The mechanism mediating fast neurotransmitter release at the calyx of Held synapse

PhD Thesis

in partial fulfillment of the requirements for the degree 'Dr. rer. nat.' in the Neuroscience Program at the Georg-August-University Göttingen, Faculty of Biology

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Declaration	
I hereby declare that my PhD thesis 'The mechanism release at the calyx of Held synapse' has been written in or sources than quoted.	5 0
Göttingen, September $16^{\rm th}$ 2008	

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Introduction

1.1 Neurotransmitter release - exemplified at the calyx of Held synapse

The calyx of Held synapse was first described in 1893 by the German anatomist Hans Held. Its rather unique features have established it as a model system early on and throughout the conceptional and technological advances in the field of neuroscience. This introduction is meant to reflect this development, to present our knowledge of the model synapse that the calyx of Held has become in relation to our general understanding of synaptic function.

1.1.1 Morphology

The first description of the calycial terminals in the auditory brain stem of cats originated from Golgi staining experiments conducted by the German anatomist Hans Held in 1893. This discovery was made at a time when the nature of connectivity within the brain was heavily disputed. Held, in support of the reticular brain theory, proposed that the axons giving rise to the calyx cross midline during development to eventually fuse onto the principle cells of the medial nucleus of the trapezoid body (MNTB). Ramon y Cajal however viewed the calyx of Held as an example to disprove the reticularists.

Further studies employing staining techniques (Harrison and Irving, 1966), electron microscopy (EM) (Lenn and Reese, 1966; Nakajima, 1971) and intraaxonal recordings combined with horseradish peroxidase staining (Friauf and Ostwald, 1988) investigated

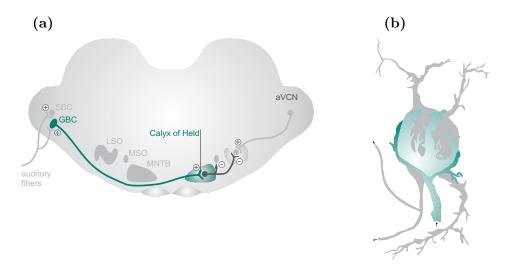


Figure 1.1: (a) Diagram of the superior olivary complex circuitry. Auditory fibers form synapses onto globular (GBC) and spherical bushy cells (SBC) in the anterior ventral cochlear nucleus (aVCN). GBC axons cross midline and give rise to the calyx of Held terminal in the medial nucleus of the trapezoid body (MNTB). Its target, the glycinergic principle cell of the MNTB, projects to the lateral superior olive (LSO) which is thought to serve as a coincidence detector for auditory signals. Plus and minus signs indicate excitatory and inhibitory synapses. (b) Illustration of the calyx of Held (green) and its corresponding principle cell (gray). Adapted from von Gersdorff and Borst (2002)

the connectivity in the auditory pathway that the calyx of Held is part of. The calycial axons are formed by globular bushy cells (GBCs) in the anterior ventral cochlear nucleus (aVCN). Consistent across mammalian species such as cat, bat and rodents, these projections form a direct connection arising in the aVCN, through the trapezoid body, across midline to branch of at the contralateral MNTB where they form large terminals, the calyx of Held, onto a principal cell (Fig. 1.1(a)). These afferents present the main input to the MNTB, however the principle cells also receive small inputs. In younger animals, axons forming two calyx terminals have been described (Kuwabara et al, 1991).

A detailed study of the morphology of a single postnatal day 9 (p9) rat calyx of Held terminal and its connecting principle cell by serial reconstruction of ultra thin sections imaged with electron microscopy was conducted by Sätzler et al. The calyx terminal had a thickness of $<1 \,\mu m$ covering $\sim 40 \,\%$ of the principle cell that was $22 \,\mu m$ and $18 \,\mu m$

in its longer and shorter diameter. They identified 554 active zones (AZs) with a mean distance of 590 nm. On average, two synaptic vesicles were docked at the active zone whereas 2800 vesicles were found within 20 nm from the active zone. By extrapolation, the total number of vesicles within the terminal was estimated to be higher than 70000 (Sätzler et al, 2002).

The calyx of Held has also been a model system for the study of synapse formation. Experiments using the established Golgi staining technique could show the developmental stages from the growth-cone bearing migrating axon to a cup-shaped terminal (around postnatal day 5) and the mature calyx terminal with its digit-like structures (seen around postnatal day 14) (Morest, 1968; Kandler and Friauf, 1993). With development, the calyx of Held also shows subcellular specializations. Rowland et al reported the atypical structural organization of puncta adherentia and mitochondria which they named mitochondria-associated adherens complex, thought to play a role in vesicle and energy supply, calcium homeostasis and synaptic stability to support high frequency transmission at this synapse (Rowland et al, 2000). Wimmer et al have shown a 'donut'-like assembly of synaptic vesicles and active zones arranged around mitochondria by confocal microscopy and electron microscopy that forms after the onset of hearing at p11–p12 (Wimmer et al, 2006).

1.1.2 Physiology

Studies that have investigated the morphology of the calyx of Held synapse by electron microscopy already proposed that is glutamatergic (Lenn and Reese, 1966; Nakajima, 1971). More direct evidence was provided by experiments mapping glutamate-like immunoreactivity in the MNTB which was most prominently found at the calyx of Held (Grandes and Streit, 1989).

Intraaxonal recordings could show that GBCs present peristimulus time histogram responses much like the primary afferents, in a first attempt to correlate electrophysiological characteristics and the morphology of cell types found in the aVCN (Friauf and Ostwald, 1988). Intracellular recordings from MNTB principle cells showed that excitatory postsynaptic potentials (EPSPs) could be blocked by the AMPA/ kainate receptor antagonists CNQX and DNQX (Banks and Smith, 1992). This was confirmed by the first whole-cell voltage clamp recordings which described the excitatory postsy-

naptic currents (EPSCs) to be mediated by AMPA and NMDA receptors with distinct kinetics (Forsythe and Barnes-Davies, 1993a). Additionally MNTB principle cells receive small excitatory and inhibitory inputs. It has been shown that inhibitory postsynaptic currents (IPSCs) are initially mediated by GABA receptors, but are driven by glycine receptors later in development (Banks and Smith, 1992; Awatramani et al, 2004).

The presynaptic channels mediating action potential (AP) firing have been characterized both electrophysiologically as well as immunohistochemically. Excitation of GBCs passes through a myelinated axon via saltatory conduction. AP generation can be completely blocked by application of the sodium channel blocker tetrodotoxin (TTX). Voltage-gated sodium channels are found in high density at nodes of Ranvier and the heminode (the unmyelinated last segment of the axon) but only a low density is found at the terminal. It has been shown that there is a developmental increase in Na_v1.6 α -subunit expression at the calyx heminode. This increase, combined with the apparent absence of sodium channels in the terminal, leads to a shortening of the AP and avoids changes in the sodium gradient that is needed to drive Ca²⁺ extrusion mechanisms. The variable expression of Na⁺ channels could therefore serve as one of the properties that enable high-frequency transmission at the calyx of Held (Leão et al, 2005).

Forsythe and Barnes-Davies have also studied the conductances mediating AP generation and identified an inward sodium current and a low- and a high-voltage activated potassium current ($I_{K,LV}$, $I_{K,HV}$) as well as a hyperpolarization-activated current (I_H) (Forsythe and Barnes-Davies, 1993b). $I_{K,LV}$ is attributed to the K_v1 family of potassium channels, $I_{K,HV}$ to the K_v3 family. Again, expression is differential between the heminode and the terminal. K_v1 channels colocalize with the $Na_v1.6$ channels in the heminode at the light microscopic level and are absent from the terminal. K_v1 channels are only found in the terminal. Apart from these delayed rectifiers, I_H and Ca^{2+} -activated K^+ channels (BK) have been described presynaptically. Their physiological role at the calyx of Held, however, remains unclear.

Various other receptors have also been found in the presynaptic terminal: mGluRs, $GABA_B$, noradrenergic $\alpha 2$ and adenosine receptors, all of which can modulate transmitter release at the calyx of Held (reviewed in Schneggenburger and Forsythe, 2006).

Calcium channels, cytosolic calcium and neurotransmitter release

Wu et al studied the contribution of $\mathrm{Ca^{2+}}$ channel subtypes to neurotransmitter release at the calyx of Held. Subtype specific toxins were used to identify the high-voltage activated P/Q-, N- and R-type channels ($\mathrm{Ca_v2.1}$, $\mathrm{Ca_v2.2}$, $\mathrm{Ca_v2.3}$) as mediators of the presynaptic $\mathrm{Ca^{2+}}$ current in animals younger than p10. In response to a single AP, P/Q-type channels account for ~50% of the measured $\mathrm{Ca^{2+}}$ -current ($\mathrm{I_{Ca}}$), N- and R-type for ~25% each. Immunocytochemical analysis of the channel distribution hinted at a differential distribution of these subtypes, with N- and R-type channels being predominantly located more distant from the release sites (Wu et al, 1999). The fraction of the $\mathrm{Ca^{2+}}$ current attributed to N- and R-type channels is reduced from p7 on and diminished by p10, leaving P/Q-type channels as the predominant subtype (Iwasaki and Takahashi, 1998; Iwasaki et al, 2000). Studies performed with α_{1A} -subtype-deficient mice, which lack P/Q-type currents, showed that the upregulation of N-type channel expression can partially compensate for the lack of P/Q channels. Interestingly, $\mathrm{Ca^{2+}}$ current facilitation was absent in these KO mice (Inchauspe et al, 2004; Ishikawa et al, 2005).

With its unusually large size, the calyx of Held serves as a good model system to study the intracellular calcium concentration ($[Ca^{2+}]_i$), especially its spatial and temporal characteristics after AP induced calcium influx, the calcium sensitivity of neurotransmitter release and the cell's mechanisms to control $[Ca^{2+}]_i$.

Borst and Sakmann measured the I_{Ca} induced by an AP waveform in whole-cell voltage clamp experiments. The recorded I_{Ca} had a Gaussian-like shape with a full width of half maximum (FWHM) of 0.36 ms with a peak of 2.5 nA and a charge integral of 0.92 pC on average at room temperature (RT). The I_{Ca} increased in amplitude and got briefer at physiological temperature (36 °C) (Borst and Sakmann, 1998).

Although single Ca^{2+} -channel conductances have not been obtained from the calyx of Held, about 25000 Ca^{2+} -channels would have to open to account for an I_{Ca} of 2.5 nA (assuming a maximal channel conductance of 0.1 pA as found for L-type channels). However, the open probability for P/Q-type channels is low and likely even lower when channel opening is governed by an AP. The total number of Ca^{2+} -channels is therefore expected to be much higher.

Direct access to the presynaptic terminal allows for dialysis of controlled amounts of Ca^{2+} -sensitive fluorescent dyes in order to measure the Ca^{2+} signal underlying release. Helmchen et al measured the resting $[Ca^{2+}]$ at the calyx of Held to be 40 nM at RT and estimated a calcium binding ratio (κ_s) of 40. This implies that 2.5 % of the Ca^{2+} ions entering the terminal after AP induced channel opening remain free after buffer equilibrium is reached. The volume averaged Ca^{2+} concentration will therefore only increase by 300 nM rather than 12 μ M (0.96 pC charge entering the terminal with a volume of $\sim 400 \, \mu m^3$). The decay time constant of the volume averaged $[Ca^{2+}]_i$ was measured to be $\sim 106 \, \text{ms}$, which represents the sum of clearance of Ca^{2+} from the cytosol (extrusion via the plasma membrane, uptake into organelles i.e. mitochondria) and unbinding from the buffer(s) (Helmchen et al, 1997). A recent study has highlighted the importance of the endogenous mobile Ca^{2+} buffer parvalbumine in shaping the $[Ca^{2+}]_i$ decay and estimated the decay time constant to be $\sim 30 \, \text{ms}$ in unperturbed calyces (Müller et al, 2007).

However, volume averaged Ca^{2+} measurements cannot reflect the complex Ca^{2+} transients that arise in the vicinity of Ca^{2+} channels. In order to probe the coupling of Ca^{2+} influx and vesicular release, exogenous buffers such as EGTA and BAPTA have been used as an assay for the diffusional distance of Ca^{2+} . These chelators can effectively reduce the diffusion of Ca^{2+} ions from the channel pore into the terminal. The mean distance an ion travels before being captured by the buffer depends on its on-binding rate (Neher, 1998). The on-binding rate of EGTA is ~160-fold smaller than BAPTA's, so that differential effects of the two buffers on transmitter release can infer information on channel-release-coupling (Naraghi and Neher, 1997). It has to be noted that Ca^{2+} indicator dyes also act as exogenous buffers. Their contribution, especially that of high affinity ones, to Ca^{2+} buffering within the cell cannot be neglected (see Neher, 2008).

Borst and Sakmann studied the effect of EGTA on neurotransmitter release at the calyx of Held. EPSC amplitudes were reduced by low millimolar concentrations of EGTA and up to 50% in $10\,\mathrm{mM}$ (Borst and Sakmann, 1996). The effectiveness of EGTA has previously been shown in chromaffin cells (Neher and Marty, 1982), whereas it is ineffective in the squid giant synapse where a very tight coupling of vesicles with single Ca^{2+} channels has been postulated (Adler et al, 1991; Augustine et al, 1991).

Cell attached recordings of the release face of the calyx-type synapse of the chick ciliary ganglion showed large Ca²⁺ conductances which were attributed to Ca²⁺ channel clusters (Stanley, 1991). Borst and Sakmann implicated that at the calyx of Held multiple Ca²⁺ channels govern the release of a single vesicle (channel domain overlap) (Borst and Sakmann, 1996). The measured EPSC amplitudes show a supralinear relationship to both peak I_{Ca} and its integral which supports the concept of overlapping domains (Borst and Sakmann, 1999). At the squid giant axon terminal, the recruitment of additional Ca²⁺ channels only leads to a linear increase in release, again illustrating the tight coupling of single channels and vesicles (Augustine et al, 1991).

There is another marked difference between the calyx and the squid giant axon. At the calyx of Held, it has been found that 1 mM BAPTA and 10 mM EGTA are similarly effective in blocking neurotransmitter release (Borst et al, 1995; Borst and Sakmann, 1996). This result cannot be explained by models that assume a homogeneous vesicle population (Naraghi and Neher, 1997), but is in line with models that assume a variable channel-vesicle distance among release sites within a single calyx (Meinrenken et al, 2002).

Transient and localized 'hot spots' of Ca^{2+} entry have been measured using n-aequorin-J, which emits blue light after binding Ca^{2+} , in squid (Llinás et al, 1992) and by using total internal reflection fluorescence microscopy (TIRFM) in retinal bipolar cells, saccular haircells (Zenisek et al, 2003) and chromaffin cells (Becherer et al, 2003). Although microscopy techniques have improved in both spatial and time resolution, there is, as of now, still no technique that would allow the exact measurement of an AP invoked $[Ca^{2+}]_i$ transient considering that it has a microdomain nature and fast kinetics (which would ideally be measured at the membrane/ site of Ca^{2+} entry alone).

Facing the difficulties in measuring the $[Ca^{2+}]_i$ transients directly, Heidelberger et al have relied on uniform elevations of $[Ca^{2+}]_i$ by flash photolysis of the Ca^{2+} chelator DM-nitrophen (DMn). The uniform and relatively long lived elevation of $[Ca^{2+}]_i$ simplifies its detection with fluorescent indicator dyes and allows to determine vesicular release rates in dependence of intracellular Ca^{2+} . The so-called 'reverse approach' can then be used to estimate the Ca^{2+} concentration that an average vesicle 'sees' before fusion by fitting the Ca^{2+} dependence of release with a kinetic model. The derived parameters of the model can then be used to infer the local $[Ca^{2+}]_i$ transient that drives an average vesicle to fuse after AP induced channel opening. Bollmann et al and Schneggenburger

and Neher have used this approach at the calyx of Held and estimated $[\mathrm{Ca^{2+}}]_i$ to rise to $10\text{--}25\,\mu\mathrm{M}$ with a half-width of $\sim\!0.5\,\mathrm{ms}$ (Bollmann et al, 2000; Schneggenburger and Neher, 2000). These numbers are much lower than the $200\text{--}300\,\mu\mathrm{M}$ indicated for the squid giant axon (Llinás et al, 1992).

Bollmann and Sakmann combined flash photolysis of DMn with the use of exogenous $\mathrm{Ca^{2+}}$ buffers and were able to create a uniform, but transient, elevation of $[\mathrm{Ca^{2+}}]_i$ (FWHM < 0.5 ms) in the terminal that could evoke a postsynaptic response similar to that of an AP, further emphasizing the role of $[\mathrm{Ca^{2+}}]_i$ in controlling the dynamics of release and the fidelity of synaptic strength (Bollmann and Sakmann, 2005).

Although the exact topology of Ca²⁺ channels and vesicles remains to be shown, multiple lines of evidence support that multiple channels contribute to the local elevations of [Ca²⁺]_i that drive fusion of a vesicle at the calyx of Held. The channel-vesicle distance is variable and can extent hundreds of nanometers (as opposed to the squid giant synapse). However, comparably low levels of [Ca²⁺]_i at the transmitter release site can support a high rate of release at the calyx of Held. This is due to the highly non-linear relationship between release and the Ca²⁺ signal driving it, so that two orders of magnitude increase in [Ca²⁺]_i from resting levels can support five orders of magnitude increase in the release rate (Lou et al, 2005).

Parameters determining synaptic strength

The first paired recording of a principle cell and its corresponding calyx terminal by Borst et al opened the possibility to tightly control and describe the parameters governing synaptic strength at the calyx of Held synapse (Borst et al, 1995). Synaptic strength can be described quantitatively with

$$PSC = N \cdot p \cdot q \tag{1.1}$$

which describes the amplitude of the postsynaptic current (PSC) as the product of the quantal size (q), the number of release sites (N) and the probability that release occurs at such a site (p) (del Castillo and Katz, 1954; Zucker, 1973). The parameter N of the binomial model is fixed, whereas the probability p can be interpreted as the joint probability for fusion of a synaptic vesicle $(p_{\rm rel.})$ and the availability of a synaptic vesicle at the release site $(p_{\rm avail.})$

This model opens the ability to separate the pre- and postsynaptic contributions to synaptic strength which can change dramatically in response to repetitive activity. This synaptic plasticity can manifest itself by an increase (facilitation) or decrease (depression) of subsequent release on various time scales, short- and long-term (reviewed in von Gersdorff and Borst, 2002). When the calyx of Held is stimulated by trains of APs at high frequency the EPSC is dominated by short-term depression (STD). Facilitation is observed under conditions of lowered release probability. These short-term changes, a characteristic attributed to all synapses, play an important role in relaying information in the nervous system. The quantitative description of N, p and q at the calyx of Held might help to provide a conceptional understanding of synaptic function when faithful transmission at high frequencies is required. All three parameters can contribute to the STD observed at the calyx of Held. Postsynaptic receptor desensitization and saturation would result in reduction in q. A reduction in Ca^{2+} -influx would cause a reduction in $p_{rel.}$, whereas the depletion of releasable vesicles would lower $p_{avail.}$ (reviewed in Schneggenburger et al, 2002).

It is important to note that the definition of 'release site' in the binomial model is ambiguous, their number N is estimated to lie between two numbers that can be determined experimentally: the number of morphological active zone (N_{az}) with a corresponding p_{az} that release occurs at such a site. On the other hand, N_{ves} describes the number of releasable vesicles with the probability p_{ves} that a given vesicle is released.

Using EPSC variance-mean analysis, an upper limit of ~600 active zones was estimated for the calyx of Held (Meyer et al, 2001; Scheuss et al, 2002). This number is comparable to the ~550 active zones that were identified by the EM study mentioned above (Sätzler et al, 2002). Trains of APs have also been used to estimate the number of releasable vesicles in the terminal. However, postsynaptic effects such as AMPA receptor desensitization and saturation manifest in a reduction of q during the train and will lead to an underestimation of N_{ves} . These postsynaptic effects can be controlled pharmacologically by adding cyclothiazide (CTZ) and kynurenic acid (Kyn) which lead to an estimate of ~900 releasable vesicles by EPSC variance-mean analysis (Scheuss et al, 2002). A single active zone is therefore, on average, used only once or twice during short trains of AP.

Pool sizes (N_{ves}) can also be estimated by non-physiological stimulations, either long lasting depolarizations (i.e. 50 ms) or flash photolysis of caged Ca^{2+} . Flash-induced

elevations of $[Ca^{2+}]_i$ to levels above 12 μ M gave a pool estimate of ~1800 vesicles when analyzing cumulative release on a 10 ms interval (Schneggenburger and Neher, 2000). Studies employing long depolarizations of the terminal to deplete the readily releasable pool (RRP) of vesicles combined with deconvolution analysis of the obtained EPSCs found N_{ves} to be ~2300 (Sakaba and Neher, 2001a,b). Even higher numbers, 3300–5200, were obtained by measurements of increases in cell capacitance in response to depolarizing stimuli of varied length (Sun and Wu, 2001). Additionally, the analysis of the covariance of EPSCs following AP trains also estimated N_{ves} to be ~2000 (Scheuss et al, 2002). The release of thousands of vesicles within a few milliseconds implies that multiple vesicles fuse at each active zone.

The quantal content (the number of quanta released in response to an AP) was estimated to be 210, 140 and 157 on average in three studies (Borst and Sakmann, 1996; Schneggenburger et al, 1999; Meyer et al, 2001). With the high number of release ready vesicles at the calyx of Held, a very low release probability p_{ves} would result which cannot explain the depression pattern observed at the calyx. Accordingly, simple depletion models fail to properly predict the experimental data (Weis et al, 1999). The model assumes a high release probability and that an invariable fraction of the releasable pool is consumed per AP. However, the decay in EPSC amplitudes during a train of APs observed at the calyx of Held is shallower than predicted by the model.

This inconsistency can be revoked if heterogeneity in the release probability is considered. Indeed, Wu and Borst and subsequently Sakaba and Neher showed that at the calyx of Held release probability is heterogeneous and that two subpools contribute equally to the pool of releasable vesicles (Wu and Borst, 1999; Sakaba and Neher, 2001a). 50% of vesicles released rapidly within a few milliseconds. Both pools possess distinct kinetic properties which allow their separation when analyzing the cumulative release in response to a sustained depolarization of the terminal. The release kinetics of the fast releasing pool of vesicles (FRP) are about 10 times faster than those of the slow releasing pool of vesicles (SRP).

Sakaba and Neher also showed that the recovery of the SRP is quick ($\tau = 200 \,\mathrm{ms}$) whereas the FRP recovers slowly. The latter is $\mathrm{Ca^{2+}}/\mathrm{calmodulin}$ dependent and can be blocked by infusing a calmodulin binding domain peptide (Fig. 1.2). Heterogeneity of release probability has been reported in various other systems by either uncaging induced $[\mathrm{Ca^{2+}}]_i$ elevations in tonic and phasic crustacean neuromuscular junctions

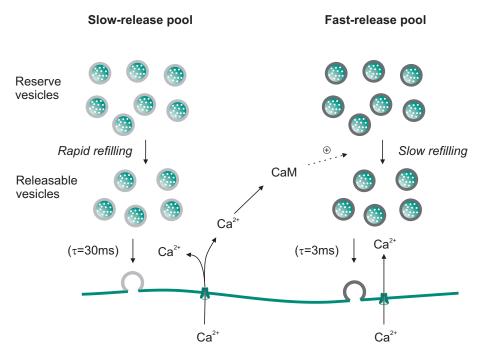


Figure 1.2: Vesicle pool dynamics at the calyx of Held. Two kinetically distinct pools can be identified at the calyx of Held when neurotransmitter release is stimulated by long depolarizations of the synaptic terminal. Vesicles from the slow releasing pool are rapidly refilled but release with a time constant \sim 10-fold slower than those from the fast releasing pool. Refilling of the latter is slow, but is accelerated by Ca²⁺/ calmodulin (CaM). Adapted from Matthews (2001).

(Millar et al, 2005), brief depolarizing steps in cultured autaptic hippocampal neurons (Rosenmund et al, 1993) and minimum stimulation in hippocampal CA1 cells (Dobrunz and Stevens, 1997).

Multiple mechanistic concepts could serve as an explanation for an apparent heterogeneity in p. Assuming that postsynaptic effects such as receptor desensitization and saturation are counteracted pharmacologically as described by Neher and Sakaba, all remaining causes should have a presynaptic origin (Neher and Sakaba, 2001). A variable Ca^{2+} channel/ vesicle topology as described above could explain this phenomenon.

Calcium-secretion coupling on a molecular level

Multiple proteins involved in neurotransmitter release have been identified in the last 30 years. Among them and attributed as key players for exocytosis are the plasma

membrane proteins syntaxin-1 and SNAP-25 as well as the vesicle protein synaptobrevin-2/VAMP (vesicle associated membrane protein). All three are members of the SNARE family of proteins (SNAREs are SNAP receptors, SNAPs are soluble NSF attachment proteins and NSF stands for N-ethylmaleimide sensitive factor).

Synaptobrevin, syntaxin and SNAP-25 form tight complexes that are generally accepted as being essential for membrane fusion. Neuronal SNARE complexes are formed by the parallel alignment of four helical SNARE motifs, one of which is contributed by synaptobrevin-2, one by syntaxin-1 and two by SNAP-25. According to the 'zippering hypothesis', the assembly is initiated by the interaction of synaptobrevin with heterodimers of syntaxin and SNAP-25, N-terminal to the SNARE motifs. The 'zippering' will then proceed towards the C-terminus. After fusion of the membranes, SNAPs-NSF serves to disassemble the SNARE complexes (reviewed in Jahn and Scheller, 2006). In this context, complexes that bring the vesicle membrane in apposition to the plasma membrane are referred to as *trans* complexes, whereas a *cis* complex resides in only one membrane.

In vitro studies employing liposome fusion assays have shown that synaptic vesicles are constitutively active fusion machines (Holt et al, 2008). Interestingly, the kinetics of fusion were solely influenced by the number of 1:1 acceptor heterodimers of syntaxin and SNAP-25 but no Ca^{2+} dependence was observed. Synaptotagmin-1, -2 and -9 are thought to present the Ca^{2+} sensors for synchronous neurotransmitter release observed in neurons (Xu et al, 2007). However, no additional increase in fusion kinetics mediated by synaptotagmin/ Ca^{2+} could be observed in *in vitro* assays (Stein et al, 2007).

This discrepancy can be attributed to multiple regulatory proteins that have been identified, in addition to the ones mentioned so far, as part of the active zone or direct interaction partners with the fusion machinery:

Munc13 (mammalian unc13) is an active zone protein thought to be essential for vesicle priming. Munc18 is a member of the Sec1/Munc18 (SM) family of proteins which are important for membrane trafficking and secretion. Although its precise role remains to be shown, Munc18 is essential for neurotransmitter release. Mice lacking Munc18-1 show normal brain development and synapse morphology, yet no transmitter release could be observed (Verhage et al, 2000). α RIMs (Rab3 interaction molecules) are Rab3 effectors in the active zone that bind Munc13 and also play a role in priming. Rab3s

are members of the Rab family, small monomeric GTPases, and exclusively found on synaptic vesicles.

Synaptotagmin is thought to be the Ca²⁺ sensor for release. It has been proposed that synaptotagmin is able to replace complexin from the SNARE complex after binding Ca²⁺. Complexin in turn is thought to act as a 'fusion clamp' that is able to stabilize a SNARE complex intermediate (reviewed in Rizo and Rosenmund, 2008).

The large number of regulatory proteins that have been identified so far illustrates that Ca²⁺ triggered exocytosis in nerve terminals critically depends on the timed interplay of multiple proteins at the active zone. As of now, this configuration cannot be reconstructed *in vitro*.

The molecular basis of vesicle fusion can, though to a limited extent, be studied at the calyx of Held as an accessible and hence well characterized model synapse. Specifically, the impact of molecular perturbations on the Ca²⁺ sensitivity of release can be studied.

Lou et al used the phorbol ester PDBu (which mimics diacylglycerols) to enhance the activity of presynaptic protein kinase C. Mapping $[Ca^{2+}]_i$ and release rates showed a potentiation of release. The increased Ca^{2+} sensitivity was interpreted as an increased 'fusion willingness' of synaptic vesicles.

Studying neurotransmitter release in genetically modified animals is difficult at the calyx of Held. Since the synapse only forms after birth, electrophysiological studies can only be done from postnatal day 6 on, an age that is not reached by many animals in which presynaptic proteins have been deleted. Synaptotagmin-2 KO animals are viable at p8–p10 and have been studied by Sun et al. The studied calyces lacked the stereotypical supralinear dependence of release on $[Ca^{2+}]_i$, but release at basal levels (< 1 μ M) was intact and slightly elevated. The remaining release was attributed to a second Ca^{2+} sensor for asynchronous release which exhibits a smaller cooperativity towards Ca^{2+} binding (Sun et al, 2007).

The accessibility of the calyx of Held for recording pipettes allows the perturbation of release by infusing peptides, antibodies and toxins into the terminal. Sakaba et al have done the latter to study the specific effect of clostridial neurotoxins onto release. Clostridial neurotoxins are zinc-endopeptidases that selectively cleave synaptobrevin, SNAP-25 and syntaxin. Seven serotypes of botulinum neurotoxin (BoNT) secreted by Clostridium botulinum have been identified (BoNT A–G) whereas C. tetani produces tetanus neurotoxin (TeNT). The BoNTs and TeNT are composed of a heavy (~100 kDa)

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and a light chain (~50 kDa) with the latter conferring the peptidase activity. The toxins cause severe to complete reduction in postsynaptic responses and severe reduction in spontaneous release at vertebrate and invertebrate synapses (reviewed in Humeau et al, 2000; Schiavo et al, 2000).

When the recombinant light-chains of either BoNT/C (which cleaves syntaxin) or TeNT (which cleaves synaptobrevin) were introduced into the terminal, an all-or-none block of release was observed. For these two toxins, no significant changes in the efficiency of Ca²⁺ to trigger release were found. BoNT/A infusion (which cleaves SNAP-25), however, lead to a 4-fold reduction in the Ca²⁺ sensitivity of release. The kinetics of release under BoNT/C are unchanged when release is triggered by long depolarizations. This is in marked contrast to TeNT, under which a preferential loss of the fast component of release could be shown. It was therefore postulated that cleavage of synaptobrevin might interfere with a 'maturation' step during which vesicles are incorporated in the vicinity of Ca²⁺ channel clusters to enable fast neurotransmitter release (Sakaba et al, 2005).

1.2 Motivation

Synapses in neural circuitry are not merely relay stations to pass on information coded in action potentials innervating their terminals. They themselves serve as modulators of information, processing units that can effectively filter or amplify the incoming signals. This modulation can be mediated on multiple time scales and the short-term effects were of key interest in the work that will be presented here. Accordingly, the strength of a synapse critically depends on the mechanisms governing this short-term plasticity during sustained high-frequency firing. How robust synapses can sustain activity will critically depend on the number of available vesicles, their release probability and the rate at which they are recruited to the plasma membrane.

Neurotransmitter release at the calyx of Held synapse, which was studied here, depresses in response to high-frequency stimulation which is attributed to the depletion of vesicles in the RRP (Rosenmund and Stevens, 1996; Dittman and Regehr, 1998). Interestingly, the vesicles found in the RRP are not released with equal probability when an AP innervates the terminal. In glutamatergic synapses, such as the calyx of

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Held and hippocampal synapses, a number of studies identified 'reluctant' or 'slow' vesicles (Wu and Borst, 1999; Hallermann et al, 2003; Moulder and Mennerick, 2005). Also, the presence of asynchronous release in response to an AP was taken as evidence for 'reluctant' vesicles (Sun et al, 2007). Multiple mechanisms could account for the apparent reluctance of vesicles to be released by an AP, two of which have gained much attention:

First, reluctant vesicles could be intrinsically different from fast releasing vesicles. This could be mechanistically attributed to i.e. a different Ca²⁺ sensor for release or an incomplete formation of the release apparatus (or in other words, a lack of fusion competence). For example, synaptotagmin-1, -2 and -9 have been proposed as mediators for fast and synchronous release, whereas an, as of now, unidentified second sensor is thought the mediate slow neurotransmitter release (Sun et al, 2007; Xu et al, 2007).

On the other hand, the coupling of vesicles to the source of the Ca^{2+} influx that triggers release, the Ca^{2+} channels, may be variable between populations of vesicles. Variable distances between Ca^{2+} channels and synaptic vesicles have been predicted for the calyx of Held in a modeling study (Meinrenken et al, 2002). In the scheme of a microdomain elevation of $[Ca^{2+}]_i$ around the channel pore, vesicles that are more distant from the channels would release only when sustained activity had led to an increase in global $[Ca^{2+}]_i$ and therefore release much slower (due to the supralinear relationship of release to Ca^{2+}).

These two hypothesis can be discerned when the increase of $[Ca^{2+}]_i$ is obtained by uniform uncaging of Ca^{2+} from a caged compound rather than being mediated by Ca^{2+} influx through Ca^{2+} channels in the plasma membrane. By making use of the differential recovery paradigm of the fast and slow component of release at the calyx of Held (Sakaba and Neher (2001a), also see Fig. 1.2), the fast releasing vesicles can be depleted so that the response of reluctant vesicles to strong depolarizations and flash uncaging can be studied in isolation and in contrast to each other.

Investigating this issue provides important implications for the study of synaptic transmission. If the former hypothesis is correct, one would expect that the choice of the Ca²⁺ sensor will determine the kinetics of transmitter release. The presence of a fast Ca²⁺ sensor would then allow for rapid release to occur, whereas a slow Ca²⁺ sensor would cause slower release of synaptic vesicles.

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If the latter hypothesis is correct, then 'molecular priming' of synaptic vesicles will not be the rate limiting step for fast neurotransmitter release, but rather the association of synaptic vesicles with Ca²⁺ channel clusters at the active zone. The degree of association would then govern the kinetics of release at different synapses.

Additionally, it has been shown that the recovery of fast releasing vesicles can be modulated in multiple ways, one of which being the differential block of this component mediated by introducing TeNT into the terminal. In this work, the implicated role of synaptobrevin in recovery of the fast component of release was studied further. The use of additional toxins and peptides served as a valuable tool to bridge the wealth of information on proteins involved in Ca²⁺-dependent exocytosis with the vesicle pool definitions that are based on the kinetic analysis of neurotransmitter release, in an attempt to identify the critical steps in vesicle fusion that mediate rapid neurotransmitter release at the calyx of Held.

Material and Methods

2.1 Slice preparation

The preparation of acute brain slices for electrophysiological recordings followed the general guidelines provided by Edwards et al (1989) and their specific application to obtain recordings from the medial nucleus of the trapezoid body (MNTB) as described by Forsythe and Barnes-Davies (1993b) and Borst et al (1995).

Postnatal day 8 to 10 (p8–10) Wistar rats were decapitated and the head was transferred into a dish containing ice cold artificial cerebrospinal fluid (aCSF) containing (in mM): 125 NaCl, 2.5 KCl, 25 NaHCO₃, 1.25 NaH₂PO₄, 25 glucose, 3 MgCl₂, 0.1 CaCl₂, 0.4 L-ascorbic acid, 3 myo-inositol, 2 sodium pyruvate (according to von Gersdorff et al (1997), modified from Borst et al (1995)). Using this cold, low Ca²⁺, saline was meant to prevent anoxia and should reduce the metabolic activity of cells.

The skull was opened by carefully cutting along the midline from the neck to the eye line. Cuts along the mediolateral axis exposed the cerebrum and the cerebellum. Using a scalpel, two cuts (one coronal and close to the eyeline, the second slightly tilted close to the rostral-dorsal base of the cerebellum, see Fig. 2.1) allowed to remove the entire cerebrum. The remaining brainstem and cerebellum could then carefully be pushed caudally and out of the skull. Any fibers still connected to the brain stem were severed with forceps. In order to facilitate the slicing, fine forceps were used to remove the pia mater and the arachnoid mater. The location of the MNTB region could be estimated upon careful inspection of the ventral brainstem surface where the

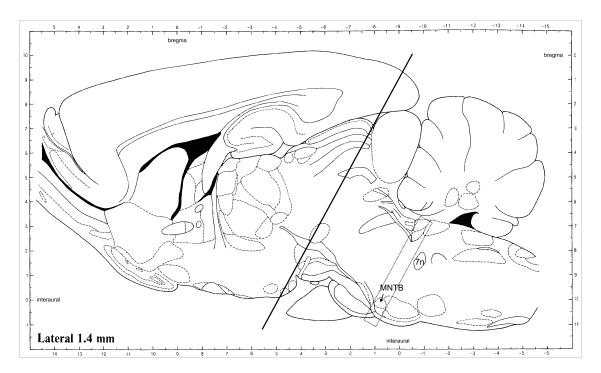


Figure 2.1: Lateral view of the adult rat brain. The diagonal line indicates the cut that was performed to separate the cerebrum from the brainstem. The latter was then mounted for slicing. The region of interest is highlighted by the dashed box and contains the medial nucleus of the trapezoid body (MNTB). While slicing, the seventh cranial nerve (7n) served as a landmark that indicated the beginning of the region of interest. Modified from Paxinos and Watson (1982).

crossing fiber bundles leave a white band which becomes more prominent with age (and increased myelination).

The preparation, containing the brain stem and the cerebellum, was then glued onto the Teflon base of a slicing chamber using cyanoacrylate glue, oriented so that the caudal side faced up and the ventral side of the brainstem faced the slicer's blade.

Transverse brain slices were made using a vibratome slicer (Integraslice 7550 MM, Campden Instruments, Leicester, UK) with the horizontal vibration set at 70 Hz and advancing $0.18 \, ^{\text{mm}}/\text{s}$. Initial slicing was performed at higher speed and thicker than the final desired slices since they did not contain the MNTB region. The 7^{th} cranial nerve (facial nerve) served as a landmark to the experimenter and was usually visible unilaterally. A slight tilt in the mounted preparation could be corrected by rotating the Teflon base. Once the 7^{th} nerve was visible, $200 \, \mu \text{m}$ thick slices were collected. The

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MNTB region was usually contained in the 2nd to 5th slice (Fig. 2.1). The exact number of slices critically depended on the size and age of the animal.

Slices containing the MNTB were transferred to an incubation chamber containing aCSF at 37 °C using a glass suction pipette and placed onto a nylon mesh. The aCSF solution was identical to the one mentioned above except that $2\,\mathrm{mM}$ CaCl₂ and $1\,\mathrm{mM}$ MgCl₂ were used. The solution was continuously bubbled with carbogen gas (95 % O₂, 5 % CO₂). At 37 °C, slices were viable for 3–4 hours.

2.2 Solutions

2.2.1 Extracellular solution

The extracellular solution for electrophysiological recordings was the same as the solution used for incubation (in mM): 125 NaCl, 2.5 KCl, 25 NaHCO₃, 1.25 NaH₂PO₄, 25 glucose, 1 MgCl₂, 2 CaCl₂, 0.4 L-ascorbic acid, 3 myo-inositol, 2 sodium pyruvate (NaCl, KCl, NaHCO₃, MgCl₂ and CaCl₂ from Merck (Darmstadt, Germany), all others from Sigma (Steinheim, Germany)). This aCSF had an osmolarity of ~310 mOsm with a pH of 7.4 when bubbled with carbogen. Depending on the type of experiment, additional chemicals were added. All experiments utilized 10 mM tetraethylammonium (TEA) chloride to block potassium channels and 1 μ M of the sodium channel blocker tetrodotoxin (TTX). Non-AMPA type glutamate channels were blocked with 50 μ M of the competitive NMDA antagonist D(-)-2-amino-5-phosphonopentanoic acid (D-AP5), effectively isolating postsynaptic AMPA receptor mediated excitatory postsynaptic currents (EPSCs).

Strong non-physiological stimulations, such as sustained depolarizations of the terminal or flash uncaging, lead to massive neurotransmitter release. 100 μM cyclothiazide (CTZ) was therefore used to prevent AMPA receptor desensitization in all experiments (Neher and Sakaba, 2001). To prevent AMPA receptor saturation, the competitive receptor antagonists γ-D-glutamylglycine (gDGG) or kynurenic acid (Kyn) were used. 1 mM Kyn reduces the mean miniature excitatory postsynaptic current (mEPSC) amplitude by 50 %. Since Kyn absorbs ultraviolet (UV) light and therefore interferes with flash experiments, 2 mM gDGG was used in these cases which reduces the mean mEPSC amplitude to 30 %. To detect mEPSC faithfully, neither Kyn nor gDGG were used when mEPSC were recorded. In the latter type of experiments, excitatory mEPSC were

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isolated by adding $30\,\mu\mathrm{M}$ bicuculline (to block GABA_A receptors) and $5\,\mu\mathrm{M}$ strychnine (as antagonist for glycine receptors) to the extracellular solution. TEA was purchased from Sigma, TTX from Alamone Labs (Jerusalem, Israel). D-AP5, CTZ, gDGG, Kyn, bicuculline and strychnine came from Tocris (Bristol, UK).

2.2.2 Intracellular solutions

The principle composition of the pipette solutions was taken from Schneggenburger and Neher (2000) and Sakaba and Neher (2001a). The stock solution used for postsynaptic recordings and presynaptic recordings that did not involve Ca²⁺ measurements contained (final concentration in mM): 100 Cs-gluconate, 10 HEPES, 5 Na₂-phosphocreatine, 4 MgATP, 0.3 NaGTP, 20 TEA-Cl and was prepared as a 4x stock. The pH was adjusted to 7.2 using CsOH (Cs-gluconate was custom-made, all others from Sigma). Additional substances could be added to this stock solution and the final volume was established by adding purified water (Milli-Q Plus, Millipore, Billerica, USA) as with all other solutions presented herein. For postsynaptic recordings, 5 mM Cs-EGTA (from Sigma, pH adjusted with CsOH) and 40 mM Cs-gluconate were added to adjust the osmolarity to ~320 Osm prior to experiments.

The presynaptic pipette solution contained (final concentration in mM): 100 Cs-gluconate, 20 HEPES, 5 Na₂ATP, 0.5 NaGTP, 0.5 MgCl₂ and 20 TEA-Cl, also prepared as a 4 x, pH adjusted, stock solution. For experiments in which transient elevations of [Ca²⁺]_i were desired, 2 mM DM-nitrophen (DMn) (Calbiochem, La Jolla, CA, USA), 1.7 mM CaCl₂, 1 mM Cs-EGTA and 0.5 mM Oregon Green BAPTA-5N (OGB-5N) (Molecular Probes, Eugene, Oregon, USA) were added. An additional 35 mM Cs-gluconate raised the osmolarity to ~320 Osm.

Step-like elevations of $[Ca^{2+}]_i$ were achieved by adding 1–3 mM DMn, 0.8–2.7 mM $CaCl_2$, 0.2 mM fura-2FF (Teflabs, Austin, TX, USA) and 30 mM Cs-gluconate.

For mEPSC measurements, fura-2FF was substituted with fura-4F (Molecular Probes) to be able to determine $[\mathrm{Ca^{2+}}]_i$ around resting levels and $< 1\,\mu\mathrm{M}$. For some experiments, free $\mathrm{Ca^{2+}}$ in the pipette solution was 'clamped' to an estimated 100 nM by adding 10 mM EGTA and 4 mM $\mathrm{CaCl_2}$.

The free Ca²⁺ concentration in solution was estimated by a computational routine kindly provided by Dr. Sørensen (Xu et al, 1997).

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Peptides and toxins: Recombinantly produced light chain botulinum neurotoxin serotype B (BoNT/B) was kindly provided by Dr. Stein and stored as a $59\,\mu\text{M}$ ($3\,\text{mg/ml}$) stock solution containing (in mM): 130 CsCl, 10 HEPES and 1 DTT at pH 7.2.

Table 2.1: Synthetic peptides

Peptide	Sequence	Notes
Rp3a(650–669)	AKGERLKHWYECLKNKDKKI	amino acids 650–669 of rat rabphilin3a
Rp3a(s650–669)	KERDKIKWECKHLLKNYGAK	scrambled version of Rp3a(650–669)
Syt2(383–401)	ATGTELRHWSDMLANPRRP	homologue of Rp3a(650–669) in rat synaptotagmin-2

All synthetic peptides were obtained from Biosyntan (Berlin, Germany) and delivered as lyophilized TFA salts. Desalting was performed on a short column packed with Sephadex G-10 (GE Healthcare, Chalfont St. Giles, UK). The Sephadex media was swelled in 165 mM CsCl, 10 mM HEPES and 1 mM DTT according to the manufacturer's guidelines. A 20 cm · 1 cm column was packed with ~10 cm of medium. Packing and sample application followed the guidelines provided in the Amersham Biosciences gel filtration handbook. 10 mg peptide were dissolved in 500 μ l buffer and eluted at the void volume of the column, collecting 500 μ l fractions. The protein concentration was determined using 1:20 dilutions of the sample in 8 M guanidinium hydrochloride in a UV spectrometer by measuring the absorption spectrum from 250–320 nm. The measured absorption was then correlated to the predicted one as calculated by the ExPASy tool ProtParam to determine the peptide concentration in a given fraction. Fractions were pooled if possible and resulted in stock concentrations of 4.58 mg/ml (1.83 mM) for Rp3a(650–669), 6.88 mg/ml (2.76 mM) for Rp3a(s650–669) and 7.3 mg/ml (3.31 mM) for Syt2(383–401).

Control experiments for both peptide and toxin experiments were performed by adding equivalent amounts of their respective buffer solutions to the intracellular solution.

2.3 Electrophysiological recordings, calcium uncaging and calcium imaging

After 40–60 min incubation time, slices were transferred to a custom build recording chamber with a glass base. A U-shaped platinum wire with parallel Nylon threads spanned across it was used to hold the slice in place. The recording chamber connected to a gravity driven perfusion system with adjustable flow rate which contained both plain extracellular solution and extracellular solution supplemented with the drugs listed in 2.2.1. After establishing a whole-cell patch clamp configuration on a postsynaptic principle cell, the bath solution was exchanged within 3–4 min for the latter solution.

Recording pipettes were made from borosilicate glass filaments (2 mm outer- and 1.4 mm inner diameter from Hilgenberg, Malsfeld, Germany) using a temperature controlled two-step vertical pipette puller (PIP 5 from HEKA, Lambrecht/Pfalz, Germany). Whole-cell voltage clamp recordings were performed at room temperature (RT) using either EPC9/2 or EPC10/2 amplifiers controlled by Patchmaster (v. 2.20, HEKA). Pipette- and other stray capacitances as well as cell capacitance were compensated using the built-in compensation mechanisms. The series resistance (R_s) for postsynaptic recordings was generally 5–10 M Ω and 8-20 M Ω for presynaptic patches. The series resistance was also compensated electronically by setting the fractional compensation such that the remaining resistance was 2.5–3 M Ω for post- and 8 M Ω for presynaptic recordings. The remaining R_s was corrected offline (see 2.4). Data were sampled at 20–50 kHz and low-pass filtered (Bessel, 6 kHz corner frequency).

2.3.1 Experimental setup

Two electrophysiological setups were used: One consisted of an Axioskop FS (Carl Zeiss, Jena, Germany) equipped with an infrared (IR) differential interference contrast (DIC) system. The slice was centered around the midline crossing point of the trapezoid body using a Zeiss PlanNEOFLUAR air objective with 2.5 x magnification. Cell pairs were selected and imaged using a 60 x water immersion objective (LUMPlan FI/IR, 0.9 N.A. from Olympus, Tokyo, Japan).

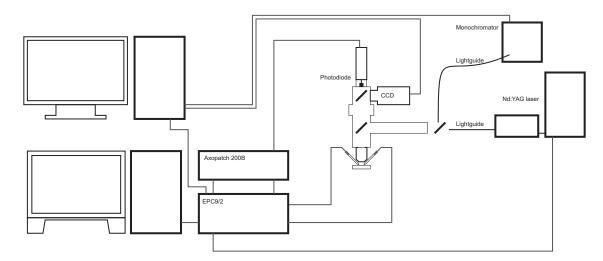


Figure 2.2: Experimental setup for UV-laser uncaging: A setup for slice electrophysiology was equipped with a photodiode for fast Ca²⁺ measurements as described by Bollmann (2001). The fluorescence signal was sampled by a computer which also controlled the UV-laser and the patch clamp amplifier. Both light sources were coupled into the epifluorescence port of the microscope via light guides. A CCD camera was employed for visual control of the electrophysiological recordings.

The second setup followed the principle design outlined above. It consists of an Olympus BX51WI microscope equipped with IR Dodt gradient contrast (DGC) illumination (Luigs & Neumann, Ratingen, Germany) and the UPlanFL N 4x and LUMPlan FI/IR 60x objectives (both Olympus).

Both setups were equipped with fluorescence imaging systems consisting of a monochromator (Polychrome V) and a 2-port condenser that coupled the light coming from the monochromator into the epifluorescence port of the microscope. Images were acquired by a CCD camera (Imago VGA). Both camera and monochromator were software controlled by TILLvisIO (all from TILL Photonics, Gräfelfing, Germany). Calcium uncaging was achieved via Xenon flash lamps that produced a brief (1–1.5 ms) flash of UV light (DP-10 or JML-C2 by Rapp OptoElectronic, Hamburg, Germany). The flash lamp was coupled into the second port of the epifluorescence condenser via a light guide. Quartz glass was used to couple both light paths into the microscope. Its reflective properties result in an attenuation of the monochromator light (passing 8% into to microscope), but 92% of the UV flash light are transmitted.

The photometric system used to measure transient elevations of [Ca²⁺]_i was adapted from Bollmann (2001). The fluorescence signal was detected using a low-noise, high sensitivity PIN photodiode (S5973-02, Hamamatsu, Hamamatsu, Japan) mounted into a Viewfinder III (TILL Photonics). The viewfinder passes 95 % of the 400–590 nm light onto the detector whereas red and IR light is directed towards the CCD camera to visualize the preparation. A rectangular aperture allowed one to limit the window from which the fluorescence signal was collected and was adjusted to match the size of the calyx of Held under investigation. An Axopatch 200B patch-clamp amplifier (Molecular Devices, Sunnyvale, CA, USA) served as a current-to-voltage converter and signal amplifier. Its signal was then fed into the EPC9/2 for data acquisition. For these experiments, a frequency-tripled Nd:YAG laser (Continuum, Santa Clara, CA, USA) was used. The peak laser output of 2 mJ was attenuated by the built-in adjustable energy attenuator and the components in the optical path so that peak amplitudes < 100 µJ were measured in the focal plane of the objective. Additional control of the laser output could be achieved by altering the timing of the quality switch (Q-switch). A shorter Q-switching time leads to reduced charging of the laser medium and therefore a decreased laser output.

2.3.2 Measurements of intracellular calcium

Ca²⁺ uncaging

Homogeneous elevations of cytosolic [Ca²⁺] were achieved by photolyzing the Ca²⁺ cage DMn (Kaplan and Ellis-Davies, 1988). The cell-impermeant form of DMn was dialyzed into the terminal via the patch pipette. Due to its high affinity for Ca²⁺ ($K_D = 5 \, \text{nM}$), DMn can be loaded with Ca²⁺ to a high degree while still maintaining low levels of free Ca²⁺ (~100 nM). The product of extinction coefficient (ε) and quantum yield (Φ) of DMn is comparatively high. Consequentially, the use of UV flash lamps allows one to elevate [Ca²⁺]_i to levels of 30 μ M and higher. DMn shows a 600000-fold reduced affinity to Ca²⁺ after photolysis ($K_D = 3 \, \text{mM}$) (reviewed in Ellis-Davies, 2007, 2008). These properties make DMn the prime candidate for uncaging studies at the calyx of Held. However, it has to be noted that DMn has a relatively high affinity for Mg²⁺ ($K_D = 2.5 \, \mu$ M). The presynaptic pipette solution used here contained only 0.5 mM

 MgCl_2 which effectively results in lowered levels of MgATP available to the cell. DMn is therefore not suitable in environments where [Mg²⁺] cannot be tightly controlled (see Ellis-Davies, 2006) or when physiological levels of MgATP are required.

To determine the purity of the DMn stock solution, a titration series with increasing levels of Ca²⁺ was made. The Ca²⁺ indicator fura-4F was used to determine the amount of Ca²⁺ that saturated the available fraction of chelating DMn by measuring the fluorescence ratio in microcuvettes. The purity was determined for each batch of DMn and varied among lot numbers from 70–80%. The error introduced by impure DMn was compensated by using proportionally more of the stock solution. All concentrations of DMn given herein reflect the effective concentration of DMn in the pipette solution.

Single wavelength dye

Experiments in which fast-decaying Ca²⁺ transients were recorded made use of the Ca²⁺ indicator dye OGB-5N which was perfused into the presynaptic terminal via the patch pipette. OGB-5N is a low affinity dye ($K_{\rm D}=20\,\mu{\rm M}$) with an excitation/ emission spectra like fluorescein (excitation maximum at 492 nm, emission maximum at 517 nm) according to the manufacturer. Faas et al measured $k_{\rm on}$ to be ~2.5 × 10⁸ M⁻¹s⁻¹ and a $k_{\rm off}$ around 8 × 10³ s⁻¹ (Faas et al, 2005).

Fluorescence traces collected via the photodiode were expressed as $\Delta F/F$ and corrected for both the dark-current of the photodiode and the background fluorescence of the preparation. The latter was determined by retracting both pipettes (excised patch configuration) after recordings were completed and placing them onto the slice in a nearby location from which a fluorescence signal was then recorded.

Dual wavelength dyes

The ratiometric Ca^{2+} indicator dyes used in this study all belong to the fura-2 family (reviewed in Takahashi et al, 1999). Fura-2 is an UV excited indicator that shifts its absorption maximum upon Ca^{2+} binding. Peak absorption for Ca^{2+} free solution is 362 nm and 335 nm at maximal $[Ca^{2+}]$ with an emission peak at ~510 nm (Grynkiewicz et al, 1985). This feature can be used to derive $[Ca^{2+}]$ from ratiometric measurements rather than relying on the increased fluorescence signal that single wavelength dyes emit

upon binding of Ca²⁺. By measuring the dye signal at two wavelengths, [Ca²⁺] values can be derived from their ratio which is essentially independent of dye concentration.

Grynkiewicz et al described the calibration equation as

$$[\mathrm{Ca}^{2+}] = K_{\mathrm{D}} \frac{R - R_{\mathrm{min}}}{R_{\mathrm{max}} - R} \left(\frac{S_{\mathrm{f2}}}{S_{\mathrm{b2}}} \right)$$

The ratio R signifies the fluorescence signal measured at two wavelengths λ_1 and λ_2 : $R = F_{\lambda 1}/F_{\lambda 2}$. S indicates a proportionality coefficient that depends on dye properties and properties of the measurement system (i.e. excitation intensity, extinction coefficient and quantum efficiency). Four constants $S_{\rm fl}, S_{\rm f2}, S_{\rm b1}, S_{\rm b1}$ describe both ${\rm Ca}^{2+}$ free and ${\rm Ca}^{2+}$ bound states at the two wavelengths.

Experimentally, values for $R_{\rm min}$, $R_{\rm max}$ and $K_{\rm eff}$ are determined by performing ratiometric measurements in solutions for which the free [Ca²⁺] is known. Taking $K_{\rm eff} = K_{\rm D} \cdot \frac{S_{\rm f2}}{S_{\rm b2}}$ simplifies the above equation to:

$$[\mathrm{Ca}^{2+}] = K_{\mathrm{eff}} \frac{R - R_{\mathrm{min}}}{R_{\mathrm{max}} - R}$$

The cytosolic $[Ca^{2+}]$ can therefore be inferred by measuring fluorescence ratios $R = F_{350\,\mathrm{nm}}/F_{380\,\mathrm{nm}}$ after the calibration parameters have been determined. The experiments presented herein made use of two derivatives of fura-2. As a rule of thumb, the affinity of a given dye to Ca^{2+} should relate to the $[Ca^{2+}]$ in question such that $0.1 \cdot K_D \leq [Ca^{2+}] \leq 10 \cdot K_D$ is satisfied. Fura-4F $(K_D = 800\,\mathrm{nM})$ was therefore used as a high affinity dye to measure $[Ca^{2+}]_i$ around resting levels and in experiments that employed weak uncaging flashes (Gee et al, 2000). Fura-2FF (originally described as fura-F by London et al and also marketed as fura-FF) is a low affinity derivative of fura-2 with two added fluorines on the BAPTA moiety (London et al, 1994). A K_D of $\sim 6-10\,\mu\mathrm{M}$ makes it the ideal dye to measure flash induced elevations of cytosolic Ca^{2+} .

In vitro calibration: The in vitro calibration was performed according to Schneggenburger (2005, chap. 53). The solutions used are based on the intracellular solution for fluorescence measurements (see 2.2.2) and were measured in microcuvettes. Since DMn absorbs UV light, it was included in the calibrations solutions to mimic the solution used for actual experiments. The composition of the solutions used for calibration of

fura-2FF is shown in Table 2.2. The calibration of fura-4F was performed according to the same principle.

Table 2.2: Solutions for *in vitro* calibration of fura-2FF. EGTA, DPTA (Fluka, Buchs, Switzerland) and DPTA/ $\mathrm{Ca^{2+}}$ was taken from stock solutions with pH 7.2 (adjusted with CsOH). The osmolarity was elevated to 320 mOsm with Cs-gluconate. The free [$\mathrm{Ca^{2+}}$] of the solution to measure $\mathrm{R_i}$ was calculated to be 10.5 $\mu\mathrm{M}$.

Substance	Final concentration [mM]
R _{min}	
intracellular solution	
Fura-2FF	0.05
DM-nitrophen	1
EGTA	10
R_i	
intracellular solution	
Fura-2FF	0.05
DM-nitrophen	1
DPTA, 50 % loaded with CaCl ₂	5
DPTA	5
R_{max}	
intracellular solution	
Fura-2FF	0.05
DM-nitrophen	1
CaCl ₂	10

2.4 Data analysis

Electrophysiological data were analyzed using Igor Pro (v. 5.0.5.7, Wavemetrics, Lake Oswego, OR, USA). Recordings were imported into Igor via the patcher's powertools routines kindly provided by Mr. Würriehausen. Remaining R_s after online correction (see 2.3) was compensated according to Traynelis (1998). In order to assess mEPSCs, a template matching algorithm as described in Clements and Bekkers (1997), kindly provided by Dr. Taschenberger, was used. Average data are presented as mean \pm standard error of the mean (SEM).

2.4.1 Deconvolution

For the experiments presented here, presynaptic release rates were inferred from EPSC recordings. This assay is only legitimate under the assumption that mEPSCs add up linearly to form the EPSC and that each measured mEPSC corresponds to a single presynaptic fusion event. As mentioned, postsynaptic receptor desensitization and saturation would violate this principle since both would lead to changes in mEPSCs during a prolonged stimulation and were therefore counteracted pharmacologically in this study. Additionally, such stimulations lead to excess glutamate in the synaptic cleft, which would be detected postsynaptically, yet does not represent additional release. This 'spill-over current' is caused by delayed clearance of glutamate and has to be subtracted from the compound EPSC prior to calculating the presynaptic release rate.

In order to derive release rates, the deconvolution method as described in Neher and Sakaba (2001); Sakaba and Neher (2001b) was used. The routine was implemented in Igor Pro and kindly provided by Dr. Neher and Mr. Würriehausen. A 'template recording' consisting of depolarizations to 0 mV of varied length was used to invoke EPSCs of variable amplitudes. The parameters for mEPSC kinetics and spill-over current were then adjusted manually, so that the spill-over current accounted for the decay phases of the EPSCs. The resulting estimate for the spill-over current was then subtracted from the EPSC. Deconvolution with the mEPSC then yields the quantal release rate. The assumed underlying mEPSC distribution was scaled according to the means taken to reduce postsynaptic non-linearities. For CTZ alone, a mean mEPSC amplitude of 30 pA was used, 15 pA for CTZ and Kyn and 10 pA for CTZ and gDGG.

The integral of the release rate derived from deconvolution yields the cumulative release of vesicles over time. For further analysis, the cumulative release was corrected for vesicles that had been recruited to the pool in the course of a given stimulus protocol (Sakaba and Neher, 2001b).

2.4.2 Kinetic release models

As mentioned in 1.1.2, kinetic models are a valuable tool to estimate release rates in response to the Ca^{2+} waveform driving vesicle fusion. Its reverse analogy allows to adjust the Ca^{2+} waveform so that the model prediction matches experimental release

rates in order to determine amplitude and time course of the Ca²⁺ signal at the Ca²⁺ sensor (Bollmann et al, 2000; Schneggenburger and Neher, 2000).

$$\mathbf{B_0} \xrightarrow{5k_{\mathrm{on}}[\mathrm{Ca}^{2+}]} \mathbf{B_1 \cdots B_4} \xrightarrow{k_{\mathrm{on}}[\mathrm{Ca}^{2+}]} \mathbf{B_5} \xrightarrow{\gamma} \text{fused}$$
 (2.1)

The model used by Schneggenburger and Neher represents a minimal sequential model describing the consecutive binding of 5 Ca²⁺ ions. The occupancy of each binding state is governed by the rate constants $k_{\rm on}$ and $k_{\rm off}$ as well as a cooperativity factor b which is necessary to account for the supralinear dependence of release on $[{\rm Ca^{2+}}]_i$ (Heidelberger et al, 1994). Fusion occurs irreversible from the fully occupied state at rate γ (see scheme 2.1). This kinetic model was translated into a set of rate equations that describe the occupancy of each binding site and implemented in Igor Pro and Matlab (v 7.6, Mathworks, Natick, MA, USA). It was solved numerically with a built-in algorithm for stiff problems available in both programs. The release rate per vesicle was determined by differentiation of the cumulative release (the 'fused' state in scheme 2.1).

Release rates predicted by a kinetic model can be compared to experimental data by either multiplying with an average pool size or by dividing the peak release rate obtained experimentally by the pool size estimate for the given cell (*release rate per* vesicle).

Although this sequential model is minimalistic in nature, it sufficiently describes the steep $\mathrm{Ca^{2+}}$ dependency of release identified at the calyx of Held. Lou et al (2005) showed however that the sequential model fails to predict release rates at $[\mathrm{Ca^{2+}}]_i$ levels $< 1\,\mu\mathrm{M}$ which have been found to be several orders of magnitude higher than predicted. This is owed to the model assumption that fusion can only occur when all binding sites have successively been occupied in scheme 2.1.

Higher release rates at basal levels of $[Ca^{2+}]_i$ can be accounted for if one allows fusion to occur from a sensor that is not fully occupied. Changes between interconvertible

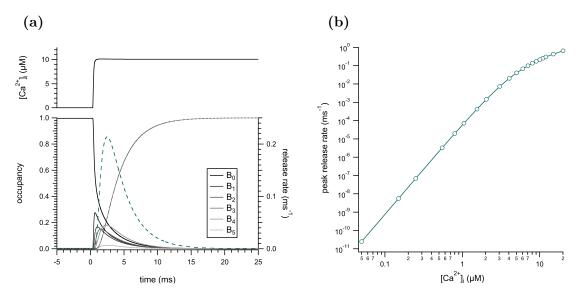


Figure 2.3: Example of the minimal kinetic release model described by Schneggenburger and Neher (2000). (a) shows the occupancy of each binding site in scheme 2.1 over time (bottom) in response to a simulated Ca²⁺ waveform after flash induced Ca²⁺ uncaging (top). The dotted line represents the 'fused' state (or cumulative release) whereas the superimposed dashed line shows its derivative, the release rate. (b): Dose-response-curve plotting peak release rates obtained from a simulation as in (a) against various levels of [Ca²⁺]_i on a double logarithmic scale. The model parameters in this example were taken from Fig. 3.4(d).

states in the absence of a regulator are a feature of an allosteric model (also called *MWC model*, reviewed in Changeux and Edelstein (1998)).

In scheme 2.2, the **B** and **C** states correspond to the 'tense' and 'relaxed' states in the MWC model. Fusion can only occur from the 'relaxed' states. All **C** states have a finite probability to lead to fusion, which can occur without binding Ca^{2+} (B_0), from partially filled states ($B_{1...4}$) or a fully occupied one (B_5 in scheme 2.2). The equilibrium constant of the reaction from states **B** to **C** increases by a factor f with each successive Ca^{2+} ion that is bound. Also, the binding affinity for Ca^{2+} is larger in the **C** states. A cyclic mechanism, without an energy input, has to conform to the demand for microscopic reversibility. The dissociation constant for agonist binding in the relaxed state is therefore multiplied by f^{-1} . As a result, going from B_0 to C_1 clockwise or counterclockwise is energetically equivalent. Fusion is then promoted irreversibly from the **C** states and limited by rate γ as in scheme 2.1.

In Lou et al (2005), a simplified allosteric model was adopted for which scheme 2.1 was extended by allowing fusion to occur irreversible from each \mathbf{B} state mediated by a single forward rate l_+ and factor f, as in scheme 2.2.

Results

3.1 Heterogeneity in release probability at the calyx of Held

Heterogeneity in release probability at the calyx of Held was first reported by Wu and Borst. They used voltage-clamp to depolarize the terminal via two action potential waveforms (APWs) with an interstimulus interval of 1 s. Both APWs were followed by a 10 ms step depolarization meant to deplete the releasable vesicles. It was shown that the second APW can only trigger a small amount of release during recovery from the induced depression. The following step however elicited a substantial excitatory postsynaptic current (EPSC). Wu and Borst concluded that vesicles are rapidly replenished after depression of the releasable pool, yet are 'reluctant' and can therefore only be released by an increase in the global $[Ca^{2+}]_i$ (Wu and Borst, 1999).

This heterogeneity in p_{ves} can also be revealed when the calyx of Held is exposed to a 50 ms step depolarization in the presence of 0.5 mM EGTA which has been shown to effectively block overlapping facilitation of release (Sakaba and Neher, 2001a). Figure 3.1 shows an example for this type of stimulation. The resulting AMPA-receptor mediated EPSC was deconvolved as described in 2.4.1 to yield the vesicular release rate. Its integral gives the cumulative release over time. Figure 3.1(b) illustrates that this kind of stimulation depletes almost all available vesicles. Accordingly, the readily releasable pool (RRP) is defined herein as comprising vesicles that can be released by strong stimulations such as sustained depolarizations of the presynaptic terminal. When the cumulative release trace is fitted with a biexponential function after correction for pool refilling, two time constants of release can be obtained. They differ roughly by a

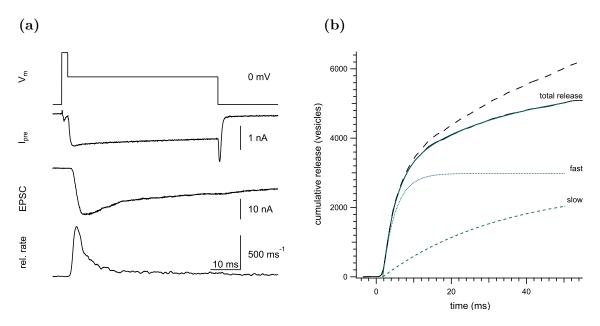


Figure 3.1: (a) A 50 ms step-depolarizations depletes the RRP at the calyx of Held. Traces shown are (from top to bottom): presynaptic membrane potential, presynaptic Ca²⁺ current, EPSC and vesicle release rate. The presynaptic terminal was depolarized to +70 mV for 2 ms and then to 0 mV for 50 ms to deplete the readily releasable pool. A pre-depolarization to +70 mV maximizes Ca²⁺ channel opening without causing influx of Ca²⁺ ions. 0.5 mM EGTA was added to the presynaptic pipette solution to facilitate the differentiation between fast and slow releasing vesicles. CTZ and Kyn were added to the presynaptic pipette solution to block postsynaptic AMPA receptor desensitization and saturation. (b) cumulative release in response to the 50 ms step-depolarization shown in (a) (dashed black line) obtained after integrating the vesicular release rate. The cumulative release was corrected for pool refilling as described in 2.4.1 (solid black line). The double exponential fit to the corrected cumulative release trace is shown in green and can be separated into the contributions of the fast releasing pool (dotted green line) and slow releasing pool of vesicles (dashed green line).

factor of 10 (2–3 ms and 20–30 ms respectively) and are attributed to the fast and slow component of release. The heterogeneity among vesicles of the RRP can therefore be ascribed to the presence of two subpools, the fast releasing pool of vesicles (FRP) and the slow releasing pool of vesicles (SRP).

Figure 3.2 shows an experiment similar to the one by Wu and Borst. A calyx terminal and its corresponding medial nucleus of the trapezoid body (MNTB) principle cell were double voltage-clamped at $-80\,\mathrm{mV}$. A square pulse to $+40\,\mathrm{mV}$ for 1.5 ms was used to mimic an action potential (AP). The subsequent 10 ms step depolarization to $0\,\mathrm{mV}$



Figure 3.2: Release evoked by AP-like stimulation is blocked when the vesicles from the fast releasing pool of vesicles have been depleted. Traces shown are (from top to bottom): Presynaptic membrane potential, presynaptic Ca²⁺ current, excitatory postsynaptic current and vesicle release rate. The inset shows an overlay of the first (gray) and second (black) EPSC response to action potential-like stimulation.

(after a 2 ms prepulse to $+70\,\mathrm{mV}$) released the fast releasing vesicles ($3 \times \tau_{\mathrm{fast}} \approx 10\,\mathrm{ms}$) corresponding to $\sim 50\,\%$ of the RRP (Sakaba and Neher, 2001a). A second AP-like pulse followed after 200 ms. This interval allowed for replenishment of the SRP but not the FRP owing to the differential recovery times of these pools. It has been shown previously that the SRP recovers fast whereas the FRP recovers more slowly (Sakaba and Neher, 2001a). A final 50 ms step depolarization was then used to release all remaining vesicles in order to obtain a faithful RRP size estimate.

Recordings were performed in the presence of the NMDA receptor blocker D-AP5, Kyn (or gDGG) and CTZ to prevent AMPA receptor saturation and desensitization. Presynaptic Ca²⁺ currents were pharmacologically isolated by TEA and TTX.

In the example given in Fig. 3.2, the first brief depolarization elicited an EPSC of $\sim 2 \,\mathrm{nA}$ whereas the second response to the AP-like stimulation was largely depressed (see inset). The lack of release in response to the second AP-like stimulus was not caused by inactivation of $\mathrm{Ca^{2+}}$ currents, because the $\mathrm{Ca^{2+}}$ charge integral of the first and second AP-like stimulus was not significantly different: the ratio of AP2/AP1

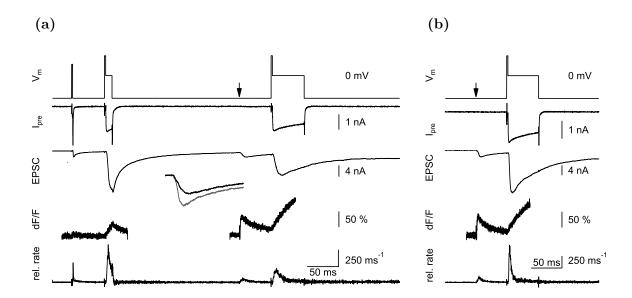


Figure 3.3: Release evoked by flash-induced transient elevations of $[Ca^{2+}]_i$ is intact after depletion of the rapidly releasing vesicles. Traces shown are (from top to bottom): Presynaptic membrane potential, presynaptic Ca^{2+} current, excitatory postsynaptic current, fluorescence signal of the Ca^{2+} indicator OGB-5N and vesicle release rate. OGB-5N is a single wavelength dye, Ca^{2+} binding to the dye is reported as the relative increase of fluorescence over the baseline fluorescence (dF/F). (a) shows the same protocol as Fig. 3.2, but flash uncaging was used instead of the second AP-like depolarization (time point of the flash indicated by an arrow). (b) shows a flash response where the preceding prepulse was omitted. Both (a) and (b) were recorded in the same cell pair as Fig. 3.2.

was $94.7\% \pm 1.5\%$ (mean \pm standard error of the mean (SEM)). As reported by Wu and Borst, a subsequent long depolarization elevates $[Ca^{2+}]_i$ sufficiently to release the 'reluctant' vesicles.

To investigate the nature of these 'reluctant' vesicles and to see whether they are less sensitive to $[Ca^{2+}]_i$, the second stimulus was substituted with a laser-induced transient, but spatially uniform, elevation of $[Ca^{2+}]_i$. If vesicles are 'reluctant' because of a loose coupling between Ca^{2+} channels and the Ca^{2+} sensor of release, uniform elevations of $[Ca^{2+}]_i$ should elicit rapid release (hypothesis 1), while AP-induced release would be small. Alternatively, if vesicles are 'reluctant' because of a reduced Ca^{2+} sensitivity for fusion, uniform elevations of $[Ca^{2+}]_i$ would only trigger slow release (hypothesis 2). Bollmann and Sakmann revised the Ca^{2+} uncaging technique described in 1.1.2 by using a pulsed ultraviolet (UV) laser in combination with the low affinity Ca^{2+} indicator

dye OGB-5N. High concentrations of endogenous buffers were used to evoke a uniform and rapidly decaying global elevation of $[Ca^{2+}]_i$ that can be measured at high sampling frequencies with a photo diode (Bollmann and Sakmann, 2005).

Figure 3.3 shows examples in which 1 mM EGTA was added to the intracellular pipette solution to induce a faster decay of $[Ca^{2+}]_i$ after flash uncaging. It has to be noted that the transients reported here were not as short as the AP transients described in Bollmann and Sakmann (2005). However, the large amounts of EGTA and BAPTA used by Bollmann and Sakmann would make it impossible to obtain RRP size estimates in response to strong depolarizing pulses, as Ca^{2+} entering through channels would readily be buffered before inducing vesicle fusion. Still, the transient flashes shown here are closer to the physiological Ca^{2+} rise invoked by an AP since they do not cause the sustained neurotransmitter release seen in response to flash uncaging designed to produce a step-like elevation of $[Ca^{2+}]_i$ (see below).

In the stimulation protocol depicted in Fig. 3.3(a) the second AP-like stimulus of Fig. 3.2 was substituted with a laser flash. The spatially uniform elevation of $[Ca^{2+}]_i$ was able to trigger release (Fig. 3.3(a)) whereas a short depolarization was not (Fig. 3.2). This difference was not caused by an overly strong flash stimulus since the flash intensity was adjusted so that a stand-alone flash elicited an EPSC (~2 nA in the example shown in Fig. 3.3(b)) comparable to one elicited by AP-like stimulation (1st AP-like stimulus response in Fig. 3.2). Experiments as shown in Fig. 3.2 and Fig. 3.3 were performed in seven cell pairs (n=7). Within a given pair, the laser intensity was adjusted such that stand-alone flashes without a depleting prepulse (control flash) evoked an EPSC of comparable amplitudes to AP-like triggered ones (Fig. 3.3(b)). The $[Ca^{2+}]_i$ elevations $(\Delta F/F)$ were similar between flashes with a prepulse and control flashes in a given cell and gave a mean ratio of 1.04 ± 0.03 across all cell pairs. In order to study the 'reluctant' vesicles alone, 10 ms prepulses were used which, on average, released $52.9\% \pm 3.6\%$ of the total RRP. The fast releasing vesicles should have been depleted by this prepulse since the contribution of fast releasing vesicles is $\sim 50\%$ (as described in Sakaba and Neher (2001a)).

The EPSC amplitudes obtained by the double AP-like stimulation protocols like in Fig. 3.2 gave an average ratio of $AP_2/AP_1 = 0.12 \pm 0.05$. This ratio is much lower than the one given by flashes after a prepulse and control flashes which was 0.78 ± 0.21 . Similar results were obtained when the comparison of the protocols of Fig. 3.2 and

Fig. 3.3 was based on the analysis of vesicular release rates obtained after deconvolution. The peak release rates resulting from either flash or brief depolarizing stimulation were divided by the pool of available vesicles yielding the release rate per vesicle. For double pulse experiments, the release rate per vesicle for the first AP-stimulus was $0.093\,\mathrm{ms^{-1}}\pm0.04\,\mathrm{ms^{-1}}$ and $0.028\,\mathrm{ms^{-1}}\pm0.019\,\mathrm{ms^{-1}}$ for the second, giving a ratio $\mathrm{AP_2/AP_1}$ of $29\,\%\pm9\,\%$. The ratio of release rates per vesicle for flashes after a prepulse and stand-alone (control) flashes was $81\,\%\pm16\,\%$. Release rates per vesicle were $0.035\,\mathrm{ms^{-1}}\pm0.008\,\mathrm{ms^{-1}}$ for control flashes and $0.022\,\mathrm{ms^{-1}}\pm0.004\,\mathrm{ms^{-1}}$ for flashes after the prepulse.

The experiment shown in Fig. 3.3 demonstrated that an intrinsic difference in the Ca^{2+} sensitivity among vesicles is not the major cause for the heterogeneity in p_{ves} observed at the calvx of Held. Instead, the experiment indicated that reluctant vesicles are slow because of loose coupling between Ca²⁺ channels and vesicles in a qualitative manner. To study the kinetics of release quantitatively, step-like elevations of [Ca²⁺]_i were used to follow the time course of release. Ca²⁺ uncaging in combination with the ratiometric Ca²⁺ indicator dye fura-2FF allows for the quantitative determination of [Ca²⁺]_i. Using 1–2 mM DM-nitrophen (DMn) and a variable intensity of the UV flash, [Ca²⁺]_i could be elevated in a step-like manner ranging from 2–30 μM. Figure 3.4(a) shows a protocol analogous to Fig. 3.2 except that the two AP-like stimuli were omitted. A 15 ms prepulse is followed by a 50 ms pool depleting pulse with an interstimulus interval of 200 ms. In order to completely rule out recovery of the fast releasing vesicles during this interval, 20 µM of calmodulin binding domain peptide were added, which has been shown to eliminate the fast phase of FRP recovery (Sakaba and Neher, 2001a). Figure 3.4(a) shows the slow rise of the EPSC during the second stimulus which reflects the release of 'reluctant' vesicles. The EPSC rose much faster when a flash induced elevation of [Ca²⁺]_i was employed as shown in Fig. 3.4(b). In this example, intracellular Ca^{2+} was elevated to levels $> 10 \,\mu\text{M}$ which is comparable to $[Ca^{2+}]_i$ during an AP and sufficient to deplete the RRP. It can be seen that a subsequent 50 ms depolarization did not lead to additional release, indicating that the RRP had been depleted by the preceding flash. The increase in EPSC risetime indicated again that slow vesicles can in fact be released rapidly, but that the Ca²⁺ concentrations reached during depolarizations do not reach high enough levels to do so.

As mentioned previously, the two components of release, fast and slow, can be separated based on their kinetic properties. The deconvolution of the EPSC with miniature excitatory postsynaptic currents (mEPSCs) yields the release rate of vesicles. Its integral describes the cumulative release of vesicles over time. The cumulative release was corrected for the recovery of slow vesicles which takes place at a rate of $\sim 10^{\text{vesicles}/\text{ms}}$ (see 2.4.1). Figure 3.4(c) shows an example of the cumulative release in response to a flash after prepulse (solid trace). The time course of release evoked by flash uncaging can be fitted with a double exponential function (gray trace) that shows the dominant fast component of release (> 80 % depleted within 2 ms), whereas the slow rise of the 2^{nd} EPSC evoked by a 50 ms step depolarization in Fig. 3.4(a) could be fitted with a single exponential function. The lack of biexponential kinetics is indicative of the depletion of fast vesicles (dashed trace in Fig. 3.4(c)).

The dominating fast component of release was most pronounced at $[Ca^{2+}]_i > 10 \,\mu\text{M}$ and showed that the release probability amongst the RRP vesicles is more homogeneous when $[Ca^{2+}]_i$ is elevated uniformly around them. The kinetics of release were analyzed in 30 cell pairs by flash photolysis. The fast component of release accounted for $87\% \pm 4\%$ of the total release in control flashes and still $84\% \pm 4\%$ in flashes that followed a prepulse. This implies that when $[Ca^{2+}]_i$ is elevated sufficiently high (> $10\,\mu\text{M}$), most of the reluctant vesicles can be released rapidly.

It remains possible that the Ca²⁺ sensitivity of release differs at lower $[Ca^{2+}]_i$. It is therefore necessary to examine the $[Ca^{2+}]_i$ -dependence of fast and slow vesicles. This was achieved by changing the UV-flash intensity which allows to achieve post-flash $[Ca^{2+}]_i$ levels of 2–30 μ M. Figure 3.4(d) shows the peak release rate per vesicle plotted against $[Ca^{2+}]_i$. The data were fitted with the 5-site secretion model described by Schneggenburger and Neher (2000) (also see 2.4.2). Only the on-binding rate of the model needed to be changed in order to describe both experimental conditions of control flashes and flashes following a prepulse respectively ($k_{\rm on} = 1.2 \times 10^8 \, {\rm M}^{-1} {\rm s}^{-1}$ for control and $k_{\rm on} = 0.9 \times 10^8 \, {\rm M}^{-1} {\rm s}^{-1}$ after prepulse, $k_{\rm off} = 9000 \, {\rm s}^{-1}$, b = 0.25, $\gamma = 8000 \, {\rm s}^{-1}$). Overall, a ~2-fold difference was observed. The same difference was observed when the time constants of the fast component in the cumulative release were compared (data not shown).

This 2-fold difference is much smaller than the 10-fold difference in release kinetics between FRP and SRP during a depolarizing pulse (Fig. 3.1) so that the spatial distribution of vesicles seems to be the predominant cause of heterogeneity in the release probability observed at the calyx of Held. This excludes the possibility that intrinsic differences in the Ca^{2+} sensitivity can cause fast and slow neurotransmitter release in wild-type animals (Sun et al, 2007). However, even when $[Ca^{2+}]_i$ was elevated uniformly by flash uncaging, some vesicles remained 'reluctant' to release. It is therefore not admissible to exclude the possibility of an additional, though smaller, intrinsic heterogeneity of vesicles. However, the protocols employed here do not allow to discern between an intrinsic heterogeneity that exists a priori of release (Sun et al, 2007) and heterogeneity that develops during release (Wölfel et al, 2007).

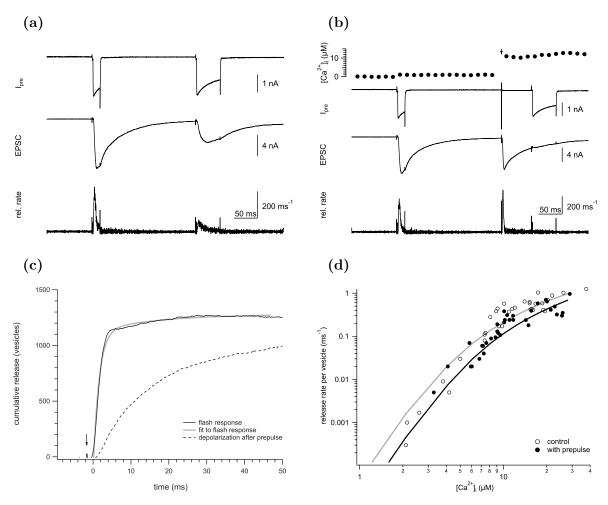


Figure 3.4: Quantitative comparison of neurotransmitter release evoked by either depolarization or flash uncaging after depletion of the fast releasing vesicles. (a) Traces shown are (from top to bottom): presynaptic Ca^{2+} current, excitatory postsynaptic current and vesicle release rate. The presynaptic terminal was depolarized to $+70\,\mathrm{mV}$ for 2 ms and then to 0 mV for 15 ms to deplete the fast releasing vesicles. The addition of calmodulin binding domain peptide and an inter stimulus interval of 200 ms allowed to study the slow releasing vesicles in isolation (Sakaba and Neher, 2001a). The following 50 ms depolarization elicited an EPSC with slow rise and decay (see release rate trace). (b) shows the same protocol as (a) except that flash uncaging was used as the second stimulus. A uniform elevation of $[Ca^{2+}]_i$ to $\sim 10\,\mu\mathrm{M}$ (top trace) evoked an EPSC with fast risetime. (c) shows the cumulative release to illustrate the difference in the release kinetics in response to the second stimulus in (a) and (b). (d) Relationship between the release rate per vesicle and $[Ca^{2+}]_i$ for flash experiments with (black circles) and without (open circles) preceding prepulse. The data were fitted with the 5-site binding model (see scheme 2.1).

3.2 The role of synaptobrevin in release at the calyx of Held

The discrimination between fast and slow releasing vesicles at the calyx of Held is based on the observation of distinct kinetic differences (Wu and Borst, 1999; Sakaba and Neher, 2001a). In order to study the slow component of release, the protocols described so far exploited the slower recovery of fast releasing vesicles and their depletion by stimuli that trigger release of $\sim 50\%$ of the RRP (the fraction attributed to be 'fast').

A selective effect on the fast releasing vesicles via a molecular perturbation of the release machinery using tetanus neurotoxin (TeNT) was described by Sakaba et al (2005). TeNT leads to a complete block of exocytosis at the calyx of Held with time, however the analysis of release kinetics revealed a preferential loss of the fast component of release when the time course of the toxin effect on release was tracked by multiple 50 ms depolarizations. It was shown that under the influence of TeNT, but not under BoNT/C, the cumulative release lost its biexponential characteristic and showed a slowing in the remaining single time constant describing the release kinetics. It was also shown, that the recovery of the fast component was slowed markedly when strong depolarizations were applied at various interstimulus intervals (ISIs), whereas recovery of the slow component was not. TeNT cleaves rat synaptobrevin-2 at the C-terminal end of the segment that participates in SNARE complex formation. However, it is not clear which region of synaptobrevin is responsible for the observed effects on release kinetics.

To further specify the role of synaptobrevin for fast release at the calyx of Held, $100\,\mu\mathrm{M}$ of the N-terminal proline-rich domain of synaptobrevin (corresponding to amino acids 1–26) was perfused into the terminal (Cornille et al, 1995). To facilitate the discrimination of the fast and slow components of release, 0.5 mM EGTA was also added (Sakaba and Neher, 2001a). As shown in Fig. 3.1, two kinetic components representing FRP and SRP can be discerned in this condition. The recovery of the two pools was probed by pairs of 50 ms depolarizations with varied interstimulus interval (Fig. 3.5). The peptide had no effect on the kinetics of release during the first depolarization (Fig. 3.5(b) and Fig. 3.5(a) black traces) whereas the release (and therefore recovery) was reduced under the influence of the peptide (gray traces). It has to be noted however that the peptide had no effect on the total amount of release whereas TeNT eventually blocks

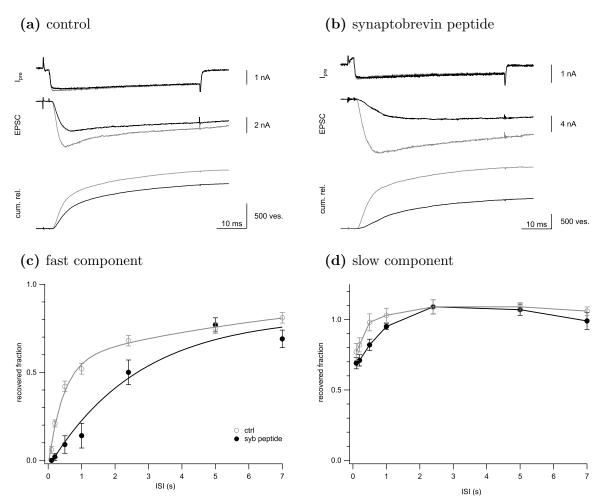


Figure 3.5: (a) & (b) Traces shown are (from top to bottom): presynaptic Ca²⁺ current, EPSC and cumulative release under control conditions and in presence of synaptobrevin proline-rich domain peptide respectively. Two successive 50 ms depolarizations were given with an inter stimulus interval of 1 s. Gray and black traces indicate the first and second stimulus. (c) & (d) show the recovered fraction of the fast and slow component of release during the second depolarizing pulse plotted against the inter stimulus interval between the sustained depolarizations as exemplified in (a) & (b). Open and filled circles indicate control and peptide experiments respectively.

release completely, indicating that the block of release is mediated at the N-terminal side of synaptobrevin but not by the proline-rich domain alone. Similar results were obtained when an antibody raised against amino acids 2–17 of synaptobrevin was used instead of the peptide.

When the recovery was analyzed for the FRP and SRP selectively, it could be shown that the reduction of release during the second pulse stems from a reduction in the recovery of the fast releasing vesicles (Fig. 3.5(c) and (d)). Figure 3.5(c) shows that this reduction was most pronounced at an interstimulus interval of 1 s. This opened the ability to study the Ca^{2+} sensitivity of release of the slow releasing vesicles again, yet under different circumstances. When the second of the 50 ms depolarizations was substituted with a flash, the vesicles released by Ca^{2+} uncaging were vesicles that were recovered within 1 s after the 1st pool depleting 50 ms pulse. The slow vesicles studied in Fig. 3.3(b) and Fig. 3.4(b) were spared during the preceding stimulation. Still, even with the difference in experimental paradigm, only a ~2-fold difference in the Ca^{2+} sensitivity of release was found (data not shown). This supports the importance of the proline-rich domain of synaptobrevin in recruitment of fast releasing vesicles without affecting the apparent Ca^{2+} sensitivity.

3.3 Cleavage of synaptobrevin alters the kinetics of fast neurotransmitter release

To further the investigation of synaptobrevin's role in vesicle fusion at the calyx of Held, $3\,\mu\mathrm{M}$ of the botulinum neurotoxin serotype B (BoNT/B) light-chain was infused into the presynaptic terminal. BoNT/B and TeNT share the same cleavage site on synaptobrevin, yet differ in their requirements for exerting their protease activity. Whereas TeNT requires the V1 copy of the SNARE motif in synaptobrevin, BoNT/B requires V2 (Foran et al (1994); Pellizzari et al (1996), also see Fig. 4.2). The difference in binding requirements was exploited in Hua and Charlton (1999) which showed that at the crayfish neuromuscular junction (NMJ), TeNT cleavage was greatly enhanced by stimulation of the terminal, whereas BoNT/B was effective even in the absence of stimulation.

The effect of various clostridial neurotoxins including TeNT at the calyx of Held was studied in Sakaba et al (2005). TeNT infusion showed a preferential loss of the fast component of release when strong depolarizing stimuli were used to assay the toxin effect. Figure 3.6 shows the effect of BoNT/B in the same experimental paradigm. The presynaptic terminal was depolarized to $0\,\mathrm{mV}$ for $50\,\mathrm{ms}$ (after a prepulse to $+70\,\mathrm{mV}$

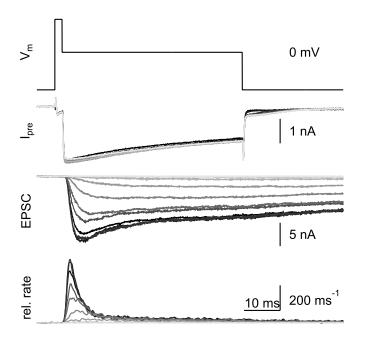


Figure 3.6: BoNT/Bcauses a complete block of neurotransmitter release at the calyx of Held. Traces shown (from top to bottom): presynaptic membrane potential, presynaptic current, EPSC and release rate. Depolarizing pulses were applied every ~60 sec, shades from dark to light show the toxin effect during the course of the experiment (\sim 12 min in this case). The presynaptic pipette solution contained 0.5 mM EGTA.

for 2 ms) roughly every 60 sec to map the toxin effect over time. The presynaptic pipette contained 0.5 mM EGTA to facilitate the differentiation between FRP and SRP. In the example shown in Fig. 3.7, neurotransmitter release was completely abolished within 12 min although the presynaptic Ca²⁺ currents were unchanged (I_{pre} in Fig. 3.6). The cumulative release shown in Fig. 3.7(a) was obtained by integration of the release rate shown in Fig. 3.6. Normalizing to the end of the depolarizing pulse revealed the preferential loss of the fast component of release (see Fig. 3.7(b)) as described for TeNT at the calyx of Held. Early into toxin action, the cumulative release shown in Fig. 3.7(b) showed biexponential kinetics (dark gray) which were eventually lost with time (light gray).

Both TeNT and botulinum neurotoxins (BoNTs) have been shown to reduce spontaneous neurotransmitter release significantly in a number of systems and species (reviewed in Humeau et al, 2000). Both BoNT/B and TeNT block miniature endplate potentials (mEPPs) at the mouse NMJ (Gansel et al, 1987) whereas TeNT has been shown to block mEPSCs in organotypic hippocampal slice cultures even when [Ca²⁺]_i was elevated by ionomycin (Capogna et al, 1997). To determine the effect of BoNT/B on spontaneous release at the calyx of Held, mEPSCs were measured while keeping

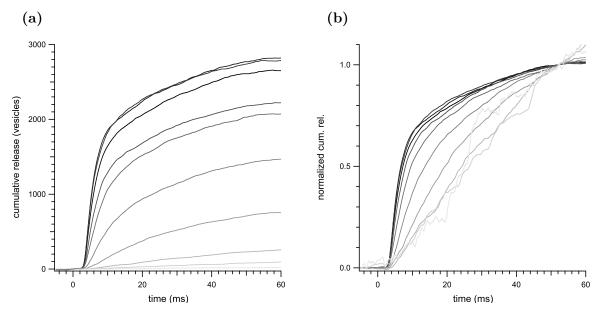


Figure 3.7: (a) Cumulative release of a BoNT/B-infused calyx of Held, obtained by integrating the release rate shown in Fig. 3.6. (b) same as (a) but normalized to the value at the end of the depolarizing pulse to illustrate the slowing of release over time. Shades from dark to light illustrate the toxin effect during the course of the experiment.

control of $[Ca^{2+}]_i$. mEPSC recordings were followed by strong depolarizing stimuli to obtain a RRP size estimate and to make sure that the toxin led to a full block of release by the end of the experiment (usually within 15 min). To allow the synapse to recover, strong stimuli were followed by ~60 sec of waiting time before mEPSCs were recorded again. Control experiments that employed the same stimulation pattern but omitting the toxin showed that both mEPSCs and EPSCs could be recorded without significant run-down in the postsynaptic response for 15–20 min. Figure 3.8(a) shows an example of spontaneous release at a slightly elevated $[Ca^{2+}]_i$ compared to resting levels. When the RRP size was still 98% compared to the one found right after break-in into whole-cell configuration, the release rate was 7×10^{-4} ms⁻¹. The RRP was reduced to 20% of its initial size after 13:30 min and no mEPSCs could be detected anymore. The reduction in RRP size was therefore paralleled by a reduction in mEPSC frequency (see Fig. 3.8(b)). Recordings of mEPSCs were performed in five cell pairs in the presence of BoNT/B and in three cell pairs in control conditions (n = 5 and n = 3 respectively).

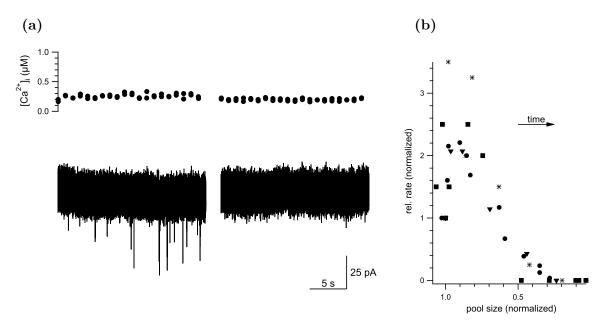


Figure 3.8: (a) Recordings of mEPSCs (bottom) with corresponding $[Ca^{2+}]_i$ (top). The presynaptic pipette solution contained 0.5 mM EGTA and fura-4F. $[Ca^{2+}]_i$ was ~200 nM in this example. Traces on the left side are 4:30 min after entering whole-cell configuration, traces on the right are after 13:30 min. (b) Plot of normalized mEPSC frequency against normalized pool size. Four cells are shown to indicate that the reduction in mEPSC frequency parallels the reduction in pool size assessed by pool depleting stimuli. It has to be noted that changes in $[Ca^{2+}]_i$ are not corrected for which explains the initial increase in the release rate observed in the early phase of the experiments.

Spontaneous quantal release could be studied in $[Ca^{2+}]_i$ ranges from ~0.5–1.5 μ M by using minimal flash uncaging and measuring $[Ca^{2+}]_i$ with the Ca^{2+} indicator dye fura-4F (Lou et al, 2005). Uncaging to low levels of $[Ca^{2+}]_i$ is particularly challenging as stable concentrations can only be obtained by counterbalancing the terminal's Ca^{2+} binding and extrusion mechanisms (readily returning $[Ca^{2+}]_i$ to basal levels) by increasing either the exposure time or frequency of monochromator illumination so that the excitation by the imaging system itself leads to incremental uncaging of Ca^{2+} . The analysis of mEPSCs frequency was therefore limited to a time window during which $[Ca^{2+}]_i$ was constant (n=4 for both BoNT/B and control condition, summarized in Fig. 3.10(a)).

To further investigate the effect of BoNT/B on neurotransmitter release at the calyx of Held, stronger flash stimuli were used to elevate [Ca²⁺]_i to the levels indicated for AP-driven neurotransmitter release (10–20 µM, Bollmann et al (2000); Schneggenburger

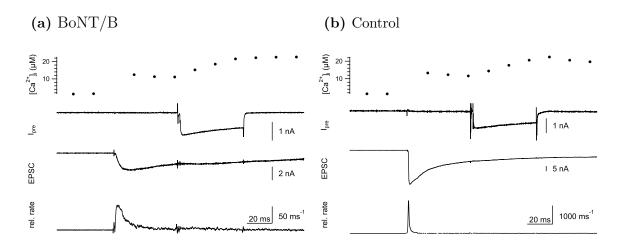


Figure 3.9: (a) shows a flash uncaging experiment in a callyx terminal infused with BoNT/B-LC. Traces shown (from top to bottom): $[Ca^{2+}]_i$ derived from fura-2FF measurement, presynaptic current, postsynaptic EPSC and release rate. (b) is analogous to (a) but under control conditions.

and Neher (2000)) and to lower levels to completely map the release rate's dependence on $[Ca^{2+}]_i$ for both control and toxin condition. As mentioned before, time constraints limit the number of flash trials that can be performed in a given cell pair. Additionally, flashes to levels > 10 μ M can induce run-down of the postsynaptic response. For these reasons, only 1–3 flash stimuli were given per cell pair (n=19 cell pairs for toxin and n=9 for control experiments).

Figure 3.9(a) shows an example where $[Ca^{2+}]_i$ was elevated to 12.4 μ M which evoked a slowly rising EPSC of ~2.5 nA. The subsequent depolarization for 50 ms did not elicit further release, indicating that the RRP had been emptied completely. A comparable flash in control conditions (13.3 μ M) led to a ~30 nA EPSC with fast rise time, again emptying the RRP completely. As with recordings of mEPSCs, BoNT/B flash experiments were done when the RRP was reduced to 5–50% of its initial size. Notably, release was slowed down severely, indicating that the Ca²⁺ sensitivity of release was reduced by BoNT/B. Figure 3.10(a) shows the observed peak release rates per vesicle for $[Ca^{2+}]_i$ levels < 1 μ M which were found to be similar for both toxin and control data. As illustrated in Fig. 3.8, spontaneous release is gradually blocked by BoNT/B but the RRP size is reduced as well. Presenting the data as release rate per vesicle (release rate normalized to the RRP size) allows to compare the control experiments to

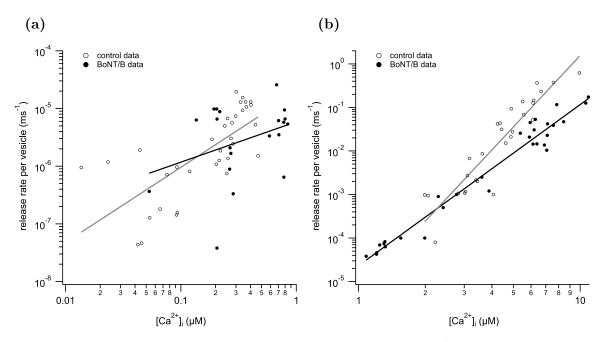


Figure 3.10: (a) Peak release rates per vesicle plotted against $[Ca^{2+}]_i$ for levels lower than $1 \,\mu\mathrm{M}$. Control and BoNT/B data were fitted with lines with slopes of 1.3 and 0.7 respectively, illustrating the shallow dependence of release on $[Ca^{2+}]_i$ at these levels. (b) Same as (a) but for $1-10\,\mu\mathrm{M}$ $[Ca^{2+}]_i$. Slopes were 5.4 and 3.7 for control and BoNT/B respectively.

the toxin ones even in the case of pronounced pool depression. Up to $5\,\mu\mathrm{M}~[\mathrm{Ca}^{2+}]_i$ no clear separation between the two datasets could be revealed. The separation is seen however at levels $> 5\,\mu\mathrm{M}$, where an up to 10-fold difference could be observed. Also, the slope of the dose-response curve is shallower under BoNT/B (Fig. 3.10(b)).

Figure 3.11 summarizes the results of mEPSC recordings and flash uncaging experiments to low and high levels of $[Ca^{2+}]_i$ for both control and toxin experiments (filled circles, n = 28 cells) and control data (open circles, n = 16 cells). Since the spontaneous release rate per vesicle is similar for low $[Ca^{2+}]_i$ levels (Fig. 3.10(a)), BoNT/B does not seem to affect the energy barrier for fusion (parameter l_+ in the allosteric model) as postulated for the effect of phorbol esters on release at the calyx of Held (Lou et al, 2005). The dashed lines in Fig. 3.11 show the predictions of the allosteric model outlined in 2.4.2. To account for the comparable release rates at low levels of $[Ca^{2+}]_i$ (Fig. 3.10(a)) but differential release rates at levels $> 5 \,\mu\text{M}$ (Fig. 3.10(b)), the number of binding steps had to be increased. The observed results could not be accounted for by changing the

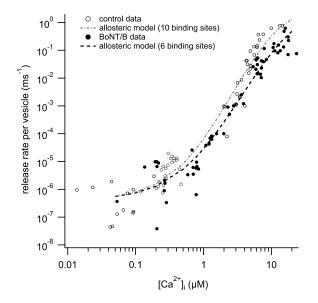


Figure 3.11: Summary plot of release rates per vesicle derived from mEPSC recordings and flash uncaging experiments plotted against intracellular calcium. Open circles show the data from recordings in cells that were infused with BoNT/B-LC and showed a 50–95% reduction in EPSC amplitude. Filled circles show recordings in control conditions. The overlaid curves show the predictions from an allosteric release model. To account for the divergent release rates between control and toxin experiments, the number of binding steps had to be altered.

parameters $l_{+/-}$, $k_{\rm on/off}$ or γ alone. A difference in $l_{+/-}$ should manifest as a difference in the release rate per vesicle at low $[{\rm Ca^{2+}}]_{\rm i}$ between control and toxin recordings, which was not observed. Changes in $k_{\rm on/off}$ shift the dose-response curve, whereas the parameter γ becomes limiting only at high $[{\rm Ca^{2+}}]_{\rm i}$ (>30 µM). The number of ${\rm Ca^{2+}}$ binding steps governs the cooperativity of the kinetic release model and its reduction was necessary to account for the shallower dose-response curve of BoNT/B data (parameters for control data: $k_{\rm on} = 3 \times 10^8 \, {\rm M^{-1} s^{-1}}$, $k_{\rm off} = 1.2 \times 10^4 \, {\rm s^{-1}}$, $l_+ = 4 \times 10^{-4} \, {\rm M^{-1} s^{-1}}$, $l_- = 4 \times 10^{-5} \, {\rm s^{-1}$

In conclusion, BoNT/B diminishes the fast component of release similarly to TeNT when the release is evoked by a step depolarization. The underlying mechanism is different, however, as TeNT blocks the coupling of vesicles to Ca^{2+} channels whereas BoNT/B reduces the Ca^{2+} sensitivity. Although the Ca^{2+} sensitivity is changed, the mechanism of action is different than the one described in Lou et al (2005), where the effect of phorbol esters was ascribed to an increased 'fusion willingness' for release governed by the parameter l_+ in a simplified allosteric model.

3.4 Rabphilin3a as a potential modulator of release at the calyx of Held

The toxins that have been shown to effect the fast component of release at the calyx of Held selectively cleave synaptobrevin. Toxins that cleave other members of the SNARE family of proteins, though mediating severe block of neurotransmitter release, did not share the differential effects on the kinetics of release. To investigate whether the fast component of release can be perturbed by other means, a peptide taken from rabphilin3a that showed a marked reduction in chromaffin cell secretion was perfused into the calyx terminal via the patch pipette. Chung et al could show that the basic COOH-terminal peptide from the C_2B domain of bovine rabphilin3a (amino acids 676–695, corresponding to amino acids 650–669 in rat rabphilin3a (Rp3a(650–669))) reduced release whereas its scrambled version and homologous peptides taken from synaptotagmin did not (Chung et al, 1998). It has also been shown that the bottom surface of the C_2B in rabphilin3a, which contains a long α -helix (Ubach et al, 1999), mediates binding to SNAP-25 (Deák et al, 2006).

Figure 3.12 illustrates the effect of Rp3a(650–669) and its homologue from synaptotagmin-2 (Syt2(383-401)) at the calyx of Held. Synthetic peptides dissolved in cesium-based solution were added to the presynaptic solution. Rp3a(650–669) was effective in blocking release at the calyx of Held at concentrations as low as 100 μ M. However, 400 μ M were used so that release could be blocked completely within 10–15 min even when the access resistances obtained in the presence of peptide were close to 20 M Ω . Within this timeframe, the contribution of run-down of release should be minimal, so that the observed block of release could solely be attributed to be a peptide effect. Figures 3.12(a), (c) and (e) show that, much like TeNT and BoNT/B, Rp3a(650–669) blocks synaptic release completely (within ~13 min in this example). Again, the fast component of release was lost early into peptide action.

The interpretation of data derived from experiments in which peptides were used as competitive antagonists is ambiguous as the concentrations used experimentally should greatly exceed those of the natively found protein they are derived from. Also, it cannot be ruled out that peptides yield secondary effects due to differential folding, dimerization or the ability to diffuse freely within the cytosol. As one control for the

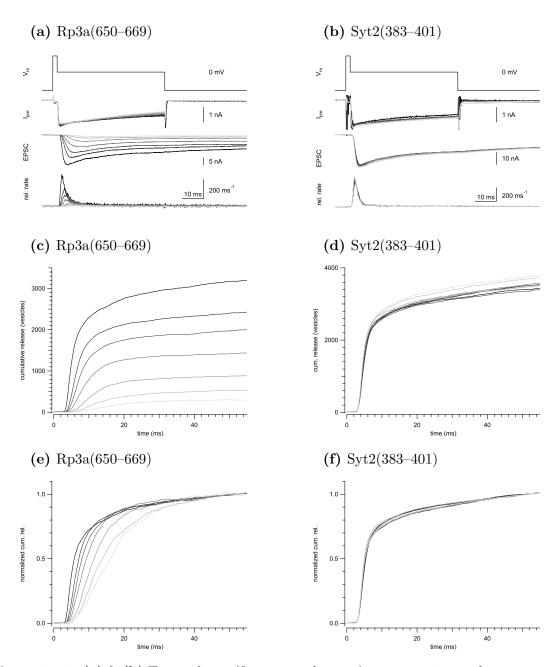


Figure 3.12: (a) & (b) Traces shown (from top to bottom): presynaptic membrane potential, presynaptic current, EPSC and release rate. Depolarizing pulses were applied every ~60 sec, shades from dark to light show the EPSCs during the course of the experiment. The presynaptic pipette solution contained 0.5 mM EGTA. (c) & (d) Cumulative release obtained by integrating the release rate shown in (a) and (b) respectively. (e) & (f): same as (c) & (d) but normalized to the value at the end of the depolarizing pulse.

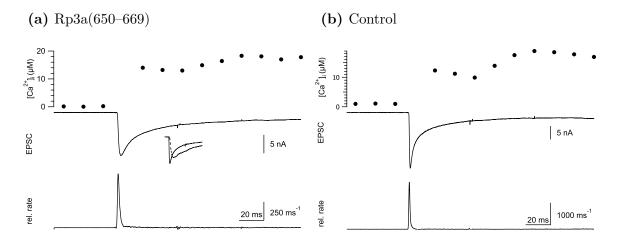


Figure 3.13: (a) Flash uncaging in a calyx of Held terminal infused with Rp3a(s650–669). Traces shown (from top to bottom): $[Ca^{2+}]_i$ derived from fura-2FF measurement, presynaptic current, postsynaptic EPSC and release rate. The inset shows an overlay of the Ca^{2+} uncaging response (solid line) compared to an EPSC evoked by a 50 ms depolarization (dashed line) in the same cell. (b) is analogous to (a) but under control conditions.

specificity of the observed effect of Rp3a(650–669), a peptide from the homologous sequence in synaptotagmin-2 (thought to be the predominant synaptotagmin isoform at the calyx of Held, Sun et al (2007)) was used. Figures 3.12(b), (d) and (f) illustrate that the synaptotagmin peptide did not alter release significantly at the same concentration, comparable access resistance and time frame as the experiments with Rp3a(650–669).

To rule out that the net charge or number of hydrophobic amino acids alone mediates the effect seen with Rp3a(650–669), its scrambled variant (Rp3a(s650–669)) as described by Chung et al was introduced into the calyx terminal. Interestingly, Rp3a(s650–669) infusion resulted in a very fast and strong inactivation of the recorded Ca²⁺ currents. The inactivation was so severe, that proper 'template protocol' (see 2.4.1) or pool depleting stimuli recordings could not be conducted. However, the EPSC amplitudes recorded in the presence of Rp3a(s650–669), though small, did not decrease over time. The effect on presynaptic Ca²⁺ currents was not described by Chung et al since they assayed the peptide effect by making use of radioactively labeled norepinephrin and digitonin to permeabalize the cells. The stability of EPSC amplitudes at the calyx of Held did however confirm that the scrambled peptide does not seem to affect vesicular release directly.

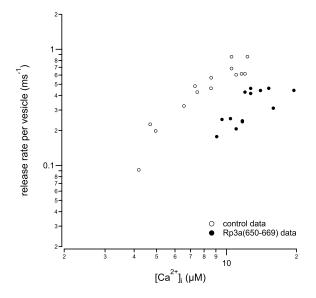


Figure 3.14: Summary plot of release rates per vesicle from flash uncaging experiments plotted against intracellular calcium. Closed circles show the data from recordings in cells that were infused with the Rp3a(650–669) peptide. Flashes were applied when the peptide had caused a 50–95% reduction in EPSC amplitude. Open circles show recordings in control conditions.

To complete the characterization of the Rp3a(650–669) effect on release at the calyx of Held, flash uncaging experiments to $[\mathrm{Ca^{2+}}]_i$ levels $> 10\,\mu\mathrm{M}$ were performed. Flash photolysis was induced when the peptide had reduced the initially observed EPSC amplitude to 50% or less. Figure 3.13 shows representative examples for both peptide and control condition. The inset in Fig. 3.13(a) indicates that the EPSC in the peptide experiment is somewhat slower compared to control, however, the peak release rate per vesicle was reduced only about 2-fold across all cell pairs. Figure 3.14 illustrates the dose-response relationship of peak release rate per vesicle to $[\mathrm{Ca^{2+}}]_i$ for cells infused with Rp3a(650–669). Since the $\mathrm{Ca^{2+}}$ sensitivity of release seemed to be altered only little when depolarization induced EPSCs were strongly reduced in amplitude and presented slowed kinetics, Rp3a(650–669) does seem to interfere with the vesicle-release coupling at the calyx of Held.

Communication between neurons in neuronal networks predominantly relies on signal transmission via chemical synapses. These sites are, however, not merely relay stations for the signals that invade the presynaptic terminal, but highly dynamic in shaping the signal that eventually passes down to the processes of the postsynaptic cell. The proficiency at which a synapse operates is dependent on the history of its activity, both on long and short timeframes. Activity dependent changes in synapses that last, at the most, a few minutes, coined short-term synaptic plasticity, present themselves in either facilitation, depression or a mixture of both in virtually all synapses and have been recognized early on (reviewed in Zucker and Regehr, 2002).

The synaptic strength of a synapse can be described by the model introduced in the introduction (eq. 1.1), namely the parameters N, p and q that shape the postsynaptic response. Seemingly a simple model, a remarkable number of processes within the cell have been found to contribute to either one of these parameters. Special interest has been taken in the attempt to pinpoint a single, the predominant, process that shapes synaptic transmission at a given synapse.

The calyx of Held, a model glutamatergic synapse, depresses when stimulated with action potential (AP) trains and the depletion of vesicles from the readily releasable pool (RRP) is seen as the main cause of short-term synaptic depression at glutamatergic synapses such as the calyx of Held (Schneggenburger et al, 1999; Wu and Borst, 1999; Bollmann et al, 2000).

The situation at the calyx of Held is complicated by the fact that the RRP consists of two subpools that were identified based on their release kinetics and recovery rate (see

Fig. 1.2, Sakaba and Neher (2001a)). The fast releasing vesicles among the RRP were released 10-fold faster than the slowly releasing vesicles under sustained depolarization of the terminal, yet took much longer to recover after depletion. These two subpools of vesicles were studied further in Sakaba (2006). It was found that synchronous release at the calyx of Held is mainly mediated by the fast releasing vesicles. Consequential, the depression observed during the synchronous phase of AP trains is a result of their depletion. The slowly releasing vesicles in turn contribute mainly to the asynchronous release phase during high frequency stimulation and are less prone to depression.

A heterogeneous RRP was also identified in glutamatergic hippocampal neurons in culture, where action potentials only drew a fifth of the vesicle pool released during sucrose application. Just as the calyx of Held, the remaining 'reluctant' vesicles could be forced to release by increasing the Ca²⁺ influx (Moulder and Mennerick, 2005), i.e. through higher frequency AP trains.

Investigating the nature of the slowly releasing or reluctant vesicles was attempted in this work. The comparison of fast and slow release then allows to clarify how fast release, responsible for AP-induced neurotransmitter release, is mediated.

It was shown that fast release could still be triggered by flash photolysis of caged $\mathrm{Ca^{2+}}$ which elevated $[\mathrm{Ca^{2+}}]_i$ to levels above 10 $\mu\mathrm{M}$ when AP evoked release was largely depressed (see Fig. 3.3(a)). A calcium concentration of roughly 10 µM is thought to represent the amplitude of the average AP-induced Ca^{2+} transient that vesicles are exposed to. The vesicles remaining after a depolarizing pulse that depleted the fast releasing pool of vesicles (FRP) therefore were 'releasable' and the observed depression in AP-evoked release was not caused by a lack of fusion competence. Furthermore, quantitative analysis of the [Ca²⁺]_i dependence of release combined with a minimal model describing the kinetics of release revealed that the intrinsic sensitivity of reluctant vesicles to Ca²⁺ is reduced at most 2-fold. Intrinsic differences between vesicles should therefore only contribute little to the 10-fold difference in release kinetics observed in response to strong depolarization of the calyx of Held terminal. The 10-fold slower release kinetics of slowly releasing vesicles observed during a long depolarizing pulse (Sakaba and Neher, 2001a) can therefore be explained by a lower [Ca²⁺]; that these vesicles are exposed to when the elevation in $[Ca^{2+}]_i$ is mediated by Ca^{2+} channels. It was estimated that slow vesicles 'see' an average Ca²⁺ transient peaking at 6-7 µM

whereas fast releasing vesicles are exposed to an average $\mathrm{Ca^{2+}}$ transient of 10–12 $\mu\mathrm{M}$ (Wadel et al, 2007).

It has to be noted that a fraction of vesicles could not be released 'fast' as 16% of vesicles released slowly even when $[Ca^{2+}]_i$ was elevated uniformly via flash uncaging. However, intrinsically slow vesicles do not seem to be a major contributor to release under more physiological conditions since this should have presented itself when transient elevations of $[Ca^{2+}]_i$ were induced (Fig. 3.3). On average, the release rate per vesicle was reduced only $\sim 20\%$ for transient flashes that followed partial pool depletion.

From this data it is reasonable to conclude that the slow vesicles are merely further away from sites were Ca^{2+} enters the cell, namely calcium channels or clusters thereof. This implies that the fusion competence of fast and slow releasing vesicles is similar. Distant vesicles will be exposed to lower Ca^{2+} concentrations due to the sharp spatial profile Ca^{2+} transients are assumed to possess (see 1.1.2). It is important to note that the slowness is indeed caused by the lowered local $[\mathrm{Ca}^{2+}]$ and not by a time delay imposed by diffusion from the channels' pores to a given vesicle. Due to the supralinear relationship of vesicle release to $[\mathrm{Ca}^{2+}]_i$ at the calyx of Held (Fig. 3.4(d)), the difference of 6–7 μ M versus 10–12 μ M average local $[\mathrm{Ca}^{2+}]$ can readily account for the 10-fold difference in kinetics observed for the two classes of vesicles.

It has to be noted that 'positional priming' of vesicles (the association of vesicles with Ca^{2+} channel clusters at the active zone) is different from the priming step in the vesicle cycle that is thought to involve proteins such as Munc13. Since the slow releasing pool of vesicles (SRP) can be released fast by homogeneous elevations of $[Ca^{2+}]_i$, vesicle docking and priming (a prerequisite for fusion to occur) do not seem to be the rate limiting steps in the exocytosis of 'slow' vesicles, but rather their positioning within the active zone and Ca^{2+} channel clusters. The remaining 2-fold difference in kinetics would then be explained by additional protein interactions ('superpriming', Schlüter et al (2004)).

The RRP contains an average of 3000 vesicles at the calyx of Held and is made up by about equal numbers of slow and fast vesicles (Sakaba and Neher, 2001a). About 500 active zones were identified by serial electron microscopy (EM) reconstruction of a p9 calyx of Held (Sätzler et al, 2002), so that on average three vesicles should be released fast and three slow at a given active zone. It was shown previously that the SRP recovers very quickly ($\tau = 200 \, \text{ms}$), whereas the FRP recovers much slower (see Fig. 3.5,

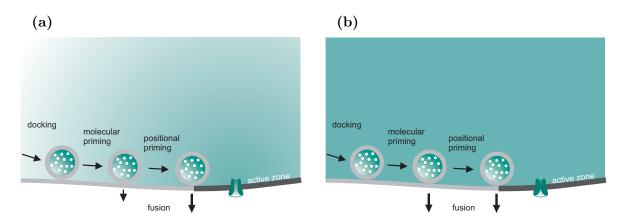


Figure 4.1: Illustration of the steps towards Ca^{2+} -triggered exocytosis of synaptic vesicles when Ca^{2+} influx is mediated by Ca^{2+} channel opening (a) or when $[Ca^{2+}]_i$ is elevated uniformly via flash uncaging (b). In (a), vesicles that are close to Ca^{2+} channels release rapidly, whereas more distant vesicles are 'reluctant' to release. These vesicles are release competent and can be released rapidly in response to uniform elevations of $[Ca^{2+}]_i$ as shown in (b).

Sakaba and Neher (2001a)). It is plausible that these two pools recover sequentially so that the FRP draws from the SRP and that this 'maturation' involves the recruitment of vesicles into close proximity of Ca²⁺ channels. Naturally, trying to understand this maturation step involves the attempt to modulate it which can be achieved by both physiological and non-physiological means.

It was shown that the recovery of the fast releasing vesicles depends on $\mathrm{Ca^{2+}/}$ calmodulin (Sakaba and Neher, 2001a) and cAMP (Sakaba and Neher, 2003a). The rate of recovery of the fast pool was described quantitatively by Hosoi et al (2007) who found a linear dependence on the global $[\mathrm{Ca^{2+}}]_i$. Interestingly, the recovery of the SRP is rather robust and could only be slowed down by latrunculin A, causing actin depolymerization, and ATP- γ S, a nonhydrolyzable analogue of ATP (Sakaba and Neher, 2003b).

Infusing the proline-rich N-terminal domain of synaptobrevin into the calyx terminal revealed a slowing of the recovery of the fast component, but did not interfere with recruitment of the slow releasing vesicles after depletion of the entire RRP (see Fig. 3.5). This indicates the involvement of the proline-rich domain of synaptobrevin in 'positional priming' of synaptic vesicles. The total amount of release, however, was not influenced by the peptide. This is in contrast to the action of tetanus neurotoxin (TeNT) which

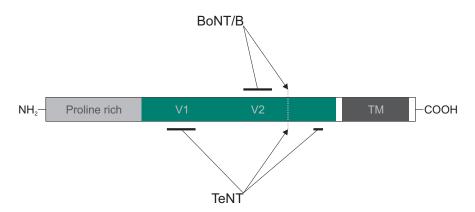


Figure 4.2: Schematic diagram of the synaptobrevin structure and its interaction with BoNT/B and TeNT. The proline-rich N-terminal domain is indicated in light gray. The coiled-coil homology domain (green) mediates the formation of the SNARE complex with syntaxin-1 and SNAP-25 and contains two copies of the SNARE motif (V1 and V2) with which the synaptobrevin-specific neurotoxins BoNT/B and TeNT selectively interact. The cleavage site for BoNT/B and TeNT is indicated by the dashed line (rat synaptobrevin-2 is cleaved at peptide bond GASQ⁷⁶FETS, whereas rat synaptobrevin-1 is not cleaved due to a variation in the cleavage site GASVFESS). The transmembrane domain (TM) of synaptobrevin is indicated in dark gray. Modified from Schiavo et al (2000).

cleaves both the proline-rich domain and the SNARE motives from synaptobrevin. At the calyx of Held, TeNT demonstrates an all-or-none effect which eventually leads to a complete block of release if sufficient time for toxin action is provided. During the early phase of toxin action, TeNT preferentially blocks the fast component of release, a first indication of synaptobrevin's role in mediating, at least partially, the incorporation of vesicles in the vicinity of Ca²⁺ channels. It is, however, not understood how the proline-rich domain mediates 'positional priming' in molecular terms.

Experiments that characterize the effect of BoNT/B on transmission at the calyx of Held confirmed the kinetic effect of synaptobrevin cleavage (see Fig. 3.7). The comparison of TeNT and BoNT/B allows interesting insights into the way the fusion machinery is assembled. Hua and Charlton (1999) made use of both toxins and studied their effect in crustacean neuromuscular junctions (NMJs). It was found that the efficiency of TeNT in cleaving synaptobrevin was greatly enhanced by nerve stimulation, whereas BoNT/B was effective even in a 'silent' synapse. Since TeNT and BoNT/B share the same cleavage site on synaptobrevin, the observed difference was accounted to the difference in the required toxin binding sites on synaptobrevin (see Fig. 4.2).

Tentral requires the N-terminal SNARE motif for binding whereas Bont/B needs the C-terminal one. Hua and Charlton concluded that a stable, partially assembled trans-SNARE complex shields the Tent binding site but leaves the Bont/B one accessible. Bont/B would therefore even be able to exert its protease activity on vesicles that are docked, primed and available for fast Ca²⁺ driven exocytosis. It has to be noted that experiments that are based on comparing stimulated and non stimulated terminals cannot be reconciled with experiments in vertebrate neurons. At the calyx of Held, the rapid rate of docking and undocking as well as spontaneous release lead to sufficient 'cycling' of vesicles to open a time window for Tentral cleavage. Accordingly, botulinum neurotoxins were found to be efficient in blocking release in the absence of stimulation (T. Sakaba, personal communication).

However, BoNT/B did show an additional effect on the kinetics of release at the calvx of Held (see Fig. 3.11). When flash photolysis was used to trigger release at the calyx of Held after BoNT/B had caused more than 50% reduction in the RRP size, lower release rates per vesicle were observed at a given $[Ca^{2+}]_i$. The reduced Ca^{2+} sensitivity was evident at $[Ca^{2+}]_i > 5 \mu M$, but no differences could be identified at lower and resting levels of $[Ca^{2+}]_i$. Attempts to describe the observed effect by an allosteric kinetic model for vesicle fusion indicate that BoNT/B changes the calcium cooperativity of the release machinery. The contribution of SNAREs to calcium cooperativity was postulated for the Drosophila NMJ (Stewart et al, 2000) previously. Yet at the cerebellar basket cell terminal, BoNT/B cleavage of synaptobrevin led to a shift in Ca²⁺ sensitivity but no change in cooperativity was observed (Sakaba, 2008). However, spontaneous events were not analyzed in the latter study which makes it difficult to attribute the observed effect to changes in K_D , l_+ or the number of binding sites in the context of the model presented here. The data herein are compatible with a model of 'zippered' vesicle fusion, which allows BoNT/B to cleave synaptobrevin at a late stage of vesicle fusion, possibly the hemifusion state directly prior to Ca²⁺ triggered exocytosis.

In summary, BoNT/B mediated cleavage of synaptobrevin after molecular priming and before Ca²⁺ triggered exocytosis presents as a change in the Ca²⁺ sensitivity of the release machinery.

Clostridial neurotoxins are not the only agents that interfere with positional priming. In this work, it was shown that a peptide taken from the C-terminal end of rabphilin3a's C₂B domain (Rp3a(650–669)) also caused a pronounced block of neurotransmitter

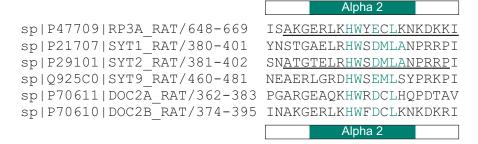


Figure 4.3: Sequence alignment of sequences containing the α-helix that has been described as a conserved feature of C_2B domains. Conserved residues within the helix are highlighted in green. The residues corresponding to the peptides Rp3a(650–669) and Syt2(383–401) used in experiments are underlined. The corresponding sequences from synaptotagmin-1, -2 and -9 as well as Doc2A/B are shown for comparison. Modified from Ubach et al (1999).

release that initially presented as a loss of the fast component of release. Infusion of Rp3a(650–669) essentially mimics the TeNT phenotype at the calyx of Held. These kinds of experiments might suffer from non specific binding of the peptide. The effectiveness of Rp3a(650–669) in blocking release was therefore validated by a homologous peptide from synaptotagmin-2 (Syt2(383–401)) and a scrambled version of the rabphilin3a peptide (Rp3a(s650–669)), neither of which caused a block of release as observed with Rp3a(650–669).

Rp3a(650–669) contains the α -helix that was described as a unique feature of C_2B domains and is thought to be involved in their Ca^{2+} independent interactions (Ubach et al, 1999). This helix is a conserved feature of C_2B domains and can hence not only be found in rabphilin3a, but also in the rat synaptotagmins and Doc2A/B. A later study identified this helix in rabphilin3a to mediate binding to SNAP-25 using NMR spectroscopy (Deák et al (2006), but see Tsuboi et al (2007)). The authors found an increased recovery from use-dependent synaptic depression in cultured neurons from rabphilin3a KO mice that was not described in previous work on synaptic transmission and plasticity in rabphilin3a deficient animals (Schlüter et al, 1999). The phenotypes described by Schlüter et al and Deák et al cannot be reconciled with results that were obtained by utilizing rabphilin3a in an 'overexpression' scheme (Chung et al, 1998; Burns et al, 1998; Tsuboi and Fukuda, 2005) and the results described herein.

Rather than interpreting the results obtained with Rp3a(650–669) at the calyx in the context of rabphilin3a's role in synaptic transmission, the focus should be set on

its interaction partner. It is plausible that the observed block of release is caused by the interaction with SNAP-25 as described by Deák et al. In order to unify the results seen with Rp3a(650–669) and those of clostridial neurotoxins on fast neurotransmitter release at the calyx of Held, a collaborative effort is currently under way to see if and where Rp3a(650–669) binds to SNAP-25.

The work presented here served to investigate the kinetics of neurotransmitter release observed at the calyx of Held synapse. The heterogeneity in release probability, or in other words: the reluctance of a subpopulation of vesicles to be released by an AP invading the synaptic terminal was attributed to the spatial profile of vesicles within the terminal which in turn highlights the importance of $[Ca^{2+}]_i$ and the Ca^{2+} transients that trigger release and the long-standing interest of scientist in studying their role in synaptic transmission and plasticity.

It has to be noted that reluctant release of vesicles can be explained by alternative mechanisms that likely contribute to the mechanism that was described here. A study at the calyx of Held described differences in the intrinsic Ca^{2+} sensitivity of vesicles as the reason for heterogeneity in p_{ves} . The observed 'submaximal' release was interpreted mechanistically as an 'a posteriori' adaptation of the release apparatus to the Ca^{2+} signal (Wölfel et al, 2007). Other studies have highlighted the importance of small changes of the Ca^{2+} influx for short-term depression which were found to be specifically important at lower stimulation frequencies (Xu and Wu, 2005).

As outlined, the recovery of vesicles for fast release can be modulated at many levels at the calyx of Held by both physiological and non physiological means. Conversely, the release machineries sensitivity to Ca^{2+} seems to be very robust so that changes are most easily induced by interfering directly with SNAREs (Sakaba et al, 2005) or the Ca^{2+} sensor (Sun et al, 2007).

Clearly, future work at the calyx of Held will attempt to unify biophysical concepts (such as 'positional priming') with those from biochemical ones ('molecular priming').

Summary

Fast and synchronous neurotransmitter release relies on the time course of calcium influx into the terminal, the availability of fusion competent vesicles and the proteins comprising the release machinery. Studies in glutamatergic synapses have identified a subpopulation of vesicles in the releasable pool that do not fuse in response to action potential (AP) driven calcium influx and release only slowly in response to sustained stimulation. The nature of these 'reluctant' vesicles, however, remained unclear.

This question was investigated at the calyx of Held, a large glutamatergic synapse in the auditory brainstem. By combining electrophysiological recordings of the preand postsynaptic compartments with calcium imaging and uncaging, it is shown that uniform elevations of the intracellular calcium concentration in the presynaptic terminal were able to elicit rapid release after the vesicles mediating fast and synchronous neurotransmitter release had been depleted. The calcium sensitivity of 'reluctant' vesicles was reduced only 2-fold which is insufficient to explain the 10-fold slower release kinetics that is observed in response to sustained depolarizations of the presynaptic terminal. It is concluded, that 'reluctant' vesicles are indeed release competent, but that their distance to calcium channel clusters at the active zone prevents them to release in response to APs. This indicates that the incorporation of vesicles into the vicinity of calcium channels at the active zone (a 'positional priming' step) is rate-limiting in fast neurotransmitter release rather than acquiring fusion competence (a 'molecular priming' step).

The accessibility of the presynaptic terminal of the calyx of Held to patch-pipettes allows one to infuse toxins and peptides to further investigate the 'positional priming' 5 Summary 63

step in exocytosis. Here, it is shown that the N-terminal of synaptobrevin is important for mediating this step, which has been indicated in a previous study based on the effect of tetanus neurotoxin (TeNT) on release at the calyx of Held. In contrast, it was found that BoNT/B, which shares the same cleavage site as TeNT but requires a different binding site on synaptobrevin, changes the intracellular calcium sensitivity of vesicles for fusion, likely by interfering with vesicle exocytosis at a step between calcium sensing and fusion.

'Positional priming' was also affected when a C-terminal peptide from rabphilin3a's C₂B domain was infused into the calyx of Held, which blocked synaptic transmission, potentially by interacting with SNAP-25. Its effect on the kinetics of release mimics that of TeNT as it preferentially blocked the fast component of release.

These results highlight the importance of calcium channel-vesicle coupling as well as the delicate interplay of multiple proteins of the release machinery in mediating fast neurotransmitter release.

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Appendix

A.1 Abbreviations

aCSF artificial cerebrospinal fluid

AMPA α-Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid

AP action potential

APW action potential waveform

aVCN anterior ventral cochlear nucleus

AZ active zone

BAPTA 1,2-Bis(2-aminophenoxy)ethane-N,N,N',N'-tetraacetic acid

BoNT botulinum neurotoxin

BoNT/B botulinum neurotoxin serotype B

CTZ cyclothiazide

D-AP5 D(-)-2-amino-5-phosphonopentanoic acid

DGC Dodt gradient contrast

DIC differential interference contrast

DMn DM-nitrophen (1-(2-Nitro-4,5-dimethoxyphenyl)-

N,N,N',N'-tetrakis[(oxycarbonyl)methyl]-1,2-ethanediamine)

EGTA Ethylene glycol-bis(2-aminoethylether)-N,N,N',N'-tetraacetic acid

EM electron microscopy

EPSC excitatory postsynaptic current

FRP fast releasing pool of vesicles (a subpool of the readily releasable pool)

FWHM full width of half maximum

GBC globular bushy cell gDGG γ-D-glutamylglycine

IR infrared

IPSC inhibitory postsynaptic current

ISI interstimulus interval

Kyn kynurenic acid

A.1 Abbreviations 75

mEPP miniature endplate potential

mEPSC miniature excitatory postsynaptic current MNTB medial nucleus of the trapezoid body

Nd:YAG neodymium-doped yttrium aluminum garnet

NMDA N-methyl-D-aspartic acid NMJ neuromuscular junction

NSF N-ethylmaleimide sensitive factor

OGB-5N Oregon Green BAPTA-5N

RRP readily releasable pool (defined as the number of docked and fusion

competent vesicles that can be released by strong depolarizing pulses or

Ca²⁺ uncaging at the calvx of Held)

RT room temperature

SEM standard error of the mean SNAP soluble NSF attachment protein

SNAP-25 synaptosomal associated protein of 25 kDa

SNARE SNAP receptor

SRP slow releasing pool of vesicles (a subpool of the readily releasable pool)

TEA tetraethylammonium TeNT tetanus neurotoxin

TTX tetrodotoxin UV ultraviolet

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