

# Supplementary Material

# 1 NA<sup>+</sup>-CA<sup>2+</sup> EXCHANGER MODEL

$$I_{NCX} = I_{NCX}^{max} \cdot \frac{1}{1 + \left(\frac{k_{dact}}{Ca_i}\right)^2} \cdot \frac{e^{\frac{\nu V_m F}{RT}} \cdot Ca_o \cdot Na_i^3 - e^{\frac{(\nu-1)V_m F}{RT}} \cdot Na_o^3 \cdot Ca_i}{Km_{Ca_i} \cdot Na_o^3 \left(1 + \left(\frac{Na_i}{Km_{Na_i}}\right)^3\right)} \cdot \frac{1}{1 + k_{sat} \cdot e^{\frac{(\nu-1)V_m F}{RT}}} + Km_{Na_o}^3 \cdot Ca_i \cdot \left(1 + \frac{Ca_i}{Km_{Ca_i}}\right) + (Km_{Ca_o} + Ca_o) \cdot Na_i^3 + Na_o^3 \cdot Ca_i$$

$$\begin{split} I_{\rm NCX}^{max} &= 4.41\,{\rm pA/pF} \\ k_{dact} &= 0.384 \times 10^{-3}\,{\rm mM} \\ Km_{Ca_i} &= 3.59 \times 10^{-3}\,{\rm mM} \\ Km_{Ca_o} &= 1.3\,{\rm mM} \\ Km_{Na_i} &= 12.29\,{\rm mM} \\ Km_{Na_o} &= 87.5\,{\rm mM} \\ nu &= 0.35 \\ k_{sat} &= 0.27 \end{split}$$

### 2 CA<sup>2+</sup> AND NA<sup>+</sup> DIFFUSION MODEL



**Figure S1.** Schematic representation of  $Ca^{2+}$  and  $Na^+$  concentrations and their across the different model compartments in a membrane domain.

#### 3 NA<sup>+</sup> MASS BALANCE

In the case of the simplified

$$\frac{dNa_i}{dt} = \frac{J_{Na_{SLmyo}}}{v_{myo}} \cdot \left( (Na_{SL}^1 - Na_i) + (Na_{SL}^2 - Na_i) \right) + \frac{(Na_i^{-1} - Na_i) + (Na_i^{-2} - Na_i)}{\tau_D}$$

where  $\tau_D = 0.6$  /s is the diffusion constant.  $Na_{SL}^1$  and  $Na_{SL}^2$  are the Na<sup>+</sup> concentrations in the SL space of the membrane domains, and  $Na_i^{-1}$  and  $Na_i^{-2}$  are the Na<sup>+</sup> concentrations in the neighbouring left and right segments, respectively.  $J_{Na_{slmoy}}$  is the net buffered Na<sup>+</sup> flux and equals  $1.6386 \times 10^{-12}$  L/ms.

Since in this study we simulate only one segment and both membrane domains are identical, the above equation simplifies to:

$$\frac{dNa_i}{dt} = \frac{J_{Na_{SLmyo}}}{v_{myo}} \cdot 2 \cdot \left(Na_{SL} - Na_i\right)$$

At the membrane domains, Na<sup>+</sup> is updated as:

$$\frac{dNa_{SL}}{dt} = I_{Na_{SL}}^{tot} \cdot \frac{C_m F}{v_{SL}} + J_{Na_{juncSL}} \cdot (Na_{junc} - Na_{SL}) + \frac{2 \cdot J_{Na_{SL}myo}}{v_{SL}} \cdot (Na_i - Na_{SL}) - Buff_{Na_{SL}}$$

$$\frac{dNa_{junc}}{dt} = I_{Na_{junc}}^{tot} \cdot \frac{C_m F}{v_{junc}} + \frac{J_{Na_{juncSL}}}{V_{junc}} \cdot (Na_{SL} - Na_{junc}) - Buff_{Na_{junc}}$$

where  $J_{Na_{juncSL}} = 1.8313 \times 10^{-14} \,\text{l/ms}$  is the Na<sup>+</sup> flux between the SL and junc spaces.  $Buf f_{Na_{SL}}$  and  $Buf f_{Na_{junc}}$  are the buffering variables of Na<sup>+</sup> in the SL and junc spaces, respectively, and are given by:

$$\frac{dBuff_{Na_{SL}}}{dt} = k_{on}^{Na} \cdot Na_{SL} \cdot (B_{Na_{SL}}^{max} - Buff_{Na_{SL}}) - k_{off}^{Na} \cdot Buff_{Na_{SL}}$$

$$\frac{dBuff_{Na_{junc}}}{dt} = k_{on}^{Na} \cdot Na_{junc} \cdot (B_{Na_{junc}}^{max} - Buff_{Na_{junc}}) - k_{off}^{Na} \cdot Buff_{Na_{junc}}$$

where  $k_{on}^{Na} = 0.0001$  and  $k_{off}^{Na} = 0.001$  are the rate constants of Na<sup>+</sup>buffer complex formation and dissociation, respectively.  $B_{Na_{SL}}^{max}$  and  $B_{Na_{junc}}^{max}$  are the maximum buffer capacities in the SL and junc spaces and are equal to 7.561 mM and 1.65 mM, respectively.

#### 4 SERCA2A MODEL

$$J_{Serca} = V_{max} \cdot \frac{\left(\frac{Ca_i}{Kmf}\right)^{nh} - \left(\frac{Ca_{SR}}{Kmr}\right)^{nh}}{1 + \left(\frac{Ca_i}{Kmf}\right)^{nh} + \left(\frac{Ca_{SR}}{Kmr}\right)^{nh}}$$

$$V_{max} = 5.3 \times 10^{-3} \,\mathrm{mM/s}$$
  
 $Kmf = 0.000\,625 \,\mathrm{mM}$   
 $Kmr = 1.0 \,\mathrm{mM}$   
 $nh = 1.787$ 

#### 5 CA<sup>2+</sup> BUFFERS

Buffers are modelled using a two-state model:

$$\frac{dC_B}{dt} = k_{on}C(B_{max} - C_B) - k_{off}C_B$$

where  $B_{max}$  is the maximum buffer capacity, C and  $Ca_B$  represent the concentration of free and bound ligand, and  $k_{on}$  and  $k_{off}$  are the rate coefficients of ion-buffer complex formation and dissociation, respectively.

Buffer	Compartment	Bmax (µM)	kon (/s)	<b>koff</b> (/ $\mu$ Ms)
Troponin C High (Ca <sup>2+</sup> )	cytosol	140E-3	2.37	3.2E-5
Troponin C High (Mg <sup>2-</sup> )	cytosol	140E-3	3E-3	3.33E-3
Troponin C Low	cytosol	70E-3	163.5	98E-3
Calmodulin	cytosol	2.4E-3	34	238E-3
Myosin (Ca)	cytosol	140E-3	13.8	4.6E-4
Myosin (Mg)	cytosol	140E-3	0.0157	5.7E-5
SR	cytosol	17.1E-3	100	6E-2
SLlow	SL	$37.4\text{E-3} \cdot \frac{v_{myo}}{v_{SL}}$	100	1.3
SLlow	junc	$4.6\text{E-4} \cdot \frac{v_{myo}}{v_{junc}}$	100	1.3
SLhigh	SL	$13.4\text{E-3} \cdot \frac{v_{myo}}{v_{SL}}$	100	3E-2
SLhigh	junc	$1.65\text{E-4} \cdot \frac{v_{myo}}{v_{junc}}$	100	3E-2
Calsequestrin	SR	$140\text{E-}3 \cdot \frac{v_{myo}}{v_{SR}}$	100	65
Na <sup>+</sup>	SL	7.561	1.0E-4	1.0E-3
Na <sup>+</sup>	junc	1.65	1.0E-4	1.0E-3

Table S1. Buffer parameters.

#### **6 INITIAL CONDITIONS**

Ionic traces	Initial concentration	Description
Ca <sub>cyt</sub>	$0.21\mu M$	Ca <sup>2+</sup> concentration in the cytosol.
Ca <sub>junc</sub>	$0.073\mu\mathrm{M}$	Ca <sup>2+</sup> concentration in the junctional domain of the SS space.
Ca <sub>sl</sub>	$0.23\mu M$	Ca <sup>2+</sup> concentration in the sarcolemmal domain of the SS space.
Ca <sub>SR</sub>	$0.25\mu\mathrm{M}$	Luminal $Ca^{2+}$ concentration in the sarcoplasmic reticulum.
Ca <sub>SRS</sub>	$0.073\mu M$	$Ca^{2+}$ concentration in the sub SR space.
Na <sub>cyt</sub>	$11.3\mathrm{mM}$	Na <sup>+</sup> concentration in the cytosol (2Hz).
Najunc	$11.3\mathrm{mM}$	Na <sup>+</sup> concentration in the junctional domain of the SS space.
Na <sub>SL</sub>	$11.3\mathrm{m}\mathrm{M}$	Na <sup>+</sup> concentration in the sarcolemmal domain of the SS space.
K <sub>cyt</sub>	$120\mathrm{m}\mathrm{M}$	K <sup>+</sup> concentration in the cytosol.
Cl <sub>cyt</sub>	$30\mathrm{m}\mathrm{M}$	Cl <sup>-</sup> concentration in the cytosol.
Ca <sub>o</sub>	$1.8\mathrm{mM}$	Extracellular $Ca^{2+}$ concentration.
Na <sub>o</sub>	$140\mathrm{m}\mathrm{M}$	Extracellular Na <sup>+</sup> concentration.
Ko	$5.4\mathrm{mM}$	Extracellular K <sup>+</sup> concentration.
Clo	$132\mathrm{m}\mathrm{M}$	Extracellular Cl <sup>-</sup> concentration.

 Table S2. Initial ionic concentrations in the different compartments of the model.

#### 7 SELECTED POPULATION

Table S3. Parameter values of the 16 models of the calibrated population.

G <sub>CaL</sub>	G <sub>CaT</sub>	$\mathrm{I}_{\mathrm{NaK}}^{max}$	I <sub>NaCa</sub>	G <sub>Na</sub>	G <sub>to</sub>	G <sub>Kr</sub>	G <sub>Ks</sub>	G <sub>K1</sub>	$I_{CaP}^{max}$	G <sub>Cab</sub>	G <sub>Nab</sub>	G <sub>Clb</sub>
0.228	0.415	2.449 E-3	1.436	0.088 E-3	0.712	0.120	0.174	0.623	0.589	0.970 E-3	0.398 E-3	6.392 E-3
0.146	0.474	1.801 E-3	1.162	0.068 E-3	0.606	0.056	0.119	0.333	0.246	0.506 E-3	0.623 E-3	7.647 E-3
0.192	0.069	2.425 E-3	1.073	0.068 E-3	0.443	0.268	0.058	0.277	0.401	1.348 E-3	0.885 E-3	5.078 E-3
0.202	0.437	1.651 E-3	3.314	0.060 E-3	0.492	0.176	0.178	0.673	0.403	1.024 E-3	0.612 E-3	8.786 E-3
0.295	0.124	2.573 E-3	1.165	0.071 E-3	0.790	0.248	0.199	0.263	0.244	0.348 E-3	0.631 E-3	5.742 E-3
0.099	0.147	1.320 E-3	0.762	0.063 E-3	0.155	0.135	0.182	0.344	0.466	1.022 E-3	0.772 E-3	4.047 E-3
0.256	0.156	1.892 E-3	3.786	0.050 E-3	0.354	0.193	0.155	0.777	0.067	1.033 E-3	0.143 E-3	6.964 E-3
0.189	0.406	1.599 E-3	1.313	0.064 E-3	0.435	0.181	0.084	0.697	0.582	0.482 E-3	0.944 E-3	5.345 E-3
0.222	0.325	2.366 E-3	2.012	0.104 E-3	0.449	0.209	0.122	0.625	0.395	1.594 E-3	0.221 E-3	5.321 E-3
0.189	0.431	2.100 E-3	1.532	0.042 E-3	0.514	0.205	0.113	0.432	0.220	1.360 E-3	0.176 E-3	7.643 E-3
0.231	0.038	1.602 E-3	3.161	0.073 E-3	0.604	0.142	0.174	0.299	0.128	0.524 E-3	0.405 E-3	7.133 E-3
0.322	0.388	2.407 E-3	3.200	0.109 E-3	0.558	0.262	0.121	0.414	0.315	0.949 E-3	1.570 E-3	8.421 E-3
0.180	0.458	1.934 E-3	1.081	0.062 E-3	0.162	0.144	0.062	0.314	0.455	0.312 E-3	0.851 E-3	7.995 E-3
0.293	0.204	2.671 E-3	1.528	0.104 E-3	0.729	0.262	0.181	0.396	0.505	0.173 E-3	1.228 E-3	7.827 E-3
0.201	0.305	1.302 E-3	1.882	0.105 E-3	0.699	0.161	0.133	0.531	0.263	0.471 E-3	0.112 E-3	8.358 E-3
0.155	0.329	1.775 E-3	3.535	0.094 E-3	0.574	0.146	0.129	0.380	0.339	1.410 E-3	1.373 E-3	3.252 E-3

#### 8 I-V CURVES

Table S4. Maximum and minimum peak current values obtained from simulated I-V curves in Fig. S2, and comparison with experimental values reported in Aslanidi et al. (2009)(1) and Muraki et al. (1995)(2). Values are in pA/pF.

		I-V curv	ves	2 Hz Pacing			
	Simulated		Experimental	Our I	model	Aslanidi model	
	Min. peak	Max. peak	Experimental	Min. peak	Max. peak	Asianui mouei	
I <sub>CaL</sub>	-13.9	-4.3	-8 (PM), -6 (CT)	-6.4	-2.1	-3.6	
I <sub>Na</sub>	-737	-291	-70	-393	-192	-120	
Ito	3.4	16.6	13 (PM), 8 (CT)	0.87	4.0	1.3	
I <sub>K1</sub>	0.88	2.6	5 (PM), 4 (CT)	0.88	2.6	0.66	
I <sub>Kr</sub>	0.45	2.1	0.67*	0.41	2.4	0.43	

\* Assuming a cell capacitance of  $50 \,\mathrm{pF}$ 



**Figure S2.** Simulated I-V curves (blue) of  $I_{CaL}$ ,  $I_{Na}$ ,  $I_{to}$ , and  $I_{K1}$  for the 16 models in the control population. The voltage-clamp protocol applied consisted of trains of 200 ms square depolarizing pulses at a cycle length of 2 s from a holding potential of -80 mV to voltages between -80 mV and 40 mV, in steps of 10 mV. Experimental I-V curves (red) are shown for comparison. In panels A and C, curves for both pectinate muscle (PM, stars) and crista terminalis (CT, squares) rabbit atrial CMs are shown (data from Aslanidi et al. (2009)). The experimental curve in panel D is normalized data, shown here for qualitative comparison purposes. The experimental curve in E assumed a cell capacitance of 50 pF.



**Figure S3.** Comparison between  $I_{CaL}$ ,  $I_{Na}$ ,  $I_{to}$ ,  $I_{K1}$ ,  $I_{Kr}$ , and  $I_{Ks}$  obtained with our control population and with the Aslanidi model(1) (CellML version) at 2 Hz pacing.

## REFERENCES

- [1]Aslanidi et al. *Mechanisms of transition from normal to reentrant electrical activity in a model of rabbit atrial tissue: Interaction of tissue heterogeneity and anisotropy*, Biophysical Journal 96(3), 798-817, 2009.
- [2]Muraki et al. *Delayed rectifier K*<sup>+</sup> *current in rabbit atrial myocytes*, Am J of Phys-Heart and Circ Phys 269(2), H524-H532, 1995.