

Calcium-Dependent Exocytosis, Biophysical Models of

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Synonyms

[Non-constitutive exocytosis](#); [Regulated exocytosis](#)

Definition

Calcium-dependent exocytosis is the biochemically controlled fusion of the bilipid secretory vesicle membrane with the bilipid cell membrane, triggered by the binding of several Ca^{2+} ions to control proteins such as synaptotagmins anchored at the interface between these two membranes. Exocytosis results in the release of vesicle contents into the extracellular space, namely, the release of neurotransmitter into the synaptic cleft in the case of neuronal synapses and neuromuscular junctions or the secretion of hormone into the bloodstream in the case of endocrine cells. Exocytosis also allows the transmembrane proteins contained in the vesicle membrane to be incorporated into the cell membrane, although such membrane protein trafficking is more characteristic of Ca^{2+} -independent, constitutive exocytosis.

Detailed Description

In synapses, neuromuscular junctions, and endocrine cells, fast Ca^{2+} -triggered exocytosis of a neurotransmitter or hormone-containing vesicle occurs primarily through the interaction of the so-called SNARE (soluble *N*-ethylmaleimide-sensitive factor attachment protein receptor) proteins spanning the synaptic and vesicle membranes with specific isoforms of the Ca^{2+} -sensitive protein synaptotagmin and with involvement of other control proteins such as complexins (Jahn and Fasshauer 2012). Despite highly specialized morphology, a similar exocytosis-triggering mechanism is found at most high-throughput ribbon sensory synapses (Cho and von Gersdorff 2012; Sterling and Matthews 2005), except for cochlear hair cells which may differ in both their molecular exocytosis machinery and Ca^{2+} sensitivity (Nouvian et al. 2011) and where exocytosis may require nonneuronal isoforms of synaptotagmin (Johnson et al. 2010) or a different Ca^{2+} sensor, otoferlin (Pangrsic et al. 2012; Roux et al. 2006). The mechanism of exocytosis of hormone-containing large dense-core vesicles in endocrine cells is very similar to the mechanisms of neurotransmitter vesicle release (Chow et al. 1992; Voets 2000), but is believed to proceed at a somewhat slower rate due to less tight morphology of the release site and more loose coupling between voltage-dependent calcium channels and release-ready vesicles (Verhage and Toonen 2007; Wu et al. 2009). However, this distinction may be specific to only certain subclasses of endocrine cells, since a close channel-vesicle coupling characteristic of that in neurons has been found in pancreatic beta cells (Barg et al. 2001).

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The dependence of the neurotransmitter or hormone release rate on Ca^{2+} concentration is known to be steeply nonlinear, as was first observed by Dodge and Rahamimoff (1967) when studying neuromuscular junction potentials at different extracellular Ca^{2+} concentrations and later confirmed in studies that varied intracellular Ca^{2+} concentration directly, using caged- Ca^{2+} release in endocrine cells, the giant calyx of Held synapse, and other synaptic terminals (Neher and Sakaba 2008). For low, subsaturating concentrations of Ca^{2+} , this nonlinear relationship can be summarized as

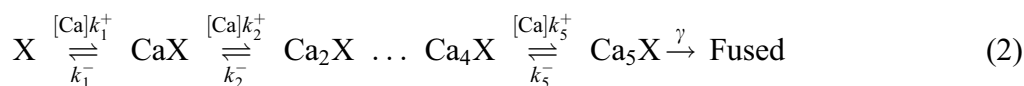
$$R([\text{Ca}]) \propto [\text{Ca}]^n \quad (1)$$

Here $[\text{Ca}]$ denotes Ca^{2+} concentration at the vesicle fusion site, R denotes instantaneous neurotransmitter release rate, and n is the intrinsic (biochemical) Ca^{2+} cooperativity of exocytosis, which varies from three to five in most preparations. Note however that a significantly less cooperative, near-linear Ca^{2+} dependence has been reported in mature auditory hair cells, possibly because of differences in molecular exocytosis sensors mentioned above (Cho and von Gersdorff 2012; Johnson et al. 2010; Nouvian et al. 2011), although it has been suggested that the linear dependence could potentially arise from the averaging across synaptic contacts (Heil and Neubauer 2010).

The release (exocytosis) rate R is usually estimated in units of vesicles per second, and measured as a membrane capacitance increase or by electrochemical detection of released molecules, using a carbon fiber electrode, but often assessed only indirectly by measuring postsynaptic currents or potentials. The steep nonlinear dependence given by Eq. 1 indicates that simultaneous binding of several Ca^{2+} ions to exocytosis control proteins, most likely isoforms of synaptotagmin, is needed for release. In fact, the biochemical cooperativity can be viewed as a lower bound on the number of Ca^{2+} -binding events required for exocytosis (Dittrich et al. 2013). Although the C2A and C2B domains of a given synaptotagmin molecule do possess a total of five Ca^{2+} ion-binding sites (Chapman 2002), not all of these sites are necessarily involved in fast (phasic) Ca^{2+} -triggered exocytosis, and therefore it is possible that the relevant Ca^{2+} -sensitive sensors could be distributed among several synaptotagmin molecules, which dimerize and bind Ca^{2+} simultaneously to trigger vesicle fusion (Mutch et al. 2011). However, most models (reviewed below) assume for simplicity a small number of Ca^{2+} -binding sites, all of which have to be occupied for vesicle fusion to occur. An alternative detailed model that assumes an excess of Ca^{2+} -binding sites and includes Monte Carlo simulation of Ca^{2+} ion diffusion and binding has recently been examined in Dittrich et al. (2013).

Sequential Ca^{2+} -Binding Model

Assuming for concreteness five distinct Ca^{2+} -binding sites comprising the putative exocytosis gate (sensor) X , the most general Ca^{2+} -sensitive exocytosis process can be described by the following reaction:



where $[\text{Ca}]$ is the Ca^{2+} concentration at the vesicle site, k_j^\pm are the binding and unbinding rates of each binding site, and the final irreversible reaction represents the actual vesicle fusion event. In a deterministic simulation, this reaction is converted to a system of ordinary differential equations, using the principle of mass action:

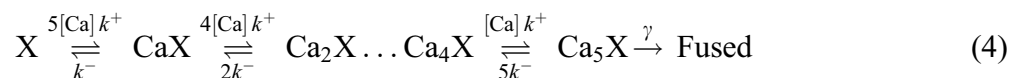
$$\begin{cases} \frac{d[X]}{dt} = -k_1^+[Ca][X] + k_1^-[CaX] \\ \dots \\ \frac{d[Ca_5X]}{dt} = k_5^+[Ca][Ca_4X] - (\gamma + k_5^-)[Ca_5X] \end{cases} \quad (3)$$

where $[Ca_nX]$ represents the fraction of exocytosis gates with n -binding sites occupied by a Ca^{2+} ion. The maximal fusion rate is given by the product $\gamma[Ca_5X]$. Note that the Ca^{2+} concentration, $[Ca]$, should be modeled independently; in the simplest case of global Ca^{2+} elevation produced by caged Ca^{2+} release, it is approximately constant, or it can be represented as a brief pulse of certain width and amplitude when modeling vesicle release produced by a train of action potentials.

The binding and unbinding rates k_j^\pm are in general distinct, and if the earlier unbinding rates are slow, a significant accumulation of partially bound states of X can result during a train of stimuli; such accumulation due to slow unbinding is the basis for the so-called bound- Ca^{2+} model of synaptic facilitation (see entry on “► [Facilitation, Biophysical Models](#)”; Bertram et al. 1996; Bornschein et al. 2013; Matveev et al. 2006).

Parallel Ca^{2+} -Binding Model

It is instructive to first consider the simplest scenario where all five release sites comprising exocytosis gate X are identical and can bind Ca^{2+} independently, leading to the following simplified version of the reaction given by Eq. 2:

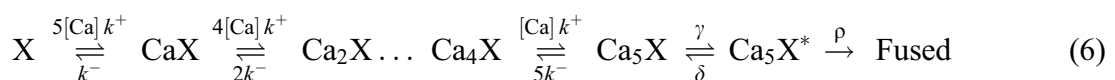


Here the final reaction representing vesicle fusion is irreversible and is triggered when all five binding sites are occupied; exocytosis proceeds at rate $R = \gamma [Ca_5X]$. Even though Eq. 4 appears to describe a series of five consecutive binding reactions, it is equivalent to five identical reactions occurring in parallel, with fusion taking place when all five sites are bound, as follows:

$$\begin{cases} Y \xrightleftharpoons[k_1^-]{[Ca]k_1^+} CaY, \quad 5CaY \xrightarrow{\gamma} \text{Fused} \\ \frac{d[CaY]}{dt} = k^+[Ca](1 - [CaY]) - k^-[CaY] \end{cases} \quad (5)$$

where $[Y] = 1 - [CaY]$ represents the fraction of unbound gates. In this case, the fusion rate is given by the product $\gamma[CaY]^5$ (identically equal to $\gamma[CaX]$ in reaction (4)), where $[CaY]$ is the bound fraction (probability) of each gate Y.

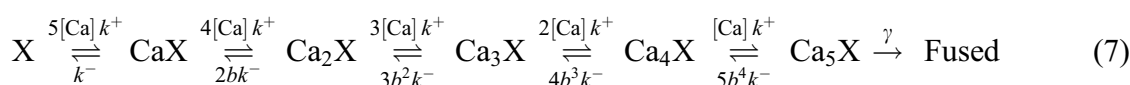
Some implementations of vesicle release process postulate the existence of an independent final release-promoting conformational transition reaction occurring after the release sensor is fully Ca^{2+} bound, as, for instance, done in the model of neurotransmitter release at the calyx of Held synaptic terminal by Bollmann and Sakmann (2005) and Bollmann et al. (2000):



Here ρ is the maximal vesicle fusion rate. In Bollmann et al. (2000), the parameter values that we found to fit well the data from the calyx of Held synaptic terminal are given by $\rho = 40 \text{ ms}^{-1}$, $k^+ = 0.3 \mu\text{M}^{-1}\text{ms}^{-1}$, $k^- = 3 \text{ ms}^{-1}$, $\gamma = 30 \text{ ms}^{-1}$, and $\delta = 8 \text{ ms}^{-1}$. The inclusion of an additional post-binding step helps to achieve more constant shape of the release time course at different amplitudes of Ca^{2+} influx, as seen in experiment (Bollmann et al. 2000; Yamada and Zucker 1992).

Cooperative Ca^{2+} -Binding Model

Some studies suggest that the Ca^{2+} -sensitive exocytosis triggers exhibit strong cooperativity, whereby the target protein undergoes a conformational change with each successive Ca^{2+} ion binding, which in turn leads to an increase in the Ca^{2+} affinity of the remaining (yet unoccupied) Ca^{2+} -binding sites. The following widely used modification of reaction given by Eq. 2 has been proposed to implement this possibility (Felmy et al. 2003; Heidelberger et al. 1994; Schneggenburger and Neher 2000):



Note that cooperative binding can be represented as either a progressive increase in forward binding rates or decrease of backward rates; Eq. 7 corresponds to the latter possibility. The cooperativity parameter should satisfy $b < 1$, with a typical value used in the literature of 0.25, indicating that the final Ca^{2+} -binding reactions are only slowly reversible. In Schneggenburger and Neher (2000) and Wolfel and Schneggenburger (2003), quantifying the Ca^{2+} dependence of vesicle release at the calyx of Held synaptic terminal, the following parameter values were obtained: $k^+ = 0.09 \mu\text{M}^{-1} \text{ms}^{-1}$, $k^- = 9.5 \text{ ms}^{-1}$, $\gamma = 6 \text{ ms}^{-1}$, and the cooperativity parameter $b = 0.25$.

Note that the use of the term “cooperativity” is ambiguous in the context of exocytosis mechanisms, since it is used to refer to two different properties of the exocytosis process: cooperativity can either refer to the number of Ca^{2+} -binding sites (as inferred from the log-log slope of the Ca^{2+} dose–response curve) or indicate that the Ca^{2+} -binding affinity of these sites is not equal and increases as the first sites become Ca^{2+} bound.

Models with Independent Sets of Ca^{2+} -Binding Sites

Several recent biophysically detailed models of exocytosis take into account the possibility that Ca^{2+} may bind to several different proteins or to different domains of the same protein, such as the C2A and C2B domains of synaptotagmin, each characterized by a distinct set of rates and cooperativity values. Examples of such models are given below in the context of more comprehensive models of exocytosis – see model schemes (16) and (20).

Exocytosis Rate at a Steady Ca^{2+} Concentration

To simulate the exocytosis rate during prolonged Ca^{2+} elevation, for instance, to reproduce caged- Ca^{2+} release experiments, it is sufficient to consider the equilibrium point of the reactions summarized above. At equilibrium, the occupancy of the Ca^{2+} -bound states is easily found by equating the right-hand side of the binding reaction to zero. For the case of independent gates, described by Eqs. 4 and 5, the equilibrium is given by

$$[\text{CaY}] = \frac{k^+[\text{Ca}]}{k^+[\text{Ca}] + k^-} = \frac{[\text{Ca}]}{[\text{Ca}] + K} \quad (8)$$

where the ratio of the backward and forward rates $K = k^-/k^+$ is referred to as the Ca^{2+} affinity or dissociation constant and is an important Ca^{2+} -sensitivity parameter: the lower the K , the higher is the affinity (sensitivity) of the release sensor. It follows from Eq. 8 that K represents the Ca^{2+} concentration at which half the sensors become bound. Since the final fusion reaction in Eq. 5 requires the binding of all five sensors, the release rate at equilibrium will be given by the 5th power of the steady-state binding occupancy of gate Y :

$$R([\text{Ca}]) = \gamma \left(\frac{[\text{Ca}]}{[\text{Ca}] + K} \right)^5 \quad (9)$$

At low Ca^{2+} concentration, this agrees with the cooperativity condition given by Eq. 1. Note that the release rate equals $1/2^5$ of its maximal value when $[\text{Ca}^{2+}] = K$.

For the case of cooperative binding, Eq. 7, the steady-state fusion rate R is given by the equilibrium value of fully bound state Ca_5X , which obeys a modified form of the Adair-Klotz-Pauling equation (Weiss 1997)

$$R([\text{Ca}]) = \frac{\gamma [\text{Ca}]^5}{[\text{Ca}]^5 + 5 [\text{Ca}]^4 b^4 K + 10 [\text{Ca}]^3 b^7 K^2 + 10 [\text{Ca}]^2 b^9 K^3 + 5 [\text{Ca}] b^{10} K^4 + b^{10} K^5} \quad (10)$$

where $[\text{Ca}]$ is the Ca^{2+} concentration at the release site. In the limit of very small values of b (strong cooperativity limit), the above expression approaches the well-known and widely used Hill function:

$$R([\text{Ca}]) \approx \gamma \frac{[\text{Ca}]^n}{[\text{Ca}]^n + K_D^n} \quad (11)$$

where $n = 5$ and $K_D \approx b^2 K$ ($b \ll 1$). The dissociation constant K_D quantifies the Ca^{2+} sensitivity of the entire 5-step process, as opposed to the sensitivity of any individual Ca^{2+} -binding site: when $[\text{Ca}] = K_D = b^2 K$, the exocytosis rate reaches its half-maximal value. Note that this is the most constrained and most important model parameter, since extensive experimental data on Ca^{2+} sensitivity has been collected at various types of synaptic terminals: in general, physiological rates of release are found when $[\text{Ca}^{2+}]$ at the vesicle location reaches the range of 10–50 μM (reviewed in Neher and Sakaba 2008), although there are reports that highly sensitive vesicle pools exist in some cells, possibly controlled by distinct isoforms of the synaptotagmin sensor with Ca^{2+} affinity of several μM (reviewed in Pedersen and Sherman 2009).

Note that the Hill functional form given by Eq. 11 should only be viewed as a crude qualitative approximation of the true Ca^{2+} dependence of exocytosis rate, even at equilibrium, and tends to be somewhat overused (Weiss 1997). This is because the agreement between Eqs. 10 and 11 becomes sufficiently accurate only for values of cooperativity parameter satisfying $b < 0.1$; in the absence of firm experimental evidence for such a strong cooperativity, the Hill function should be avoided. Assuming a more moderate but still strong cooperativity corresponding to the typically used value $b = 0.25$, fitting the data to a Hill function would lead to a gross underestimate of the true number of Ca^{2+} -binding sites, as illustrated in Fig. 1. This underestimate is mostly due to the poor match of the saturating part of the Hill curve and can be greatly improved by omitting the data from saturating levels of $[\text{Ca}^{2+}]$. Therefore, the most model-independent way of estimating the cooperativity factor n is to only fit the log-log slope of the experimentally obtained Ca^{2+} sensitivity curve *below* the saturation inflection, which is often done in practice.

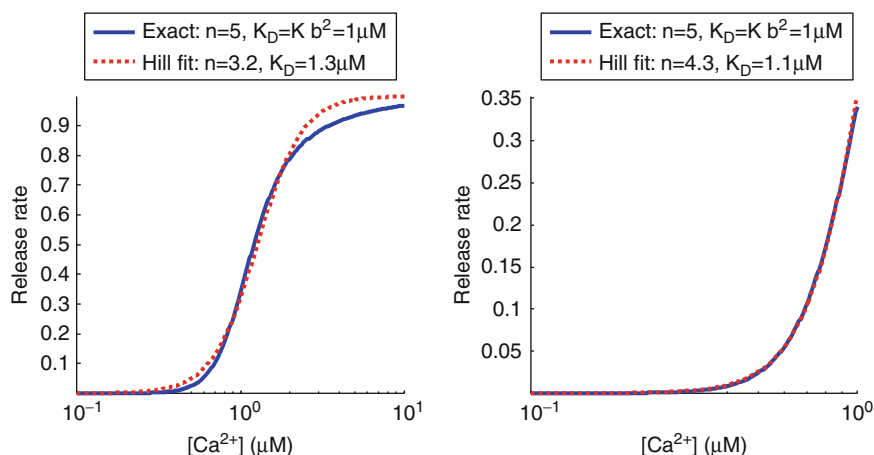


Fig. 1 Hill function fit underestimates true biochemical cooperativity and affinity of a 5th-order Ca²⁺-binding reaction (Eqs. 7 and 10), even in the case of moderately strong cooperativity parameter ($b = 0.25$). Note that the low quality of the fit is mostly a result of poor matching of the saturating part of the curve by the Hill function: the cooperativity is more accurately predicted if only the non-saturating part of the data is considered (*right panel*)

Cumulative Release During Prolonged Ca²⁺ Elevation

The Ca²⁺ dependence of release rate can be used to quantify the total cumulative release with prolonged Ca²⁺ elevation, as measured by the total membrane capacitance increase or the total amount of released neurotransmitter. In the case of prolonged and approximately steady Ca²⁺ elevation of duration Δt , the amount of fused vesicles is given by Quastel et al. (1992):

$$F([\text{Ca}]) = F_{\max} [1 - \exp(-R_{\text{eq}}([\text{Ca}]) \Delta t)] \quad (12)$$

where F_{\max} represents the total amount of available exocytosis resources (say, vesicles) and $R_{\text{eq}}([\text{Ca}])$ represents the equilibrium reaction rate at Ca²⁺ concentration [Ca] attained at the vesicle location. The exponential term can be interpreted as the probability of release failure. For pulses raising [Ca²⁺] to subsaturating levels, given Eq. 1 we have $R \approx k[\text{Ca}]^n$; therefore (Quastel et al. 1992)

$$F([\text{Ca}]) = F_{\max} [1 - \exp(-k [\text{Ca}]^n \Delta t)] \quad (13)$$

However, if [Ca²⁺] is high enough to saturate the release sensor during the long depolarizing pulse, the total amount of released neurotransmitter will become limited by the duration of stimulation only:

$$F = F_{\max} [1 - \exp(-R_{\max} \Delta t)] \quad (14)$$

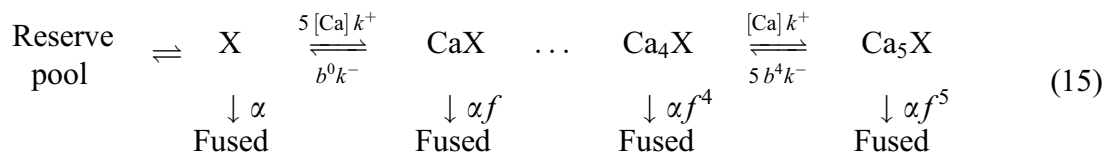
As was explained above, Eqs. 12 and 13 only apply to the situation where [Ca²⁺] is elevated for a long time relative to the kinetics of the Ca²⁺-binding sites, as, for instance, during whole-cell [Ca²⁺] elevation produced by Ca²⁺ uncaging experiments. If on the other hand [Ca²⁺] is understood as the *peak* concentration achieved in the vicinity of the vesicle during a brief action potential, the above steady-state results would overestimate the true release and could only serve as an upper bound on release rate (Shahrezaei and Delaney 2005). The “true” neurotransmitter release rate would be given by a solution to the differential equation (5), with [Ca²⁺] representing the time-dependent Ca²⁺ concentration at the vesicle release site, which has to be modeled independently,

using simulations of Ca^{2+} entry through Ca^{2+} channels, diffusion, and binding to intracellular Ca^{2+} buffers.

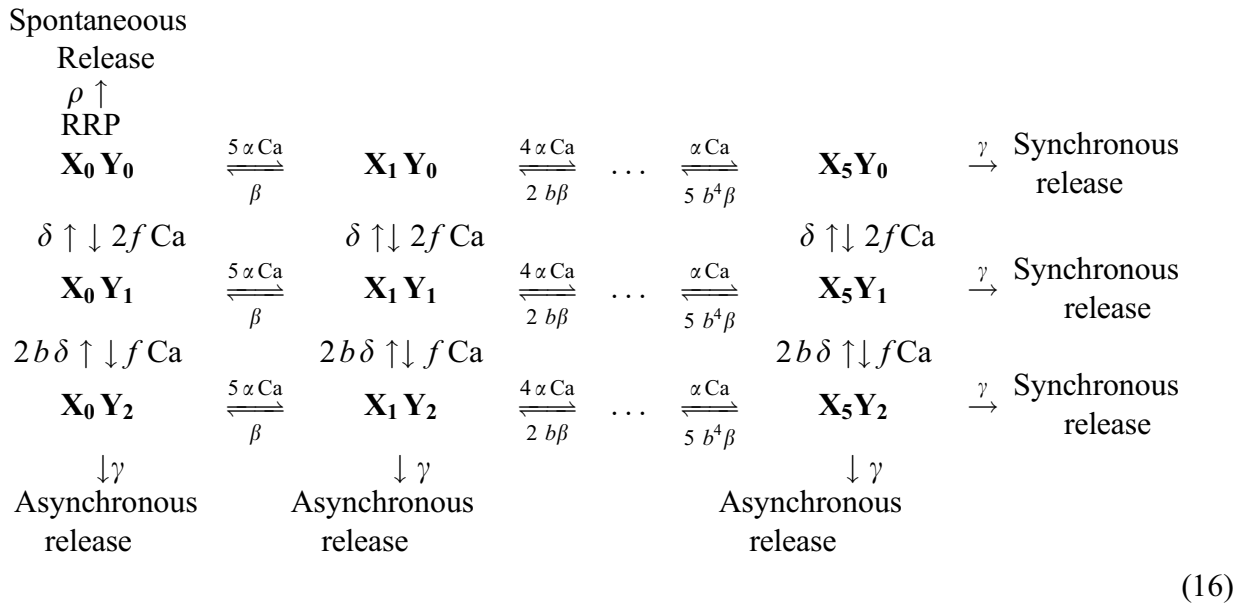
Release at Low Ca^{2+} Concentration: Asynchronous and Spontaneous Transmission

At very low Ca^{2+} concentration, depolarization-evoked exocytosis is more stochastic and less synchronized with the stimulus, and its rate decreases, becoming comparable to the spontaneous transmission observed in the absence of stimulation. This delayed component of exocytosis is termed asynchronous or delayed exocytosis and at some synapses represents a significant contribution to the overall neurotransmitter release under physiological conditions. A prominent feature of asynchronous release is that it seems to exhibit a lower, almost linear Ca^{2+} dependence (Lou et al. 2005; Sun et al. 2007). As will be discussed further below, there is some debate whether the distinct components of synaptic transmission are all manifestations of the same Ca^{2+} -dependent process or are in fact caused by distinct vesicle pools released via independent mechanisms (Chung and Raingo 2013; Smith et al. 2012). However, it should be noted that the mechanisms of asynchronous release may vary across distinct types of synapses (Kaesler and Regehr 2014).

The so-called allosteric model proposed by Lou et al. (2005) explains the observed decrease in apparent cooperativity of Ca^{2+} action at low Ca^{2+} concentration by the presence of a slow conformational change of the release proteins, leading to a second route of vesicle fusion, which becomes progressively less likely at higher Ca^{2+} concentration. This is implemented by the following scheme with a reverse-cooperativity parameter $f < 1$:



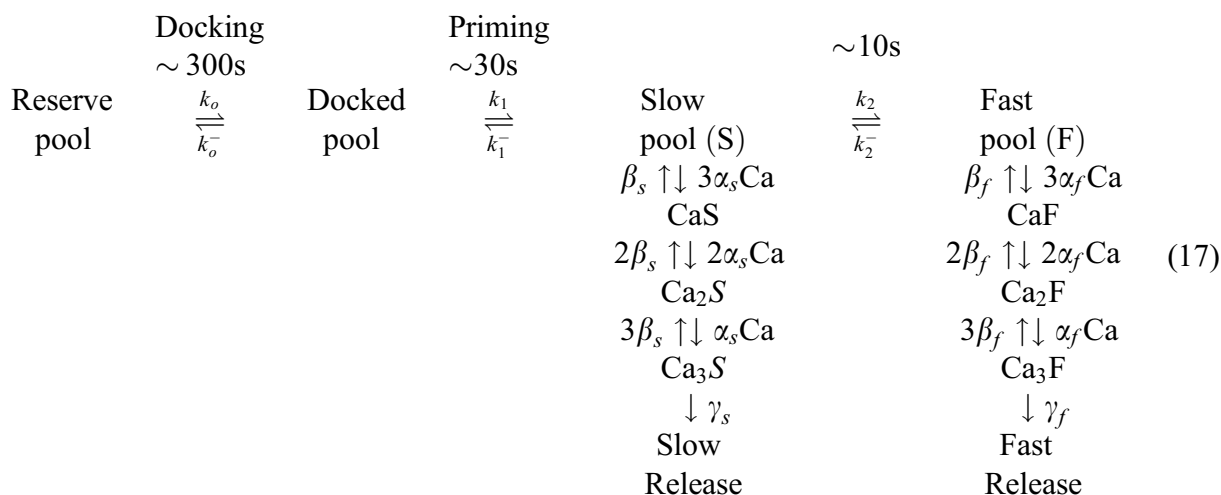
A more detailed model proposed by Sun et al. (2007) and referred to as the *dual-sensor* model provides an improved quantitative description of release at low Ca^{2+} concentrations, since it reproduced more accurately the latency of synaptic response at very low levels of Ca^{2+} . This model assumes two independent Ca^{2+} sensors acting in parallel and each triggering a distinct mode of neurotransmitter release:



Here the synchronous release involves a sensor “X” with Ca^{2+} cooperativity of five, while a different and independent sensor “Y” is responsible for asynchronous release and possesses a cooperativity of two. Note that in this model implementation, both sensors bind Ca^{2+} cooperatively, with equal cooperativity parameter b . The existence of two distinct sensors is supported by recent evidence that in some synapses asynchronous release requires a special non-synaptotagmin sensor Doc2 (Yao et al. 2011). However, the mechanisms of asynchronous and spontaneous release are still under debate and may vary across synaptic types (Chung and Raingo 2013; Kaeser and Regehr 2014; Smith et al. 2012).

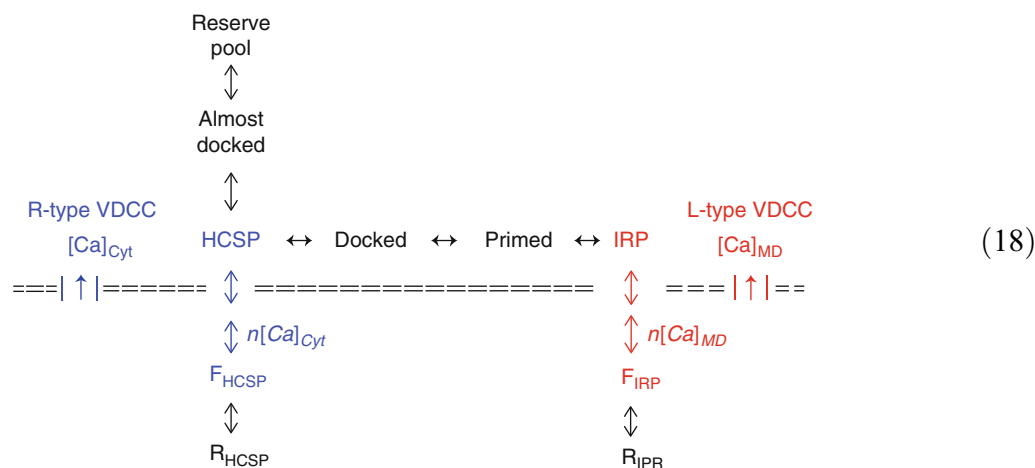
Multiple Vesicle Pool Models

Apart from the intrinsic heterogeneity in Ca^{2+} affinity and speed of exocytosis due to the existence of distinct release pathways guiding exocytosis of each vesicle, an additional explanation for the observed heterogeneous components of vesicle release is the existence of independent pools of vesicles with variable degree of “preparedness” for exocytosis. This heterogeneity is especially pronounced in endocrine cells, which exhibit a fast-decaying initial phase of release followed by a second, more slowly decaying component, but such behavior is also found in other high-throughput neuronal synapses such as the calyx of Held synaptic terminal and in ribbon synapses (Neher 2012; Neher and Sakaba 2008; Pedersen and Sherman 2009; Verhage and Toonen 2007). The identity of such distinct vesicle pools is currently under debate: they could be distinguished either by the variations in the distance between vesicles and corresponding voltage-gated Ca^{2+} channels (“positional priming”: Wadel et al. 2007) or by differences in the arrangement of the molecular machinery needed for exocytosis (“molecular priming” and “super-priming” (Chung and Raingo 2013; Lee et al. 2013; Sorensen 2004; Verhage and Toonen 2007)). Most quantitative models of exocytosis that include multiple releasable vesicle pools are based on the two-pool model of Heinemann et al. (1993), Voets (2000), and Voets et al. (1999) that was first put forward to quantify two temporal components of secretory vesicle release in adrenal chromaffin cells. Here is a recent implementation of such a two-pool model explored by Sorensen (2004):



In this model exocytosis proceeds with cooperativity of three from both the slow (“sustained”) and fast synchronous pools, which are replenished via two preparatory steps, docking and priming, from the reserve pool.

There are indications that in some cells, the slowly releasing pool may in fact have a much higher sensitivity to Ca^{2+} , in the range of several μM rather than 10 s of μM , which allows it to participate in exocytosis despite a much greater separation from the Ca^{2+} channels compared to the immediately releasable, fast pool (reviewed in Pedersen and Sherman 2009). The delay in the release of the highly sensitive vesicles is explained by the longer diffusional distance and therefore longer time required for Ca^{2+} concentration to reach μM range far from the Ca^{2+} channels. Such so-called highly Ca^{2+} -sensitive pool (HCSP) has been described in adrenal chromaffin cells (Yang et al. 2002), rod photoreceptor ribbon synapses (Thoreson et al. 2004), and insulin-secreting beta cells (Yang and Gillis 2004). A recent multiple-pool model that includes the HCSP pool along with the lower-affinity immediately releasable pool (IRP) and implements a deterministic bi-domain model of $[\text{Ca}^{2+}]$ dynamics can be found in Pedersen and Sherman (2009). This model is based on an earlier model of Chen et al. (2008), and it accurately predicts the characteristic biphasic release of insulin from pancreatic beta cells, with the first phase of secretion mostly due to the docked vesicles that are rapidly converted to immediately releasable pool (IRP) and exocytosed in response to local “microdomain” $[\text{Ca}^{2+}]$ ($[\text{Ca}]_{\text{MD}}$) entering the cell through L-type Ca^{2+} channels, whereas the second, delayed phase of insulin release builds up more slowly due to the recruitment of the reserve vesicles into the HCSP and gradual accumulation of cytosolic Ca^{2+} ($[\text{Ca}]_{\text{Cyt}}$) entering through R-type Ca^{2+} channels:



Here Ca^{2+} -dependent transitions are indicated in color, according to the primary source of Ca^{2+} for the corresponding transition. The R-type and L-type voltage-dependent Ca^{2+} channels (VDCCs) contribute respectively to the cytosolic and the microdomain pools of Ca^{2+} , $[\text{Ca}]_{\text{Cyt}}$, and $[\text{Ca}]_{\text{MD}}$ (the diffusional exchange between these two Ca^{2+} pools is not shown). The Ca^{2+} -dependent exocytosis steps lead to the two “fused” states, F_{HCSP} and F_{IRP} , which in turn feed into the final released insulin states R_{HCSP} and R_{IRP} through a final Ca^{2+} -independent pore-expansion transition.

Although the existence of multiple vesicle pools can also be considered as a potential explanation for asynchronous release (Chung and Raingo 2013), it is likely that intrinsic sources of heterogeneity in release properties captured by the allosteric and the dual-sensor models (see above) are also present in each vesicle (Kaeser and Regehr 2014; Sun et al. 2007). For instance, experiments that disrupt specific molecular components of release machinery indicate that at some synapses synchronous and asynchronous release can be independently manipulated and that asynchronous release is mediated by a distinct Ca^{2+} sensor, Doc2 (Yao et al. 2011), and possibly involves a distinct isoform of the vesicle SNARE protein VAMP4 (Raingo et al. 2012). Further, both the synchronous and the delayed components are observed with global (whole-terminal) Ca^{2+} elevations that activate vesicles in all pools. Therefore, both sources of heterogeneity, the ones intrinsic to each vesicle and the extrinsic ones (distinct pools), seem to be relevant in many synapses (Kaeser and Regehr 2014). In order to take into account all factors of vesicle release heterogeneity and predict more accurately the exocytosis rate under low Ca^{2+} conditions, one can combine the dual-pool model with the allosteric model of Lou et al. (2005) or the dual-sensor model of Sun et al. (2007). This was done, for instance, by Wolfel and Schneggenburger (2003), who considered allosteric Ca^{2+} binding for both pools, as in reaction (15) (neglecting however the exchange between the two pools due to the short timescales considered in that work). Finally, it should be noted that the separation of releasable vesicles into only two pools instead of a larger set of pools or a continuum of states is most probably a simplification, but this level of detail is sufficient to accurately quantify release under most physiological conditions. Including more than two pools would most likely lead to an underdetermined model and to data over-fitting.

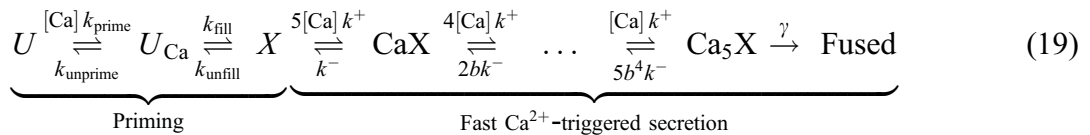
Models with Ca^{2+} -Dependent Vesicle Priming

Vesicle docking and priming steps in reactions (17)–(18) most likely include multiple molecular processes (involving Munc18 and other proteins) and morphological steps (e.g., positional priming), the identity and sequence of which is still under investigation (Lee et al. 2013; Verhage and Toonen 2007). Experimental evidence suggests that at least some of these vesicle priming steps are Ca^{2+}

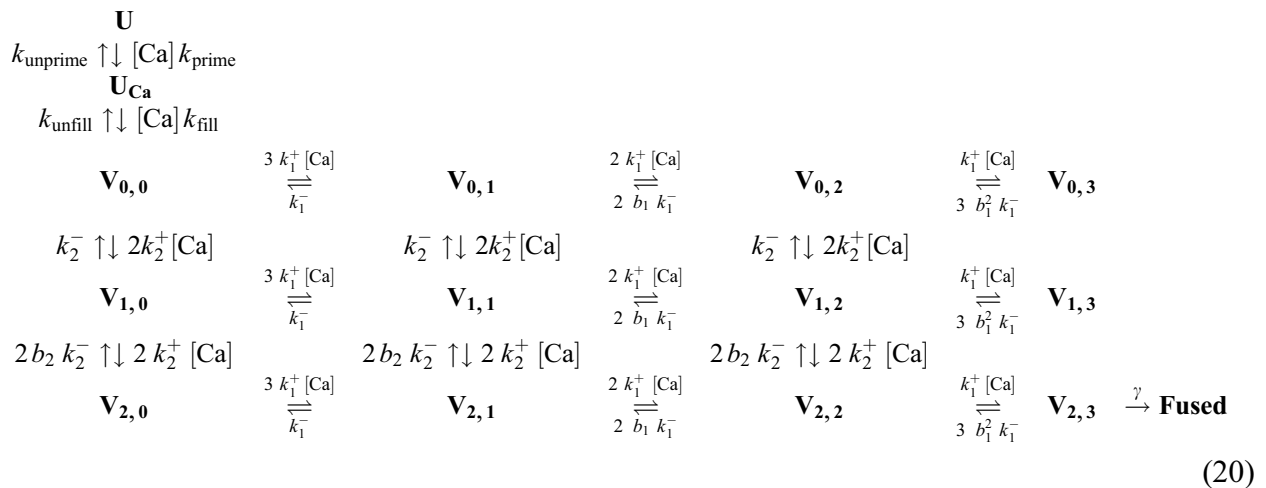
dependent in many synapses, albeit with a lower Ca^{2+} cooperativity compared to exocytosis itself, with a near-linear dependence on intracellular Ca^{2+} concentration (Neher and Sakaba 2008). Depending on the balance between the different vesicle pools at resting Ca^{2+} , the Ca^{2+} -dependent priming can manifest itself in two different properties of short-term synaptic plasticity (Zucker and Regehr 2002):

1. Under conditions of low initial release-ready vesicle pool size, Ca^{2+} -dependent priming would cause the secretion rate to increase during stimulation as a result of an increase in vesicle “mobilization” to the release-ready, primed pool, which can serve as a mechanism of short-term synaptic facilitation (Dittman et al. 2000; Millar et al. 2005; Pan and Zucker 2009; Worden et al. 1997)(see also encyclopedia entry on “► Facilitation, Biophysical Models”).
2. Under conditions of high vesicle release probability, Ca^{2+} -dependent priming would manifest itself through activity-dependent acceleration of recovery from short-term synaptic depression (Dittman and Regehr 1998; Hosoi et al. 2007; Stevens and Wesseling 1998; von Ruden and Neher 1993; Wang and Kaczmarek 1998).

The simplest model of release that includes a Ca^{2+} -dependent priming step can be obtained by modifying reaction (7): for instance, Millar et al. (2005) considered the following scheme (see also Bornschein et al. (2013) and Sakaba (2008) for a slightly modified version of this scheme):



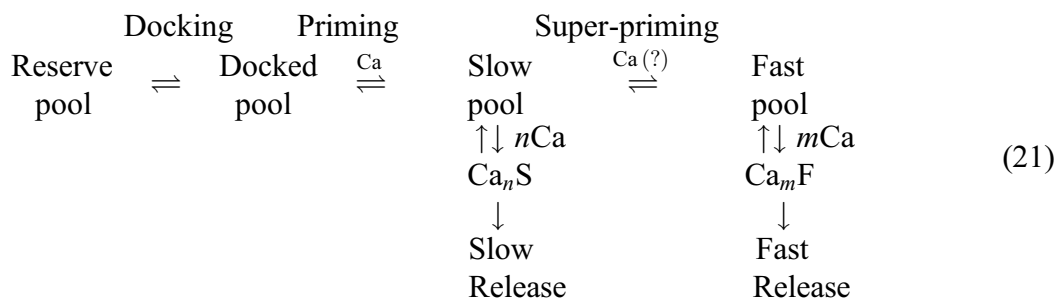
The study of Millar et al. (2005) also considered a more detailed version of this model that takes into account the independence of Ca^{2+} binding to C2A and C2B domains of synaptotagmin, leading to the following modification of the above (see also Pan and Zucker 2009):



Here the horizontal state transitions represent the binding of the three C2A sites of synaptotagmin, while the vertical transitions correspond to the binding of the C2B sites, which is assumed to be independent of the Ca^{2+} binding of the C2A domain. Both domains are assumed to bind Ca^{2+} cooperatively (with cooperativity parameter $b_1 = b_2 = 0.5$). These models were successful in

explaining the difference between release properties of the facilitating “tonic” and depressing “phasic” crustacean neuromuscular junctions (Millar et al. 2005; Pan and Zucker 2009).

Vesicle priming has also been considered in the context of more detailed models that include multiple releasable vesicle pools. In such models, the priming step is usually assumed to occur upstream of the process of exchange between the slow-releasing and the fast-releasing vesicle pools. Adding the priming step to scheme (17) leads to the following class of models (Verhage and Toonen 2007):



In this scheme, “super-priming” refers to the conversion of the slowly releasable pool to the fast releasable pool, the nature and the Ca^{2+} dependence of which is currently unknown. Note that the use of the term “super-priming” is model dependent; recently it has been used to describe a newly identified additional kinetic step in the priming of the fast pool at the calyx of Held synaptic terminal (Lee et al. 2013). The possible Ca^{2+} dependence of the conversion of the slow pool to the fast pool is unknown; it is possible that this conversion may even be retarded by Ca^{2+} at some synapses (Neher and Sakaba 2008). At the calyx of Held, the contribution of the fast component of release is observed to increase with increasing Ca^{2+} levels, at the expense of the slow component (Wolfel et al. 2007). An explanation of this observation was suggested that the slow pool may be small at rest but increases due to activity-dependent conversion of vesicles from the fast pool and that this conversion is less pronounced at high Ca^{2+} levels, so it is not able to reduce the fast pool at more intense stimulation levels (Neher and Sakaba 2008).

Comprehensive Biophysical Models of Ca^{2+} -Dependent Exocytosis

Several particularly detailed recent modeling studies include simulations of Ca^{2+} dynamics and combine together many of the mechanisms reviewed above to build a more comprehensive model of vesicle exocytosis:

1. The model of Dittrich et al. (2013) is one of the most complete recent models of exocytosis that includes stochastic simulations of Ca^{2+} ions diffusion, buffering, and binding to multiple synaptotagmin molecules and takes into account most recent data on the copy number of synaptotagmin molecules and SNARE complexes per vesicle (Chapman 2002; Han et al. 2004; Mutch et al. 2011). This is the first modeling study that considered the possibility that only a subset of synaptotagmin sites have to be Ca^{2+} bound to trigger exocytosis, and it examines the effective Ca^{2+} cooperativity of exocytosis as an emergent characteristic of such partial binding of a subset of Ca^{2+} -binding sites.
2. The study of Pan and Zucker (2009) builds a comprehensive model of release and short-term plasticity at tonic and phasic crustacean neuromuscular junctions, based on the scheme (20) reviewed above but adds positional priming, leading to a transition scheme describing vesicle exchange between the following distinct vesicle pools: reserve, docked, primed channel-

detached, and primed channel-attached. One of the main aims of this study was to build a model that most fully accounts for the properties of short-term synaptic facilitation observed at crustacean tonic neuromuscular junctions, as well as the properties of short-term depression and asynchronous release at phasic neuromuscular junctions.

The latter model of Pan and Zucker (2009) is a more detailed example of a class of comprehensive models of Ca^{2+} -dependent exocytosis that go beyond the simulation of vesicle release in response to a single presynaptic depolarization, aiming to build a comprehensive model of short-term changes in synaptic transmission strength in response to repeated stimulation. Such modeling work is reviewed in more detail in the entry “► [Facilitation, Biophysical Models.](#)”

Spatial Coupling Between Vesicle and Ca^{2+} Channels

Building a comprehensive model of Ca^{2+} -dependent exocytosis at a particular class of synaptic terminal requires accurate description of Ca^{2+} dynamics at the release site, which in turn requires taking into account the latest data on the spatial coupling between voltage-dependent Ca^{2+} channels and synaptic vesicles (Eggermann et al. 2012; Gentile and Stanley 2005; Moser et al. 2006; Oheim et al. 2006; Stanley 1997). The geometric arrangement of channels and vesicles at the release site has been recently a subject of intensive modeling effort, using both stochastic simulation of Ca^{2+} diffusion and buffering (Bennett et al. 2004; Glavinovic and Rabie 2001; Nadkarni et al. 2012; Scimemi and Diamond 2012; Shahrezaei et al. 2006; Shahrezaei and Delaney 2005) as well as deterministic solution of continuous reaction–diffusion equations (Bucurenciu et al. 2008; Ermolyuk et al. 2013; Matveev et al. 2009, 2011; Meinrenken et al. 2002, 2003; Schmidt et al. 2013; Zucker and Fogelson 1986) and steady-state approximations of Ca^{2+} distribution near an array of open Ca^{2+} channels (Bertram et al. 1999; Coggins and Zenisek 2009).

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