SUPPLEMENTAL MATERIAL

Model Stimulation Protocol

Table S1: Model Protocol Values

Model Parameters

Canine Models

Table S2: Parameters varied in Fox Model

Fox Model is based on experimental data collected from canine midmyocardial cells.

Notes

*The scaling factor for I_{Kr} is not formally a maximal conductance, since this is multiplied by $\sqrt{K_0/4}$ and can therefore be greater than this value. For a constant value of extracellular $[K^+]$, changing this factor served to scale the current.

Table S3: Parameters varied in Hund Model

Hund Model is based on data from canine epicardial cells.

Notes

*The scaling factors for I_{K1} and I_{Kr} are not formally maximal conductances, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. For a constant value of extracellular [K+], changing this factor served to scale the current.

[†]The scaling factor for I_{Ks} is not formally a maximal conductance, since it is multiplied by a function of intracellular Ca^{2+} and can therefore be greater than this value.

‡The units for this maximal conductance are mistakenly listed as uA/uF in the original manuscript.

Parameter	Definition	Baseline value*
INaB	Fast Na ⁺ Channel	1
INabB	Background Na+Channel	1
INaLB	LateNa ⁺ Channel	1
ICaLB	L-type Ca ²⁺ Channel	1
ICabB	Background Ca ²⁺ Channel	
Ito1B	Transient outward K ⁺ Channel	
Ito2B	Calcium-activated Cl Channel	
IKrB	Rapid delayed rectifier K+Channel	
IKsB	Slow delayed rectifier K+Channel	
IK1B	Inward rectifier K ⁺ Channel	
IKpB	Plateau K ⁺ Channel	
INaKB	Na ⁺ -K ⁺ pump current	
INaCaB	Na^+ -Ca ²⁺ exchange current	
IrelB	SR Ca ²⁺ release	
IleakB	Passive SR Ca ²⁺ leak	
IupB	SR Ca ²⁺ uptake (SERCA)	
IpCaB	Sarcolemmal Ca ²⁺ pump current	
ICIB	Background Cl channel	

Table S4: Parameters varied in Heijman Model

Notes

*These variable names were present in the online version of the Heijman Model: <http://rudylab.wustl.edu/research/cell/code/AllCodes.html>. The original code had (1-IB) in front of each current equation with each baseline IB term equal to zero. We modified the code to impose variability by removing the (1-IB) with just IB and replacing its baseline value with 1.

Guinea Pig Models

TableS5: Parameters varied inLivshitzModel

Notes

- P_{Ca} , P_{Na} , P_{K} values were retrieved from http://rudylab.wustl.edu/research/cell/lrd2.htm and are different from the ones in the paper.
- [†]The scaling factors for I_{K1} and I_{Kr} are not formally maximal conductances, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. For a constant value of extracellular $[K^+]$, changing this factor served to scale the current.

 $*$ The scaling factor for I_{Ks} is not formally a maximal conductance, and is based on a function of intracellular [Ca²⁺] and can therefore be greater than this value. $\overline{G}_{Ks} = 0.433 +$ 0.2598

$$
1 + (38e - \frac{6}{[Ca^{2+}]i})^{1.4}
$$

Table S6: Parameters varied in Devenyi Model

Notes

*The scaling factors for I_{K1} and I_{Kr} are not formally maximal conductances, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. For a constant value of extracellular [K+], changing this factor served to scale the current.

 t The scaling factor for I_{Ks} is not formally a maximal conductance, and is based on a function of intracellular [Ca²⁺] and can therefore be greater than this value. $G_{Ks} = 0.433 + 0.2598$

$$
\frac{0.2556}{1+(38e-\frac{6}{[Ca^{2+}]i})^{1.4}}
$$

Rabbit Model

Table S7: Parameters varied in Shannon Model

Notes

*These values were reported incorrectly in the paper, and were taken from the source code: https://somapp.ucdmc.ucdavis.edu/Pharmacology/bers/

 \dagger The scaling factor for I_{Ks} is not formally a maximal conductance, since it is multiplied by a function of intracellular Ca^{2+} and can therefore be greater than this value.

[‡]The scaling factors for I_{K1} and I_{Kr} are not formally maximal conductances, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. For a constant value of extracellular $[K^+]$, changing this factor served to scale the current.

 V_{max} is 2.2 umol/L cytosol/sec in the paper but was converted to A/F as used in the source code: https://somapp.ucdmc.ucdavis.edu/Pharmacology/bers/

Human Models

All human models are based on experimental data from endocardial cells.

Parameter	Definition	Baseline value
G_{Na}	Maximal Na ⁺ conductance	14.838 mS/µF
G_{bNa}	Background Na ⁺ conductance	$2.9e-4$ mS/ μ F
G_{Cal}	Maximal L-type Ca ²⁺ current permeability	1.75e-4 $\rm cm^3/\mu F \, s^{-1}$
G_{bCa}	Background Ca ²⁺ conductance	$5.92e-4$ mS/ μ F
G_{to}	Maximal transient outward K ⁺ conductance	0.073 mS/ μ F
$G_{\rm Kr}$	Rapid delayed rectifier K ⁺ scaling factor [*]	0.096 mS/ μ F
G_{Ks}	Maximal slow delayed rectifier K ⁺ conductance	0.245 mS/ μ F
G_{K1}	Inward rectifier K ⁺ current scaling factor [*]	5.405 mS/ μ F
G_{pK}	Maximal plateau K ⁺ conductance	0.0146 mS/ μ F
P_{NaK}	Maximal Na ⁺ -K ⁺ pump current	1.362 pA/pF
k_{NaCa}	Maximal Na ⁺ -Ca ²⁺ exchange current	1000 pA/pF
a_{rel}	SR Ca ²⁺ release scaling factor	$16.464 \mu M \text{ ms}^{-1}$
c_{rel}	SR Ca ²⁺ release scaling factor	8.232 µM ms-1
$V_{\rm leak}$	Passive SR leak scaling factor	8e-5 ms ⁻¹
V_{maxup}	Maximal rate of SR Ca ²⁺ uptake (SERCA)	$0.425 \mu M \text{ ms}^{-1}$
G_{pCa}	Maximal sarcolemmal Ca ²⁺ pump current	0.025 pA/pF

Table S8: Parameters varied in TT04 Model

Notes

*The scaling factors for I_{Kr} and I_{K1} are not formally a maximal conductance, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. For a constant value of extracellular [K+], changing these factors serves to scale the current.

TableS9: Parameters varied in TT06 Model

Notes

*The scaling factors for I_{Kr} and I_{K1} are not formally a maximal conductance, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. For a constant value of extracellular [K+], changing these factors serves to scale the current.

 \dagger Additionally, V_{rel} = 0.102 ms⁻¹, as per the source code available at http://wwwbinf.bio.uu.nl/khwjtuss/ SourceCodes/HVM2/, rather than 40.8 mM ms-1 as inthe manuscript.

Table S10: Parameters varied in O'Hara Model

Notes

- *The scaling factors for I_{K1} and I_{Kr} are not formally maximal conductance, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. Changing this factor scales the current at all values of extracellular [K+] while maintaining the dependence on this variable.
- [†]The scaling factor for I_{Ks} is multiplied by a function of intracellular [Ca2+]. This value is therefore not precisely the current's maximal conductance.
- ‡ $\overline{I}_{\text{NaK}}$ was not labeled in the original paper and have been given these names to keep to terminology consistent.
- $$P\alpha$$ Parameters controlling the magnitude of SR Ca²⁺ release through the ryanodine receptor (RyR) and through SERCA are an introduced unitless multiplier with baseline value equals to 1.00.

Parameter	Definition	Baseline value
G_{Na}	Maximal Na ⁺ conductance	23 mS/uF
$G_{\underline{N}a\,\underline{B}}$	Background Na ⁺ conductance	$0.597x10^{-3}mS/uF$
p_{Ca}	Maximal L-type Ca ²⁺ current permeability*	2.7e-4cm/sec
$G_{\underline{Ca_{B}}}$	Background Ca ²⁺ conductance	5.513e-4 uA/uF
$G_{\text{to_slow}}$	Maximal slow transient outward K ⁺ conductance	0.0376mS/uF
$G_{\text{to} \text{ Fast}}$	Maximal fast transient outward K ⁺ conductance	0.0014 m S/uF
	Maximal rapid delayed rectifier K ⁺ current	0.035 mS/uF
g_{Kr}	conductance ²	
	Maximal slow delayed rectifier K ⁺ current	0.0035 mS/uF
$g_{\rm ks\,sl} = g_{\rm ks\,junc}$	conductance	
G_{K1}	Maximal inward rectifier K ⁺ conductance ^{†,‡}	0.35 mS/uF
g_{kp}	Maximal plateau K ⁺ conductance	0.002 mS/uF
$G_{Cl_{Ca}}$	Maximal Ca ²⁺ -activated Cl-conductance	0.0548 mS/uF
G_{CLB}	Background Cl conductance	9e-3 mS/uF
$I_{\text{Na}_{K}}$	Maximal Na ⁺ -K ⁺ pump current	1.8 uA/uF
I_{NCX}	Maximal Na ⁺ -Ca ²⁺ exchange current	4.5 uA/uF
k_{s}	SR Ca ²⁺ release	$25 \,\mathrm{ms}^{-1}$
$K_{\underline{\text{leak}}}$	Passive SR Ca ²⁺ leak scaling factor [‡]	5.348e-6 ms ⁻¹
$V_{\text{max}}_{\frac{\text{SR}}{\text{Cap}}}$	Maximal rate of SR Ca ²⁺ uptake (SERCA)	5.3114e-3 mM/ms
I _{PMCA}	Maximal sarcolemmal Ca ²⁺ pump current	0.0673 uA/uF

Table S11: Parameters varied in Grandi Model

Notes

 p_{ca} = 2.7 x 10⁻⁴ cm/sec, as per the source code available at

https://somapp.ucdmc.ucdavis.edu/Pharmacology/bers/, rather than 1.215 x 10-4cm/sec as in the manuscript.

[†]The scaling factors for I_{K1} and I_{Kr} are not formally maximal conductance, since each is multiplied by $\sqrt{K_0/5.4}$ and can therefore be greater than this value. Changing this factor scales the current at all values of extracellular [K+] while maintaining the dependence on this variable.

 G_{K1} and K_{leak} were not labeled in the original paper and have been given these names to keep to terminology consistent.

Population Distributions

Table S12: Data collected during Population Runs

Table S13: Population Statistics

Figure S1. Procedure to determine EAD Threshold

AP waveforms of beats 91-100 of the O'Hara Human Model under calcium perturbation at 1 Hzpacing rate. EAD appearance was not consistent at every beat. Thus, in order to address this variability, beats 91-100 were analyzed and the first I_{cal} Factor that caused an EAD between beats 91-100 was considered the threshold for triggering EADs in the model. This procedure was repeated for each model. The O'Hara Model displayed an EAD first on the 91st beat at an I_{Cal} Factor of 15.2. This analysis was replicated in each model.

Figure S2. AP Clamp Simulation Procedure

This represents the AP Clamp procedure performed in the O'Hara Model. The baseline model's AP waveform was split into two epochs: the depolarized period when membrane potential was more positive than -75 mV and the resting period when membrane potential was more negative than -75 mV. The two epochs were either stretched or compressed using the *interp1* function in MATLAB to make the AP longer or shorter while keeping the total duration constant at 1000 ms. This procedure was replicated in the remaining nine models.

The same simulations performed in Figure 1 on the O'Hara model were replicated two additional human models (TT04 and Grandi). (A/F) Only a low I_{ks} version for the TT04 model and a high I_{ks} version for the Grandi model were developed because both of these models at baseline have very high and low levels of I_{ks} at baseline, respectively. Thus, increasing the levels of I_{κ} in the TT04 and decreasing the levels of I_{κ} in the Grandi would not provide further information. The TT04 Low I_{ks} version Grandi High I_{ks} by decreasing or increasing (20 fold) IKs and adjusting IKr to produce an identical APD. (B/G) IKs and IKr waveforms of the different versions of the same model. The shaded regions represent the integrated currents (area under the curve, or AUC) during the APs. The current AUCs were used to calculate the fraction of repolarizing current contributed by (C/H) IKs (IKs Fraction) in each model. (D/I) Using each of the model variants as the baseline model, model populations (300 cells each) were generated by randomly varying parameters. Distributions of APD show that greater IKs promotes less variability within a population. (E/J) APD Spread was calculated for each of the populations to quantify the variability within the APD distributions.

The same simulations performed in Figure 2 on the O'Hara model were replicated two additional human models (TT04 and Grandi). I_{cal} was progressively increased in each model to induce EADs and arrhythmic behavior. I_{cal} factor refers to the increase in channel permeability coefficient (1.0 equals control level).(A) APs simulated in the baseline version of TT04 with I_{cal} augmentation (8.1 and 16.1 times). (B) APs simulated in the low TT04 model with the same I_{Cat} perturbations. (C) Both versions of the TT04 model were run under a wide range of I_{Cat} factors and plotted against APD. The end of each line represents the last factor before an EAD formed. (D) APs simulated in the baseline version of Grandi with I_{cal} augmentation (1.6 and 2 times). (E) High I_{ks} version of Grandi simulated under the same conditions. (F) Both versions of the Grandi model were run under a wide range of I_{cat} factors and plotted against APD.

Figure S5. Baseline AP and I_{Kr}/I_{Ks} Waveforms

AP waveforms and slow & rapid delayed rectifier K^{*} currents in seven additional ventricular

Figure S6. Population Variability across multiple species.

Histograms of AP duration of a population of 300 model variants generated by imposing random variability.

Aps and ICaL waveforms of each model with increasing I_{cat} current. IThe blue curves show the baseline I_{cal} whereas the black dashed curves plot I_{cal} just before the first EAD occurred. The gray shaded regions are the area under the curve (AUC). Plots of Shannon and TT06 represent the 91st beat, plot of Fox is 100th beat, plot of Hund is the 99th beat, and plots of Heijman and Devenyi are the 92nd beat, while Livshitz is the 93rd beat.

Figure S8. Inducing proarrhythmic behavior through constant inward current inject across multiple species. AP waveforms of each model with increasing constant inward current injection (from blue to red) until proarrythmic behaviors were present. The bar graph represents the level of current injected in order to produce the failed APs in red in each model.

(A) The scaling factor (I_{cat} factor) by which the calcium current was increased does not correlate with the levels of Iks. Since the baseline I_{cal} magnitude differed between models, the I_{cal} factor did not provide a fair basis of comparison. To compare suscep bility to EADs, we integrated the I_{cal} waveform at the level just under the EAD threshold as depicted in Figures 4 & 5. (B) The percent change in APD between the APD when 0.1 A/F inward current was injected into the models and the baseline APD was calculated and compared with the levels of I_{κ} . This shows that models with low I_{κ} are much more suscep ble to APD prolonga on when an arbitrary inward current is injected as compared to models with high I_{κ} .

The AP Clamp simulation performed in Figure 6 was replicated in all the models. The x-axis refers to the factor by which the APD was increased. The y -axis represents the integrated currents (area under the curve, AUC) of each K⁺ currents, normalized to the integrated current of the baseline AP. As APD increases, there is a much greater relative increase in I_{ks} compared with I_{ks} across all models. The x marked lines represent I_{κ} and the circle marked lines represent I_{κ} . The inset in each figure represents the normalized current at the last time point in each figure. Please note that the TT04, TT06, and the Grandi models had longer baseline APDs, thus increasing the APD by a factor of 3 was not possible.

Figure S11. Contribution of individual substrates to changes in APD Spread

We individually blocked the phosphorylation of each of the eight PKA targets to see how this contributes to the change in APD Spread. We simulated the same population of 300 cells under the ten conditions. The no ISO population contained thirteen APs that formed EADs, which were removed from each population.