

## A Simple Model of Transmitter Release and Facilitation

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We describe a model of synaptic transmitter release and presynaptic facilitation that is based on activation of release sites by single  $\text{Ca}^{2+}$  microdomains. Facilitation is due to  $\text{Ca}^{2+}$  that remains bound to release sites between impulses. This model is inherently stochastic, but deterministic equations can be derived for the mean release. The number of equations required to describe the mean release is small, so it is practical to use the model with models of neuronal electrical activity to investigate the effects of different input spike patterns on presynaptic facilitation. We use it in conjunction with a model of dopamine-secreting neurons of the basal ganglia to demonstrate that transmitter release is greater when the neuron bursts than when it spikes continuously, due to the greater facilitation generated by the bursting impulse pattern. Finally, a minimal form of the model is described that is coupled to simple models of postsynaptic receptors and passive membrane to compute the postsynaptic voltage response to a train of presynaptic stimuli. This form of the model is appropriate for neural network simulations.

### 1 Introduction ---

Several models have been proposed for synaptic transmitter release and presynaptic facilitation, the process whereby impulse-evoked release is enhanced for up to several hundred milliseconds if preceded by one or more conditioning impulses (Magleby, 1987). In the models of Zucker and Fogelson (1986) and Yamada and Zucker (1992), release sites and  $\text{Ca}^{2+}$  channels are spatially separated, and facilitation is due primarily to residual free  $\text{Ca}^{2+}$  left over from previous impulse-induced  $\text{Ca}^{2+}$  channel openings, so the diffusion of free  $\text{Ca}^{2+}$  is important. In an alternative model (Stanley, 1986; Bertram, Sherman, & Stanley, 1996), each transmitter release site is colocalized with a single  $\text{Ca}^{2+}$  channel. Release is then activated by the microdomain of  $\text{Ca}^{2+}$  surrounding the channel mouth, eliminating the need to perform  $\text{Ca}^{2+}$  diffusion calculations. This model is intended to describe release evoked by short depolarizations, such as action potentials, during which the opening of multiple  $\text{Ca}^{2+}$  channels in the neighborhood of a release site is unlikely.

In the single-domain model, each release site has four independent  $\text{Ca}^{2+}$  binding sites or gates, each of which must be bound for release to occur. Kinetics of the gates are graded from slow unbinding, high affinity to fast unbinding, low affinity. Facilitation is due to the accumulation of  $\text{Ca}^{2+}$  bound to one or more of the gates rather than to residual free  $\text{Ca}^{2+}$ . This model was motivated by the findings that release sites can be activated by the  $\text{Ca}^{2+}$  microdomain under a single channel (Stanley, 1993) and that there is a step-like frequency dependence of facilitation associated with a steplike decline of  $\text{Ca}^{2+}$  cooperativity (Stanley, 1986). There is also considerable evidence that residual free  $\text{Ca}^{2+}$  does not play an important role in facilitation (Stanley, 1986; Blundon, Wright, Brodwick, & Bittner, 1993; Winslow, Duffy, & Charlton, 1994), although it does appear to play a role in other forms of enhancement that last for several seconds (augmentation) or several minutes (posttetanic potentiation) following a high-frequency conditioning tetanus (Delaney, Zucker, & Tank, 1989; Delaney & Tank, 1994).

## 2 The Mathematical Model

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The binding of  $\text{Ca}^{2+}$  to the  $j$ th gate obeys a first-order kinetic scheme:



where  $B_j$  and  $U_j$  are the normalized ( $B_j + U_j = 1$ ) concentrations or probabilities of bound and unbound gates and  $\text{Ca}$  is the concentration of free  $\text{Ca}^{2+}$  at the release site. The binding rates ( $k_j^+$ , in  $\text{msec}^{-1}\mu\text{M}^{-1}$ ) and unbinding rates ( $k_j^-$ , in  $\text{msec}^{-1}$ ) were chosen to produce steps in the facilitation curve:  $k_1^+ = 3.75 \times 10^{-3}$ ,  $k_1^- = 4 \times 10^{-4}$ ,  $k_2^+ = 2.5 \times 10^{-3}$ ,  $k_2^- = 1 \times 10^{-3}$ ,  $k_3^+ = 5 \times 10^{-4}$ ,  $k_3^- = 0.1$ ,  $k_4^+ = 7.5 \times 10^{-3}$ ,  $k_4^- = 10$ . The  $k_j^-$  vary from small to large, resulting in a wide range of unbinding time constants ( $1/k_j^-$ ): 2.5 sec, 1 sec, 10 msec, and 0.1 msec, respectively. The temporal evolution of  $B_j$  is described by

$$\frac{dB_j}{dt} = -(k_j^- + k_j^+ \text{Ca}) B_j + k_j^+ \text{Ca}, \quad j = 1, 2, 3, 4. \quad (2.2)$$

The opening and closing of a  $\text{Ca}^{2+}$  channel is a stochastic process, with transition rates that depend on membrane voltage ( $V$ ). The quantity  $\text{Ca}$  is therefore a random variable. In Bertram et al. (1996), a Monte Carlo procedure was used to handle the effects of stochastic channel activity on release. Although effective, this approach is computationally demanding, and a better approach was subsequently developed (Bertram & Sherman, in press). Here, Fokker-Planck-type partial differential equations were derived for the probability distributions of bound gates under open and closed

$\text{Ca}^{2+}$  channels. Ordinary differential equations for the expected values of these distributions—that is, the mean concentrations of bound gates of each type—were then derived. Although the four gates are physically independent, they are statistically correlated through their common dependence on the domain  $\text{Ca}^{2+}$  concentration. Therefore, expected or mean release is not simply the product of the mean bound concentrations of each gate type; it depends on the mean concentrations of all gate configurations, resulting in a system of 30 differential equations. However, it was shown that the correlation among gates is removed and the number of equations necessary to compute release reduced to 4 if it is assumed that the gates respond to the *average* domain  $\text{Ca}^{2+}$  concentration—the domain  $\text{Ca}^{2+}$  concentration averaged over the entire population of  $\text{Ca}^{2+}$  channels. Using perturbation analysis, it was shown that the error introduced by this approximation is typically small.

Taking the expected values of both sides of equation 2.2, and replacing  $E[\text{Ca } B_j]$  by  $E[\text{Ca}] E[B_j]$  (the average domain  $\text{Ca}^{2+}$  approximation), we get

$$\frac{d\sigma_j}{dt} = -(k_j^- + k_j^+ \overline{\text{Ca}}) \sigma_j + k_j^+ \overline{\text{Ca}}, \quad j = 1, 2, 3, 4, \quad (2.3)$$

where  $\sigma_j$  is the expected value of  $B_j$  and  $\overline{\text{Ca}} = E[\text{Ca}]$  is the average domain  $\text{Ca}^{2+}$  concentration, equal to the product of the fraction of open channels ( $m$ ) and the domain  $\text{Ca}^{2+}$  concentration at an open channel ( $\text{Ca}_{\text{open}}$ ). The latter is assumed to be proportional to the  $\text{Ca}^{2+}$  influx through an open channel ( $i(V)$ ):  $\text{Ca}_{\text{open}} = -A i(V)$ . Assuming that the  $\text{Ca}^{2+}$  channel has a single open and a single closed state, the fraction of open channels satisfies

$$\frac{dm}{dt} = \alpha_m (1 - m) - \beta_m m \quad (2.4)$$

where  $\alpha_m, \beta_m$  are the channel opening and closing rates, respectively. Expected release is then

$$E[R] = \sigma_1 \sigma_2 \sigma_3 \sigma_4. \quad (2.5)$$

The fourth gate, unlike the others, has fast unbinding kinetics, so that  $\sigma_4$  changes rapidly with changes in  $\overline{\text{Ca}}$ . Thus, the number of differential equations needed to compute  $E[R]$  is further reduced by one by assuming that  $\sigma_4$  is in equilibrium with  $\overline{\text{Ca}}$ .

### 3 Example: The Dopamine Neuron

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Our release model can be used with any model of membrane voltage or even with experimental voltage recordings. As an example, we use it in

conjunction with a model of in vitro electrical activity of the dopamine neuron of the basal ganglia (Li, Bertram, & Rinzel, 1996). A periodic bursting pattern, high-frequency spiking followed by a long period of silence, is produced (see Fig. 1B) in the presence of the glutamate agonist N-methyl-D-aspartate (NMDA). In the absence of NMDA, the cell spikes continuously at a lower frequency (see Fig. 1A). Mean release evoked by the continuous spiking pattern shows some facilitation (see Fig. 1E), due to the slow growth of  $\sigma_1$  and  $\sigma_2$  (see Fig. 1C). However, facilitation generated by the bursting pattern is much greater (see Fig. 1F), consistent with in vivo experimental findings (Gonon, 1988). In this case  $\sigma_3$  increases during the first portion of the active phase, while  $\sigma_1$  and  $\sigma_2$  increase throughout the active phase (see Fig. 1D), facilitating release within an active phase. Between active phases  $\sigma_3$  decays back to its baseline value, but the burst period is short enough (800 msec) that the active-phase increases in  $\sigma_1$  and  $\sigma_2$  are not entirely lost. Therefore,  $\sigma_1$  and  $\sigma_2$  accumulate over several bursts, and release evoked by later bursts is greater than that evoked by earlier ones.

#### 4 A Minimal Model

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To demonstrate how the transmitter release model can be used in neural network simulations, it is simplified to a form that can be used with integrate-and-fire neurons. This model is coupled to simple models of postsynaptic transmitter receptors and passive membrane to generate a postsynaptic voltage response. Following Destexhe, Mainen, & Sejnowski (1994a) we assume that each impulse leads to the release of a square pulse of transmitter of 1 msec duration. In contrast to Destexhe et al., we assume that the size of the transmitter pulse evoked by each stimulus is not constant. The magnitude of the initial pulse is  $T^{(1)}$ , while the magnitude of each subsequent pulse is the product of  $T^{(1)}$  and the presynaptic facilitation. Facilitation is computed by solving equation 2.3 for  $\sigma_1$ ,  $\sigma_2$ , and  $\sigma_3$  (or just one of the three for the simplest model that exhibits facilitation) assuming that each presynaptic stimulus leads to a square pulse of  $\overline{Ca}$  lasting 1 msec. (With 10 mM external  $Ca^{2+}$ ,  $\overline{Ca}$  summed over an impulse is  $63 \mu M$ , using the Hodgkin-Huxley equations [Hodgkin & Huxley, 1952] to generate the impulse. This value may be used as the magnitude of the  $\overline{Ca}$  pulse, scaled linearly to reflect any changes in the external  $Ca^{2+}$  concentration.) Facilitation contributed by the  $j$ th gate is then the ratio of  $\sigma_j$  at the end of the  $n$ th pulse to the value at the end of the first pulse.

We assume that the facilitating transmitter pulses act on a simple two-state postsynaptic receptor (Destexhe et al., 1994a), with the fraction of bound receptors ( $a$ ) given by

$$\frac{da}{dt} = \alpha_a T(1 - a) - \beta_a a, \quad (4.1)$$

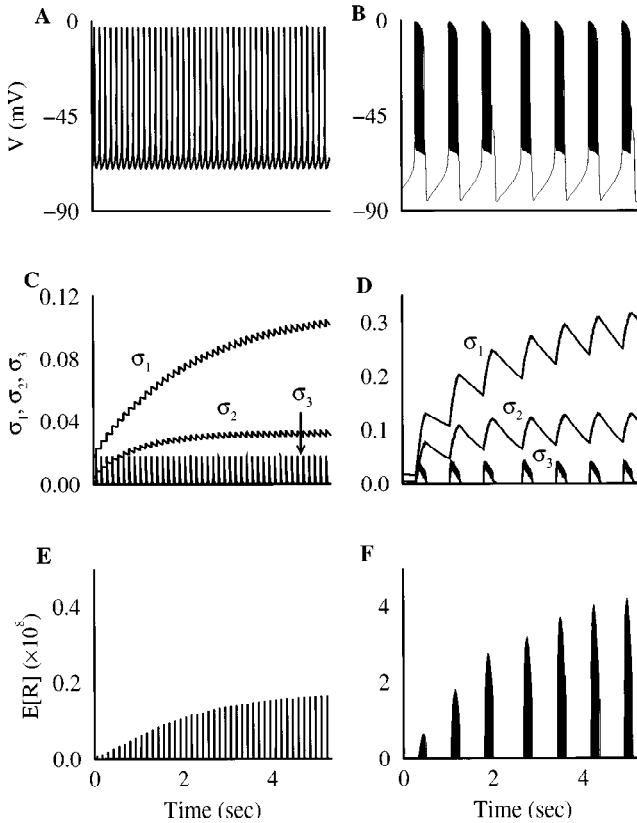


Figure 1: Expected release (see equations 2.3–2.5) evoked by a dopamine neuron model. In the presence of NMDA, the cell generates bursts of action potentials (B); otherwise, it spikes continuously (A). Release is much greater when the cell bursts (E, F; notice the different scales) due to the greater accumulation of bound gates (C, D;  $\sigma_3$  has been multiplied by 30). The single-channel  $\text{Ca}^{2+}$  current is given by a Goldman-Hodgkin-Katz expression,  $i(V) = 1.45V[\text{Ca}_{\text{ex}}/(1 - \exp(0.07V))]$ , where  $\text{Ca}_{\text{ex}} = 1 \text{ mM}$  is the external  $\text{Ca}^{2+}$  concentration. We use  $A = 0.1 \mu\text{M fA}^{-1}$  and  $\alpha_m = 3.1 e^{V/10}$ ,  $\beta_m = e^{-V/26.7}$  to compute  $\overline{\text{Ca}}$ . Both  $\alpha_m$  and  $\beta_m$  are based on squid giant synapse data (Llinás, Steinberg, & Walton, 1981) modified for the higher temperature at which dopamine neurons typically operate ( $37^\circ\text{F}$ ).

where  $T$  is the transmitter concentration and  $\alpha_a$  and  $\beta_a$  are the binding and unbinding rates, respectively. Postsynaptic membrane voltage ( $V_{\text{post}}$ ) is computed with the passive membrane equation

$$\frac{dV_{\text{post}}}{dt} = -(I_{\text{mem}} + I_{\text{syn}}) / C_{\text{mem}} \quad (4.2)$$

where  $C_{\text{mem}}$  is the membrane capacitance,  $I_{\text{mem}} = \bar{g}_{\text{mem}} (V - V_{\text{mem}})$  is the passive membrane current, and  $I_{\text{syn}} = \bar{g}_{\text{syn}} a (V - V_{\text{syn}})$  is the synaptic current. Note that since both  $\bar{Ca}$  and  $T$  appear as square pulses in equations 2.3 and 4.1, the equations have piecewise exponential solutions (see Destexhe et al., 1994a).

In Figure 2 this minimal model is used to compute the postsynaptic voltage response to a short 100-Hz stimulus train. The synapse facilitates throughout the train, producing consistently larger transmitter pulses (see Fig. 2A) and postsynaptic voltage depolarizations (see Fig. 2B). Combined with the relatively slow membrane dynamics, this leads to a substantial rise in the average postsynaptic voltage (see Fig. 2B, solid curve). In contrast, when facilitation is excluded by assuming that the size of each transmitter pulse is the same, the average postsynaptic voltage exhibits only a small rise, associated with the slow membrane dynamics (see Fig. 2B, dashed curve).

## 5 Discussion

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Computational models of synaptic transmission typically concentrate on postsynaptic mechanisms, assuming explicitly or implicitly that the amount of neurotransmitter released by each impulse is the same (Rall, 1967; Destexhe et al., 1994a). However, a key feature of the transmission process is that the number of transmitter molecules released by an impulse depends on the history of presynaptic electrical activity. The model discussed here is one of several intended to describe presynaptic transmitter release and short-term presynaptic enhancement (Zucker & Fogelson, 1986; Parnas, Dudel, & Parnebas, 1986; Yamada & Zucker, 1992). In addition to the differences discussed earlier, these other release models differ from the present one in the structure of their release sites. In Zucker and Fogelson (1986), it was assumed that the kinetics of  $\text{Ca}^{2+}$  binding to a release site are instantaneous, so that the rate of release is proportional to some power of the  $\text{Ca}^{2+}$  concentration at the release site. This model was later modified to include finite kinetic rates (Yamada & Zucker, 1992), but because the modified model includes only one slow  $\text{Ca}^{2+}$  binding site, facilitation depends largely on residual-free  $\text{Ca}^{2+}$ , contrary to a growing body of experimental data (Stanley, 1986; Blundon et al., 1993; Winslow et al., 1994). In addition, this model does not account for the multiple time scales of facilitation observed experimentally (Stanley, 1986; Magleby, 1987). Finally, the model by Parnas et al. (1986)

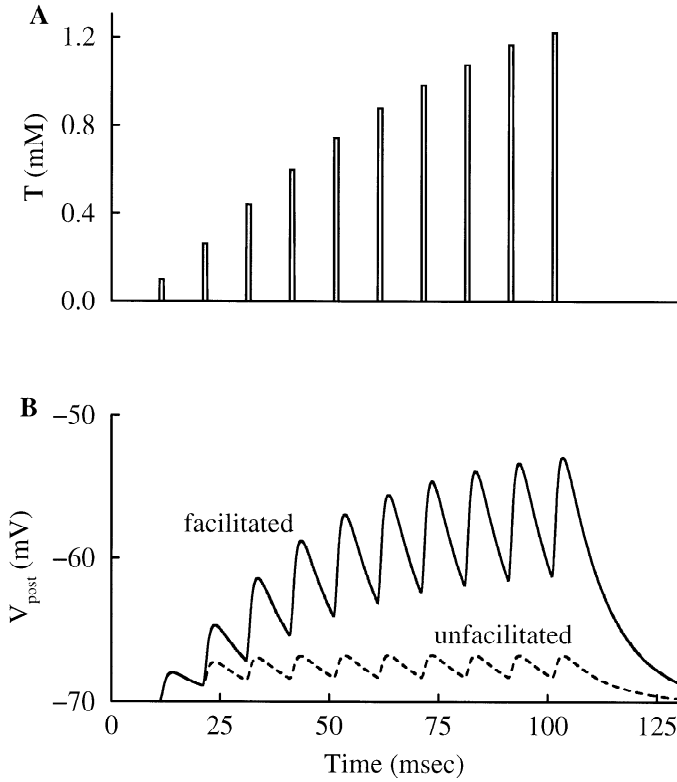


Figure 2: The minimal model of presynaptic facilitation is combined with simple models of postsynaptic binding and passive membrane to describe the postsynaptic voltage response to a short 100-Hz impulse train (see equations 2.3, 4.1, and 4.2). (A) Each stimulus elicits a 1-msec square pulse of transmitter ( $T$ ). The magnitude of the first pulse is assumed to be 0.1 mM. The magnitude of subsequent pulses grows as the synapse facilitates. (B) The increased magnitude of  $T$  results in an increased postsynaptic voltage response with each stimulus, leading to a significant rise in the average postsynaptic voltage (solid line). In the absence of facilitation, the average voltage equilibrates at a lower level (dashed line). Postsynaptic parameter values are:  $\alpha_a = 2 \text{ msec}^{-1} \text{ mM}^{-1}$ ,  $\beta_a = 1 \text{ msec}^{-1}$ ,  $\bar{g}_{\text{mem}} = 0.1$ ,  $\bar{g}_{\text{syn}} = 0.2 (\mu\text{M cm}^{-2})$ ,  $V_{\text{mem}} = -70$ ,  $V_{\text{syn}} = 0$  (mV), and  $C_{\text{mem}} = 1 \mu\text{F cm}^{-2}$ .

postulates an explicit voltage dependence in the release mechanism, a feature that has not been supported by experimental data (Landò & Zucker, 1994).

The present model also belongs to the family of kinetic models used to describe a wide variety of neuronal mechanisms (see Destexhe, Mainen, & Sejnowski, 1994b, for an overview). These include voltage-gated and ligand-gated ion channels (Hodgkin & Huxley, 1952; Destexhe et al., 1994b); IP<sub>3</sub>-sensitive Ca<sup>2+</sup> channels in the membrane of the endoplasmic reticulum (De Young & Keizer, 1992); postsynaptic receptors and channels (Standley, Ramsey, & Usherwood, 1993; Destexhe et al., 1994a,b); and presynaptic transmitter release sites (Parnas et al., 1986; Yamada & Zucker, 1992; Destexhe et al., 1994b; Bertram et al., 1996). In Destexhe et al. (1994b) a nonfacilitating kinetic model of transmitter release was coupled to kinetic models of postsynaptic receptor binding to provide a description of the complete synaptic signal transduction process. As demonstrated in Figure 2, the release model discussed in this article can be applied in a similar manner, introducing the key feature of presynaptic facilitation to the signal transduction scheme.

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