

Nuclear FRAP Model

Model definition

I propose the following parameterization for the Nuclear FRAP model.

tPB_i : time of PB i, with $1 \leq i \leq \text{number of FRAPs}$

Δt : time between consecutive data points

$n[t]$: number of nuclear molecules at time t

$c[t]$: number of cytoplasmic molecules at time t

$effVn$: nuclear effective volume, that is where the molecules can diffuse.

$effVc$: cytoplasmic effective volume.

$effVfrac = effVc / effVn$: cytoplasmic to nuclear effective volume ratio

kI : Import rate (has volume units)

kE : Export rate (has volume units)

PBn : fraction of nuclear fluorescence photobleached in each FRAP

PBc : fraction of cytoplasmic fluorescence photobleached in each FRAP

$\delta PBn1$: extra PBn of first FRAP as compared to following FRAPs

$\delta PBc1$: extra PBc of first FRAP as compared to following FRAPs

$PBfrac = PBc/PBn$: ratio of cytoplasmic to nuclear photobleaching

$n0$: number of nuclear fluorescent proteins at the start of the experiment

$c0$: number of cytoplasmic fluorescent proteins at the start of the experiment

$tot0 = n0 + c0$: total number of fluorescent proteins at the start of the experiment

n_i : number of nuclear fluorescent molecules just after the PB i

c_i : number of cytoplasmic fluorescent molecules just after the PB i

i: PB index. $i=0$ if $(t < tPB_1)$, $i=1$ if $(t \geq tPB_1 \& t < tPB_2)$, ...

j: time course index, identifies all data point acquired in the same “movie” or file.

$geomVn$: nuclear geometric volume, relevant for microscopy

$geomVc$: cytoplasmic geometric volume, relevant for microscopy

$geomVfrac = geomVc / geomVn$: cytoplasmic to nuclear geometric volume ratio

γ : proportionality constant between molecule concentration in $geomV$ and fluorescence signal, assuming perfect focus.

$focusN[t]$: function describing the nuclear focus during the experiment

$focusC[t]$: function describing the cytoplasmic focus during the experiment

$auto0n$: non photobleachable nuclear autofluorescence amount (in units of fluorescence signal)

$auto0c$: non photobleachable cytoplasmic autofluorescence amount (in units of fluorescence signal)

$auto1n$: photobleachable nuclear autofluorescence at the start of the experiment amount (in units of fluorescence signal)

$auto1c$: photobleachable cytoplasmic autofluorescence at the start of the experiment amount (in units of fluorescence signal)

$autoPBn$: fraction of nuclear photobleachable autofluorescence photobleached in each FRAP

$autoPBc$: fraction of cytoplasmic photobleachable autofluorescence photobleached in each FRAP

the “recovery” model after each FRAP is given by the ODEs

$$\left\{ \begin{array}{l} n' [t] == -kE \frac{n[t]}{effVn} + kI \frac{c[t]}{effVc} , \\ c' [t] == kE \frac{n[t]}{effVn} - kI \frac{c[t]}{effVc} , n[tPB_i] == n_i , c[tPB_i] == c_i \end{array} \right\}$$

that have analytic solution

$$n[t] = \frac{(c_i + n_i) \text{effVn } kI + (n_i \text{effVc } kE - c_i \text{effVn } kI) e^{\left(-\frac{kE}{\text{effVn}} - \frac{kI}{\text{effVc}}\right) t}}{\text{effVc } kE + \text{effVn } kI}$$

$$c[t] = c_i + n_i - n[t]$$

Assuming that at the start of the experiment the system is in steady state we have

$$n0 = \frac{\text{tot0 effVn } kI}{\text{effVc } kE + \text{effVn } kI}$$

$$c0 = \frac{\text{tot0 effVc } kE}{\text{effVc } kE + \text{effVn } kI}$$

The initial conditions after each frap are given by

$$n[tPB_i] = \begin{cases} n[tPB_i - \Delta t] (1 - PB_n - \text{deltaPBn1}) & i = 1 \\ n[tPB_i - \Delta t] (1 - PB_n) & i > 1 \end{cases}$$

$$c[tPB_i] = \begin{cases} c[tPB_i - \Delta t] (1 - PB_c - \text{deltaPBc1}) & i = 1 \\ c[tPB_i - \Delta t] (1 - PB_c) & i > 1 \end{cases}$$

Note that we allow for the first PB to produce a larger drop in fluorescence than the subsequent ones, as this is what we observe experimentally. The model has to be solved in chunks of time. (we could get an analytical expression, but it would be quite messy).

$$\text{if } t < tPB_1$$

$$n[t] = \frac{\text{tot0 effVn } kI}{\text{effVc } kE + \text{effVn } kI}$$

$$c[t] = \frac{\text{tot0 effVc } kE}{\text{effVc } kE + \text{effVn } kI}$$

$$\text{if } t \geq tPB_i \ \& \ t < tPB_{i+1} \left\{ \begin{array}{l} n_i = \begin{cases} n[tPB_i - \Delta t] (1 - PB_n - \text{deltaPBn1}) & i = 1 \\ n[tPB_i - \Delta t] (1 - PB_n) & i > 1 \end{cases} ; \\ c_i = \begin{cases} c[tPB_i - \Delta t] (1 - PB_c - \text{deltaPBc1}) & i = 1 \\ c[tPB_i - \Delta t] (1 - PB_c) & i > 1 \end{cases} ; \\ n[t] = \left((c_i + n_i) \text{effVn } kI + (n_i \text{effVc } kE - c_i \text{effVn } kI) e^{\left(-\frac{kE}{\text{effVn}} - \frac{kI}{\text{effVc}}\right) t} \right) / \\ (\text{effVc } kE + \text{effVn } kI) ; \\ c[t] = c_i + n_i - n[t] ; \end{array} \right\}$$

A second part of the model relates to the “microscopy effects”. The observed fluorescence is proportional to the concentration of molecules in the “geometrical volume”, not the “effective volume”. (I’m assuming the fluorescence signal has been corrected for imaging PB). Taking into account the autofluorescence and the focus, the expression for the fluorescence level as a function of $n[t]$ and $c[t]$ is given by

$$Fn[t] = \text{focusN}[t] \left(\gamma \frac{n[t]}{\text{geomVn}} + \text{auto1n} (1 - \text{autoPBn})^i + \text{auto0n} \right)$$

$$Fc[t] = \text{focusC}[t] \left(\gamma \frac{c[t]}{\text{geomVc}} + \text{auto1c} (1 - \text{autoPBc})^i + \text{auto0c} \right)$$

To avoid identifiability issues, I adimensionalized the equations to get rid of some parameters. First I define the adimensional rate parameters by dividing by the cytoplasmic effective volume

$$\kappa I = \frac{kI}{\text{effVc}} \Rightarrow kI = \kappa I \text{effVc}$$

$$\kappa E = \frac{kE}{\text{effVc}} \Rightarrow kE = \kappa E \text{effVc}$$

Using this in the equation for $n[t]$

$$n[t] = \frac{(c_i + n_i) \kappa I + (n_i \kappa E \text{effVfrac} - c_i \kappa I) e^{(-\kappa E \text{effVfrac} - \kappa I) t}}{\kappa E \text{effVfrac} + \kappa I};$$

where

$$\text{effVfrac} = \frac{\text{effVc}}{\text{effVn}}$$

for the focus we have that $0 \leq \text{focusN} \leq 1$ and $0 \leq \text{focusC} \leq 1$, but because we can't know when we are in perfect focus, I define the relative focus by dividing by the nuclear focus at time zero

$$\phi n[t] = \frac{\text{focusN}[t]}{\text{focusN}[0]}$$

$$\phi c[t] = \frac{\text{focusC}[t]}{\text{focusN}[0]}$$

$$Fn[t] = \phi n[t] \text{focusN}[0] \left(\gamma \frac{n[t]}{\text{geomVn}} + \text{auto1n} (1 - \text{autoPBn})^i + \text{auto0n} \right)$$

$$Fc[t] = \phi c[t] \text{focusN}[0] \left(\gamma \frac{c[t]}{\text{geomVc}} + \text{auto1c} (1 - \text{autoPBc})^i + \text{auto0c} \right)$$

Second, I adimensionalize the fluorescence variables and parameters, or really I just express the amounts in terms of nuclear fluorescence in the nuclear focus of time zero.

$$\eta[t] = \gamma \text{focusN}[0] \frac{n[t]}{\text{geomVn}} \Rightarrow n[t] = \eta[t] \frac{\text{geomVn}}{\gamma \text{focusN}[0]}$$

$$\xi[t] = \gamma \text{focusN}[0] \frac{c[t]}{\text{geomVn}} \Rightarrow c[t] = \xi[t] \frac{\text{geomVn}}{\gamma \text{focusN}[0]}$$

$$T0 = \gamma \text{focusN}[0] \frac{\text{tot0}}{\text{geomVn}}$$

$$\alpha 1n = \text{focusN}[0] \text{auto1n}$$

$$\alpha 1c = \text{focusN}[0] \text{auto1c}$$

$$\alpha 0n = \text{focusN}[0] \text{auto0n}$$

$$\alpha 0c = \text{focusN}[0] \text{auto0c}$$

plugging this into the fluorescence equations we get

$$Fn[t] = \phi n[t] \left(\eta[t] + \alpha 1n (1 - \text{autoPBn})^i + \alpha 0n \right)$$

$$Fc[t] = \phi c[t] \left(\frac{\xi[t]}{\text{geomVfrac}} + \alpha 1c (1 - \text{autoPBc})^i + \alpha 0c \right)$$

where

$$\text{geomVfrac} = \frac{\text{geomVc}}{\text{geomVn}}$$

I'll assume that the focus for the nucleus and the cytoplasm is constant during a movie. Otherwise, there are identifiable problems between the focus drift parameters and the kinetic parameters of the model.

$$\phi_n[t] = \phi_{n_j}$$

$$\phi_c[t] = \phi_{c_j}$$

where $t \in \text{movie}_j$. I will use second order polynomials evaluated at the time of the PB to model the fluctuations in the focus.

Noting that $\phi_n[0]=1$ we have

$$\phi_{n_j} = 1 + \alpha_{n1} (t_{PB_j} - t_{PB_1}) + \alpha_{n2} (t_{PB_j} - t_{PB_1})^2$$

$$\phi_{c_j} = \alpha_{c0} + \alpha_{c1} (t_{PB_j} - t_{PB_1}) + \alpha_{c2} (t_{PB_j} - t_{PB_1})^2$$

So in summary, noting that the equations for n and c are linear and therefore don't depend on the units they are expressed in, we have

$$\begin{aligned} & \text{if } t < t_{PB_1} \\ & \quad \eta[t] = \frac{T_0 \kappa I}{\kappa E \text{effVfrac} + \kappa I} \\ & \quad \xi[t] = \frac{T_0 \kappa E \text{effVfrac}}{\kappa E \text{effVfrac} + \kappa I} \\ & \text{if } t \geq t_{PB_i} \ \& \ t < t_{PB_{i+1}} \left\{ \right. \\ & \quad \eta_i = \begin{cases} \eta[t_{PB_i} - \Delta t] (1 - \text{PBn} - \text{deltaPBn1}) & i = 1 \\ \eta[t_{PB_i} - \Delta t] (1 - \text{PBn}) & i > 1 \end{cases} ; \\ & \quad \xi_i = \begin{cases} \xi[t_{PB_i} - \Delta t] (1 - \text{PBc} - \text{deltaPBc1}) & i = 1 \\ \xi[t_{PB_i} - \Delta t] (1 - \text{PBc}) & i > 1 \end{cases} ; \\ & \quad \eta[t] = \frac{(\xi_i + \eta_i) \kappa I + (\eta_i \kappa E \text{effVfrac} - \xi_i \kappa I) e^{(-\kappa E \text{effVfrac} - \kappa I) t}}{\kappa E \text{effVfrac} + \kappa I} ; \\ & \quad \xi[t] = \xi_i + \eta_i - \eta[t] ; \left. \right\} \\ & \text{Fn}[t] = \phi_{n_j} (\eta[t] + \alpha_{1n} (1 - \text{autoPNn})^i + \alpha_{0n}) \\ & \text{Fc}[t] = \phi_{c_j} \left(\frac{\xi[t]}{\text{geomVfrac}} + \alpha_{1c} (1 - \text{autoPBc})^i + \alpha_{0c} \right) \\ & \phi_{n_j} = 1 + \alpha_{n1} (t_{PB_j} - t_{PB_1}) + \alpha_{n2} (t_{PB_j} - t_{PB_1})^2 \\ & \phi_{c_j} = \alpha_{c0} + \alpha_{c1} (t_{PB_j} - t_{PB_1}) + \alpha_{c2} (t_{PB_j} - t_{PB_1})^2 \end{aligned}$$

So the free parameters I have to use in the fit are

- 1) $\kappa E \text{effVfrac}$ = combined $\kappa E \times \text{effVfrac}$ to avoid identifiable issues
- 2) κI
- 3) T_0
- 4) PBn
- 5) PBc (or PBfrac)
- 6) deltaPBn1
- 7) deltaPBc1
- 8) geomVfrac
- 9) α_{1n}
- 10) autoPNn
- 11) α_{0n}
- 12) α_{n1}
- 13) α_{n2}
- 14) α_{1c}
- 15) autoPBc

- 16) $\alpha 0c$
- 17) $ac0$
- 18) $ac1$
- 19) $ac2$

To be compatible with previous notation, I go back to latin letters, BUT KEEPING THE INTERPRETATION OF THE GREEK LETTERS!

$n \equiv \eta$
 $c \equiv \zeta$
 $tot0 \equiv T0$
 $kI \equiv \kappa I$
 $kE \equiv \kappa E$
 $Vfrac \equiv effVfrac$
 $kEVfrac \equiv \kappa EeffVfrac$
 $auto1n \equiv \alpha 1n$
 $auto0n \equiv \alpha 0n$
 $auto1c \equiv \alpha 1c$
 $auto0c \equiv \alpha 0c$

so we get the following equations (assuming $\delta PBc1 = \delta PBn1 \cdot PBfrac$)

$$\text{if } t < tPB_1$$

$$n[t] = \frac{tot0 \cdot kI}{kEVfrac + kI}$$

$$c[t] = \frac{tot0 \cdot kEVfrac}{kEVfrac + kI}$$

$$\text{if } t \geq tPB_i \text{ \& } t < tPB_{i+1} \left\{ \right.$$

$$n_i = \begin{cases} n[tPB_i - \Delta t] (1 - PBn - \delta PBn1) & i = 1 \\ n[tPB_i - \Delta t] (1 - PBn) & i > 1 \end{cases} ;$$

$$c_i = \begin{cases} c[tPB_i - \Delta t] (1 - (PBn + \delta PBn1) \cdot PBfrac) & i = 1 \\ c[tPB_i - \Delta t] (1 - PBn \cdot PBfrac) & i > 1 \end{cases} ;$$

$$n[t] = \frac{(c_i + n_i) \cdot kI + (n_i \cdot kEVfrac - c_i \cdot kI) \cdot e^{(-kEVfrac - kI) \cdot t}}{kEVfrac + kI} ;$$

$$c[t] = c_i + n_i - n[t] ; \left. \right\}$$

$$Fn[t] = focusN_j \left(n[t] + auto1n (1 - autoPNn)^i + auto0n \right)$$

$$Fc[t] = focusC_j \left(\frac{c[t]}{geomVfrac} + auto1c (1 - autoPBc)^i + auto0c \right)$$

$$focusN_j = 1 + an1 (tPB_j - tPB_1) + an2 (tPB_j - tPB_1)^2$$

$$focusC_j = ac0 + ac1 (tPB_j - tPB_1) + ac2 (tPB_j - tPB_1)^2$$

With free parameters for the fit

- 1) $kEVfrac$
- 2) kI
- 3) $tot0$
- 4) PBn
- 5) $PBfrac$
- 6) $\delta PBn1$
- 7) $geomVfrac$

- 8) auto1n
- 9) autoPBn
- 10) auto0n
- 11) an1
- 12) an2
- 13) auto1c
- 14) autoPBc
- 15) auto0c
- 16) ac0
- 17) ac1
- 18) ac2

For example, if we assume

PBfrac = 0, no cytoplasmic PB
 deltaPBn1 = 0, no extra PB in first FRAP
 autoPBn = 1, all photobleachable nuclear autofluorescence PB in first FRAP
 auto0n = 0, no non-photobleachable nuclear autofluorescence
 auto1c = 0, no photobleachable cytoplasmic autofluorescence
 autoPBc = 0, no cytoplasmic PB
 ac0 = 1, initial cytoplasmic focus equivalent to initial nuclear focus
 ac1 = 0, no cytoplasmic focus drift
 ac2 = 0, no cytoplasmic focus drift

resulting in 9 free parameters for the fit

- 1) kEVfrac
- 2) kI
- 3) tot0
- 4) PBn
- 5) geomVfrac
- 6) auto1n
- 7) an1
- 8) an2
- 9) auto0c

Model Fit

The parameters kEVfrac, kI and geomVfrac are scanned in log scale. Therefore I define the parameters

$\log_{10_kEVfrac} = \log_{10}(\text{kEVfrac})$
 $\log_{10_kI} = \log_{10}(\text{kI})$
 $\log_{2_geomVfrac} = \log_2(\text{geomVfrac})$

which are scanned lineally.

geomVfrac is not well defined by the fluorescence data alone. I has identifiability problems with tot0. Chanching the value of geomVfrac influences the value of kI and kEVfrac, as it changes the nuclear fraction. Therefore, we decided to obtain an independent estimate of geomVfrac, based on the cell's images. This can be done for each cell, or use a distribution for the population. We implemented a "ridge regression", where the cost function penalizes departures from the estimated geomVfrac. To do this, we define two new parameters.

$\log_{2_geomVfrac_mean} = \text{best estimation for } \log_{2_geomVfrac}$
 $\log_{2_geomVfrac_sd} = \text{standard deviation for the estimation of } \log_{2_geomVfrac}$

Note that the distribution of geomVfrac is assumed to be lognormal. Experimental data supports this assumption. The cost function is defined basically as in Cedersund and Roll 2008

$$\mathbf{v}(\mathbf{p}) = \sum_i \sum_j \frac{(\mathbf{y}_i[\mathbf{t}_j] - \hat{\mathbf{y}}_i^M[\mathbf{t}_j])^2}{\sigma_i^2[\mathbf{t}_j]} + \sum_k \alpha_k h_{\text{pen}}[\mathbf{p}_k - \mathbf{p}_k^g]$$

where $h_{\text{pen}}[x] = x^2$ and $\alpha_k = \frac{1}{\sigma_k^2}$

In this case there is only one variable over which to penalize ($k=1$), namely log2_geomVfrac . So we have

$$\mathbf{p}_1^g = \text{log2_geomVfrac_mean}$$

$$\alpha_1 = \frac{1}{\sigma_1^2} = \frac{1}{(\text{log2_geomVfrac_sd})^2}$$

the cost function is therefore

$$\mathbf{v}(\mathbf{p}) = \sum_i \sum_j \frac{(\mathbf{y}_i[\mathbf{t}_j] - \hat{\mathbf{y}}_i^M[\mathbf{t}_j])^2}{\sigma_i^2[\mathbf{t}_j]} + \frac{(\text{log2_geomVfrac} - \text{log2_geomvfrac_mean})^2}{(\text{log2_geomVfrac_sd})^2}$$

this statistic is assumed to follow a chi square distribution.