Mathematical models of protein trafficking in neurons

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Paul C Bressloff Mathematical models of protein trafficking in neurons



- Proteins must be transported long distances along dendrites and axons
- Protein transport crucial in synapse formation and plasticity
- Certain neurodegenerative diseases involve a breakdown in protein transport

Protein receptor trafficking and synaptic plasticity

- Cable theory of receptor trafficking
- Motor transport of mRNA

Dendritic spines



- Most excitatory synapses are located in small mushroom–like appendages called dendritic spines which are rich in actin.
- Associated neurotransmitter is glutamate (Glu).

Glutamate (Glu) receptors



- Two major types of Glu receptor: AMPA and NMDA
- Binding of Glu receptors opens AMPA ion channels leading to depolarization
- Binding of Glu + sufficient depolarization opens NMDA ion channels leading to influx of calcium

Long-term potentiation/depression



- High frequency (100 Hz) stimulus induces LTP
- Low frequency (1Hz) stimulus induces LTD

Separation of time-scales



- Induction: Calcium signal via NMDA induces action of kinases and phosphotases
- Expression: (De)Phosphorylation of AMPAR receptors regulates their trafficking and their conductance state
- ► Maintenance: Persistent changes require protein synthesis

Regulation of AMPAR trafficking



 A major expression mechanism for LTP/LTD is a Ca²⁺-induced change in the number of synaptic AMPARs

Various mechanisms of receptor trafficking



- 1. Newly synthesized proteins inserted in cell surface
- 2. Lateral surface diffusion along dendrite
- 3. Surface entry into spine
- 4. Recycling with intracellular pools/binding to scaffolding proteins
- 5. Motor transport of mRNA and local synthesis of AMPAR



(Choquet and Triller, Nat Rev. Neurosci. 2003)

- Sub-µm latex bead is bound to a receptor using ligands (antibodies or scaffolding proteins) and imaged using lasers
- GFP tags reveal regions of high receptor concentration coincide with confinement domains (red)

Inactivation and recovery of surface receptors



Passafaro et al (Nature 2001)

- ► Transfect AMPAR with an immunoflourescent tag (HA/T)
- Reduce temperature to temporarily stop trafficking
- Treat neuron with thrombin to eliminate surface staining of HA/T
- Washout thrombin and return to normal temperatures: surface expression of HA/T recovers in 30 min.

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- AMPA receptor
- scaffolding protein



Kinetic equations

$$a\frac{dR}{dt} = \omega_{+}U - \omega_{-}R - kR + \sigma^{rec}C - \alpha[Z - Q]R + \beta Q$$
$$a\frac{dQ}{dt} = \alpha[Z - Q]R - \beta Q$$
$$\frac{dC}{dt} = -\sigma^{rec}C - \sigma^{deg}C + kR + \sigma$$

- \blacktriangleright U is dendritic receptor concentration at spine boundary
- Z is concentration of scaffolding proteins
- ► *a* is area of synapse

Adiabatic approximation

$$Q(t) = rac{lpha Z R(t)}{lpha R(t) + eta}$$

Reduced single-spine model

$$a\frac{dR}{dt} = \omega_{+}U - \omega_{-}R - kR + \sigma^{rec}C$$
$$\frac{dC}{dt} = -\sigma^{rec}C - \sigma^{deg}C + kR + \sigma$$

Strength of a synapse

$$W = a[g_R R + g_Q Q]$$

where g_R, g_Q are conductances of free and bound AMPARs

Multi-spine diffusion-trapping model³



- Consider a dendritic cable of circumference / and length L.
- Somatic flux J_s at x = 0. Reflecting BC at x = L

³Bressloff et. al. PRE 2007, SIAM J. Appl. Math. 2008

1D Continuum model⁴

- Assume uniform concentration around circumference of cable
- Represent fluxes between spines and cable using Dirac delta functions:

$$\frac{\partial U}{\partial t} = D \frac{\partial^2 U}{\partial x^2} - \rho(x) [\omega_+ U(x,t) - \omega_- R(x,t)],$$

with

$$\rho(x) = \frac{1}{2\pi I} \sum_{j=1}^{N} \delta(x - x_j)$$

Continuum approximation: take ρ(x) to be a smooth function so that we obtain a 1D cable equation.

⁴Earnshaw and Bressloff J. Comp Neurosci 2008, Bressloff 2009



Transverse current

$$J(x,t) = \omega_+ U(x,t) - \omega_- R(x,t)$$

Basal parameter values

$$k = 10^{-3} \mu m^2 s^{-1}, \sigma^{\text{rec}} = 10^{-3} s^{-1}, \sigma^{\text{rec}} = 10^{-5} s^{-1}, D = 0.1 \mu m^2 s^{-1}$$

Steady-state cable equation

• Uniform spine density $\rho_0 = N/LI$:

$$\frac{d^2 U}{dx^2} - \frac{\rho_0 \overline{\omega}}{D} [U - \overline{R}] = 0$$

with

$$\overline{\omega} = rac{\omega_+ k(1-\Lambda)}{\omega_- + k(1-\Lambda)}, \quad \overline{R} = rac{\omega_-}{\omega_+} rac{\Lambda \sigma}{k(1-\Lambda)}, \quad \Lambda = rac{\sigma^{
m rec}}{\sigma^{
m rec} + \sigma^{
m deg}}.$$

Solution is

$$U(x) = ZJ_S \frac{\cosh(\gamma[x-L])}{\sinh(\gamma L)} + \overline{R}$$

where Z is characteristic impedance of cable and $\xi \equiv \gamma^{-1}$ is a diffusive space constant:

$$\gamma = \sqrt{\frac{\rho_0 \overline{\omega}}{D}}, \quad Z = \frac{1}{ID\gamma}.$$



- Diffusion from soma is insufficient to supply distal synapses with receptors
- Source at soma favors proximal synapses. Synaptic democracy requires non-uniform spines

 Given solution U(x) to cable equation, the free synaptic receptor concentration is

$$R(x) = rac{\omega_+ U(x) + \Lambda \sigma}{\omega_- + k(1 - \Lambda)}$$

The steady-state synaptic weight is then

$$W(x) = a(x) \left[g_R R(x) + g_Q \frac{\alpha(x) Z(x) R(x)}{\alpha(x) R(x) + \beta(x)} \right]$$

- Spine parameters can be classified according to whether or not they have a nonlocal effect on steady-state receptor numbers mediated by diffusion
- Synaptic parameters are purely local: rates of binding/unbinding, number of binding sites, area of synapse
- Parameters of constitutive recycling are nonlocal: rates of exo/endocytosis, local production and degradation
- If the expression of LTP/LTD involves changes in number of binding sites, then receptor diffusion unlikely to mediate heterosynaptic plasticity



- Increasing local production rate σ for spines between $90\mu m$ and $110\mu m$ leads to a global increase in receptor numbers.
- Increasing degradation rate σ^{deg} leads to a global reduction in receptor numbers



 Increasing rate of endocytosis k or decreasing rate of recycling σ^{rec} for spines between 90μm and 110μm leads to a global decrease in receptor numbers. Response to a time-dependent surface current source I(x, t) switched on at t = 0:

$$A(x,t) = \int_0^t \int_0^L \mathcal{G}_A(x,y;t-t')\mathcal{I}(y,t')dydt'$$

for A = U, R, C and A(x, 0) = 0.

- ► Green's function G_A(x, x₀; t) is probability density that at time t a single labeled receptor is at position x and in state A given that at t = 0 it was injected into dendritic surface at x₀
- Can be generalized to a time-dependent modification of constitutive recycling induced by synaptic activity.

▶ Plots of Green's functions for $x_0 = L/2$:



• Dependence of \mathcal{G}_U on diffusive coupling ω .



• Let $\mathcal{P}_A(t)$ denote probability to be in state A at time t:



$$\mathcal{P}_{A}(t) = \int_{0}^{L} \mathcal{G}_{A}(x, L/2; t) dx$$

"Sum-over-trips" on dendritic trees



- Each branch is uniform
- All terminal nodes are closed
- Continuity of receptor concentration at branch nodes
- Conservation of current at branch nodes

Use Laplace transforms on *j*th branch

$$rac{d^2 \widetilde{U}_j}{dx^2} - \Xi_j(s) \widetilde{U}_j(x,s) = -rac{\widetilde{\mathcal{I}}_j(x,s)}{D}$$

Perform the rescalings

$$x \to X = \gamma_j(s)x, \quad L_j \to \mathcal{L}_j(s) = \gamma_j(s)L_j$$

with $\gamma_j(s) = \sqrt{\Xi_j(s)}$ so that

$$rac{d^2\widetilde{U}_j}{dX^2} - \widetilde{U}_j(X,s) = -rac{\widetilde{\mathcal{I}}_j(X,s)}{D}$$

At each branch node

$$\widetilde{U}_i(0,s)=\widetilde{U}_j(0,s)$$

for all pairs (i, j) radiating from the node and

$$\sum_{j} z_{j}(s) \left. \frac{\partial \widetilde{U}_{j}}{\partial X} \right|_{X=0} = 0, \quad z_{j}(s) = l_{j} D \gamma_{j}(s),$$

where the sum is over all branches *j* connected to the node.At all terminal modes

$$\left.\frac{\partial \widetilde{U}_j}{\partial X}\right|_{X=\mathcal{L}_j}=0.$$

General solution is of the form

$$\widetilde{U}_i(X,s) = \sum_{j\in\mathcal{T}} \int_0^{\mathcal{L}_j(s)} G_{ij}(X,Y;s) \widetilde{\mathcal{I}}_j(Y,s) dY$$

The Green's function G_{ij}(X, Y; s) satisfies the homogeneous equation

$$rac{d^2 G_{ij}(X,Y;s)}{dX^2} - G_{ij}(X,Y;s) = -rac{1}{D}\delta_{i,j}\delta(X-Y),$$

with the same boundary conditions as $\widetilde{U}_i(X, s)$ for fixed j, Y.

▶ Using the "sum-over-trips" method⁵ it can be shown that

$$G_{ij}(X,Y;s) = \sum_{ ext{trips}} \mathcal{A}_{ ext{trip}}(s) \mathcal{G}_{\infty}(\mathcal{L}_{ ext{trip}}(i,j,X,Y,s)),$$

where $G_{\infty}(X)$ is Green's function for an infinite cable:

$$G_{\infty}(X)=\frac{\mathrm{e}^{-|X|}}{2D},$$

and $\mathcal{L}_{trip}(i, j, X, Y, s)$ is the length (in rescaled coordinates) of a path along the tree starting at the point X on branch *i* and ending at the point Y on branch *j*.

Sum restricted to a special class of paths or trips

⁵Abbott et. al. 1991; Bressloff et. al. 1996; Coombes et. al. 2007, Bressloff 2009

Definition:

A trip from (X, i) to (Y, j) may start out in either direction along branch *i* but it can subsequently change direction only at a branch or terminal node. A trip is always reflected back at a terminal node, whereas at a branch node it may be transmitted to another branch or reflected back. A trip may pass through the points (X, i)and (Y, j) an arbitrary number of times as long as it starts at (X, i) and ends at (Y, j).



For each trip, the associated amplitude ${\cal A}_{trip}$ is calculated according to the following rules:

- 1. Initially take $\mathcal{A}_{\mathrm{trip}}(s) = 1$.
- For every branch node at which the trip passes from an initial segment m to a different segment n, n ≠ m, multiply A_{trip}(s) by a factor 2p_n(s)
- 3. For every branch node at which the trip is reflected back along the same segment *m*, multiply $A_{trip}(s)$ by a factor $2p_m(s) 1$
- 4. For every closed (open) terminal node, multiply $\mathcal{A}_{\mathrm{trip}}(s)$ by a factor +1 (-1)

The factor p_m is given by

$$p_m(s)=\frac{z_m(s)}{\sum_n z_n(s)},$$

where sum is over all branches n radiating from branch node.

- 1. How is receptor trafficking **regulated** following the induction of **synaptic plasticity**?
- 2. How are **stable** synaptic memories maintained given **rapid protein turnover**? Roles of protein synthesis and actin cytoskeleton?
- 3. What is the effect of **intrinsic noise** (small receptor numbers) on receptor trafficking and plasticity?

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Protein synthesis required for persistent LTP⁶



- (A) Single train of tetanic stimulation produces decremental potentiation
- Early-phase LTP insensitive to translational inhibition (anisomycin) and transcriptional inhibition (actinomyosin-D)
- (B) Repeated stimulation induces persistent potentiation that is sensitive to both translational and transcriptional inhibition

⁶Kelleher et. al. Neuron 2004

Motor transport of mRNA⁷

- Newly transcribed mRNA granules are transported into the dendrite by kinesin motors on microtubules.
- Following synaptic activation, mRNA is localized to spines by the actin-based myosin



⁷Bramham and Wells Nat. Rev. Neurosci. 2007



- Motile mRNA particles in cultured hippocampal cells exist in 3 states: oscillatory, anterograde and retrograde
- KCL-induced depolarization increases anterograde motion



- Many problems in nature require search for a randomly hidden target: foraging animals, a protein searching for a specific target site on DNA, microtubular transport of mRNA
- An efficient stochastic search strategy is to alternate between (A) a slow motion (diffusive), intensive search phase and (B) a fast (motor-assisted) ballistic non-search phase

⁸Loverdo et. al. Nat. Phys. 2008



- Particle injected into dendrite at x = 0 and t = 0. Fixed hidden target (dendritic spine) at x = X.
- Particle exists in 3 states: anterograde with velocity v₊, retrograde with velocity -v₋, and stationary.
- In stationary state particle can be absorbed at a rate k if within a distance a of target
- Absorbing boundary at x = L

⁹Bressloff and Newby 2009

- ▶ Let p_m(x, t) denote probability density for anterograde (m = +), retrograde (m = −) and stationary (m = 0) states
- Master equation of the form

$$\begin{aligned} \frac{\partial p_{+}}{\partial t} &= -v_{+} \frac{\partial p_{+}}{\partial x} + \alpha p_{0} - \beta_{+} p_{+} \\ \frac{\partial p_{-}}{\partial t} &= v_{-} \frac{\partial p_{-}}{\partial x} + \alpha p_{0} - \beta_{-} p_{-} \\ \frac{\partial p_{0}}{\partial t} &= -2\alpha p_{0} + (\beta_{+} p_{+} + \beta_{-} p_{-}) - k\chi(x - X)p_{0} \end{aligned}$$

where $\chi(x) = 1$ if |x| < a and is zero otherwise.

Initial and boundary conditions:

$$v_{-}p_{-}(0,t) = v_{+}p_{+}(0,t), \quad p_{-}(L,t) = 0, \quad p_{m}(x,0) = \delta(x)\delta_{m,+}$$

Bias in anterograde direction

$$\beta_+/v_+ < \beta_-/v_-$$

Probability of finding target after time t is

$$\gamma(t) = k \int_{t}^{\infty} \int_{X-a}^{X+a} p_0(x,\tau) dx d\tau$$

 Define hitting probability Π and conditional MFPT T according to

$$\Pi = \gamma(0), \quad T = \int_0^\infty \frac{\gamma(t)}{\gamma(0)} dt$$

- Determine hitting probability Π and MFPT T by solving backwards equation or using Laplace transforms
- Optimization problem: find parameter values that minimize search time and maximize hitting probability?

Plots of Π and T for β₋ = 1s⁻¹ (solid black), β₋ = 2.5s⁻¹ (dashed) and unidirectional (gray)



- Determine minimal MFPT for a given hitting probability
- For finite β₋ (partially biased) there exists a minimum of T as a function of α
- Unidirectional transport gives smaller MFPT than bidirectional transport





- Unidirectional search and n + 1 identical targets
- Hitting probability of finding most downstream target is

$$\widehat{\Pi} = (1 - \Pi)^n \Pi$$

• $\widehat{\Pi}$ is maximized when

$$\beta_{max}(\alpha) = \frac{\ln\left(1+\frac{1}{n}\right)v}{2ka}(\alpha+k).$$

The maximum hitting probability is

$$\widehat{\Pi}_{max} = \frac{n^n}{(n+1)^{n+1}}.$$



Quasi-steady-state approximation

- Fix units of space and time: x = 1mm and $t = 10^4 s$
- Then $v_{\pm} = \mathcal{O}(1)$ and transition rates $lpha, eta_{\pm} \gg 1$
- Introduce small parameter

$$\varepsilon = \frac{1}{\beta_-} + \frac{1}{\beta_+} + \frac{1}{\alpha}$$

and set $\mathbf{a} = \varepsilon \alpha, \mathbf{b}_{\pm} = \varepsilon \beta_{\pm}$

Have linear reaction-hyperbolic system

$$\begin{aligned} (\partial_t + v_+ \partial_x)p_+ &= \frac{1}{\epsilon}(-b_+p_+ + ap_0)\\ (\partial_t - v_- \partial_x)p_- &= \frac{1}{\epsilon}(-b_-p_- + ap_0)\\ (\partial_t + k(x))p_0 &= \frac{1}{\epsilon}(b_+p_+ + b_-p_- - 2ap_0) \end{aligned}$$

Using projection methods one can show that

$$p_0(x,t)\sim rac{q(x,t)}{a}, \quad p_\pm(x,t)\sim rac{q(x,t)}{b_\pm}$$

where q satisfies advection-diffusion equation

$$\frac{\partial q}{\partial t} = -\frac{k(x)}{a}q - v_{\text{eff}}\frac{\partial q}{\partial x} + D_{\text{eff}}\frac{\partial^2 q}{\partial x^2}$$

where

$$v_{
m eff}=\gamma_{-}\equivrac{1}{b_{+}}-rac{1}{b_{-}}$$

$$D_{ ext{eff}} = \epsilon \left(rac{1-\gamma_-}{b_+^2} + rac{1+\gamma_-}{b_-^2}
ight).$$

• If $k \equiv 0$ then one can write the solution in the form

$$p_m(x,t) = \eta_m Q_{\varepsilon}\left(\frac{x-vt}{\sqrt{\varepsilon}},t\right)$$

Can prove that¹⁰

$$Q_arepsilon\left(s,t
ight)
ightarrow q(s,t)$$
 as $arepsilon
ightarrow \infty$

where q(s, t) is solution of heat equation.

¹⁰cf. Reed et. al. 1990, Friedman and Hu 2007

- AMPAR trafficking: Berton Earnshaw (Utah), Michael Ward (UBC)
- mRNA transport: Jay Newby (Utah)
- Movies: Mike Ehlers (Duke University)
- National Science Foundation



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