was observed. For 15 trabeculae exposed to 1 μM Verapamil, the absolute change in
177 APD_{90} (ΔAPD_{90}) was computed. From there, the relative change in APD_{90} was computed
178 with the following equation:

$$\% \Delta APD_{90} = \Delta APD_{90} / APD_{90, \text{baseline}}, \quad (4)$$

179 $APD_{90, \text{baseline}}$ referring to the baseline APD_{90} .

180 The scatter plot of ΔAPD_{90} and $\% \Delta APD_{90}$ against baseline APD_{90} is plotted in
181 Figure F. The ΔAPD_{90} and $\% \Delta APD_{90}$ showed not correlated with baseline APD_{90} .
182 Consistent results were obtained for all drugs and all drug concentrations.

183 Therefore, normalising ΔAPD_{90} would not improve the understanding of drug effect.
184 Furthermore, in the clinic, changes in QT are measured in absolute, average prolongation
185 of the QT interval by more than +5 ms being the limit of tolerance (ICH, 2006). At the
186 cellular level, ΔAPD_{90} is more directly linked with the safety marker than $\%\Delta\text{APD}_{90}$. As
187 a conclusion, the ΔAPD_{90} was used for the rest of this study.

188 To investigate further the importance of using ΔAPD_{90} over $\%\Delta\text{APD}_{90}$ in this study
189 the results of the main text were repeated using the relative APD_{90} change expressed as a
190 percentage of the baseline APD_{90} ($\%\Delta\text{APD}_{90}$), instead of ΔAPD_{90} . The relative APD_{90}
191 change was computed in each trabecula as:

$$\Delta\%\text{APD}_{90} = 100 \times \Delta\text{APD}_{90} / \text{APD}_{90, \text{baseline}}. \quad (5)$$

192 The experimental $\%\Delta\text{APD}_{90}$ is plotted in Figure G. The cubic surface was similar to
193 the cubic surface observed in Figure 4, granted the scalings differed between the two
194 figures.

195 Predictions of $\%\Delta\text{APD}_{90}$ were also computed for the 11 AP models, and plotted in
196 Figure H. Very similar trends were observed in 2-D maps for ΔAPD_{90} and $\%\Delta\text{APD}_{90}$.
197 Therefore, the choice of using ΔAPD_{90} over $\%\Delta\text{APD}_{90}$ did not impact the interpretation
198 of the results presented in this study.

199 5 Model predictions vs observations

200 Model predictions are plotted against experimental observations in Figure I. It enables
201 an extra visualisation of the model performances, similar to Figure 6 in the main text.
202 The same observations can be made as described in the main text. The ORd-like models
203 capture the strong APD_{90} -prolonging perturbations, but they largely overpredict the
204 APD_{90} response to compounds inhibiting both I_{Kr} and I_{CaL} . The TP-like models are less
205 prone to overprediction, but they are unable to predict strong APD_{90} prolongation.

206 **6 The TdP risk metric Q_{net} as a function of I_{Kr} and** 207 **I_{CaL} inhibition**

208 The CiPA initiative was established with the objective of developing an *in silico* model
209 classifying drugs into three TdP risk categories (low, intermediate, high risk), providing a
210 more specific safety assessment than the I_{Kr} -centric guideline (Sager *et al.*, 2014). One
211 popular candidate, Q_{net} , relies on the net charge flux over the repolarisation phase of one
212 AP computed with the ORd-CiPA model (Li *et al.*, 2019). Q_{net} is defined as the integral
213 of the net currents that are active in the repolarisation phase over one AP at 0.5 Hz pacing
214 after 1000 pre-paces defined as:

$$Q_{\text{net}} = \int_{0 \text{ ms}}^{2000 \text{ ms}} (I_{\text{Kr}} + I_{\text{CaL}} + I_{\text{Ks}} + I_{\text{NaL}} + I_{\text{K1}} + I_{\text{to}}) dt. \quad (6)$$

215 In this section, we compare Q_{net} as a function of I_{Kr} and I_{CaL} inhibition with experimental
216 ΔAPD_{90} measurements.

217 As with the AP models, Q_{net} was computed with the ORd-CiPA model for $101 \times 101 =$
218 10,201 combinations of I_{Kr} and I_{CaL} inhibition. The reduction of I_{Kr} was modelled by
219 applying a multiplying factor to the maximal conductance of I_{Kr} , as in the main text,
220 although in the original methods of Li *et al.*, the I_{Kr} inhibition by drugs is modelled
221 with the dynamic hERG binding model (Li *et al.*, 2017). Note that Li *et al.* classified
222 compounds into the TdP risk categories based on the average Q_{net} computed at 1–4
223 times their maximal effective free therapeutic concentration. Nevertheless, a qualitative
224 interpretation of the 2-D map remains possible.

225 Pixels of the 2-D map were colored based on the TdP risk category corresponding to
226 Q_{net} obtained with the ORd-CiPA model after inhibition of the ionic currents. As in (Li
227 *et al.*, 2019), Q_{net} values greater than $0.0671 \mu\text{C} \cdot \mu\text{F}^{-1}$ were classified as low risk (green),
228 Q_{net} between $0.0581 \mu\text{C} \cdot \mu\text{F}^{-1}$ and $0.0671 \mu\text{C} \cdot \mu\text{F}^{-1}$ as intermediate risk (blue), and Q_{net}
229 lower than $0.0581 \mu\text{C} \cdot \mu\text{F}^{-1}$ as high risk.

230 The resulting 2-D map is shown in Figure J.

231 Interestingly, the decrease in Q_{net} (increase in TdP risk) induced by I_{Kr} inhibition is

232 mitigated by I_{CaL} inhibition, with a higher sensitivity to I_{CaL} inhibition than ΔAPD_{90}
233 predicted by the ORd-CiPA model. For example, 50% I_{Kr} inhibition and 75% I_{CaL}
234 inhibition yields a Q_{net} value classified into the low TdP risk category, while the predicted
235 ΔAPD_{90} is +73 ms.

236 The shape of the 2-D map of Q_{net} is similar to that of the 2-D map of ΔAPD_{90}
237 predicted by the TP-M model (Figure 5 in the main text). Furthermore, qualitatively
238 similar mitigation of I_{Kr} inhibition by I_{CaL} inhibition was observed between Q_{net} predictions
239 and ΔAPD_{90} observed experimentally (Figure 4 in the main text).

240 **7 Stability of AP markers after 25 minutes of steady** 241 **pacing**

242 In *ex vivo* experiments, a steady 1 Hz pacing was applied to trabeculae (Figure 1). The
243 stability of the AP was qualitatively assessed by the experimenter, based on APD_{90} , AP
244 amplitude, and RMP. If stability was not achieved after 25 min, the steady pacing was
245 prolonged. The evolution of these AP markers is plotted in Figure K for a representative
246 trabecula, for the last 2 min of steady pacing after stabilisation.

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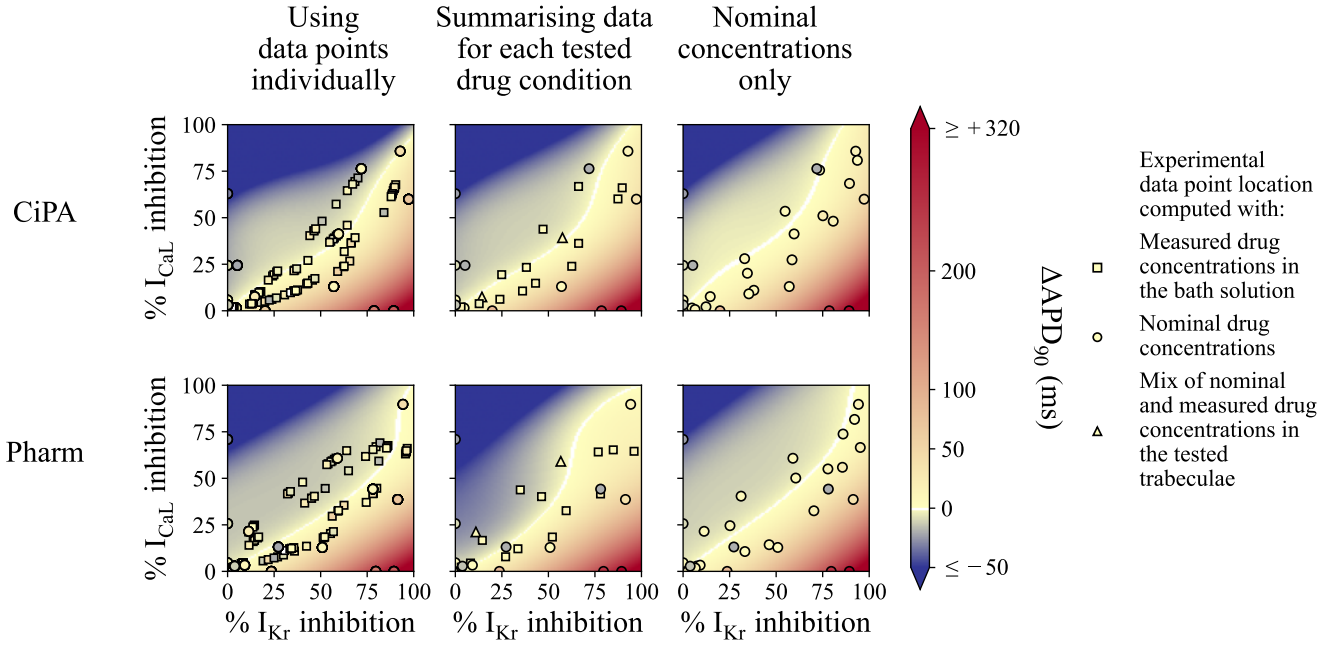


Figure D: Impact of reporting drug concentrations as measured in the bath solution or as nominal on the cubic surface and on the interpretation of the results in this study. Experimental ΔAPD_{90} measured ex-vivo under various drug conditions in human ventricular trabeculae, as a function of I_{K_r} and I_{CaL} inhibition and cubic surface approximating the experimental data points in the background. Each data point was placed with the current inhibition computed with Eq. 1 in the main text from drug concentrations and the drug IC_{50} for I_{K_r} and I_{CaL} (Table 2 in the main text). **Left:** ΔAPD_{90} is reported with one point per tested drug condition per trabecula. The cubic surface was fitted to all the data points for all the trabeculae. **Middle:** A single point is plotted per tested drug condition. The data for drug concentration and drug-induced ΔAPD_{90} was averaged the trabeculae tested with the same nominal concentration. **Right:** A single point is plotted per tested drug condition, using the nominal drug concentration to compute the drug-induced current inhibition and subsequent location on the map. ΔAPD_{90} was averaged similarly to the middle panel. When measured drug concentrations were available only for some trabeculae tested with the same nominal drug concentration, the data point was plotted as a **triangle**.

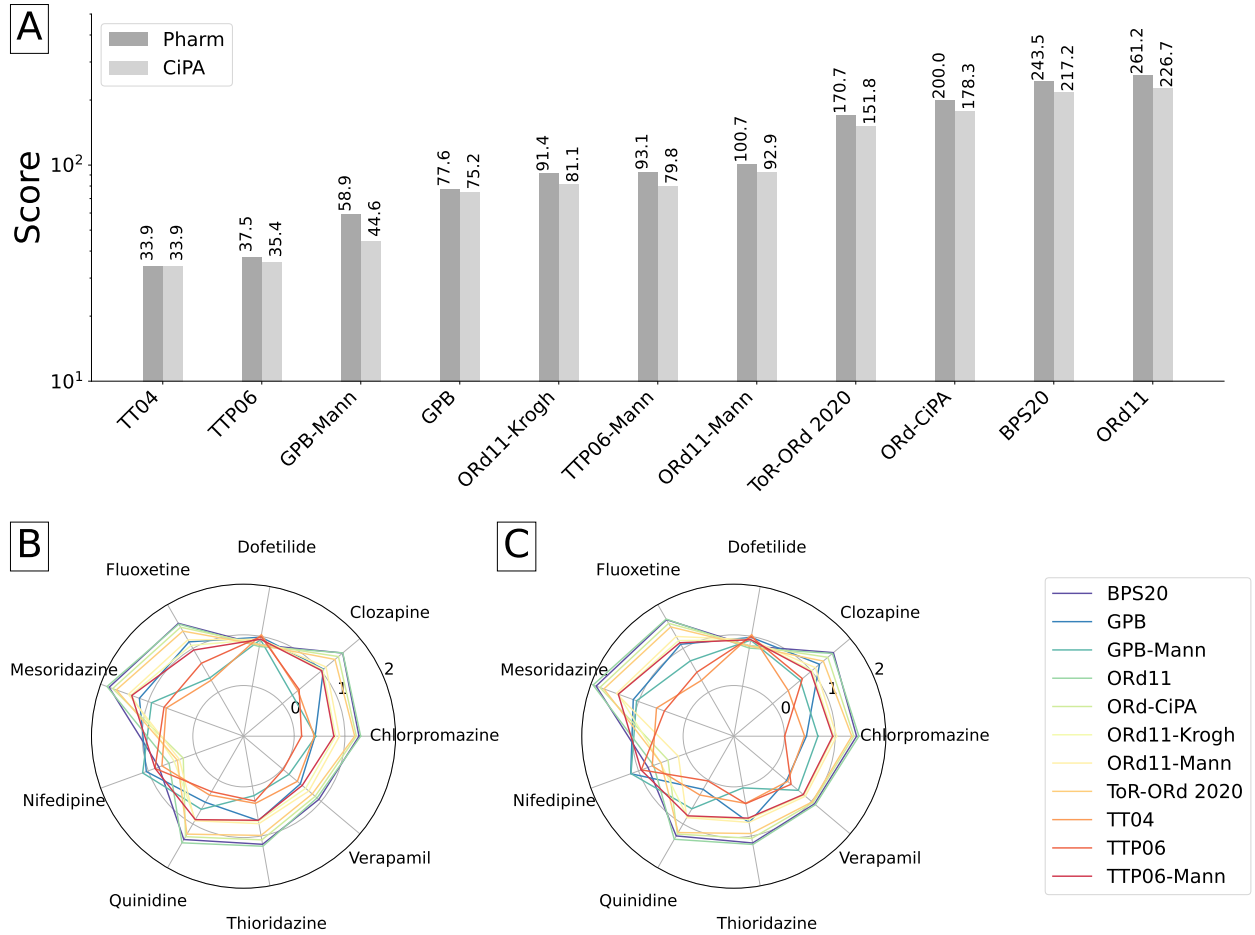


Figure E: Comparison of the abilities of human ventricular AP models to reproduce the APD_{90} response to I_{Kr} and I_{CaL} inhibition observed *ex vivo*, using only nominal drug concentrations as inputs for simulations. The error metric was computed from Eq. 2 in the main text. **A**: The error measure was summed over all the drugs used in this study, when using the Pharm and CiPA protocols to compute the reduction of ionic currents by drugs. For each model, two bar plots were plotted, to compare the predictive power of models with the Pharm (left bar) and the CiPA (right bar) datasets. **B and C**: Detail of the error measures associated to each of the drugs with the CiPA and Pharm datasets, respectively, for each model. The \log_{10} of the error measure is plotted along the radial-axis.

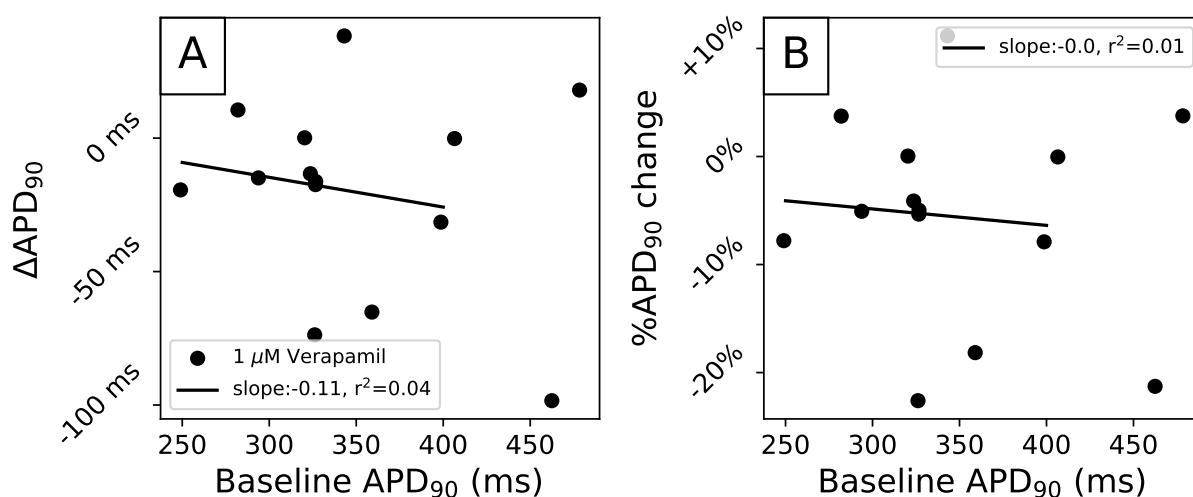


Figure F: Correlation between baseline APD_{90} and Verapamil-induced perturbations, measured as absolute change in APD_{90} (A) and as relative change in APD_{90} (B). No correlation was found.

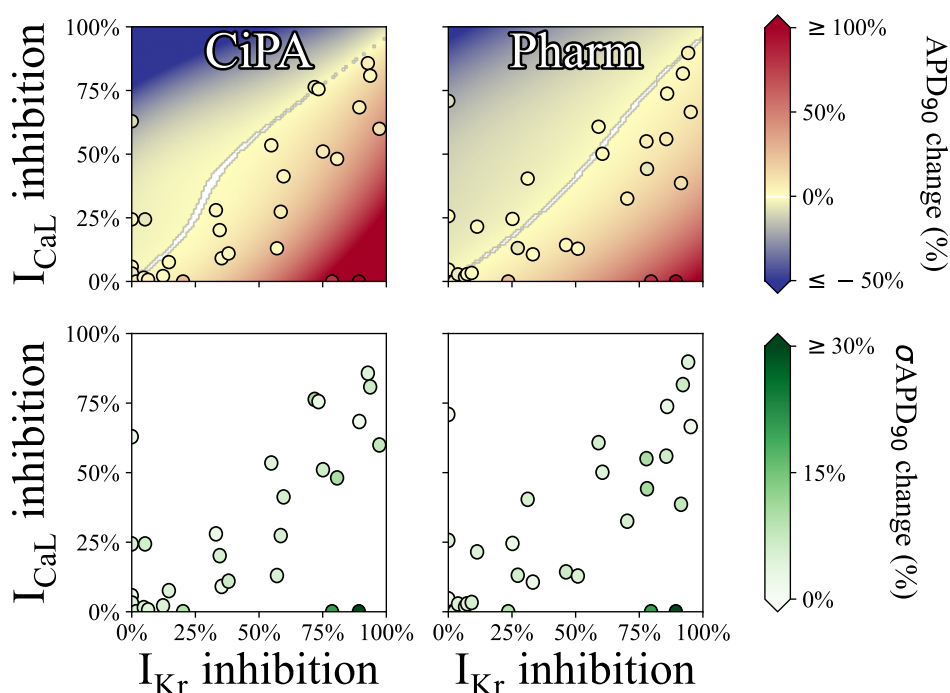


Figure G: Experimental $\% \Delta APD_{90}$ measured ex-vivo under various drug conditions in human ventricular trabeculae, as a function of I_{K_r} and I_{CaL} inhibition and cubic surface approximating the experimental data points in the background. I_{K_r} and I_{CaL} inhibition were computed using the Hill equation (Eq. 1 in the main text), with the CiPA (left) and Pharm (right) datasets (Table 2 in the main text) and nominal drug concentrations (Table 1 in the main text).

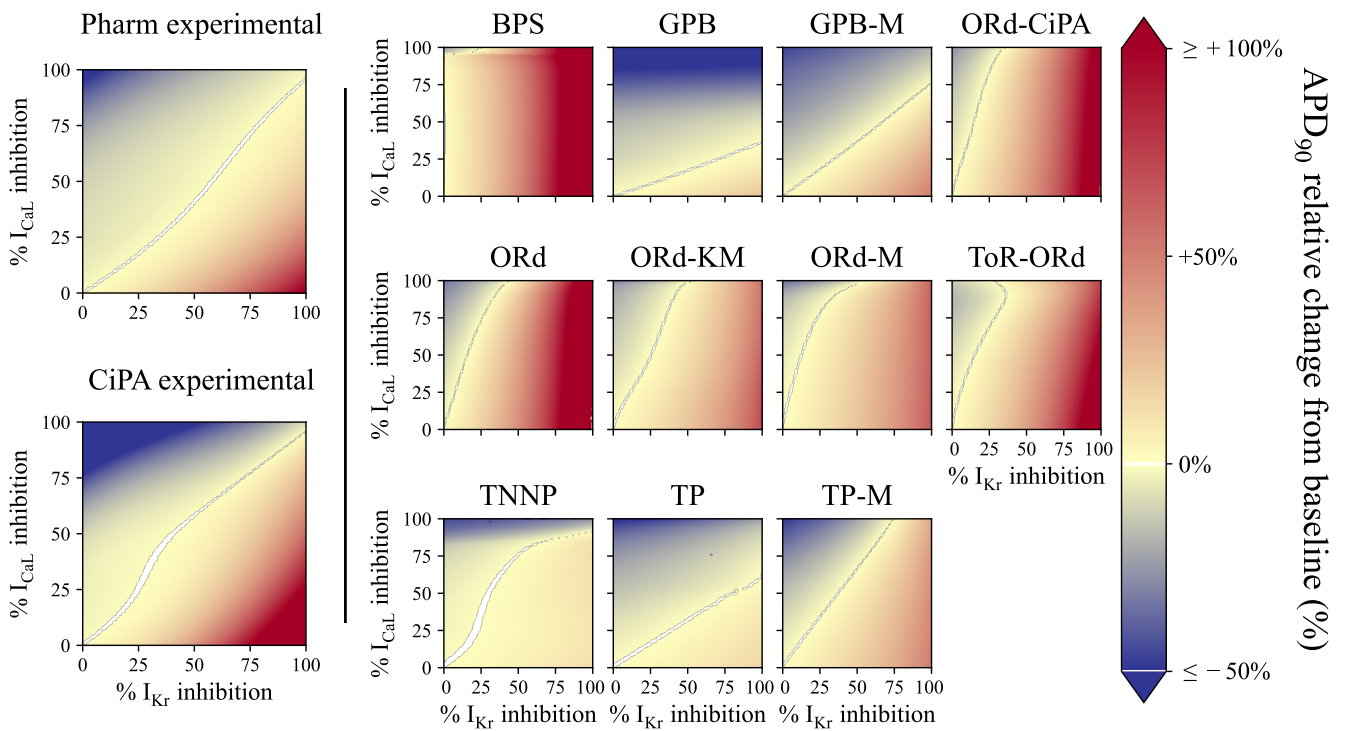


Figure H: 2-D maps of simulated $\% \Delta \text{APD}_{90}$ after I_{CaL} and I_{Kr} inhibition. The colour scale indicates shortening of APD_{90} (i.e., $\Delta \% \text{APD}_{90} < 0$ ms) for colours towards dark blue, and APD_{90} prolongation (i.e., $\Delta \% \text{APD}_{90} > 0$ ms) for colours towards red. ΔAPD_{90} values below -50 ms and above $+200$ ms were set to dark blue and red, respectively, for better visualisation. For I_{Kr} and I_{CaL} inhibition leading to $-1 \text{ ms} < \Delta \text{APD}_{90} < +1 \text{ ms}$, the pixel is coloured in white.

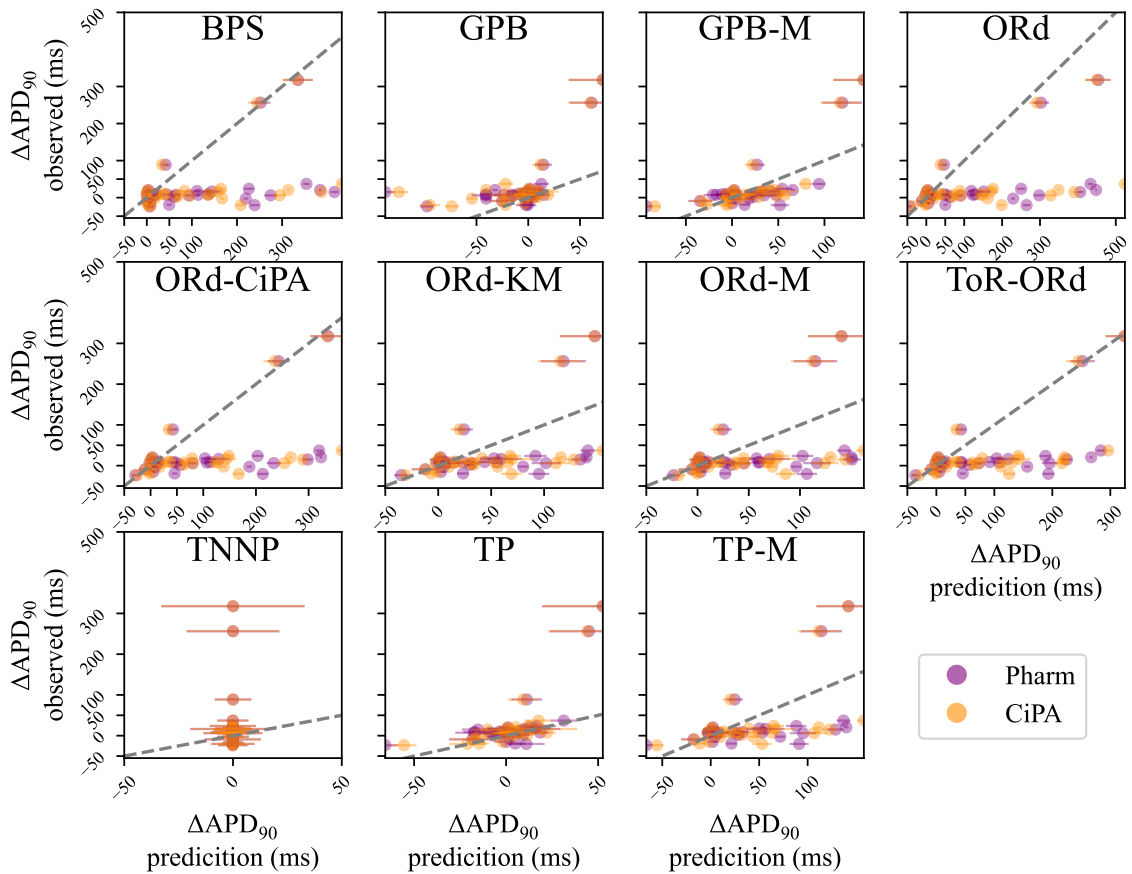


Figure I: Observed versus prediction plot for the tested models, using the CiPA and Pharm datasets (orange and purple, respectively). The experimental SEM was highlighted with horizontal error bars. The identity line is highlighted with the dashed line.

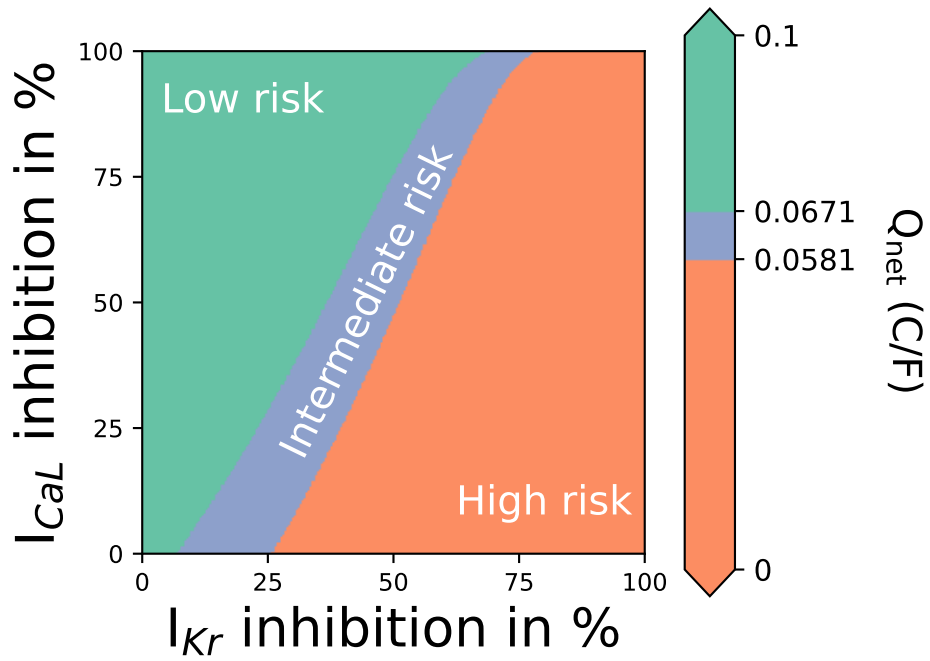


Figure J: Q_{net} computed with the ORd-CiPA model, for various combinations of I_{Kr} and/or I_{CaL} inhibition. The Torsade risk thresholds are defined in [Li *et al.* \(2019\)](#). Note that I_{Kr} inhibition was computed as plain reduction of the maximal conductance of I_{Kr} (Section 2.4.1 in the main text) instead of the dynamic hERG binding model used in the original study ([Li *et al.*, 2019](#)).

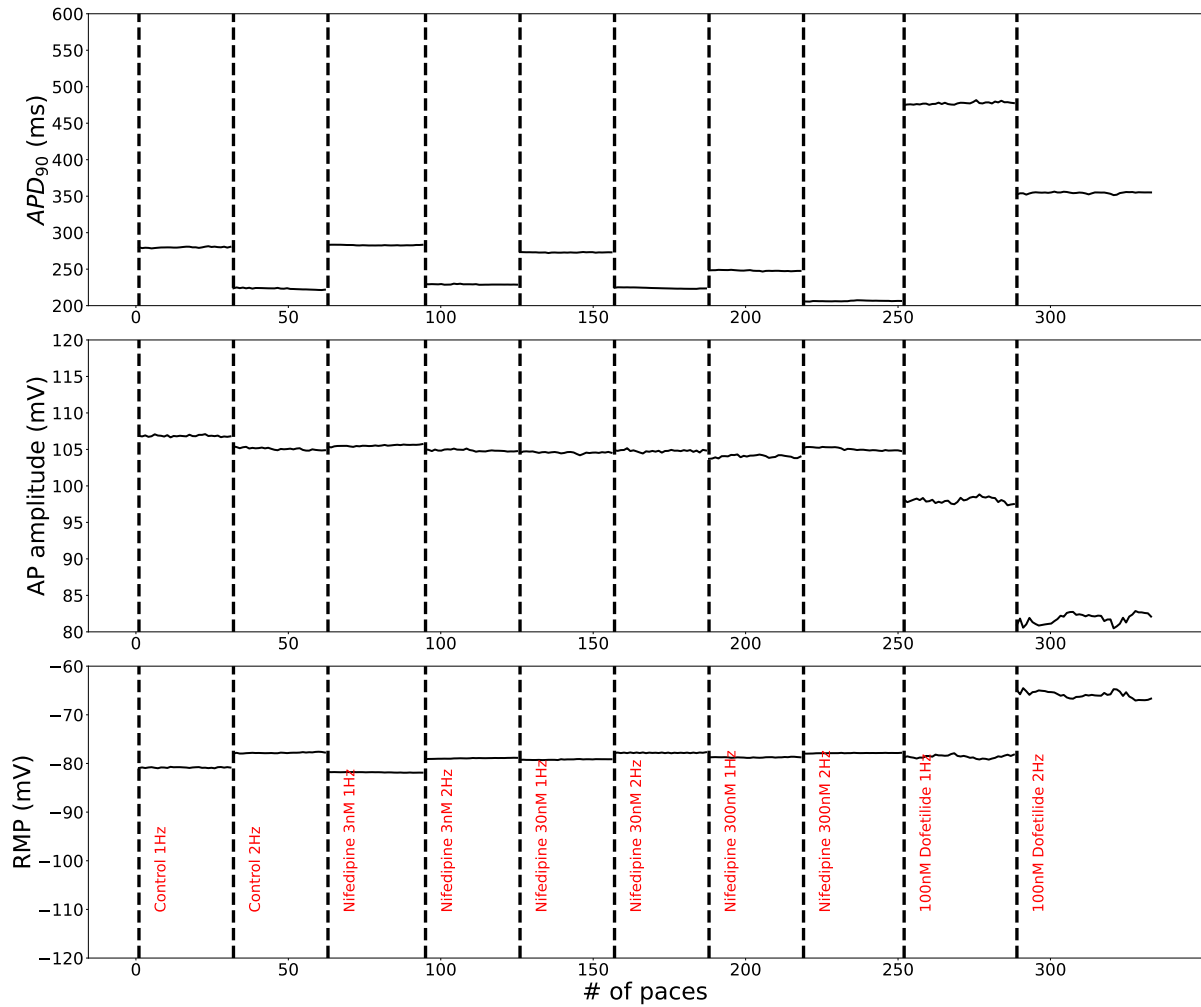


Figure K: Stability of AP markers observed in a representative trabecula, after 25 min of steady 1 Hz pacing or 3 min of 2 Hz pacing. For each condition, only the last 2 min are plotted.