

# A basic model of calcium homeostasis in non-excitable cells

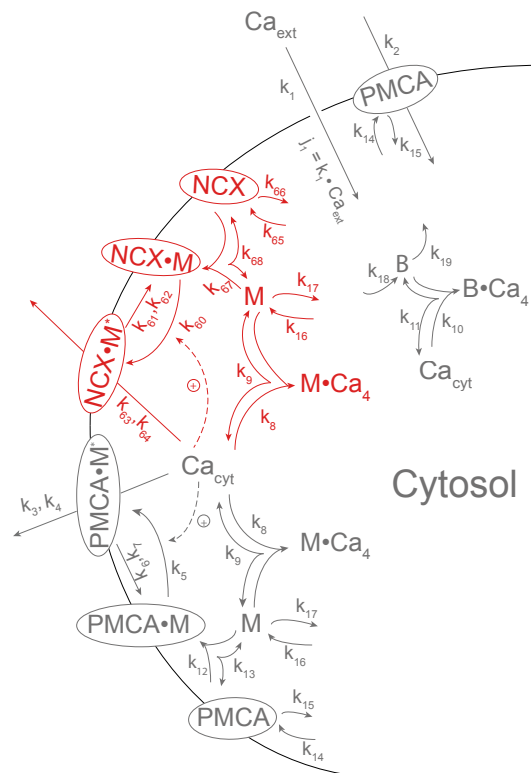
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## Supporting Information S4 Program

Rate equations, parameters, initial concentrations, and calculations for Figs 7C and 7D

Calculations in Figs 7C,D are based on the model shown in Fig S1.



**Figure S1. Model including both PMCA and NCX for cytosolic  $Ca^{2+}$  extrusion.** The added NCX part is outlined in red.

The rate equations are:

$$\begin{aligned}
\frac{dCa_{ext}^{2+}}{dt} = & k_1 \times Ca_{ext}^{2+} - k_2 \times Ca_{cyt}^{2+} \times PMCA - k_3 \times \frac{(PMCA \bullet M^*) \times Ca_{cyt}^{2+}}{k_4 + Ca_{cyt}^{2+}} \\
& - 4 \times k_8 \times M \times Ca_{cyt}^{2+} + 4 \times k_9 \times (M \bullet Ca_4) - 4 \times k_{10} \times B \times Ca_{cyt}^{2+} \\
& + 4 \times k_{11} \times (B \bullet Ca_4) - \frac{k_{63} \times Ca_{cyt}^{2+} \times (NCX \bullet M^*)}{k_{64} + Ca_{cyt}^{2+}}
\end{aligned} \tag{S1}$$

$$\frac{d(PMCA \bullet M^*)_{ext}}{dt} = k_5 \times Ca_{cyt}^{2+} \times (PMCA \bullet M) - \frac{k_6 \times (PMCA \bullet M^*)}{k_7 + (PMCA \bullet M^*)} \tag{S2}$$

$$\frac{dM}{dt} = k_{16} - k_{17} \times M - k_{12} \times M \times PMCA + k_{13} \times (PMCA \bullet M) \tag{S3}$$

$$+ k_9 \times (M \bullet Ca_4) - k_8 \times Ca_{cyt}^{2+} \times M + k_{68} \times (NCX \bullet M) \tag{S4}$$

$$- k_{67} \times M \times NCX \tag{S5}$$

$$\frac{d(M \bullet Ca_4)}{dt} = k_8 \times Ca_{cyt}^{2+} \times M - k_9 \times (M \bullet Ca_4) \tag{S6}$$

$$\frac{dB}{dt} = k_{18} - k_{19} \times B - k_{10} \times B \times Ca_{cyt}^{2+} + k_{11} \times (B \bullet Ca_4) \tag{S7}$$

$$\frac{d(B \bullet Ca_4)}{dt} = k_{10} \times B \times Ca_{cyt}^{2+} - k_{11} \times (B \bullet Ca_4) \tag{S8}$$

$$\begin{aligned}
\frac{d(PMCA \bullet M)}{dt} = & k_{12} \times PMCA \times M - k_{13} \times (PMCA \bullet M) \\
& + \frac{k_6 \times (PMCA \bullet M^*)}{k_7 + (PMCA \bullet M^*)} - k_5 \times Ca_{cyt}^{2+} \times (PMCA \bullet M)
\end{aligned}$$

$$\frac{dPMCA}{dt} = k_{14} - k_{15} \times PMCA - k_{12} \times PMCA \times M + k_{13} \times (PMCA \bullet M)$$

$$\frac{d(NCX \bullet M^*)}{dt} = k_{60} \times (NCX \bullet M) \times Ca_{cyt}^{2+} - \frac{k_{61} \times (NCX \bullet M^*)}{k_{62} + (NCX \bullet M^*)}$$

$$\begin{aligned} \frac{d(NCX \bullet M)}{dt} &= k_{67} \times M \times NCX - k_{68} \times (NCX \bullet M) \\ &\quad - k_{60} \times NCX \bullet M \times Ca_{cyt}^{2+} + \frac{k_{61} \times (NCX \bullet M^*)}{k_{62} + (NCX \bullet M^*)} \end{aligned}$$

$$\frac{dNCX}{dt} = k_{65} - k_{66} \times NCX - k_{67} \times M \times NCX + k_{68} \times (NCX \bullet M)$$

The time steps during LSODE integration is  $1.0 \times 10^{-2}$ s, with 5 phases having intervals of 48s, 144s, 210s, 148s, and 290s.

The rate constants/parameters are given as:

$k_1$ , phase 1 =  $1 \times 10^{-4} \text{s}^{-1}$ ,  $k_2 = 0.0 \text{s}^{-1}$ ,  $k_3$ , phase 1 =  $5.0 \times 10^3 \text{s}^{-1}$ ,  $k_4 = 1.2 \mu\text{M}$ ,  $k_5 = 16.0 \mu\text{M}^{-1} \text{s}^{-1}$ ,  $k_6 = 8.0 \times 10^{-3} \mu\text{M/s}$ ,  $k_7 = 1.0 \times 10^{-6} \mu\text{M}$ ,  $k_8 = 2.5 \text{s}^{-1}$ ,  $k_9 = 5.0 \text{s}^{-1}$ ,  $k_{10} = 1.0 \times 10^2 \mu\text{M}^{-1} \text{s}^{-1}$ ,  $k_{11} = 80.0 \text{s}^{-1}$ ,  $k_{12} = 1.0 \times 10^{-2} \mu\text{M}^{-1} \text{s}^{-1}$ ,  $k_{13} = 1.0 \times 10^{-1} \mu\text{M}^{-1} \text{s}^{-1}$ ,  $k_{14} = 0.0 \mu\text{M/s}$ ,  $k_{15} = 0.0 \text{s}^{-1}$ ,  $k_{16} = 0.0 \mu\text{M/s}$ ,  $k_{17} = 0.0 \text{s}^{-1}$ ,  $k_{18} = 0.0 \mu\text{M/s}$ ,  $k_{19} = 0.0 \text{s}^{-1}$ ,  $k_{60} = 16.0 \mu\text{M}^{-1} \text{s}^{-1}$ ,  $k_{61} = 8.0 \times 10^{-3} \mu\text{M/s}$ ,  $k_{62} = 1.0 \times 10^{-6} \mu\text{M}$ ,  $k_{63} = 1 \times 10^5 \text{s}^{-1}$  (all 5 phases),  $k_{64} = 1.0 \times 10^2 \mu\text{M}$ ,  $k_{65} = 0.0 \mu\text{M/s}$ ,  $k_{66} = 0.0 \text{s}^{-1}$ ,  $k_{67} = 1.0 \times 10^{-2} \mu\text{M}^{-1} \text{s}^{-1}$ ,  $k_{68} = 1.0 \times 10^{-1} \text{s}^{-1}$ ,  $k_1$ , phase 2 =  $1.4 \times 10^{-2} \text{s}^{-1}$ ,  $\alpha$ , phase 2 =  $4.5 \times 10^{-2} \text{s}^{-1}$ ,  $k_3$ , phase 2 =  $5.0 \times 10^3 \text{s}^{-1}$ ,  $\beta$ , phase 2 =  $200 \text{s}^{-1}$ ,  $k_1$ , phase 3 =  $1 \times 10^{-4} \text{s}^{-1}$ ,  $k_3$ , phase 3 =  $5.0 \times 10^3 \text{s}^{-1}$ ,  $k_1$ , phase 4 =  $1.4 \times 10^{-2} \text{s}^{-1}$ ,  $\alpha$ , phase 4 =  $4.5 \times 10^{-2} \text{s}^{-1}$ ,  $k_3$ , phase 4 =  $5.0 \times 10^3 \text{s}^{-1}$ ,  $\beta$ , phase 4 =  $200 \text{s}^{-1}$ ,  $k_1$ , phase 5 =  $0.0 \text{s}^{-1}$ ,  $\alpha$ , phase 5 =  $0.0 \text{s}^{-1}$ ,  $k_3$ , phase 5 =  $0.0 \text{s}^{-1}$ ,  $\beta$ , phase 5 =  $0.0 \text{s}^{-1}$ .

Initial concentrations (all in  $\mu\text{M}$ ):  $Ca_{cyt}^{2+} = 1.0652 \times 10^{-1}$ ,  $PMCA \bullet M^* = 3.7450 \times 10^{-4}$ ,  $Ca_{ext}^{2+} = 2 \times 10^3$ ,  $M = 9.4988$ ,  $M \bullet Ca4 = 5.0588 \times 10^{-1}$ ,  $B = 1.7785 \times 10^2$ ,  $B \bullet Ca4 = 23.679$ ,  $PMCA \bullet M = 4.6826 \times 10^{-3}$ ,  $PMCA = 4.9428 \times 10^{-3}$ ,  $NCX \bullet M^* = 3.7450 \times 10^{-4}$ ,  $NCX \bullet M = 4.6836 \times 10^{-3}$ ,  $NCX = 4.9428 \times 10^{-3}$ .

### Compiling CAREG64v6.f

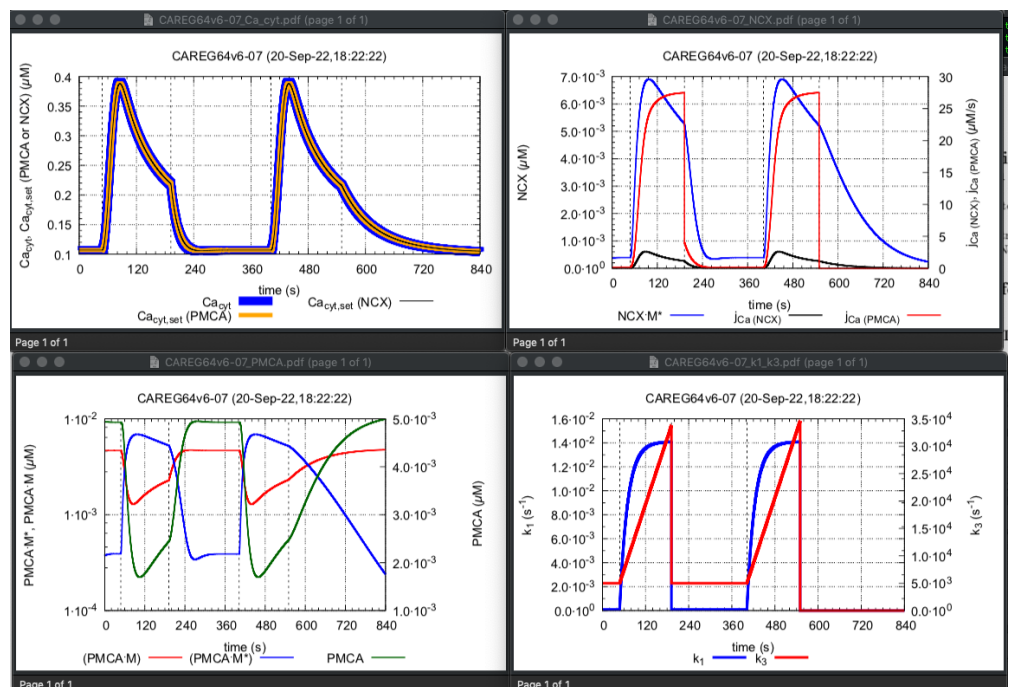
We have compiled **CAREG64v6.f** with 64 bits encoding using the Mac Terminal and Windows Cmd prompt together with Absoft's ProFortran compilers for Mac OSX and Windows:

Mac OSX: `f77 -o careg64v6 -m64 -O2 CAREG64v6.f libV77.a`  
 Windows: `f77 -o careg64v6.exe -m64 -O2 CAREG64v6.f vms.lib`  
 with the (enclosed) 64 bits libraries `libV77.a` and `vms.lib`.

## Running the model

If gnuplot and Perl are installed, on Mac OSX one runs the script file `careg64v6.sh` by using the terminal command `./careg64v6.sh`. On Windows computers you use the Cmd prompt and type `careg64v6.cmd`. The numerical output will be created, i.e. concentrations in file `CAREG64v6-07.txt` and fluxes in file `CAREG64v6-07_fluxes.txt`. `CAREG64v6-07` is our run identifier (which can be changed in the input file `CAREG64v6.INP`).

The following four graphs (Fig S2) (in pdf format) will be generated and should be opened automatically by the default pdf viewer.



**Figure S2.** Graphs created when running `careg64v6.sh` or `careg64v6.exe` with Perl and gnuplot installed.  $j_{Ca}$  (PMCA) and  $j_{Ca}$  (NCX) are the velocities by which PMCA and NCX pump calcium out of the cell.

If Perl or gnuplot are not installed you execute the provided binary files `CAREG64v6` (Mac) or `CAREG64v6-07.exe`, which will generate the concentration and flux data files `CAREG64v6-07.txt` and `CAREG64v6-07_fluxes.txt`. Graphs can then be created by a custom graphing program if it allows to plot numerical column data. The graphs in Fig S2 involve the following columns:

File `CAREG64v6-07.txt`:  
 column 1: time (s)  
 column 2:  $Ca_{cyt}$  ( $\mu M$ )  
 column 3:  $(PMCA \bullet M^*)$  ( $\mu M$ )  
 column 10:  $Ca_{cyt, set}$  (PMCA) ( $\mu M$ )

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column 11:  $\text{Ca}_{\text{cyt, set}} (\text{NCX}) (\mu\text{M})$   
column 27:  $\text{PMCA} \bullet \text{M} (\mu\text{M})$   
column 29:  $\text{PMCA} (\mu\text{M})$   
column 30:  $k_3 (1/\text{s})$   
column 31:  $k_1 (1/\text{s})$   
column 32:  $(\text{NCX} \bullet \text{M}^*) (\mu\text{M})$

File `CAREG64v6-07_fluxes.txt`:  
column 1: time (s)  
column 4:  $j_{\text{Ca}} (\text{PMCA}) (\mu\text{M}/\text{s})$   
column 14:  $j_{\text{Ca}} (\text{NCX}) (\mu\text{M}/\text{s})$