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Synaptic vesicle undocking induces low frequency depression

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This **fundamental** work substantially advances our understanding of short-term plasticity mechanisms by providing evidence for release-independent low-frequency synaptic depression that reflects a model that proposes a redistribution of vesicles within the readily releasable pool, via a reduction in docking site occupancy due to vesicle undocking. The evidence supporting this model is **convincing**, with rigorous modeling, electrophysiological data and computational analysis. The work will be of broad interest to cellular neuroscientists and synaptic physiologists.

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Abstract

Synaptic depression is often interpreted as reflecting depletion of the readily releasable pool (RRP) following exocytosis. Such a mechanism predicts little or no depression at low stimulation frequency, as RRP replenishment should then offset the loss of vesicles by exocytosis. Nevertheless, in several types of mammalian central synapses, repetitive presynaptic stimulation at low frequency (< 5 Hz) elicits synaptic depression (low frequency depression, or LFD). In the present work we count the number of synaptic vesicles released at individual active zones to study the RRP and its replenishment during LFD. Contrary to depletion models of synaptic depression, we find that LFD does not depend on previous SV consumption. We find that LFD displays a long recovery time course (tens of seconds) when challenged by isolated stimulations but is immediately reversed by a high frequency train. We suggest that LFD results from undocking, a shift between two classes of synaptic vesicles organized sequentially inside the RRP (replacement vs. docked vesicles) in favor of the upstream (replacement) state. While undocking is apparent hundreds of milliseconds after a stimulation, calcium dependent docking takes only a couple of milliseconds, explaining the fast LFD recovery when stimulating at high frequency. Consistent with the undocking model, we find that double presynaptic stimulations alleviate LFD as they favor vesicular docking and RRP replenishment. Finally, we expand our model to explain how stimulation frequency shapes short-term synaptic depression, changing from depression at low frequency to a facilitation-depression sequence at medium or high frequency trains.

Introduction

Short term synaptic plasticity (STP) governs changes in the amplitude of synaptic signals following recent neuronal activity. These changes are often rapid and large, and they potently shape information transfer across synapses. Two main components of STP are synaptic facilitation and synaptic depression (Zucker and Regehr, 2002 [↗](#)).

Synaptic depression occurs after a burst of high frequency presynaptic firing and associated synaptic release (Betz, 1970 [↗](#); Castillo and Katz, 1954 [↗](#); Charlton et al., 1982 [↗](#)). This link with previous presynaptic activity led to the view that synaptic depression is largely due to a decrease in the size of the readily releasable pool (RRP) of synaptic vesicles (SVs), following the loss of SVs by exocytosis (Kavalali, 2006 [↗](#); Regehr, 2012 [↗](#); Zucker and Regehr, 2002 [↗](#)). In addition, a

number of other mechanisms including desensitization of postsynaptic receptors (Rossi et al., 1995 [↗](#); Trussell et al., 1993 [↗](#)), activation of presynaptic inhibitory receptors (Redman and Silinsky, 1994 [↗](#)), decreased voltage-dependent calcium entry (Xu and Wu, 2005 [↗](#)), decreased SV release probability (Wölfel et al., 2007 [↗](#); Wu and Borst, 1999 [↗](#)), heterogeneous release probabilities among release sites (Neher, 2015 [↗](#); Trommershäuser et al., 2003 [↗](#)), and non-linear presynaptic calcium buffers (Bolshakov et al., 2019 [↗](#)), contribute to synaptic depression in various proportions depending on the preparation.

In more recent years, several studies have uncovered a form of synaptic depression that may differ from the RRP depletion pattern. In invertebrate synapses (*Aplysia* (Doussau et al., 2010 [↗](#)); crayfish (Silverman-Gavrila et al., 2005 [↗](#))), as well in mammalian central synapses (hippocampal CA3 to CA1 (Abrahamsson et al., 2007 [↗](#)) and CA3-CA3 (Saviane et al., 2002 [↗](#)) synapses, climbing fiber to Purkinje cell synapses (Rudolph et al., 2011 [↗](#)); parallel fiber to Purkinje cell synapses (Doussau et al., 2017 [↗](#)); calyx of Held (Lin et al., 2022 [↗](#); Müller et al., 2010 [↗](#))), repetitive stimulation at low frequency (0.1 Hz to 5 Hz) induces synaptic depression. This form of synaptic depression (low frequency depression, or LFD) does not require a previous period of intense presynaptic activity, so that it is unclear whether it can be explained by the classical RRP depletion hypothesis, or by any of the alternative mechanisms mentioned above.

Recent studies suggest that the RRP is made up of a heterogeneous group of SVs, and that the relevant parameter for synaptic plasticity is not the RRP size as such, but rather the proportion between two components of the RRP (primed vs. unprimed or superprimed SVs; loosely vs. tightly docked SVs; or docked vs. replacement SVs) (Blanchard et al., 2020 [↗](#); Doussau et al., 2017 [↗](#); Eshra et al., 2021 [↗](#); Fukaya et al., 2023 [↗](#); Kim et al., 2024 [↗](#); Kobbersmed et al., 2020 [↗](#); Koppensteiner et al., 2022 [↗](#); Lin et al., 2022 [↗](#), 2025 [↗](#); Miki et al., 2020 [↗](#), 2016 [↗](#); Taschenberger et al., 2016 [↗](#)). These results open new perspectives for the interpretation of STP (Neher, 2023 [↗](#); Neher and Brose, 2018 [↗](#); Pulido and Marty, 2018 [↗](#); Schmidt, 2019 [↗](#); Silva et al., 2021 [↗](#)).

Electron microscopy studies of hippocampal synapses in culture showed that, following a single presynaptic action potential (AP) stimulation, the distance between SVs and the active zone (AZ) membrane varies as a function of time in a complex manner (Kusick et al., 2020 [↗](#); Ogunmowo et al., 2025 [↗](#)). Following the immediate loss of SVs by exocytosis, the number of SVs directly in contact with the AZ membrane rapidly increases (within ~15 ms), reflecting the movement of SVs from a cytosolic location to the membrane ('calcium-dependent docking'). Within ~100 ms, SVs move back into the cytosol ('calcium-dependent undocking'). In the following several seconds, SVs partially move forward again to the AZ membrane as the system returns to its basal state. While SV docking has been associated with facilitation (Miki et al., 2016 [↗](#); Schmidt, 2019 [↗](#); Silva et al., 2021 [↗](#)), the functional role of subsequent undocking has remained unclear. Here we explore the possibility that undocking could underly LFD.

Testing the undocking hypothesis of LFD necessitates a functional assay of various subclasses of RRP SVs, a notoriously difficult task (Neher, 2015 [↗](#)). Our laboratory has developed a method to monitor the release of individual SVs at single AZ synapses between parallel fibers (PFs) and molecular layer interneurons (MLIs) in cerebellar slices ('simple synapse recording' (Malagon et al., 2016 [↗](#))). Simple synapse recording has provided models of synaptic facilitation and of high frequency depression based on changes in occupancy of docking sites and associated SV groups inside and outside the RRP (Miki et al., 2016 [↗](#); Tanaka et al., 2021 [↗](#); Tran et al., 2022 [↗](#)). In the present work, we suggest that LFD reflects a specific SV re-equilibration inside the RRP, as well as an inhibition of SV entry into the RRP. We further propose a general model of synaptic depression which explains differences between low and high stimulation frequencies.

Results

Biphasic dependence of paired-pulse ratio on inter-stimulus interval

PF-MLI synapses are considered facilitating (Atluri and Regehr, 1998). In the present work, we ask whether single PF-MLI synapses may turn depressing at low frequency. Using simple synapse recordings at PF-MLI junctions, short trains of 4 presynaptic APs were applied at various inter-stimulus intervals (ISIs) in a scrambled order. Each train was separated from the next by a 10 s-long time interval, considered long enough to reset the system (Tran et al., 2023) (Fig. 1A). The resulting EPSCs were deconvolved using the mean quantal EPSC as kernel to yield the numbers of released SVs per AP (s_i , where i stands for AP number) (Malagon et al., 2016). Results of an example experiment are shown in Fig. 1 B-C. Fig. 1C, top shows average EPSC traces for the first AP (average s_1) and for the second AP at various ISI values. Fig. 1C, bottom shows, for each ISI, the ratio of mean released SV numbers for AP #2 over the corresponding mean for AP #1 (paired pulse ratio; PPR: blue dots; results for AP #3-4 are shown in Supplementary Fig. 1). In this experiment, the PPR defined in this manner displayed values > 1 for ISIs ranging from 10 to 100 ms, and values < 1 for ISIs ranging from 200 to 1600 ms. In group results, the PPR was consistently > 1 for ISI values of 100 ms or less, and < 1 for ISI values of 800 or 1600 ms (Fig. 1D). A biexponential fit to the data (red curve in Fig. 1D) intersected the PPR = 1 line for an ISI near 300 ms. These results show that the synapse changes from facilitating to depressing depending on the ISI. They differ from the single facilitation component previously reported for this synapse (Atluri and Regehr, 1998). This discrepancy presumably stems from differences in experimental conditions (room temperature, stimulation of multiple presynaptic PFs and 2 mM external Ca^{2+} concentration in the previous work, vs. near-physiological temperature, single presynaptic stimulation and 3 mM external Ca^{2+} here; see below). Here, the PPR is the sum of a fast facilitation component and of a slow depression component ($\text{PPR} = 1 + A_{\text{fast}} \exp(-t/\tau_{\text{fast}}) - A_{\text{slow}} \exp(-t/\tau_{\text{slow}})$, with $A_{\text{fast}} = 0.94$, $\tau_{\text{fast}} = 230$ ms, $A_{\text{slow}} = 0.30$, $\tau_{\text{slow}} = 2100$ ms).

Low frequency depression (LFD) can be modelled as reflecting a decrease in DS occupancy but no change in RS occupancy or in IP size

We previously developed a sequential 2-step model of docking at PF-MLI synapses (Miki et al., 2018, 2016; Tran et al., 2022) (Fig. 2A). In this model, called replacement site/docking site (RS/DS) model hereafter, each AZ contains a fixed number of docking units (DUs), with a mean value of 4 DUs per AZ (Miki et al., 2017). Each DU comprises two SV binding sites: a docking site (DS), which is directly attached to the AZ membrane, and an associated replacement site (RS). The RS can accommodate an SV regardless of whether the associated DS is occupied or not, meaning that a DU can simultaneously bind two SVs (Silva et al., 2024). The RRP is the sum of the SVs bound to all DUs (RS and DS) in the same AZ. High frequency AP trains induce a flow of SVs from an upstream intermediate pool (IP) to the RRP. Once in the RRP, SVs transit from the RS to the DS, and eventually undergo exocytosis (downward red arrows in Fig. 2A).

The RS/DS model of Fig. 2A has been used to interpret phenomena of STP during and after high frequency AP trains. In response to stimulation at 100 or 200 Hz, the s_i plot displays an initial increase up to $i = 2-4$, reflecting synaptic facilitation, followed by a gradual decrease reflecting synaptic depression (Miki et al., 2016; Tran et al., 2022). According to the model, the DS occupancy, δ , increases after a stimulus over its basal value, following Ca^{2+} -dependent movement of SVs from RS to DS. This explains the initial facilitation. Subsequent synaptic depression reflects a δ decrease following the gradual exhaustion of IP SVs and consequently, a decrease in the flow of SVs into the RRP. Recovery requires the refilling of the IP, and it occurs on a time scale of hundreds of ms.

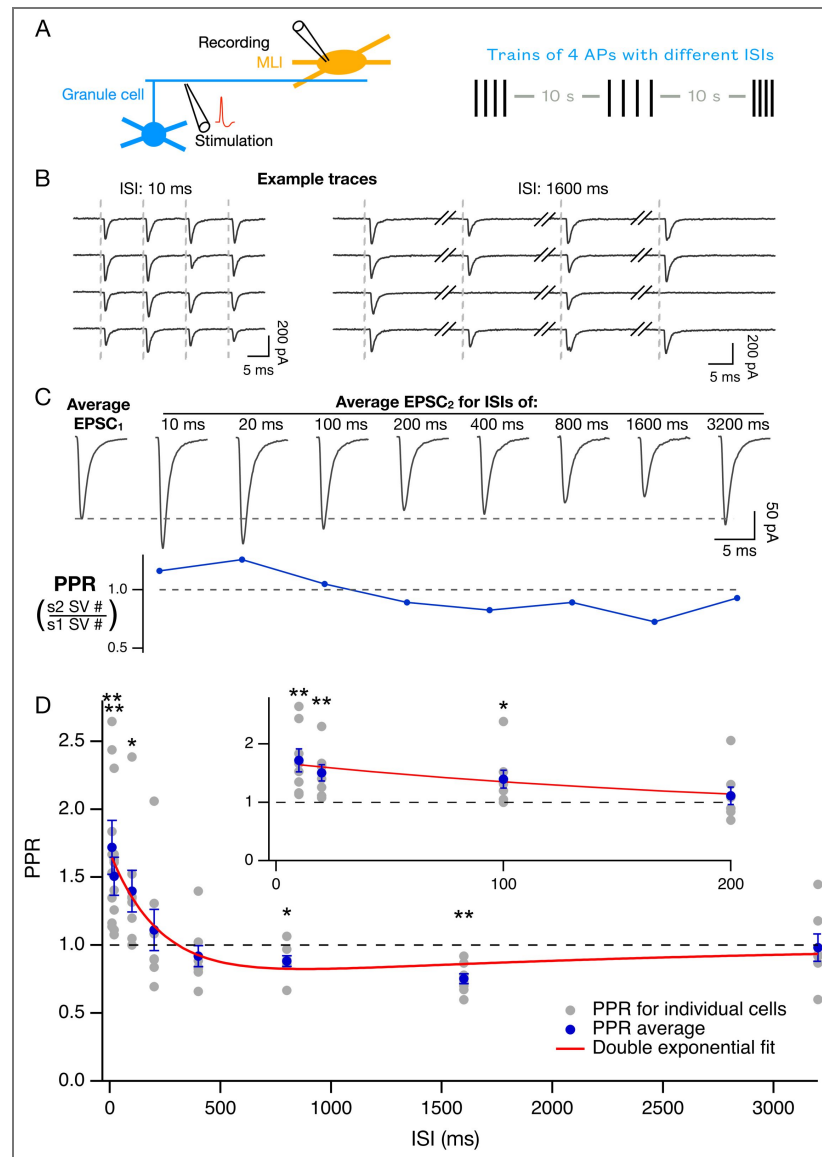


Figure 1. Dependence of paired-pulse ratio on inter-stimulus interval

A: Experimental protocol. To obtain simple synapse recordings, individual granule cell axons were stimulated with an extracellular pipette located in the granule cell layer, and EPSCs were recorded in a postsynaptic MLI. Presynaptic stimulations involved trains of 4 action potentials with various inter-stimulus intervals (ISIs), and with 10 s inter-train intervals. **B:** Example recording showing responses to trains using 10 ms ISIs (left) and 1 600 ms ISIs (right). Stimulation times indicated by dotted vertical lines. **C, Top:** Average EPSC in response to AP #1 (1st trace, labelled 'average EPSC1') followed by average responses to AP #2 for ISIs varying from 10 ms to 3200 ms, from the same experiment (20 repetitions for each ISI). **C, Bottom:** A measure of the paired-pulse ratio, PPR, calculated from the ratio of the mean numbers of released SVs for AP #2 over the average release for AP #1 from the same data. Numbers of SVs released by individual presynaptic stimulations were determined by deconvolution, using the mean quantal EPSC as kernel. **D:** Plot of PPR as a function of ISI (same analysis as in **C, bottom**: grey dots, individual cells (means from > 20 repetitions for each cell); blue dots and associated error bars, means ± sem from n = 8 cells). The red curve is a double exponential fit to the data with $A_{fast} = 0.94$, $\tau_{fast} = 230$ ms, $A_{slow} = 0.30$, $\tau_{slow} = 2100$ ms. **Inset:** Blow-up of results for ISIs of 200 ms or less. * and ** indicate data points that differ from PPR = 1 with p < 0.05 and p < 0.01 respectively.

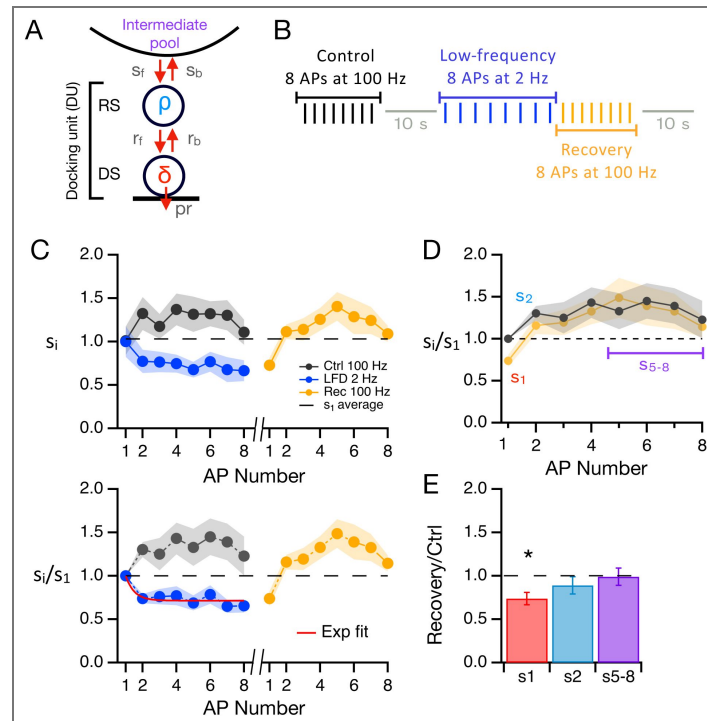


Figure 2. LFD reflects a decrease in docking site occupancy.

A: Sequential 2-state docking model. SVs coming from the intermediate pool (IP) transit through a replacement site (RS, with occupancy ρ) and an associated docking site (DS, with occupancy δ) before release. Taken together, one RS and one DS constitute one docking unit (DU). s_f , s_b , r_f and r_b represent transition rates as indicated. p_r represents the release probability of a docked SV following an AP. **B:** Experimental protocol. To probe the state of SV pools in the synapse, trains of 8 APs at 100 Hz were applied either in isolation (control train), or after a depressing train of 8 APs at low stimulation frequency (recovery train). **C, Top:** Plots (means $\pm$ sem from $n = 7$ cells; > 20 repetitions for each cell) of s_i for the control train (black), for the low frequency train (2 Hz, blue), and for the recovery train (yellow). Dotted line: Average s_1 value from control and low frequency trains. **C, Bottom:** Same, after normalization with respect to s_1 value. Exponential fit with asymptotic value = 0.69 ± 0.05 and time constant = 0.87 AP # ($= 435$ ms; red curve). **D:** Superimposed s_i plots for control (black) and recovery (yellow). Comparison of the SV release between control and recovery trains allows us to evaluate the occupancy of the RRP and of the IP at the end of the low frequency train. SV numbers in response to the 1st (s_1), 2nd (s_2), and last (s_{5-8}) APs respectively report changes in δ (red), in ρ (blue), and in the IP size (purple). **E:** Ratios of s_i values between recovery 8-AP train and control train. s_1 is significantly reduced, indicating a decrease in δ , but neither s_2 nor s_{5-8} are changed, indicating no change in ρ or in IP size.

To explore the mechanisms underlying STP as a function of AP frequency, we alternated 8-AP stimulations at 100 Hz and at 2 Hz. Immediately after the 2 Hz train, we repeated the 100 Hz train, before repeating the entire protocol again after a pause of 10 s, to allow for synapse recovery (Fig. 2B). According to our previous work (Tran et al., 2022), and within the framework of the RS/DS model, we can estimate the occupancy of the RRP and the intermediate pool (IP) during the low frequency stimulation by comparing the s_i curves of the 100 Hz trains before (control) and immediately after (recovery) the 2 Hz train. Under the hypothesis that the release probability of docked vesicles (p_r) remains constant throughout the AP train, changes in δ , in RS occupancy (ρ) and in IP size are reflected by changes in the number of released SVs for AP #1 (s_1), for AP #2 (s_2), and for the average of AP #5-8 (s_{5-8}) respectively. This analysis was derived under conditions of elevated release probability. Therefore, in most of our experiments, we used an increased external Ca^{2+} concentration (3 mM) to facilitate the interpretation of the results in terms of changes in RRP and IP occupancy; results obtained under physiological external Ca^{2+} concentration conditions will be reported below.

The results of these experiments are shown in Fig. 2C-E. In keeping with the results of Fig. 1, when stimulating at 2 Hz in 3 mM external Ca^{2+} concentration, a clear synaptic depression was observed (PPR = 0.74 ± 0.05 , paired one-tailed t-test, $p = 0.0027$, $n = 7$; blue traces in Fig. 2C; absolute SV numbers in Fig. 2C top, and values normalized to control s_1 value in Fig. 2C bottom). The s_i curve for 2 Hz trains could be approximated to an exponential decay with an asymptote of 0.69 ± 0.05 (paired one-tailed t-test, $p = 0.0032$, $n = 7$) and a time constant of 0.87 ISIs (435 ms; red curve in Fig. 2C bottom). We then compared release during control and recovery trains to investigate the origin of the synaptic depression (Fig. 2D). The value of s_1 at the onset of the recovery train replicated the stable low s_i values observed at the end of the low frequency train. It displayed a decrease with respect to the s_1 value of control trains (to 70 % of this control; $p = 0.02$, one-tailed paired t-test; Fig. 2C bottom). As s_1 is a proxy for δ under the present experimental conditions, this result suggests that LFD involves a decrease in δ , as previously suggested for high or moderate frequency depression (HFD) (Borges-Merjane et al., 2020; Lin et al., 2022; Miki et al., 2016; Tanaka et al., 2021). Strikingly, s_i values during the recovery train caught up with control values immediately after the first stimulation, so that neither s_2 nor s_{5-8} values differed from those of control 100 Hz trains ($s_2 = 1.30 \pm 0.09$ for control and 1.16 ± 0.10 for recovery; $p > 0.05$; $s_{5-8} = 5.05 \pm 0.68$ for control and 5.02 ± 0.63 for recovery; $p > 0.05$, one-tailed paired t-test; compare yellow and black s_i curves in Fig. 2D; summary graph in Fig. 2E). Within the framework of the RS/DS model, these results suggest that neither ρ nor the IP size were decreased at the end of the low frequency train. Taken together, the results suggest similarities and differences between the mechanisms of synaptic depression at high frequency, as studied before, and at low frequency, as studied here. In both cases, the RS/DS model suggests that synaptic depression is accompanied by a decrease in δ . However, while ρ and IP pool size accompany the δ decrease during synaptic depression at high stimulation frequency (Tanaka et al., 2021; Tran et al., 2022), only δ is changed during LFD.

LFD as a function of stimulation frequency and external calcium concentration

Having documented LFD at 2 Hz stimulation frequency and in 3 mM external Ca^{2+} concentration as shown in Fig. 2, we next examined the results of similar experiments when changing the stimulation frequency to 1 Hz, or when reducing the external Ca^{2+} concentration to near physiological concentration (1.5 mM). When stimulating at 1 Hz, LFD was obtained (Supplementary Fig. 2A, blue triangles; fractional response at steady state: 0.79 ± 0.08 , paired one-tailed t-test, $p = 0.0037$, $n = 5$). As was the case with 2 Hz stimulations, the s_i curve recovered immediately after the first stimulus during the recovery train (yellow curves in Supplementary Fig. 2A). Next, we examined synaptic responses at 2 Hz in 1.5 mM external calcium, finding no depression (Supplementary Fig. 2B). At this calcium concentration, synapses facilitate more than at 3 mM (compare control black normalized traces in Fig. 2C bottom in 3mM Ca^{2+} , and Supplementary Fig. 2B-C right in 1.5 mM Ca^{2+}), potentially causing a longer-lived facilitation

t would mask the depression. To test this theory, we performed the same protocol at 0.5 Hz in 1.5 mM Ca^{2+} . While the PPR was not significantly < 1 (PPR = 0.83 ± 0.11 , $n = 6$; $p = 0.21$, paired two-tailed t-test), the rest of the train led to a reduced number of released SVs compared to control s_1 (fractional response at steady state: 0.69 ± 0.05 , $n = 6$; $p = 0.00058$, paired one-tailed t-test), indicating LFD. An exponential fit to the 0.5 Hz data (red curve in Supplementary Fig. 2C, right) indicated a τ of 1.8 stimuli (3.6 s), slower than that observed for higher calcium concentrations at 2 Hz. The recovery curve (yellow, Supplementary Fig. 2C) showed a full recovery to control values once high frequency stimulation was resumed.

In conclusion, LFD can occur at 0.5 Hz, 1 Hz or 0.5 Hz; it occurs at near-physiological external Ca^{2+} concentration (1.5 mM), but a lower frequency is required compared to 3 mM Ca^{2+} (0.5 Hz instead of 2 Hz). These experiments further confirm that LFD is immediately reversed upon stimulation with high frequency.

Modelling the dependence of PPR on ISI

The facilitation/depression curve of Fig. 1 matches the docking/undocking sequence reported in flash-and-freeze experiments (Kusick et al., 2020). Therefore, we next asked whether the RS/DS model (Fig. 2A) could account for the dependence of the PPR on ISI on the basis of δ changes (constant p_r hypothesis). The sequence of events following an AP is shown in Fig. 3. Immediately after an AP, docked SVs were released with a probability p_r , resulting in a drop in δ (grey area, Fig. 3A). We modelled subsequent SV movements based on transition rates r_b , r_f , s_b , s_f (Fig. 2A; (Lin et al., 2022; Miki et al., 2018)) with appropriate time-dependent changes (Supplementary Fig. 3, Table 1, and Materials and Methods). This revealed a time window comprised between 10 and 200 ms when δ (red curve) was larger than its resting value of 0.6 (dotted red line, Fig. 3A), while ρ (blue curve) was lower than its resting value of 0.9 (dotted blue line, Fig. 3A). This time period corresponds with the facilitating part of the PPR vs. ISI plot (Fig. 3B). Due to the low ρ value, SV entry into the RRP (IP to RS transition) was facilitated. We call this the 'RS gate open' configuration (green shade in Fig. 3A). By contrast, during the depressing part of the curve (Fig. 3B, ISI values comprised between 500 ms and 10 s), δ was smaller than its resting value while ρ recovered near its resting value, and the 'RS gate' was closed (an SV cannot move into the RS if it is already occupied; red shade in Fig. 3A). The predicted PPR plot (red curve in Fig. 3B) is proportional to the δ curve of Fig. 3A under the assumption of constant p_r . It displays a good fit to the data (black dots), showing that the model can account for the facilitation/depression sequence of Fig. 1 based on changes in DS occupancy.

Modelling synaptic output in response to AP trains at high or low frequency

Next, we examined whether the same model could account for the results of Fig. 2 (LFD for 8-AP trains and following recovery). To this end, we made concatenations of the previous model over consecutive time segments (Materials and Methods). Data and simulations of 8-AP trains at high frequency (Fig. 4A), low frequency (Fig. 4B), and recovery trains at high frequency (Fig. 4C) illustrate the consequences of the time-dependent changes of DU occupancy shown in Fig. 3. Upper panels (Fig. 4A, B and C) show s_i plots together with simulation results, indicating that the model can reproduce the main features of Fig. 2C data. Lower panels (Fig. 4A, B and C) show plots of δ and ρ as a function of AP number. At high stimulation frequency (Fig. 4A), δ increases and remains high throughout an 8-AP train, explaining facilitation (red curve). Meanwhile, ρ decreases below its resting level (blue curve). Because SV movement from the IP into the RRP requires a free RS, this ρ decrease facilitates RRP replenishment. In effect, the RS gate opens at AP #2 and remains open throughout the 8AP train (Fig. 4A, bottom; Fig. 4D, green shade). At low stimulation frequency, a totally different pattern of changes is observed (Fig. 4B; Fig. 4D, blue shade). δ decreases at AP #2 and remains low throughout the train, explaining LFD. ρ remains close to its resting value of 0.9, so that the RS gate is closed before each AP of the train (Fig. 4B, bottom; note that this figure shows DU parameters just before each AP, and that it

Figure 3. Simulations of s , δ and ρ curves after an AP stimulation

A, up: Model depiction of an AZ with 3 DUs at various time periods before and after an AP stimulation. After release, SVs transition from RS (blue; some of these SVs are placed sidewise as the 2-step model cannot specify the exact location of the RS) to DS (white) within tens of ms (transient docking), then after hundreds of ms they undock, before eventually returning to their basal state. **A, bottom:** Simulated time course of δ (red curve) and ρ (blue curve) before and after a presynaptic AP (at time 0). Resting δ and ρ values indicated by dotted lines. During transient docking, δ is high and ρ is low, so that RRP replenishment is allowed (RS gate open, green shade). During undocking, δ is low and ρ is high, so that RRP replenishment is blocked (RS gate closed, red shade). **B:** PPR curve as a function of ISI (black dots, from Fig. 1) together with simulation results (red curve).

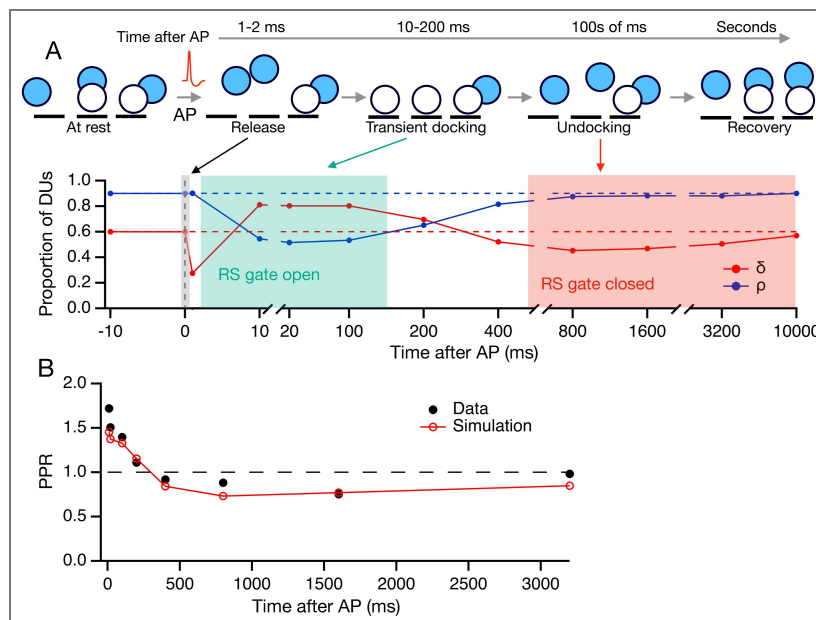


Table 1.

Simulation parameters	Fig. 3	Fig. 4	Transition Probabilities	Fig. 3	Fig. 4
N	3.8	3	EE	0.7	0.4
p_r	0.55	0.55	EU	0.15	0.31
ρ	0.9	0.9	ED	0.15	0.29
δ	0.6	0.6	UU	0.2	0.2
r	0.8	0.8	UD	0.6	0.47
s	0.4	0.6	UF	0.2	0.33
			DD	0.7	0.4
			DF	0.3	0.6

does not display the time-dependent DU changes during ISIs illustrated in Fig. 3 (blue). These results show that both at high and at low frequency, the proportions of DU states that are reached after the first ISI are kept without major changes throughout the 8-AP train.

The simulation of the recovery train used the same transition parameters as the control train. The initial ρ and δ were obtained from simulation values at the end of LFD (Fig. 4B, bottom blue). As can be seen by comparing Fig. 4C, bottom blue with Fig. 4A, bottom blue, DUs behave identically to control from AP #2 to AP #8, in agreement with the data. Thus, our model predicts that a synapse where δ has decreased following LFD can return to its basal state within a single ISI of 10 ms when challenged with a high frequency stimulation (Fig. 4D blue, yellow shade). If a high frequency train is continued the synapse enters the HFD loop (Fig. 4D blue, green shade).

LFD does not require previous SV release events

The results and simulations of Figs. 2 (blue)-4 (blue) suggest that LFD reflects a drop in δ , but they leave the question open as to whether the δ decrease is due to exocytosis, to SV return to RS (undocking), or to both. To better appreciate the share of each of these two mechanisms in LFD, we re-examined LFD experiments depending on the value of s_1 . If s_1 is nil, the first theory predicts no synaptic depression in response to the 2nd AP, because there was no prior loss of SVs by exocytosis. Conversely, a failure of release after AP #1 should not prevent synaptic depression according to the second theory. We set aside EPSC sequences during LFD where the 1st AP resulted in release failure ($s_1 = 0$; labelled with a star in Fig. 5A blue). We pooled together results obtained with ISI values of 500, 800 and 1600 ms, as the PPR did not appear to depend on the ISI value in this range (PPR = 0.74 ± 0.05 , 0.88 ± 0.04 , and 0.75 ± 0.04 for ISIs of 500 ms, 800 ms and 1600 ms respectively). We calculated separately mean s_2 values for the failure traces (noted $\langle s_2 \rangle_{s_1=0}$) as well as overall mean s_2 values (noted $\langle s_2 \rangle_{all}$). We obtained two separate LFD values (LFD_{fail} and LFD_{all}) by dividing $\langle s_2 \rangle_{s_1=0}$ and $\langle s_2 \rangle_{all}$ with the overall s_1 mean, $\langle s_1 \rangle_{all}$. With this normalization procedure, the normalization factor was the same for successes and for failures. The two LFD values were significantly < 1 ($p = 0.004$ for LFD_{fail} and $p < 0.0001$ for LFD_{all}, one-tailed, one-sample Wilcoxon rank test). Strikingly, we found that the two LFD values were similar (LFD_{fail} = 0.84 ± 0.05 vs LFD_{all} = 0.79 ± 0.03 ; $p = 0.24$, unpaired onetailed Wilcoxon rank test; Fig. 5B, left blue), indicating that the value of the LFD does not depend on whether the 1st AP leads to a failure or to a response.

Next, we expanded the above analysis by sorting traces displaying successful EPSCs as a function of s_1 . Experimental LFD values were < 1 in all cases: 0.79 ± 0.07 for $s_1 = 1$; 0.75 ± 0.10 for $s_1 = 2$; 0.73 ± 0.12 for $s_1 > 2$; respective one-tailed p values for LFD < 1 : 0.002, 0.01 and 0.03 (Fig. 5B, right blue). We finally asked whether these results were compatible with our undocking model. Simulation results using the model of Fig. 3 (blue) gave: LFD_{fail} = 0.84; LFD_{all} = 0.73; LFD _{$s_1=1$} = 0.77; LFD _{$s_1=2$} = 0.70; LFD _{$s_1>2$} = 0.59. These values are close to experimental data (Fig. 5B blue, black dots vs. bars). This agreement is remarkable since the model predicted LFD values without any parameter adjustment. Notably, the finding of identical experimental and simulated LFD_{fail} values (0.84 in both cases), is compatible with the undocking mechanism of Fig. 3 (blue) for LFD.

When comparing the evolution of DUs after SV release vs. after failure in the RS/DS model, it appears that the difference in DS occupancy resulting from release in the first case is compensated in a time frame on the order of 10 ms due to the high value of the RS $\rightarrow$ DS transition rate. Because of this short recovery time, DUs that have released display very similar δ (before the 2nd AP) to those that have failed to release after the 1st AP, leading to almost identical s_2 values in the two cases.

In conclusion, the above results indicate that LFD occurs independently of the amount of SV release following AP #1. They support the hypothesis that LFD reflects a shift of SV occupancy favoring RSs over DSs (undocking), rather than RRP depletion caused by exocytosis.

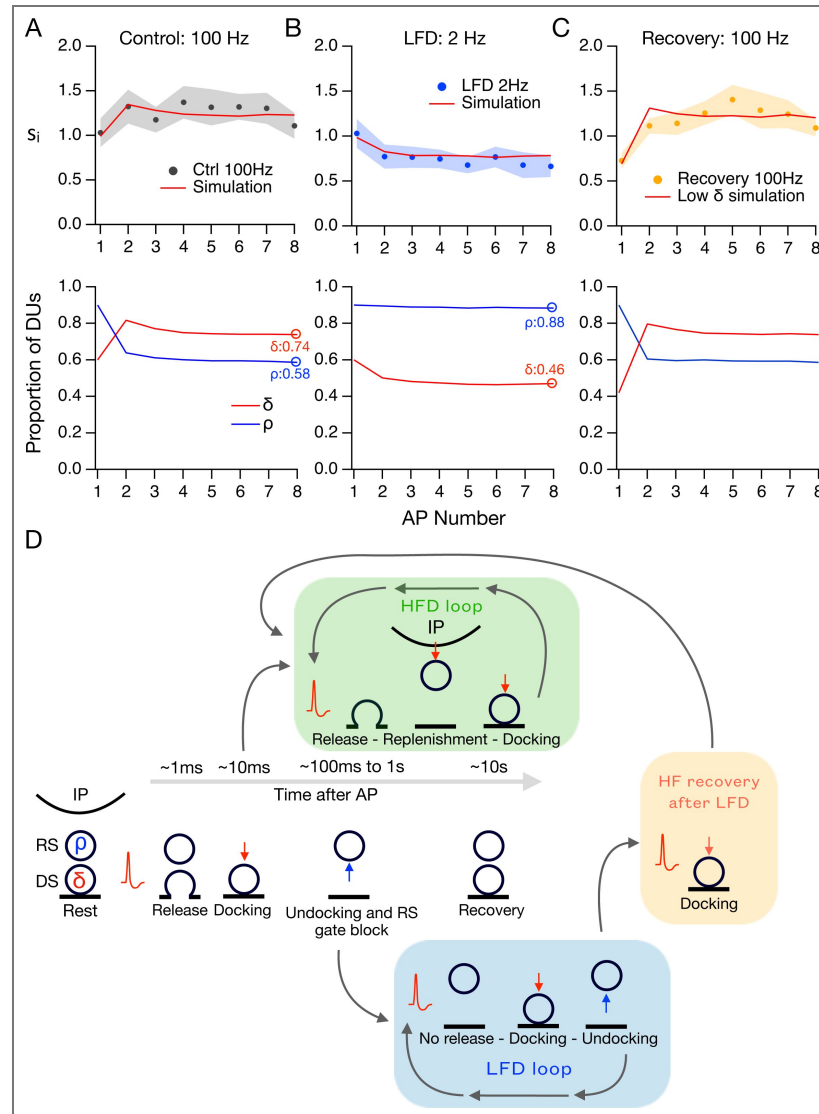


Figure 4. Simulation of SV movements during HFD and LFD

A, upper panel: s_i curve during control runs with 8 APs at 100 Hz (black dots: data from Fig. 2; red curve: simulated results; model parameters in Table 1). **A, lower panel:** simulated values of δ (red) and p (blue) observed just before AP stimulations, as a function of AP #. **B:** Same as **A**, but during 2 Hz 8-AP stimulations (data from Fig. 2). Note the behavior of δ and p compared to **A**, here p remains stable throughout the train, while δ undergoes a sharp initial decrease. At the end of the train there is a low δ value and a high p value for 2 Hz trains, while at the end of a 100 Hz 8-AP train (**A**) there is a relatively high δ and a low p . **C:** Same as **A**, but during recovery (100 Hz; data from Fig. 2). The initial δ and p values for the recovery train simulation are the end-of-train values from the 2 Hz simulation. **D:** Proposed SV movements during high-frequency depression (HFD) and LFD. **Central sequence:** Proposed timeline of SV movements inside a DU following an AP stimulation. **HFD loop:** Proposed sequence of SV movements during HFD. The DU exits the main timeline at 10 ms after the AP to enter a loop with repetitive high frequency AP stimulation, which results in high rates of exocytosis and RRP replenishment, as the RS gate is open (green shade). Eventually, SV depletion leads to depression. **LFD loop:** proposed sequence of SV movements during LFD. Here the DU exits the main timeline as the RS gate is closed, and it undergoes an idle docking-undocking cycle at each ISI (blue shade). **HF recovery after LFD** proposed sequence of SV movements during high frequency recovery train after LFD (yellow shade). The SVs that were in the RS can dock if high frequency is applied at any point during an LFD train; if further stimuli are applied, the DU is transferred to the HFD loop (green shade) after the 2nd AP.

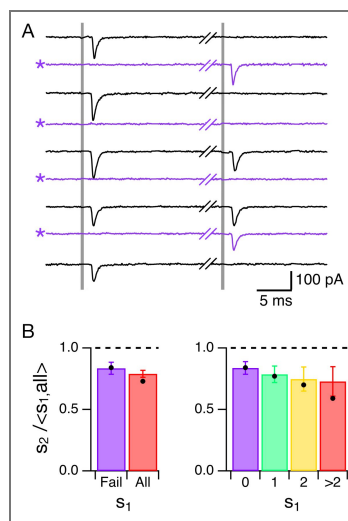


Figure 5. RRP depletion does not cause LFD.

A: Exemplar traces from an LFD experiment, illustrating responses to 1st and 2nd AP (AP times indicated by vertical grey lines) during low-frequency trains. Black traces show EPSCs including a success in response to the 1st AP, while purple traces (marked with a star) show EPSCs when there was a failure in response to the 1st AP. **B, left:** Group data showing a similar extent of LFD (mean $\pm$ sem of LFD across cells; $n = 7$ cells for 500 ms; 8 cells for 800 and 1600 ms) when there was a failure in response to the 1st AP (purple bar; m. $\pm$ sem) compared to the corresponding data taken from all traces (red bar). The LFD value was obtained by calculating the ratio of the mean numbers of released SVs for the second AP over the average release for the first AP for all trials (failures and successes). **B, right:** Same analysis, except that now LFD values are calculated separately for each s_1 value. Dots indicate predictions from the model of Fig. 3.

Long AP trains at low frequency reveal two phases of LFD occurring on widely different time scales

Earlier studies showed a slow onset LFD during long trains at low frequency (Abrahamsson et al., 2007; Doussau et al., 2010; Rudolph et al., 2011). We therefore searched for an additional slow component LFD at PF-MLI synapses. In each experiment, a series of control trains (8-APs @ 100 Hz) was presented first. Next, a long low frequency train was applied (200 APs @ 2 Hz in Fig. 6A), followed by a recovery train (50 or 100 APs @ 100 Hz; Fig. 6B; protocol in Fig. 6A, inset). Numbers of released SVs per AP (s_i) were normalized with respect to the s_1 value obtained for the control trains. During the long train, a plot of the normalized s_i (indicated as $s_i/\text{ctrl } s_1$) vs AP # displayed an initial decrease followed by a gradual depression (Fig. 6A). We call this initial drop and the subsequent linear decrease the first and second phase of LFD respectively. A linear fit to these data (red line in Fig. 6A) indicated a first phase depression (within the first 10 APs) to $75 \pm 2\%$ of the control, consistent with the previous results obtained with paired stimulation and short train protocols (Figs. 1 and 2). The second phase of the depression exhibited a slope of $-0.15 \pm 0.02\%$ per AP, such that the value reached for the linear fit at the end of the 200-AP train was 45 % of the control. A similar LFD was seen at 1 Hz with a first phase drop to $51 \pm 2\%$ of control and a slope of $-0.05 \pm 0.02\%$ reaching 40 % of the control at the end of the 200-AP train ($n = 17$ trials from 6 cells).

We next investigated δ , ρ and IP pool size at the end of long low frequency trains with the previous $s_1/s_2/s_5-s_8$ approach. As expected, the first response of the recovery train was severely depressed compared to s_1 of control trains (recovery $s_1 = 37.6 \pm 20.2\%$ of control s_1), but s_i plots during control and recovery trains were indistinguishable for i values of 2 or more (s_2 ratio: 1.01 ± 0.20 ; s_{5-8} ratio: 1.19 ± 0.23 ; Fig. 6C-D), suggesting again a drop in δ without changes in either ρ or IP size.

The $s_1/s_2/s_5-s_8$ analysis assumes a constant p_r . To test this hypothesis, we determined the parameters of the RS/DS model by fitting responses to control 8-AP trains (Fig. 6E). Next, responses to the first 8 APs of the recovery train were predicted by keeping all parameters the same as control except for either δ or p_r . In two different simulations, δ and p_r were determined to fit the s_1 point of the recovery curve (Fig. 6F, red and black curves respectively). While the δ decrease prediction closely matches the data (Fig. 6F, yellow curve), the p_r decrease prediction fails to simulate the recovery train. Altogether, these results suggest that the slow phase of LFD illustrated in Fig. 6A, like the rapid phase shown in Fig. 2C, reflects a δ decrease without significant changes in p_r , in ρ or in IP size.

In these experiments the delay between the LFD train and the recovery train was variable. A plot of the value of s_1 for the recovery plot as a function of the time interval since the end of the LFD train is shown in Supplementary Figure 4. This plot shows that recovery s_1 remains depressed after rest periods of up to 1 min after the end of the LFD train. These results suggest that, while SV re-equilibration within the RRP occurs very rapidly under elevated calcium concentration, as presumably occurs during the recovery trains, they occur very slowly under resting calcium concentration.

Finally, a full recovery train (50 or 100 APs @ 100 Hz) elicited a deep synaptic depression (high frequency depression, or HFD; Fig. 6B), similar in extent and in time course to the depression observed for stand-alone trains at high frequency. An exponential fit to these data (red curve in Fig. 6B) displayed a time constant of 412 ms (41.2x the ISI) and an asymptotic value of 13% of the control s_1 value. This type of depression, unlike LFD, was previously suggested to involve a decrease in ρ and in IP size in addition to δ (Tran et al., 2022).

LFD is not due to stimulation failures

One tacit assumption of our analysis so far is that LFD reflects a decreased synaptic response to presynaptic stimulations. We next asked whether LFD could instead originate in erratic stimulation failures. This possibility was plausible since the failure rate increased markedly during the second phase of LFD (Supplementary Figure 5A). To test it, we compared the changes

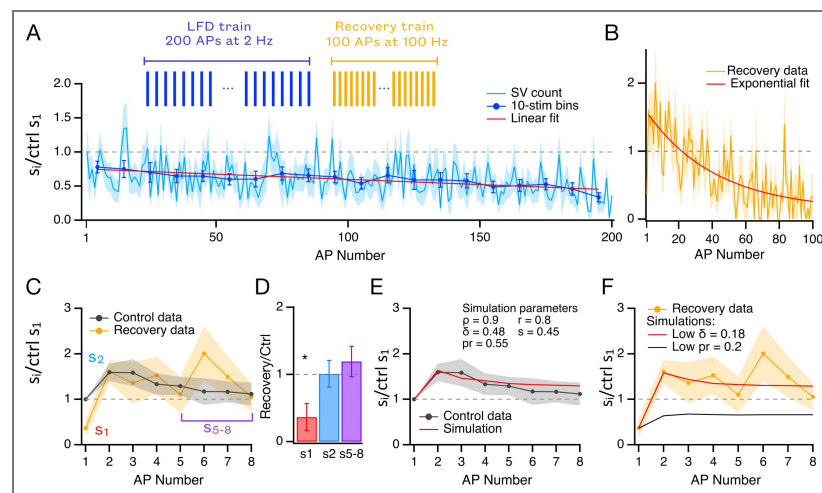


Figure 6. Prolonged low frequency trains produce a gradual decrease in docking site occupancy.

A: A long AP train at low frequency (200 APs @ 2 Hz) was followed by a long recovery train (50-100 APs @ 100 Hz; experimental protocol in **insert**). Normalized plot of s_i during long low frequency train, showing an initial rapid depression (as in Fig. 2 [B](#), notice the reduction in release for the first binned data point) followed by a gradual depression during the entire train duration (light blue: mean $\pm$ sem of individual trials; dark blue dots and associated error bars: binned data for 10 consecutive APs; red: linear fit to the data; $n = 16$ trials, 8 cells). **B:** Time course of synaptic depression during the recovery train, with exponential fit (red; time constant: 41.2 ISIs, or 412 ms). $n = 6$ trials, 3 cells for 50 APs, 6 trials, 4 cells for 100 APs. **C:** A series of control trains (8-APs @ 100 Hz, separated by 10 s-long inter-train intervals) were recorded before the long AP train to establish the characteristics of the synapse. Normalized s_i plots for this control and the first 8-APs of the recovery 100 Hz train are compared here. s_1 , s_2 and s_{5-8} are indicated in the plot. **D:** The s_1 ratio (red) between recovery and control trains is < 1 , indicating a decrease in δ at the end of the low frequency train. By contrast, neither s_2 (blue) nor s_{5-8} (purple) are changed, suggesting no change in p or in IP size. **E:** Simulation of control trains shown in **C** (black dots: average normalized data; red curve: simulation). **F:** Simulation of recovery trains shown in **C** (yellow dots: average normalized data). Decreasing the release probability from $p_r = 0.55$ to $p_r = 0.2$ without changing the other simulation parameters fails to provide a satisfactory fit of recovery data (black curve), while decreasing the DS occupancy from $\delta = 0.48$ to $\delta = 0.18$ without changing the other simulation parameters provides a satisfactory fit of the recovery data (red curve).

in the distributions of released SV numbers predicted from two LFD models. In the first model, during LFD, each DU releases less SVs when stimulated, presumably due to a reduction in δ . In the second model, a fraction of the stimulations fails to elicit a presynaptic AP during LFD, but the probability of release for each stimulated synapse remains constant (**Materials and Methods**). Both models were constrained to reproduce the observed mean number of failures during LFD. Observed distributions of released SV numbers during LFD were compatible with the decreased δ model but not with the stimulation failure model (Supplementary Figure 5 [↗](#)). We conclude that stimulation failures cannot account for LFD, consistent with the above interpretation that LFD reflects a drop in δ , as well as with previous results in other preparations (Saviane et al., 2002 [↗](#); Silverman-Gavrila et al., 2005 [↗](#))

LFD is not blocked by postsynaptic BAPTA

At synapses between PFs and Purkinje cells, low frequency stimulation leads to a gradual synaptic depression that has been attributed to a transsynaptic mechanism (Casado et al., 2000 [↗](#)). Since the simple synapse method is based on SV counts rather than EPSC amplitude, there is no doubt that the expression of LFD at PF-MLI synapses is presynaptic. Nevertheless, as AMPAR activation elevates the postsynaptic Ca^{2+} concentration (Soler-Llavina and Sabatini, 2006 [↗](#)), we included BAPTA (10 mM) in the MLI recording solution to test the possibility of a retrograde LFD mechanism that would be driven by postsynaptic Ca^{2+} . We applied a long 2 Hz train (100 APs @ 2 Hz) followed by a short 8-AP recovery train (experimental protocol in Supplementary Figure 6 [↗](#), insert). The result is very similar to Fig. 6 [↗](#) data (compare Supplementary Figure 6 [↗](#) to Fig. 6A [↗](#) and C [↗](#)) with a first phase drop to $65 \pm 3\%$ of the control and a second phase slope of $-0.2 \pm 0.05\%$ per AP. In this case recovery trains were applied immediately following the LFD train (whereas waiting times of 10 s of s were included in the experiments of Fig. 6 [↗](#)), and the recovery is identical to that observed in Fig. 6C [↗](#), suggesting likewise a reduction in δ with no associated change in p or IP size.

The similarity of results with and without a postsynaptic calcium buffer is consistent with a purely presynaptic mechanism for LFD.

LFD is not due to decreased Ca^{2+} transients

Our recovery results and simulations suggest that LFD is due to a decrease of δ , not of p_r (Fig. 6F [↗](#)). In line with this notion, earlier work reported no change in Ca^{2+} transients at PFMLI synapses for paired stimulations at short ISIs (Brenowitz and Regehr, 2007 [↗](#); Malagon et al., 2020 [↗](#); Miki et al., 2016 [↗](#)). Nevertheless, it seemed possible that a different situation would prevail with multiple stimulations and/or at longer ISIs.

To explore potential changes of presynaptic Ca^{2+} signaling during AP trains, we preloaded individual granule cells with the calcium dye OGB6F (1-2 mM in pipette solution; duration of preloading: 90 s; (Rebola et al., 2019 [↗](#))). After allowing diffusion of the calcium dye in the axon, we imaged single PF varicosities using two-photon laser microscopy (Fig. 7A [↗](#), and **Supplementary Information**), and we measured Ca^{2+} -dependent changes in fluorescence during 50-AP train stimulations at 1 Hz, or 200-AP train stimulations at 2 Hz, using stimulation conditions similar to those of the simple synapse experiments of Fig. 6 [↗](#). To avoid photodamage and photobleaching, imaging was interrupted after the 5th AP in a train, and it was resumed for the last 5 APs of the train. Traces were averaged from 3-7 trials in each varicosity to improve the signal-to-noise ratio (Fig. 7A2 [↗](#)). Averaged traces across varicosities revealed little changes in Ca^{2+} -dependent fluorescence transients during trains either at 1 Hz (Fig. 7B [↗](#)) or at 2 Hz (Fig. 7C [↗](#)). DF/F_0 values for 10 varicosities from 8 granule cells at 1 Hz were: $19.1 \pm 3.9\%$ for the 1st AP; $20.4 \pm 2.8\%$ for the 2nd AP; $19.3 \pm 2.8\%$ for the last AP; for 9 varicosities from 6 granule cells at 2 Hz: $19.4 \pm 2.9\%$ for the 1st AP; $17.9 \pm 1.6\%$ for the 2nd AP; $16.4 \pm 1.9\%$ for the last AP; $p > 0.05$ in 1 vs. 2 and 1 vs. last, paired comparisons at both frequencies, onetailed Wilcoxon Rank test (Fig. 7D [↗](#)). These data indicate that Ca^{2+} -dependent fluorescence transients were the same, within the limits of sensitivity of our measurements, in response to the first, the second, or the last AP in a train. In the same data set, we found no difference between basal fluorescence levels at the end vs.

the beginning of the trains at 1 Hz (respectively, 107 ± 12 a. u. vs. 103 ± 10 a. u.; $p > 0.05$, one-tailed Wilcoxon paired rank test; black markers in Fig. 7E1 [↗](#)), but we found an augmentation of basal fluorescence at the end of the trains compared to the pre-stimulation value at 2 Hz (118 ± 10 a. u. vs. 111 ± 9 a. u.; $p = 0.026$, one-tailed Wilcoxon paired rank test; blue markers in Fig. 7E1 [↗](#)). At 2 Hz, after the end of the train, the fluorescence trace relaxed to near its pre-stimulation value with a time constant of 750 ms, confirming that the basal Ca^{2+} concentration had increased during the train (Fig. 7E2 [↗](#)).

In conclusion, we find no evidence of decreased Ca^{2+} transients during trains, either at 1 Hz or at 2 Hz. This is in line with our earlier conclusion that p_r changes do not contribute strongly to LFD. We find a gradual increase of pre-stimulation Ca^{2+} concentration with 200 APs at 2 Hz, but not with 50 APs at 1 Hz. This effect may contribute to the second phase of LFD at 2 Hz.

Doublet stimulations inhibit LFD

Our model predicts a high ρ value during LFD that inhibits DU replenishment by preventing the transition from the IP to the RS (RS gate closed). We next tested the prediction that during a low frequency train, ρ remained elevated while δ decreased (Fig. 4B [↗](#)), by giving two successive stimuli with a 10 ms interval instead of single stimuli (doublet stimulations; Fig. 8A-B [↗](#)). According to the $s_1/s_2/s_{5-8}$ analysis, while the time course of the first response, s_1 , should give the evolution of δ , the time course of the second response, s_2 , should give the evolution of ρ (Tran et al., 2022 [↗](#)). With the same experiments, we intended to test whether doublet stimulations would open transiently the RS gate and thus interfere with the time course of LFD, as suggested by earlier work at the PF-Purkinje cell synapse (Doussau et al., 2017 [↗](#)).

We performed these experiments with a longer than usual ISI (5 s) to provide ample time for SVs to equilibrate after each stimulus, either singlet or doublet. Control single AP train stimulation experiments at 0.2 Hz revealed LFD (0.75 ± 0.11 ; paired one-tailed t-test, $p = 0.037$, $n = 8$; exponential fit of first 12 normalized si points had a time constant of 3.6 ISIs, or 18 s). We then examined the extent of LFD in response to the first 20 doublet stimulations at low frequency. We found that s_1 values were reduced compared to control (normalized s_1 value during 20 first trials at 0.2 Hz: 0.80 ± 0.10 ; $p = 0.008$, two-tailed, one-sample t-test; Fig. 8C [↗](#)). These results indicate that doublet stimulations at 0.2 Hz produced significant LFD when considering the response to the 1st stimulus. By contrast, s_2 values were not affected (s_2 values normalized to control s_1 : 1.18 ± 0.11 for control 100 Hz trains; 1.11 ± 0.10 during 20 first doublet trials; $p > 0.05$, two-tailed, one-sample t-test; Fig. 8C [↗](#)). These results are consistent with our previous conclusion that LFD involves a decrease in δ without significant changes in ρ .

Next, we compared the plots of normalized released SV numbers as a function of AP number (i) for singlet stimulations (s_{is} ; light blue), for the 1st AP of doublet stimulations (s_{id1} ; dark blue), and for the 2nd AP of doublet stimulations (s_{id2} ; orange; Fig. 8 D-E [↗](#)). In each case, the curves could be approximated to linear decay curves as a function of i ($y = \alpha - \beta i$), but the line parameters differed for the 3 curves. Values of α were similar for s_{id1} and for s_{is} (0.79 ± 0.07 and 0.88 ± 0.11 respectively; $p = 0.53$, two-tailed, two-sample t-test). This indicates that the first phase of LFD occurs with a similar amplitude for singlets and for doublets. On the other hand, values of β were markedly smaller for s_{id1} than for s_{is} ($\beta_{d1} = 0.0004 \pm 0.0014$ vs. $\beta_s = 0.0063 \pm 0.0018$; $p = 0.011$, two-tailed, two-sample t-test; Fig. 8D [↗](#) and Fig. 8E, top [↗](#)). These results suggest that the second phase of LFD is less pronounced for doublet stimulations than for singlet stimulations, with a slope ratio near 15-fold. The results for d_2 can be interpreted as a return of ρ values close to control at the end of each 5 s ISI throughout the train (orange curve in Fig. 8E, bottom [↗](#); $\alpha_{d2} = 0.97 \pm 0.04$; $\beta_{d2} = 0.001 \pm 0.001$), which suggests that the RS gate was closed at the end of each ISI after transiently opening during the preceding doublet.

Overall, the responses to the first stimulations in doublets display less LFD than singlets, and the responses to the second stimulations in doublets display no LFD at all. The results are consistent with the notion that LFD involves a closing of the RS gate, which is transiently alleviated by doublet stimulations.

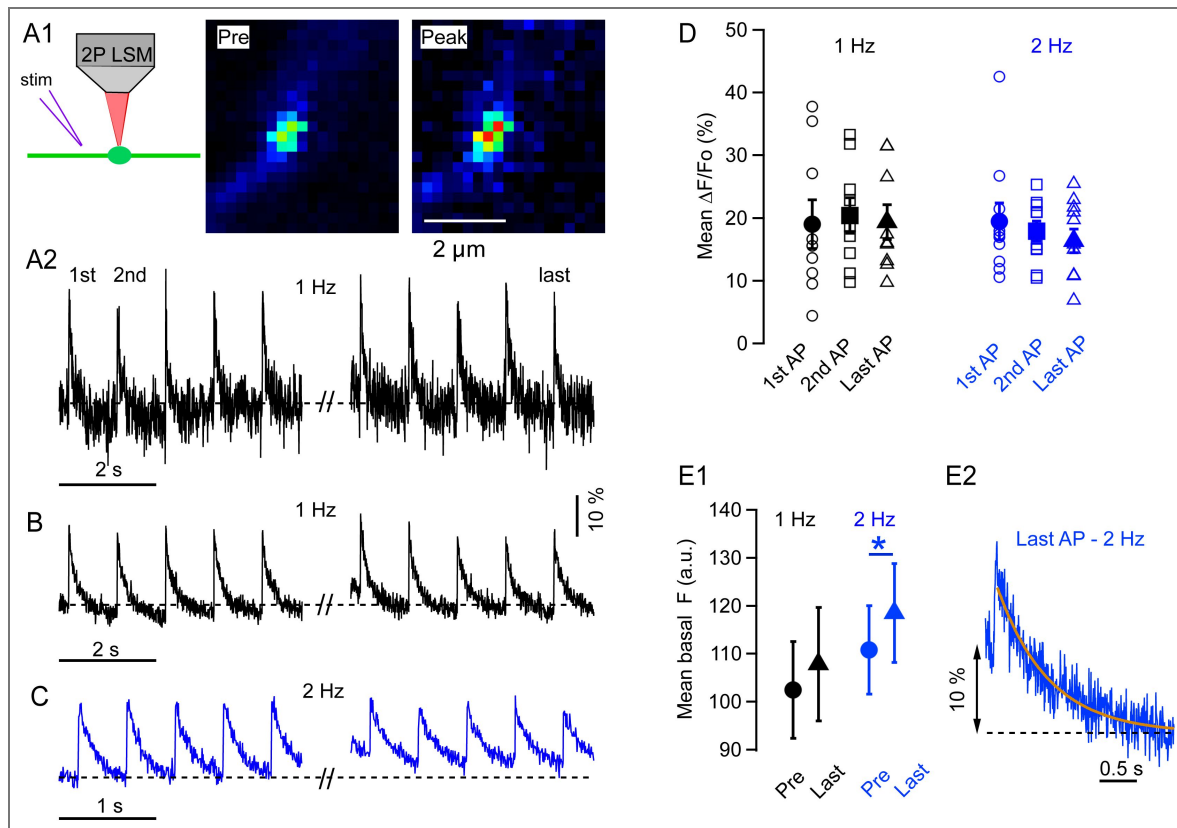


Figure 7. Presynaptic calcium transients in single PF varicosities

A1, left: Diagram of the experimental recording. **A1, center:** Image of an OGB-6F loaded PF varicosity before stimulation. **A1, right:** Same, at the peak of the response to a single presynaptic AP. **A2:** Average Ca^{2+} -dependent fluorescence transients ($\Delta F/F_0$ traces from $n = 5$ trials; see Methods) registered in response to the 5 first APs (left trace) and to the 5 last APs (right trace) in response to a train of 50 APs at 1 Hz, from the same varicosity as in A1. Imaging was interrupted between the two traces to avoid photodamage and photobleaching. **B:** Average traces from a group of 10 varicosities from 8 cells at 1 Hz. **C:** Average traces from another group of 9 varicosities from 6 cells, in response to the first and last 5 APs from a train of 200 APs at 2 Hz. **D:** Mean Ca^{2+} -dependent fluorescence transients (closed symbols: means across varicosities, with attached bars representing $\pm$ the sem; open symbols: values for individual varicosities) are the same for the 1st, 2nd, and last AP in a train (left: 1 Hz stimulation; black; right: 2 Hz stimulation; blue). **E1-E2:** Evolution of the basal fluorescence level during and after a long AP train. **E1:** Initial and final basal fluorescence levels are not statistically different for long trains at 1 Hz (left, black), while there is a significant increase at the end of trains at 2 Hz compared to its initial value (right, blue). **E2:** Return of the $\Delta F/F_0$ fluorescence trace to its initial baseline (mean from 5 varicosities) after the end of long 2 Hz trains indicates a shift of the basal fluorescence level by 11 % near the end of the train (yellow curve: exponential fit to the decay, with time constant 750 ms).

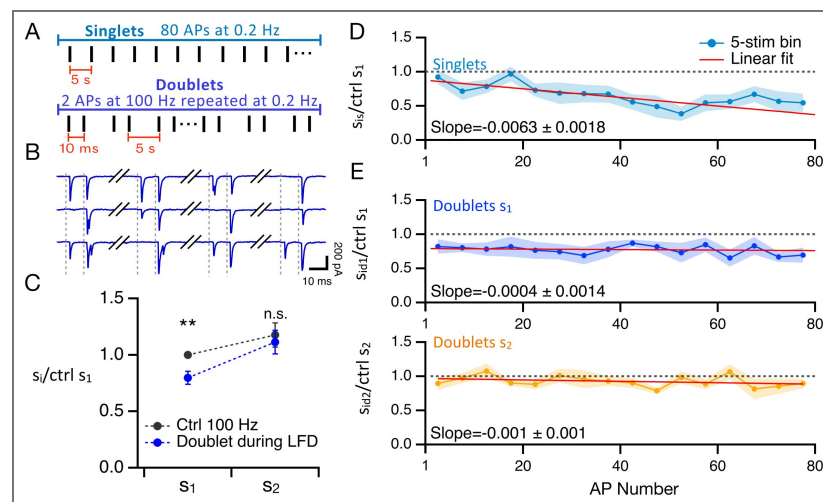


Figure 8. LFD during doublet vs. singlet train stimulations

A: Stimulation protocols. Either an 80-AP train of single stimulations ('singlets') or a sequence of 80 double stimulations 10 ms apart ('doublets') were applied at 0.2 Hz. Having 2 stimuli provides an evaluation of ρ and δ during LFD by comparing release during control and after AP# 1 or AP# 2 during doublets (s_{id1} and s_{id2} , respectively). **B:** Example traces of a doublet experiment. **C:** During the first 20 doublet stimulations, the ratio between s_{id1} and ctrl s_1 is < 1 , indicating a decrease in δ ; however, s_{id2} is not significantly different to ctrl s_2 indicating that the occupancy of the replacement sites (ρ) remains the same as control during LFD. **D:** Normalized plot of s_i during singlet trains (s_{is}) showing a 2-component depression (compare to Fig. 3C). The initial component is followed by a gradual steep depression during the entire train duration (mean $\pm$ sem of individual trials, normalized with respect to the mean s_1 value obtained during preliminary control trains; blue dots and associated error bars: binned data for 5 consecutive APs; red, linear fit to the data; 8 trials from 5 cells). **E:** Normalized plot of s_i during doublet trains (mean $\pm$ sem of individual trials, normalized with respect to the mean value obtained during preliminary control trains; dots and associated error bars: binned data for 5 consecutive APs; red, linear fit to the data; 9 trials from 6 cells). **Top:** SV release for the first AP of each doublet (s_{id1}) displays a rapid LFD component but no significant slow component. **Bottom:** SV release for the second AP of each doublet (s_{id2}), showing no significant deviation from the control value.

Discussion

In the present work, we use the SV counting method at simple PF-MLI synapses (Malagon et al., 2016) to investigate cellular mechanisms underlying LFD. Our results reveal similarities, but also significant differences, between these mechanisms and those previously described for moderate or high frequency synaptic depression (Lin et al., 2022; Tran et al., 2022). Based on our results and on our simulations using the RS/DS model, we propose that LFD reflects a decrease in δ following SV undocking. We also explain how the mechanisms of synaptic depression may vary as a function of stimulation frequency.

PPR as a function of ISI

A traditional view of synaptic physiology has been that synapses may be either facilitating or depressing. This view is clearly an oversimplification, however, as many synapses display a combination of facilitating and depressing properties (Dittman et al., 2000). Here we show that PF-MLI synapses exhibit PPR values > 1 for ISI values comprised between 10 and 200 ms, and PPR values < 1 for ISI values comprised between ~ 500 ms and ~ 1500 ms (Fig. 1). Importantly, the shift from facilitation to depression is apparent not only in group results (Fig. 1D), but also in individual experiments (Fig. 1C). A similar mix of facilitation at short ISI and depression at long ISI has been described in several central synapses (Chiu and Carter, 2024; Doussau et al., 2017). Following our analysis of simple synapse recordings, and in view of zapand-freeze results in hippocampal cultures, we propose that facilitation and LFD are linked together in a docking-undocking sequence following AP stimulation (Kusick et al., 2020).

SV movements during HFD and LFD

Based on the present study and on earlier work (Kusick et al., 2020; Miki et al., 2016; Tran et al., 2022), we propose a global model for HFD and LFD as outlined in Fig. 4D. If an AP train is presented with short ISIs (~ 10 ms), a rapid sequence of release/docking/replenishment occurs (open RS gate). The arrival of a new AP induces forward movement of SVs (from DS to exocytosis, from RS to DS, and from IP to RS). If a docked SV fails to be released, it does not undock, because a new SV rapidly occupies the corresponding RS. These events lead to facilitation at first. IP depletion eventually reduces SV influx into the RRP, such that the IP size becomes rate limiting at the end of the train. This accounts for HFD (HFD loop in Fig. 4D, green shade). If an AP train is presented with longer ISIs (0.5 s to 5 s), by contrast, release and replenishment are reduced since many DUs have their DS empty and their RS occupied at the time of the AP (closed RS gate). During the closed RS gate period, ρ is ~ 0.9 , so that the entry rate of SVs from the IP into the RRP, which is proportional to $(1-\rho)$, is reduced 10-fold by RS occupation. By comparison, ρ is ~ 0.5 when the RS gate is open. The larger ρ value results in a net 5-fold slower entry rate when the RS gate is closed compared to when it is open. With longer ISIs, SVs move from the RS to the DS after each AP, but they have the time to move back to the RS before the next AP. Therefore, DUs tend to undergo an idle docking/undocking loop without release, leading to LFD (LFD loop in Fig. 4D, blue shade).

AP induced docking and undocking happen in overlapping time frames. In Fig. 1 the highest amount of docking is seen at ~ 10 ms ISI (it is also present at 5 ms: (Miki et al., 2016; Silva et al., 2024)). Undocking on the other hand is maximal more than 1 s after the AP. For ISIs in between there is a mix of docking and undocking. According to this model, the PPR recovery observed at ISIs of ~ 200 ms is only apparent because in this period undocking is still ongoing.

When stimulation is switched to high frequency after LFD, SV release recovers within 10 ms to the same level as during high frequency control trains (Fig. 2D; Fig. 6C). According to the RS/DS model, this happens because the SVs that are in the RS dock with each AP and are still in the DS when a second AP is presented rapidly (HF recovery in Fig. 4D, yellow shade). This would explain why the s_2 of a recovery train is equivalent to the control value. If high frequency stimulation is continued the synapse will behave identically to a naïve synapse for the remainder of the train (Fig. 2D; Fig. 6C) and it will eventually undergo HFD (Fig. 6B; HFD loop in Fig. 4D, green shade).

It has long been recognized that synaptic activity enhances RRP replenishment (Neher and Sakaba, 2008). While a Ca^{2+} -dependent rate of SV entry into the RRP has been invoked as underlying mechanism, other explanations cannot be ruled out (Miki et al., 2020; Ritzau-Jost et al., 2018; Ritzau-Jost et al., 2014). Transient opening of the RS gate, as suggested here, offers a novel mechanism explaining activity-dependent RRP replenishment.

LFD may involve a specific δ decrease rather than across-the-board RRP decrease

Our conclusion that LFD results from a δ decrease is largely based in our analysis of high frequency recovery trains. While this approach is model-dependent, as it assumes the RS/DS model of Fig. 2A, our conclusion is supported by two additional lines of evidence. First, we find that LFD occurs even after failures, and that the extent of LFD does not depend significantly on s_1 (Fig. 5). These findings argue against the standard RRP depletion mechanism. By contrast, the docking-undocking sequence of Fig. 4D (main sequence and LFD loop) accounts for our results as it predicts that the DU state at the end of a long ISI depends little on the response to the previous stimulation (simulation in Fig. 5B). Secondly, we find that LFD is alleviated when using doublet rather than singlet stimulations (Fig. 8). Again, this result is inconsistent with an across-the-board RRP depletion model, but it can be readily explained by the RS/DS model. During LFD with singlet stimulations, the RS/DS model predicts a gradual δ decrease accompanied by a high ρ (Fig. 3). Doublet stimulations change this pattern in two ways. Firstly, they transiently lift the RS block that prevents RRP refilling with singlet stimulations, leading to enhanced RRP replenishment. The second AP of the doublet arrives in the period where SVs are still present in the DS, releasing a fraction of them, and promoting SV movement from the RS to the DS. This means that SVs coming from the IP are allowed, in turn, to bind the now-free RS. Secondly, doublet stimulations prevent SV undocking. This happens as the incoming SV at the RS locks the SV that is bound to the DS in its docked position. The combination of these two effects leads to an inhibition of the slow phase of LFD during doublet stimulations. Of note, this complex RS/DS interaction is only possible because the RS/DS model allows simultaneous occupation of both RS and DS by distinct SVs (Silva et al., 2024).

Altogether, the present LFD model resembles previous studies assuming two different SV states inside the RRP, and a decrease in the percentage of primed/docked SVs during synaptic depression (Doussau et al., 2017; Tanaka et al., 2021; Tran et al., 2022; Lin et al., 2025). It differs from these earlier studies in proposing a role of RS in gating SV movements during LFD.

While our model assumes that the δ decrease associated with LFD is primarily due to the return from DS to RS (undocking), we cannot exclude alternative mechanisms. SVs could leave the DS following exocytosis, or they could unbind and diffuse away in the cytosol. A previous study suggested that delayed release is insufficient to account to the δ decrease accompanying the end of facilitation (Miki et al., 2018), making the first option unlikely, but it remains possible that during LFD, part of the δ decrease is due to SV escape into the cytosol independently of the DS→RS transition.

Alternative presynaptic depression mechanisms

Simple synapse recording and deconvolution analysis give us a direct estimate of the number of SVs released, bypassing any possible postsynaptic effects. Expression of LFD as described in the present work is therefore presynaptic. Furthermore, our BAPTA experiments (Supplementary Fig. 6) argue against a participation of the postsynaptic compartment to LFD induction, so that both induction and expression of LFD are likely presynaptic. We consider here other presynaptic mechanisms than undocking that have been proposed to underlie LFD. A p_r reduction has been suggested (Wölfel et al., 2007; Wu and Borst, 1999), either resulting from a direct reduction in Ca^{2+} current (Xu and Wu, 2005) or from the buffering effect of calretinin (Bolshakov et al., 2019). The rapid reversal obtained during recovery trains in the present work is not consistent with either of these hypotheses, as both Ca^{2+} channel inhibition and a buffering effect would require hundreds of ms to seconds to recover. Furthermore, the results and simulations shown in

Fig. 6F suggest a change of δ but not of p_r during LFD. Finally, our Ca^{2+} measurements in presynaptic varicosities do not suggest a difference in Ca^{2+} entry following repetitive low-frequency stimulation although there is a change in basal calcium during long trains that could be linked to the second, slow phase of LFD (Fig. 7C). Of note, Ca^{2+} measurements have also been performed in relation to LFD in neocortical L4-L2/3 synapses with synaptic depression and no change in Ca^{2+} signal for paired stimuli was reported at ISIs of 500 ms (Chiu and Carter, 2024). In *Aplysia*, silencing of docking sites was suggested as the reason behind synaptic depression (Doussau et al., 2010), however a decrease in N would not predict the rapid return of high frequency release when recovery trains are applied after LFD.

In conclusion, it is unlikely that a p_r or N reduction underlies LFD. Overall, a δ reduction best explains LFD and its fast recovery.

Onset and offset kinetics of LFD

In some studies, LFD has been reported to develop almost immediately, after one or a few stimulations (Chiu and Carter, 2024; Rudolph et al., 2011) whereas in others it develops much more slowly as a function of stimulation number (Abrahamsson et al., 2007; Doussau et al., 2017). In the present work we observe an immediate reduction in synaptic output (Fig. 2C) followed by a very slow phase which does not reach a steady state after 200 APs at 2 Hz (Fig. 6A) and 80 APs at 0.2 Hz (Fig. 8D). After a short train, in which only the initial phase is observed, and after a long train, where both phases are present, we find a reduction in DS occupancy but no changes in RS or IP occupancy (Fig. 2E and 6D). This leads us to propose that LFD comprises two phases with different onset kinetics.

We find that LFD fails to recover significantly following a rest period of up to 1 minute (Supplementary Fig. 4). These results are consistent with earlier studies showing that for sparse stimulations, LFD recovery occurs on a time scale of minutes (Abrahamsson et al., 2007; Doussau et al., 2010; 2017). If, however, a high frequency train is presented, the synapse recovers from LFD almost instantaneously, within one single ISI (range 5-20 ms (Doussau et al., 2010; 2017); present work, Fig. 2C, 6C and 8C). As seen in Fig. 4D, the high frequency recovery loop after LFD is identical to the HFD loop: presenting new APs at short ISI does not leave enough time for undocking and immediately brings the synapse back to the HFD mode. Altogether, LFD places the synapse in a depressed state that is stable if stimulation is absent or infrequent, but that is immediately reversed if a high frequency train is presented. In our model, this dichotomy occurs because the SV-RS association is stable at rest, when presynaptic Ca^{2+} concentrations are low (RS gate closed) but is unstable during high frequency AP trains, when presynaptic Ca^{2+} concentrations are elevated (RS gate open). In contrast to LFD recovery, HFD recovers within seconds following the end of a high frequency train, reflecting the gradual replenishment of the IP (Tran et al., 2022).

Frequency-independent synaptic depression

In the present work, we observe similar extents of LFD at the end of the first phase for stimulation frequencies of 0.2 Hz (0.75 ± 0.11), 1 Hz (0.79 ± 0.08), and 2 Hz (0.69 ± 0.05). These results indicate that at PF-MLI synapses, the first phase of LFD results in a largely frequency-independent steady state synaptic output in the frequency range 0.2 Hz to 2 Hz. It has been likewise noted in earlier studies that the extent of synaptic depression at steady state is largely independent of stimulation frequency in a certain frequency range, albeit in a higher range than in the present work (10-100 Hz at Purkinje cell to deep cerebellar nuclei synapses (Turecek et al., 2016); 5-50 Hz at the calyx of Held (Lin et al., 2022)). These results have been interpreted either by a compensation between an increased facilitating component and an increased depressing component at higher frequencies (Turecek et al., 2017, 2016), or by a time limitation on calcium-dependent movement of SVs from a primed to pre-primed location, so that this movement is regulated by AP number rather than by time (Lin et al., 2025).

Molecular mechanisms of docking and undocking

The molecular mechanisms of docking and undocking are under intense scrutiny. Consistent with a role in AP-dependent docking, synaptotagmin 7 (syt7) is associated to facilitation in different preparations (Huson and Regehr, 2020 [↗](#); Jackman et al., 2016 [↗](#); MacDougall et al., 2018 [↗](#)). Using zap-and-freeze electron microscopy in hippocampal cultures revealed that syt7 KO showed no rapid docking ~10 ms after an AP, which was translated into a depressing phenotype when trains of APs are applied (Wu et al., 2024 [↗](#)). In neocortical synapses, a comparison of PPR curves in WT and syt7 KO showed no facilitation at short ISIs and a deeper depression at longer ISIs in the KO, with the same recovery time frame (5 s) (Chiu and Carter, 2024 [↗](#)), supporting a role of syt7 in AP-dependent docking, but not in undocking or baseline docking dynamics. Recently, intersectin-1 and endophilin A1 were reported to stabilize SVs in the replacement site (RS) (Ogunmowo et al., 2025 [↗](#)), indicating a role in docking/undocking and possibly in LFD. Munc-13 is another likely candidate for docking (Kusick et al., 2022 [↗](#)), as its deletion reduces the number of docked SVs and changes their relative distance to the plasma membrane (Imig et al., 2014 [↗](#); Siksou et al., 2009 [↗](#)). In the calyx of Held, RRP recovery after high frequency facilitation and depression depends on both Munc13 and synaptotagmin 3 (syt3; Weingarten et al., 2022 [↗](#)). Modeling suggested that both proteins are required to explain AP-dependent docking.

In conclusion, multiple proteins have been proposed as candidates for AP-dependent docking; in our preparation, Munc-13, syt3 and 7 are likely candidates. By contrast, the molecular mechanisms behind undocking remain largely unknown. It is likely that some of the STP heterogeneity arises from variations in protein expression and regulation amongst synapses, so that the docking/undocking kinetics are not the same. However, LFD is present in multiple preparations, and may be a general mechanism.

Functional implications

In vivo MLI recordings show very infrequent EPSCs in the absence of PF input (< 1 Hz), while sensory stimulation results in short EPSC trains at high frequency (Chadderton et al., 2004 [↗](#); Jörntell and Ekerot, 2003 [↗](#)). LFD as described here likely contributes to lower the frequency of background EPSCs, as it inhibits the responses to infrequent presynaptic APs. However, in view of the rapidity of the post-LFD recovery, full EPSC responses are likely restored as soon as a high frequency train of presynaptic APs is presented. This should increase the signal-to-noise ratio in a burst vs. background signaling mode and should improve the sensitivity of the cerebellar circuitry to external inputs. As many brain neurons display low background firing rates (< 1 Hz) *in vivo* (Buzsáki and Mizuseki, 2014 [↗](#); Margrie et al., 2002 [↗](#)), a similar signal-to-noise improvement could occur in various brain regions following LFD.

Materials and Methods Experimental model

C57BL/6 mice (12–16 days old; either sex) were used for preparation of acute brain slices. Animals were purchased from Janvier Laboratories (RRID: MGI:2670020 [↗](#)). They were housed and cared for in accordance with guidelines of Université Paris Cité (approval no. D 75-06-07; animal rearing service of the BioMedTech Facilities, INSERM US36, CNRS UMS2009).

Slice preparation

200-μm thick sagittal slices were prepared from the cerebellar vermis as follows. Mice were decapitated under anesthesia and the cerebellar vermis was carefully removed. Dissections were performed in ice-cold artificial cerebrospinal fluid (ACSF) which contained the following (in mM): 130 NaCl, 2.5 KCl, 26 NaHCO₃, 1.3 NaH₂PO₄, 10 glucose, 1.5 CaCl₂, and 1 MgCl₂. Slices were cut using a vibratome (VT1200S; Leica) and incubated in ACSF at 34°C for at least 60 min before being used for experiments.

Synaptic electrophysiology

Whole-cell patch-clamp recordings were obtained from MLIs (comprising both basket and stellate cells). The extracellular solution contained (in mM): 130 NaCl, 2.5 KCl, 26 NaHCO₃, 1.3 NaH₂PO₄, 10 glucose, 3 CaCl₂ (1.5 in the experiments where it is stated), and 1 MgCl₂ (pH set to 7.4 with 95% O₂ and 5% CO₂). The internal recording solution contained (in mM): 144 K-gluconate, 6 KCl, 4.6 MgCl₂, 1 EGTA (ethylene glycol- bis tetraacetic acid), 0.1 CaCl₂, 10 HEPES (4- (2- hydroxyethyl)- 1- piperazineethanesulfonic acid), 4 ATP- Na, 0.4 GTP- Na (pH 7.3, 300 mosm/l). Gabazine (3 μM) was included in all experiments to block GABA_A receptors. Recording temperature was 32–34°C. To establish a single PF–MLI connection, puffs using a glass pipette filled with the high K⁺ internal solution were applied in the granule cell layer to identify a potential presynaptic granule cell. The same pipette was then used for extracellular electrical stimulation of the granule cell, with minimal stimulation voltage (stimulus voltage was higher than threshold by ~5 V; range: 10–40 V; 150 μs duration). If the resulting EPSCs, particularly those that occurred after a short stimulation train, were homogeneous, then it was likely that only one presynaptic cell was stimulated. Recordings were only accepted as a simple synapse recording after analysis if the following three criteria were satisfied (Malagon et al., 2016 [DOI](#)): (1) the EPSC amplitude of the second release event in a pair was smaller than that of the first, reflecting activation of a common set of receptors belonging to one postsynaptic density; (2) all the EPSC amplitudes followed a Gaussian distribution with a coefficient of variation < 0.5; and (3) the number of release events during the baseline recording was stable.

Quantification and statistical analysis

Offline analysis of electrophysiological data and statistical analysis was performed using Igor Pro (WaveMetrics). Data are expressed as mean ± standard error of the mean (SEM). Error bars in all graphs indicate SEM. Details of statistical tests are described in the text or figure legends. Student's t-test was used when the distribution was Gaussian; Wilcoxon Rank test was used otherwise. One-tailed tests were used when the sign of the effect was predicted before performing the experiments; two-tailed tests were used otherwise. Statistical significance was accepted when $p < 0.05$. The number of trials and cells (some cells have multiple trials) for each experiment is indicated by a lowercase n. Cells were derived from different animals. Each experiment started with control recordings of 8-AP trains @ 100 Hz, with 10 s inter-train intervals. At least 25 trials were performed for this initial control run before proceeding to specific stimulation protocols. For PPR experiments (Fig. 1 [DOI](#)) 4-AP trains were applied with varying ISIs (20, 100, 200, 400, 800, 1600 and 3200 ms) in a scrambled order, with a rest period of 10 s between trains (Fig. 1A, right [DOI](#)); only synapses with minimum 20 repetitions of this protocol were considered for analysis. For Fig. 2 [DOI](#), 8-AP trains at alternating frequencies (100 Hz and 2 Hz) were applied. A 100 Hz recovery train was applied immediately after the end of each 2 Hz train (see Fig. 2B [DOI](#)); a minimum of 20 repetitions of the protocol were obtained for each synapse. Since the value of s_1 should be independent of train frequency (the synapse cannot predict the frequency at which the train will be given), values of s_1 were averaged across frequencies in each synapse. Therefore in Fig. 1D [DOI](#), the s_1 SV # used for calculation of the PPR was averaged across 8 frequencies. In Fig. 2 [DOI](#), the s_1 value used for normalization is the average of s_1 values for the control 100 Hz train and for the 2 Hz train at each train cycle. For these two protocols the repetitions of each synapse were averaged together, and synapse results were averaged for group results. For Fig. 6 [DOI](#) a 200-AP train was applied at low-frequency (2 Hz), followed by a 50-AP or 100-AP high-frequency (100 Hz) recovery train (Fig. 6A [DOI](#)). The time interval between the last AP in the LFD train and the 1st AP in the awakening train varied in the experiments (Supplementary Fig. 4 [DOI](#)). Synapses were considered with minimum one full trial (low frequency + recovery), the maximum number of trials performed in one synapse was 4. Trials were averaged for group results. For Fig. 8 [DOI](#), 80AP trains were applied at 0.2 Hz with either a single or a double stimulus (Fig. 8A [DOI](#)), trials were averaged together for group results. In 5 cells singlets and doublets were applied sequentially. For all protocols a new

control run was performed when possible at the end of the experiment to verify functional stability of the synapse. For each set of experiments the numbers of repetitions are stated in the figure legend.

Decomposition of EPSCs

The time of occurrence and the amplitude of individual release events were determined based on deconvolution analysis, as described previously (Malagon et al., 2016). In brief, for each synapse, an mEPSC (miniature excitatory postsynaptic current) template was obtained from delayed release events recorded after the end of AP trains. This template was fitted with a triple-exponential function with five free parameters (rise time, peak amplitude, fast decay time constant, slow decay time constant, and amplitude fraction of slow decay). These five parameters were then used for deconvolution of the template and of individual data traces, producing a narrow spike (called spike template) and sequences of spikes, respectively. Next, each deconvolved trace was fitted with a sum of scaled versions of the spike template, yielding the timing and amplitude of each release event. The amplitude was further corrected for receptor saturation and desensitization, using the exponential relationship between individual amplitudes and the time interval since the preceding release events. Events that were at least 1.7 times larger than the average mEPSC were split into two (Miki et al., 2017). s_1 was determined as the number of SVs released within 5 ms after the first AP of a train. s_2 was the number of SVs released within 5 ms after the second AP and s_{5-8} was the cumulative number of SVs released 5ms after each stimulus for AP# 5 to 8. As explained in Tran et al. (2022) (Tran et al., 2022), these 3 parameters are proxies respectively for the occupancy of DS (δ), the occupancy of RS (ρ), and the IP size.

Calcium measurements

To perform Ca^{2+} imaging of single PF varicosities we pre-loaded granule cells with a calcium indicator. A whole-cell recording was made with a K^+ gluconate-based solution containing 166 K gluconate, 4.1 MgCl_2 , 9.9 HEPES-K, 0.36 Na-GTP and 3.6 Na-ATP, 1-2 mM of the calcium probe Oregon Green BAPTA 6F. The neurons were held in I-clamp mode with currents of -5 to -30 pA injected to keep the average resting potential close to -90 mV for 1.5 to 2 minutes after which the pipette was removed. Obtaining an outside-out patch with > 5 GW resistance indicated successful pipette removal, and the granule cell was left for approximately 15 minutes to allow for dye diffusion into the axonal compartment. After this waiting time, a 2photon laser scanning microscope (Tan et al., 1999) was used to identify an axonal varicosity. A glass pipette with tip diameter around 1 μm , filled with extracellular saline (in mM: 145 NaCl, 2.5 KCl, 10 K-HEPES, 2 CaCl_2 , 1 MgCl_2), was introduced into the slice at the depth of the varicosity and placed 10-20 μm from the axon. The stimulation protocol consisted of 1 train of 8 pulses (100 μs duration, 10 to 40 V) delivered at 100 Hz, followed by either 50 stimulations at 1 Hz or 200 at 2 Hz and a new 8 pulse train at 100 Hz applied 30 seconds after the end of the long train. This cycle was performed 3 to 7 times. Fluorescence was acquired at dwell time of 5 ms/frame through raster scans of 5 by 5 μm fields encompassing the axonal varicosity. For each long stimulation train that was applied, we recorded two fluorescence traces. One trace covered the first 5 APs of the train (trace duration: 5 s at 1 Hz stimulation frequency, and 2.5 s at 2 Hz stimulation frequency). We next interrupted laser illumination and scanning (during 40 s at 1 Hz, and during 95 s at 2 Hz) to minimize photobleaching and photodamage. We resumed two-photon imaging near the end of the train, collecting another trace covering the last 5 APs. Time dependent fluorescence signals were analyzed in the pixels encompassing the varicosity and quantified in terms of changes relative to pre-stimulus values ($\Delta F/F_0$ expressed in %) with software written in the IGOR-Pro programming environment (Wavemetric, Lake Oswego, OR, USA). To avoid possible errors due to tissue movement, focusing was performed twice: just before the first trace, and again before the second trace. To estimate changes in baseline Ca^{2+} concentration, we compared basal fluorescence averages (calculated just before the 1st AP in each trace) for early and late traces across trials for all experiments. This revealed an apparent (non-significant) increase by 4 % of the basal fluorescence at 1 Hz stimulation frequency, and a significant 7 % increase at 2 Hz stimulation frequency (Fig. 7E1). In view of this baseline shift (F_{Inc}), the second $\Delta F/F_0$ trace needed a

correction. In this trace, the basal fluorescence used to calculate DF/F_0 was corrected according to $F_{0corr} = F_{0app} - F_{Inc}$, leading to the relation $DF/F_{0corr} = DF/F_{0app} * (1 + F_{Inc}/F_0)$ where DF/F_{0app} and DF/F_{0corr} are the apparent and corrected values of DF/F_0 for the late trace, and the ratio F_{Inc}/F_0 is 4 % at 1 Hz and 7 % at 2 Hz. For each train of stimuli, $\Delta F/F_0$ values were averaged over 5 to 7 repetitions at each varicosity and subsequently data from all varicosities were pooled together to generate the average signal as a function of time from which mean peak values were extracted.

Model and simulation of SV docking and release

Monte Carlo simulation of two-step SV docking and release (Fig. 3B) was performed using Igor Pro as described previously (Miki et al., 2016; Silva et al., 2024; Tran et al., 2022). The model is created by assuming a fixed number (N) of docking units (DUs) with 2 binding sites for each DU (RS and DS; Fig. 2A), each of which can be empty or occupied by an SV. The probabilities of initial occupancy for DS and RS are δ and ρ , respectively. The probability of SV release after one AP of an occupied DS is p_r . SV transitions within the DU are characterized by transition probabilities s_f , s_b , r_f , r_b .

For each series of experiments, control runs (usually 30 trials) were initially performed with stimulations of 8 APs at 100 Hz, interspersed by 10 s-long rest periods. We used previously developed procedures to determine from these data the parameters N, δ (the mean DS occupancy before the 1st AP), ρ (the mean RS occupancy before the 1st AP), as well as the probabilities s_f , r_f and p_r , all of which were assumed constant throughout the AP train. In this part of the simulation, the back transitions s_b and r_b were neglected. The simulation yielded the numbers of released SVs (s_i) as a function of AP#. These numbers were calculated as $s_i = N \delta_i p_r$ where δ_i is the DS occupancy just before AP# i (Scheuss and Neher, 2001). Of note, there were some variations from one set of experiments to another regarding these parameters, reflecting differences between synapses (Table 1). Similar observations have been made at calyx of Held synapses (Lin et al., 2025).

DS and RS replenishment are not independent of each other. A DS can only be filled if an SV is present in the RS, and a downward transition from the RS can only take place if the DS is empty. Likewise, a transition from DS to RS will only happen if the DS is occupied and the RS empty. These considerations are taken into account in the Monte Carlo model. To better understand how the DUs as a whole changed after an AP and during trains, we considered four DU ‘states’ and we followed them throughout the simulations: DUs were “full” when both RS and DS were occupied; “up” when RS was occupied while DS was empty; “down” when RS was empty while DS was occupied; “empty” when both RS and DS were empty (Supplementary Fig. 3B). The initial δ and ρ values were used to estimate the initial DU states, keeping in mind the following relations: 1) $\delta = \%down + \%full$, 2) $\rho = \%up + \%full$, and 3) $\%down + \%up + \%full + \%empty = 1$. The DU state was tracked alongside δ and ρ for each simulated AP.

The first AP was then simulated by releasing from the DS that were occupied (DU states full and down). Release resulted in an increase of % up and a simultaneous fall of % full (Supplementary Fig. 3B, grey shade).

Below 10 ms, rate constants are largely determined by the large amplitude, fast decaying Ca^{2+} signal occurring near voltage-dependent Ca^{2+} channels (‘ Ca^{2+} nanodomain’). The first 10 ms after an AP were simulated with short time intervals (50 μ s) while following ρ , δ and the DU states. This timeframe only takes into account forward transitions (r_f and s_f), which were determined from fitting the 100 Hz control data of each dataset, as described before (Miki et al., 2016; Silva et al., 2024). This simulation was then used to establish the overall DU changes in the first 10 ms following an AP. For example, EE was defined as the probability that a DU that was empty just after release would be again empty 10 ms after the AP; EU the probability that a DU that was empty just after release would be up 10 ms after the AP, and so forth (Table 1). These probabilities were used to simulate the initial 10 ms of SV replenishment during an ISI for low frequency train simulations. The first 10 ms after release are dominated by the calcium-dependent movement of SVs from the RS to the DS (r_f ‘transient docking’) which results in an increase in % down and a decrease in % up (see Supplementary Fig. 3B).

After 10 ms, the rate constants depend on the low amplitude, slowly decaying Ca^{2+} signals averaged over the entire varicosity ('volume-averaged Ca^{2+} '). To calculate DU state changes at times > 10 ms after the AP, time intervals for the calculations were changed from 50 μs to 10 ms. This change of time interval was justified because SV movements slow down markedly as a function of time since the AP. It resulted in a marked reduction in calculation time. The transition between the two time domains occurring at 10 ms explains the slight differences between data and simulation points at 10 vs 20 ms in Fig. 3B. Whereas transition probabilities were assumed to remain constant during the first 10 ms, they were assumed to change for each subsequent 10 ms time interval as specified in Supplementary Figure 3A. In this part of the simulation all transitions were considered, including the back transitions s_b and r_b . Note that the rate r_b has a delay before the start of the time-dependent exponential curve and that s_f remains constant throughout the ISI. Between 10 and 100 ms after the AP, up and down state proportions were stable, with a low % empty and up DUs and a high % down and full DUs (Supplementary Figure 3B; note time scale change at 10 ms). Next, SVs moved from DS to RS (undocking, with rate r_b), resulting in a gradual increase in % up and a decrease in % down (Fig. 3B). As described in the main text, this movement causes a temporary block of replenishment as ρ is high (RS gate closed). DUs gradually returned to their basal state on a time scale on the order of 10 s.

To model DU states during AP trains at low frequency, high time resolution modeling at times 0-10 ms after each AP (time domain 1) was followed by low time resolution modeling starting at 10 ms after the AP and ending at the end of each ISI (time domain 2). Models for consecutive APs were concatenated. For each binding site, the simulation started by attributing a site status to DS and RS. This was done at the beginning of each simulation trial by obtaining a pseudo-random value from the binomial distribution with a mean probability equal to the starting value of δ or ρ , for DS and RS respectively. DU states, δ and ρ were then followed step by step during the entire AP train.

Supplementary Fig. 3C depicts the changes in DU state during the simulations of high and low frequency 8-AP trains (Fig. 2 and Fig. 4). The proportion of DUs in each of the full/up/down/empty states is shown alongside ρ and δ just before each AP. During high frequency trains (**left**) an increase in % down (purple continuous curve) at the expense of a decrease in % up (green continuous curve) and % full (orange continuous curve) drives the rise in δ . During low frequency trains (**center**) the trend is the opposite, with undocking causing an increase in the % up (green), mirrored by a reduction in δ . This trend is mainly driven by a drop of % full (orange). DU changes during recovery trains (**right**) reproduce those obtained during control (**left**).

The kinetic parameters were adjusted by hand to find a good fit and to achieve a gradual return to the basal state for long ISIs. The simulation parameter values for the second domain were optimized according to the average release observed in the LFD trains as well as to the awakening train with the final δ and ρ values of the LFD train.

The steady state values of the transition rates in Fig. 3A are heavily constrained by the data within the RS/DS model. By contrast, the time dependent components of r_f , r_b and s_b , reflecting Ca^{2+} - and time-dependent SV movements, are less strictly constrained, due to the limited set of experimental data on the dependence of LFD on ISI.

The states and occupancies were stored for each simulation trial and averages were obtained at the end of the simulation. For the simulations shown, calculations were done 10,000 times for each ISI and resulting averaged values are displayed.

Our LFD simulations reached steady state after a few APs. They are a good model for the first phase of LFD, but they do not account for the second phase of LFD. Modelling the slow phase of LFD would presumably involve changes in rate constants of the model as a function of AP number; such models fall outside the scope of the present work.

Failure analysis

For this analysis we focused on the first component of LFD and more specifically, on the value of s_2 . Results were taken from experiments with ISIs of 500 ms (Fig. 2), 800 ms and 1.6 s (Fig. 1). The trains were sorted according to whether the 1st AP failed or was successful (Fig. 5A; failures

in purple, successes in black). The ratio of the released SV number after AP# 2 (s_2) and the average of the total released SV number after AP# 1 ($\langle s_1 \rangle$) was calculated for the s_1 failures and for the s_1 successes separately. One synapse had no failures; it was excluded from this analysis. Both traces with successes and failures show depression, as evidenced by a $s_2/\langle s_1 \rangle$ (PPR) lower than 1, indicating that loss of SVs is not the main mechanism of LFD (Fig. 5B, left [link](#)).

This analysis was extended by sorting the trains according to the number of SVs released for s_1 . Though the data shows a trend of deeper depression with increased release during the 1st AP, the difference is not significant (Fig. 5B, right [link](#)), indicating again that while release may play a role it is not the main mechanism. Next, we evaluated if the simulations replicated this characteristic of the data. The simulation trials were sorted the same way as the data and the PPRs calculated for each case (Fig. 5B [link](#); see black dots). The simulation also shows a significant depression for successful trials, with a somewhat stronger correlation between the amount of depression and the initial s_1 value than experimentally observed. Overall, the simulation reproduces well the results of the experiments.

Binomial analysis and predictions

According to the binomial distribution, the probability that s_i , the number of SVs released after AP number i , equals k is $p(k) = P_i^k (1 - P_i)^{N-k} N!/(k!(N - k)!)$, where P_i is the release probability per docking site ($P_i = \delta_i p_r$) and N is the number of docking sites (Malagon et al., 2016 [link](#)). The values of N and P were obtained by minimizing the summed squared deviations between the binomial prediction and data. This fitting was carried out for the control s_1 data of the synapses in Fig 7 [link](#), giving the distribution $p_{\text{ctrl}}(k)$. LFD was evaluated in a similar manner, considering each of the stimuli in the train as a separate event. The distributions of released SV numbers for 2 examples of control s_1 data is shown in black lines in Supplementary Fig. 5A [link](#) and red lines show the LFD data distributions, $p_{\text{LFD}}(k)$.

We tested the case of the depression in LFD being the result of an increase in stimulus failures, with a constant probability of release. In this case the probability of failures during LFD, $p_{\text{LFD}}(0)$, would be larger than the control probability of failures, $p_{\text{ctrl}}(0)$, but the rest of the distribution would retain the same shape for the LFD and control distributions: $p_{\text{LFD}}(k) / p_{\text{ctrl}}(k) = \text{constant}$ for $k = 1$ to N .

Considering that for each distribution, the sum of $p(k)$ from 0 to N is 1, we obtain: $p_{\text{LFD}}(k) = p_{\text{ctrl}}(k) \times [1 - (p_{\text{LFD}}(0) - p_{\text{ctrl}}(0)) / (1 - p_{\text{ctrl}}(0))]$ for $k = 1$ to N .

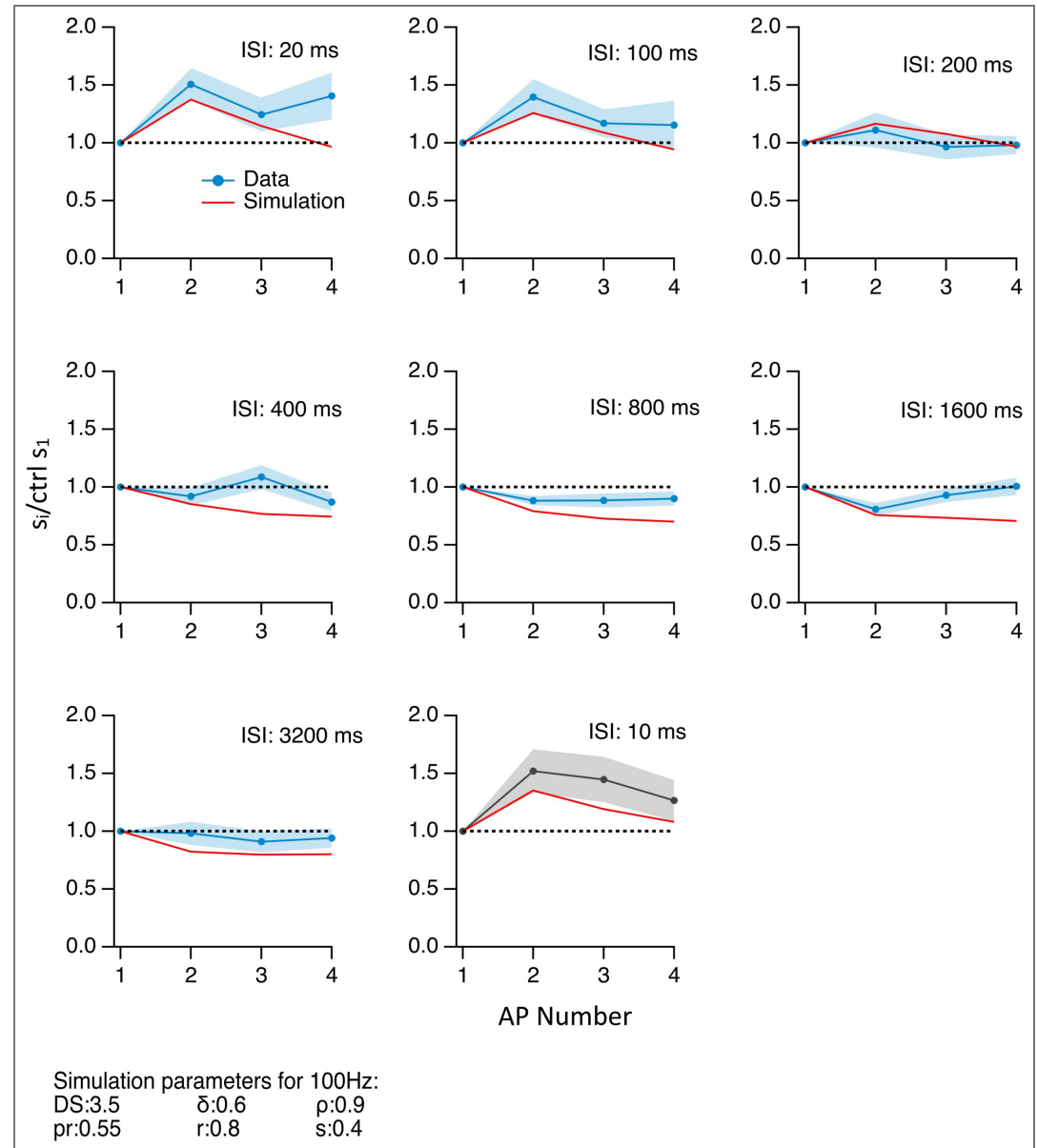
The dotted blue line in the examples of Supplementary Fig. 5A [link](#) show the result of this prediction.

If, instead, LFD was the result of a decrease in probability of release then the $p(k)$ of the LFD data would follow the distribution: $p_{\text{LFD}}(k) = (P_{\text{LFD}})^k \times (1 - P_{\text{LFD}})^{N-k} N!/(k!(N - k)!)$

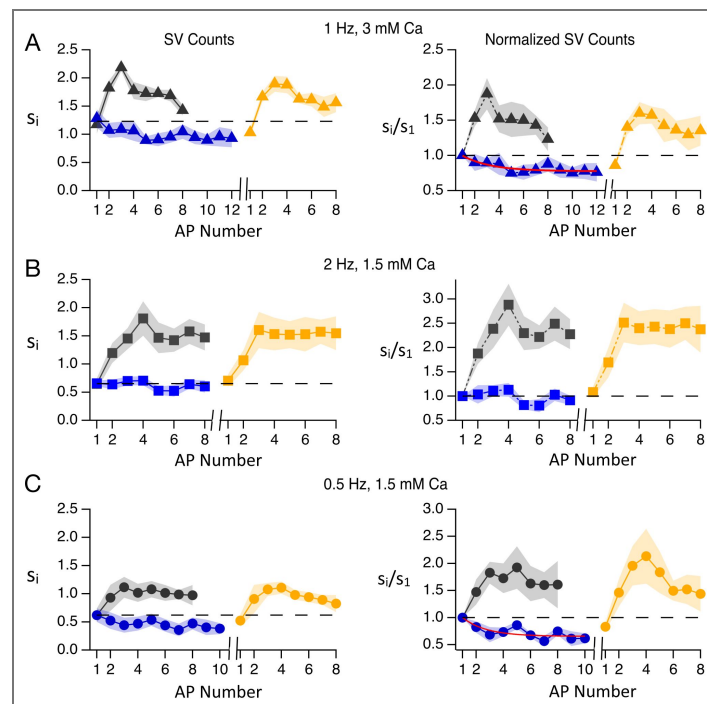
where N is the number of DS (assumed to be 5 in the example shown) and P_{LFD} is the P calculated from the LFD distribution: $P_{\text{LFD}} = 1 - p_{\text{LFD}}(0)^{1/N}$. The dotted purple lines in the examples of Supplementary Fig. 5A [link](#) show the result of this prediction. The LFD data (red line) was then compared to both predictions. In both examples the decreased P prediction (purple) is much closer to the data than the prediction for increased stimulus failure (blue). To quantify this difference residuals were calculated for each point in the plot for both predictions. The group data for the residuals of each prediction is shown in Supplementary Fig. 5B, left [link](#). The right panel depicts the average of the residuals. The prediction of a decrease in P is much closer to the data, compared to the prediction for an increase in the number of stimulus failures.

This analysis shows that an increase in stimulus failures does not account for the distribution of events in the LFD trials and that a decrease in P explains this phenomenon better.

Supplementary Material

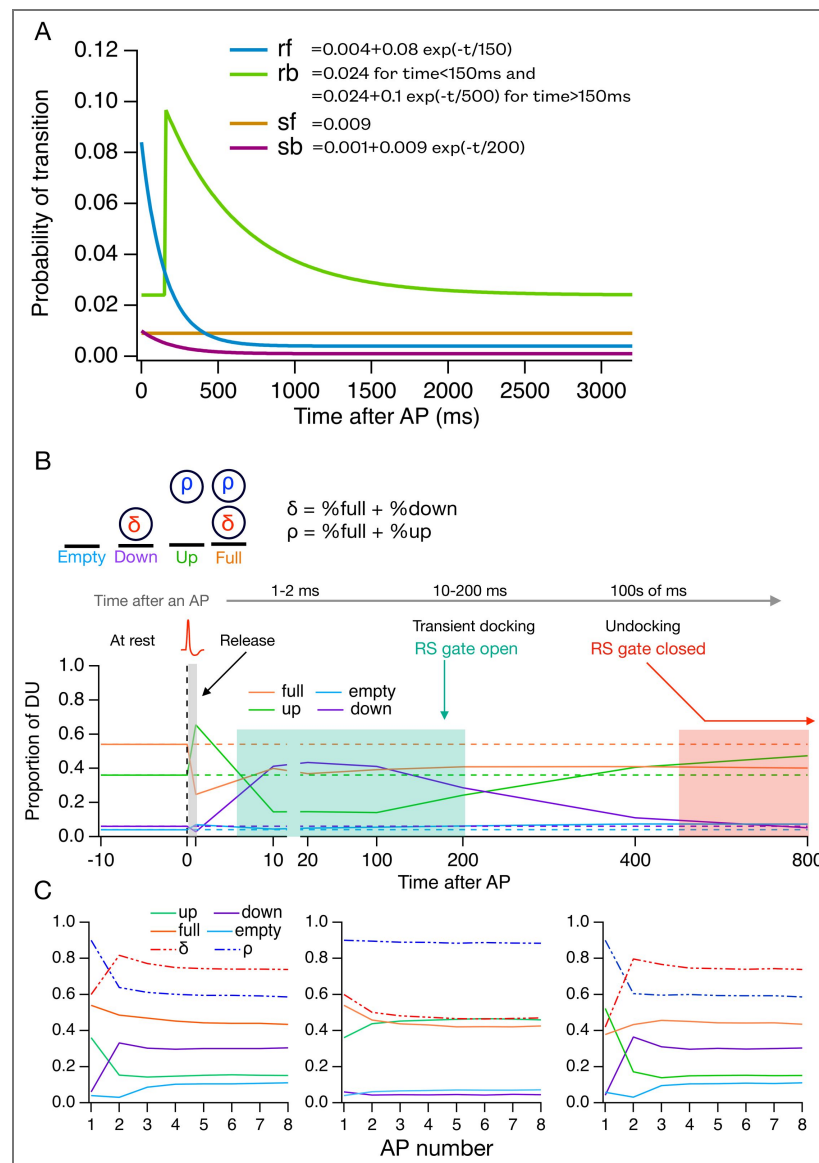


Supplementary Figure 1. Trains of 4 successive EPSCs with different ISIs Each panel shows numbers of released SVs in response to 4 consecutive APs at different ISI values. These numbers are normalized with respect to the mean s_1 value, calculated across all ISI values (dotted lines). Data (dots: mean values; shaded areas: $\pm$ sem) are in blue for ISI values of 20 to 3200 ms, and in black for control 8-AP trains with an ISI of 10 ms. Red curves show simulations based on the model of Fig. 3.



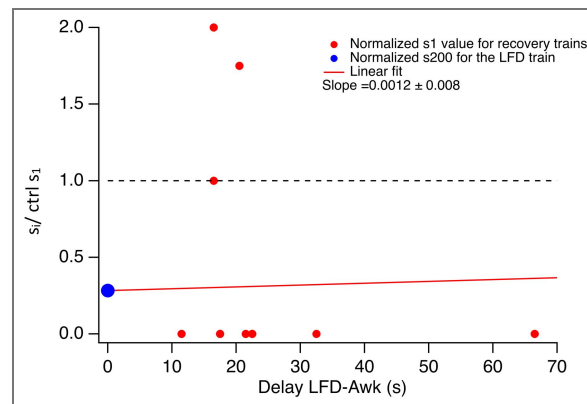
Supplementary Figure 2. LFD at various ISI and external calcium concentration values

Same experiments as in Fig. 2 at varying stimulation frequencies during low frequency trains (**A**: 1 Hz, $n = 5$; **B**: 2 Hz, $n = 6$; **C**: 0.5 Hz, $n = 6$) and varying external calcium concentrations (**A**: 3 mM; **B** and **C**: 1.5 mM). Red curves show exponential fits (**A**: asymptote: 0.79 ± 0.08 ; $\tau = 1.92 \pm 0.55$ ISIs, or 1.92 s; **C**: asymptote: 0.69; $\tau = 1.8$ ISIs, or 3.6 s).



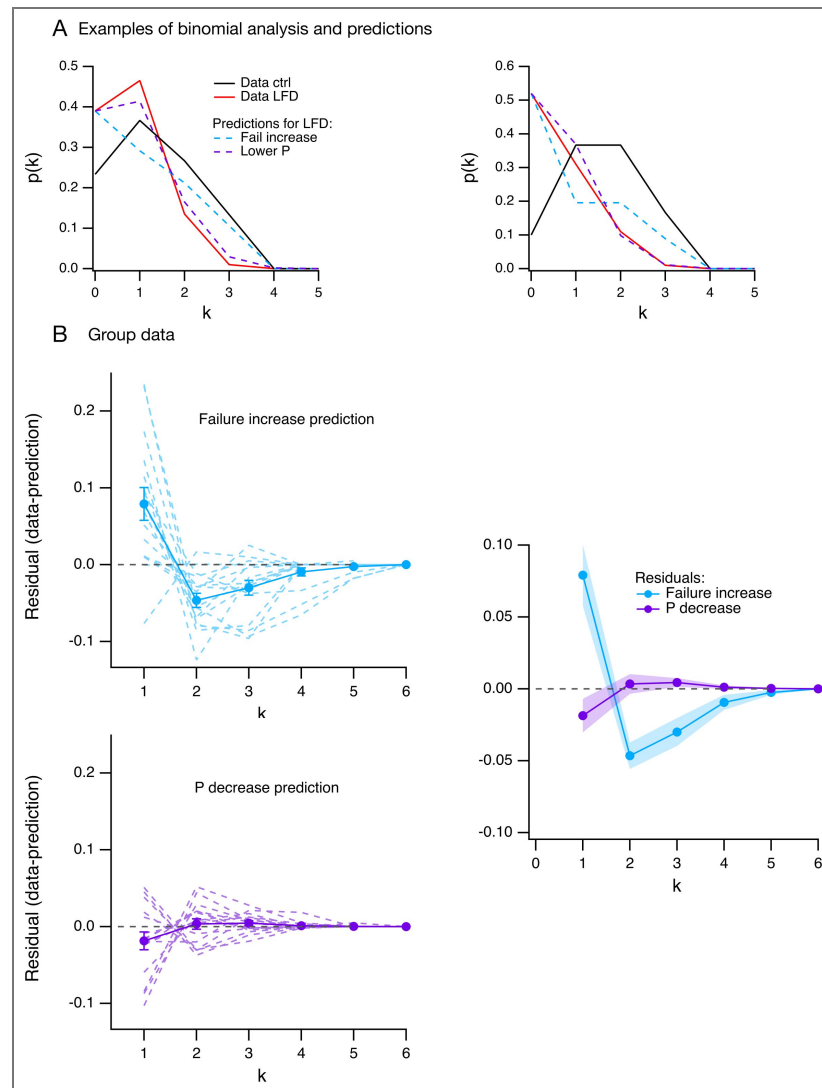
Supplementary Figure 3. Time dependent transition rates during LFD

A: Time dependent transition rates (calculated per 10 ms time bins) are shown for the 4 SV transitions of the model in Fig. 2A. Note that the first 10 ms period is excluded from this plot (see the corresponding transition probabilities in Table 1). s_f is stable as a function of time, while r_f , r_b and s_b follow exponential functions of time (t , in ms). In the case of r_b , the exponential starts with a delay of 150 ms. **B:** Following an AP, SV movements were described using 4 different DU states: “full” when both RS and DS were occupied; “up” when RS was occupied while DS was empty; “down” when RS was empty while DS was occupied; “empty” when both RS and DS were empty. Release resulted in an increase of % up and a simultaneous fall of % full (gray shade). The next 10 ms following stimulation were dominated by the calcium-dependent movement of SVs from the RS to the DS (r_f , ‘transient docking’) resulting in an increase in % down and a decrease in % up. In subsequent 10 ms time segments, SV movements were calculated based on the time-dependent changes in r_f , r_b , s_f and s_b shown in A. Between 10 and 100 ms after the AP, up and down state proportions were stable, with a low % empty and up DUs and a high % down and full DUs (green shade, RS gate open). Next, SVs moved from DS to RS (undocking, with rate r_b), resulting in a gradual increase in % up and a decrease in % down (red shade, RS gate closed). **C:** DU states before individual APs for 8-AP trains. **Left:** Control. **Center:** LFD 2 Hz. **Right:** Recovery.



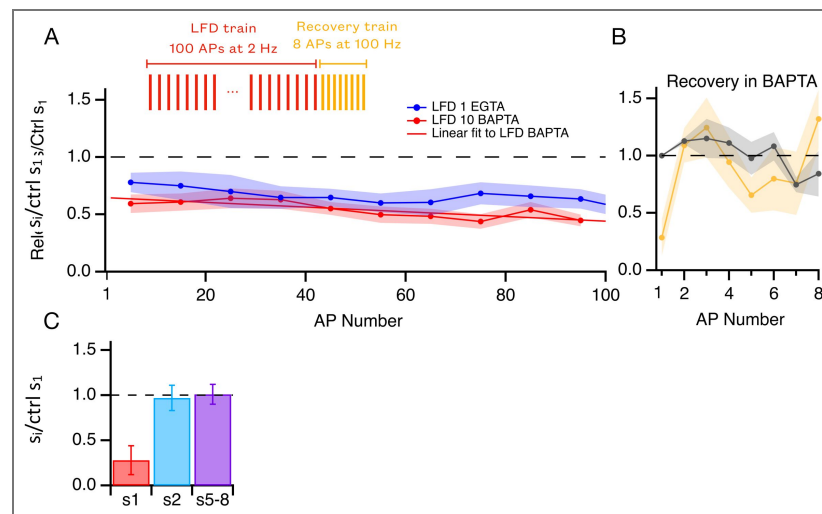
Supplementary Figure 4. Lack of recovery of LFD over tens of s after AP train

The time interval between the last AP in the LFD train and the 1st AP in the awakening train varied in the experiments presented in Fig. 7. These differences were used to analyze the time it takes for synapses to recover after a long low-frequency train. This figure shows the normalized s_1 value for recovery trains following long LFD trains (200 APs @ 2 Hz), as a function of the time duration intervening between the end of the LFD train and the onset of the recovery train. Failures were observed in a majority of the trials. The 0-delay point (blue dot) corresponds to the mean s_{200} value at the end of the LFD train. The red line is a linear fit of the data that is constrained to pass through the 0-delay point. The data indicate no recovery for up to 60 s.



Supplementary Figure 5. Binomial analysis of released SV numbers during prolonged LFD

A: Exemplar experiments. In each case, the distribution of released SV numbers (k) in response to the 1st AP during control runs (8 APs @ 100 Hz; black curves) is shown together with the distribution of released SV numbers during LFD (red curves; obtained from two of the experiments shown in Fig. 7). Next, we calculated the binomial distributions in two different models of reduced responsiveness. Both models assumed unchanged N values. In one model (blue dotted curves), the reduced response was assumed to reflect global failure at the AZ level for some stimulations, while other stimulations were assumed to be as effective as in the control. This model mimics situations such as unwanted stimulation failures during LFD. In the other model, a homogeneous reduction of P ($= \delta * p_i$; the release probability per DU) was assumed (purple dotted curves). Both models were constrained to account for the failure rate observed during LFD. In both experiments, the lower P curve is much closer to the experimental results (red curves for $k = 1$ to 3) than the failure increase curve. **B:** Group data analysis ($n = 16$ trials from 8 cells). **Left:** Residuals (dotted lines: individual LFD trials; dots and associated SEM, means across trials) calculated from the comparison of the two models with LFD data (**top:** increased failure model; **bottom:** decreased P model). **Right:** Superimposition of mean residual curves for the two models, showing that the lower P model is a much better representation of the data than the failure model.



Supplementary Figure 6. Prolonged low frequency trains with postsynaptic BAPTA

BAPTA (10 mM) was added to the internal solution used for MLI recording, keeping all other experimental conditions as before. **A:** A long AP train at low frequency (100 APs @ 2 Hz) was followed by a short 8-AP recovery train (experimental protocol in **insert**). Normalized plot of s_i during long low frequency train (red curve showing binned data for 10 consecutive APs), showing a two-component depression with an initial rapid depression ($65 \pm 3\%$ of the control) followed by a gradual depression during the entire train duration (slope = $-0.2 \pm 0.05\%$ per AP; red: linear fit to the data; $n = 12$ trials, 3 cells). The results resemble those with 1 mM EGTA shown in Fig. 6 (reproduced here as a dark blue curve). **B:** A short (8-AP) 100 Hz recovery train was applied immediately at the end of the low-frequency train. Normalized s_i plots for control and the recovery 100 Hz trains are compared here. **C:** The s_1 ratio between recovery and control trains is < 1 , indicating a decrease in δ at the end of the low frequency train while neither s_2 (blue) nor s_{5-8} (purple) are changed, suggesting no change in p or in IP size. The similarity of results with and without postsynaptic BAPTA supports a presynaptic mechanism for LFD.

Data availability

The Data that supports the finding of this study will be made available in Dryad.

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Peer reviews

Reviewer #1 (Public review):

Summary:

In this work, the authors investigate the mechanisms of low-frequency synaptic depression at cerebellar parallel fiber to interneuron synapses using unitary recordings that allow direct quantification of synaptic vesicle release. They show that sparse stimulation can induce robust synaptic depression even in the absence of substantial vesicle consumption, and that this depressed state is rapidly reversed when stimulation frequency is increased. To account for these observations, the authors propose a model in which low-frequency depression reflects a redistribution of vesicles within the readily releasable pool, in particular a reduction in docking site occupancy due to vesicle undocking.

Strengths:

I found the experimental work to be of high quality throughout. The use of simple synapse recordings to count individual vesicle release events is particularly powerful in this context and allows questions to be addressed that are difficult to approach with more conventional approaches. The demonstration that low-frequency depression can occur independently of prior vesicle release, together with the rapid recovery observed during high-frequency stimulation, places strong constraints on possible underlying mechanisms and represents a clear strength of the study.

The modeling framework is clearly laid out and helps organize a broad set of observations across stimulation frequencies. Several of the experimental tests appear well motivated by the model, including the recovery train experiments, the analysis of failures, and the use of doublet stimulation. Taken together, the data provide a coherent phenomenological description of low-frequency depression and its relationship to vesicle availability within the readily releasable pool.

Weaknesses:

The major concerns raised during the initial review have been successfully addressed through substantial revisions of both the manuscript and the presentation of the model. The distinction between experimental observations and model-based interpretation is now considerably clearer, and the discussion more appropriately reflects the explanatory nature of the modeling framework.

<https://doi.org/10.7554/eLife.110329.2.sa3>

Reviewer #2 (Public review):

Summary:

Silva and co-workers exploit their previously established methods of analyzing release events at single parallel fiber to molecular layer interneuron synapses. They observed synaptic depression at low transmission frequencies (< 5 Hz) which rapidly recovers during high-frequency transmission. Analysis of the time course of low-frequency depression revealed an initial rapid and a slow linearly increasing time course. Strikingly, the initial depression occurred even in the absence of proceeding release arguing against vesicle depletion as the underlying mechanism.

Strengths:

The main strength of the study is the careful demonstration of an interesting synaptic phenomenon challenging the classical vesicle-centered interpretation of synaptic depression.

Weaknesses:

There are no weaknesses.

<https://doi.org/10.7554/eLife.110329.2.sa2>

Reviewer #3 (Public review):

Summary:

The manuscript builds on the observation that, at some synapses, low-frequency stimulation causes synaptic depression which can be reversed by subsequent high-frequency stimulation. Such low-frequency depression (LFD) cannot be easily explained by the depletion of a single vesicle pool. Here, Silva and colleagues propose a model of activity-dependent vesicle trafficking to explain LFD at synapses between cerebellar granule cells and molecular layer interneurons.

Strengths:

Overall, LFD is interesting and worthy of examination, and the authors provide new experimental results that are of the high quality expected from this group.

Weaknesses:

The study proposes a novel model of vesicle trafficking that is not explained by known biological mechanisms, and the manuscript does not adequately compare or discuss alternative models.

I have several concerns about how the authors interpret the data. First, the manuscript's primary conceptual advance is the idea that LFD involves vesicle undocking, rather than depletion. However, most experiments were performed under conditions that promote vesicle depletion (3 mM extracellular Ca^{2+}). When experiments were repeated in physiological Ca^{2+} , there appeared to be little or no LFD (stats are not provided). Second, the RS/DS/DU/undocking model, though not outside the realm of possibility, is not readily explained by known mechanisms and is only loosely supported by experimental findings. Third, when simulating LFD, the authors do not compare alternative models and use inappropriate language to imply that a model fit represents the truth (e.g. "the finding of identical experimental and simulated values confirms that the undocking mechanism accounts for LFD"). Finally, the model is presented in an overly complicated manner. The sheer amount of terms and nomenclature makes the manuscript confusing and difficult to read. Overall, the manuscript would benefit from added experiments and more statistics, a better justification and evaluation of the model, and more nuanced language.

Comments on revised version.

I appreciate the authors' detailed responses to my initial review of the manuscript. My suggestions reflect a sincere attempt to improve the clarity and accuracy of the paper. I disagree with some of the authors conclusions and their responses to my comments, but do not wish to make any further suggestions. The authors are entitled to their own views.

A final note: If the authors want to refrain from making definitive statements on underlying cellular mechanisms, they should consider amending the title of the manuscript.

<https://doi.org/10.7554/eLife.110329.2.sa1>

Author response:

The following is the authors' response to the original reviews.

Public Reviews:

Reviewer #1 (Public review):

Summary:

In this work, the authors investigate the mechanisms of low-frequency synaptic depression at cerebellar parallel fiber to interneuron synapses using unitary recordings that allow direct quantification of synaptic vesicle release. They show that sparse stimulation can induce robust synaptic depression even in the absence of substantial vesicle consumption, and that this depressed state is rapidly reversed when stimulation frequency is increased. To account for these observations, the authors propose a model in which low-frequency depression reflects a redistribution of vesicles within the readily releasable pool, in particular, a reduction in docking site occupancy due to vesicle undocking.

Strengths:

I found the experimental work to be of high quality throughout. The use of simple synapse recordings to count individual vesicle release events is particularly powerful in this context and allows questions to be addressed that are difficult to approach with more conventional approaches. The demonstration that low-frequency depression can occur independently of prior vesicle release, together with the rapid recovery observed during high-frequency stimulation, places strong constraints on possible underlying mechanisms and represents a clear strength of the study.

The modelling framework is clearly laid out and helps organize a broad set of observations across stimulation frequencies. Several of the experimental tests appear well-motivated by the model, including the recovery train experiments, the analysis of failures, and the use of doublet stimulation. Taken together, the data provide a coherent phenomenological description of low-frequency depression and its relationship to vesicle availability within the readily releasable pool.

We thank the Reviewer for his positive assessment of our work.

Weaknesses:

While the experimental results are strong, the manuscript would benefit from rebalancing the strength of the mechanistic conclusions drawn from the modelling in light of its limitations. The framework is clearly useful and provides a coherent interpretation of the data, but it is not uniquely constrained by the experimental observations, and alternative models or interpretations could plausibly account for the findings. The use of different model regimes concatenated across time, with substantially different parameter values, highlights the abstract nature of the approach. For these reasons, the model seems best presented as one plausible explanatory framework rather than a definitive biological mechanism. Clarifying the distinction between data-driven observations and model-based inferences would help readers assess which conclusions are strongly supported and which remain more speculative.

The interpretation of the Ca²⁺-related experiments would benefit from more cautious wording. The absence of detectable changes in presynaptic Ca²⁺ signals does not exclude more localized or subtle Ca²⁺-dependent mechanisms, and conclusions regarding Ca²⁺ independence should therefore be framed accordingly. In addition, while low-frequency depression is still observed at reduced extracellular Ca²⁺, these experiments appear less diagnostic of the specific model-derived mechanism emphasized elsewhere in the manuscript - namely, a selective reduction in docking-site occupancy - and should be discussed with appropriate qualification in the text.

Concerning Ca^{2+} signals, the Reviewer is right. While we found no change in Ca^{2+} signalling apart from a slow Ca^{2+} accumulation during long trains at 1 Hz, the possibility of an undetected change cannot be excluded. We have added a word of caution in this direction on p. 11. Concerning the 1.5 mM Ca^{2+} experiments, the Reviewer presumably alludes to the first recovery train (yellow) point in Supplementary Fig. 2C. This is also the last point (s11) of the slow train at 0.5 Hz because no delay at all was interposed between the slow train and the recovery train. We have now included one more experiment (with a present total number $n = 6$), and we have corrected Fig. S2C accordingly. In the new version the depression measured for s4-s10 vs s1 during the 0.5 Hz trains is 0.69 ± 0.05 ($p = 0.00058$, paired one-tail t-test). The ratio of the s1 value of the recovery train compared to control s1 is 0.83 ± 0.08 ($p = 0.028$, paired one-tail t-test).

Major points:

(1) Clarify and qualify mechanistic claims derived from the model.

Throughout the manuscript, changes in model parameters are at times described as if they directly reflected underlying physiological mechanisms. As a result, the conceptual distinction between experimentally observed phenomena, model-derived variables, and biological interpretation is not always clear. Several conclusions in the Results and Discussion are phrased as mechanistic statements, although they rest on assumptions intrinsic to the modelling framework. The authors should systematically review the text and explicitly distinguish between (i) experimentally observed changes in synaptic responses and (ii) inferences about vesicle docking states or transitions within the model.

In particular, statements implying that vesicle undocking is the mechanism underlying low-frequency depression should be rephrased to reflect that this is an interpretation within the proposed framework rather than a uniquely demonstrated biological process. For example, statements such as "Low-frequency depression is caused by synaptic vesicle undocking" should be replaced with formulations such as "Within the framework of our model, low-frequency depression is accounted for by a redistribution of synaptic vesicles away from docking sites" or "Our results are consistent with a model in which changes in vesicle docking-state occupancy contribute to low-frequency depression."

A particularly problematic example is the statement that "these experiments further confirm that LFD only involves a decrease in δ , without accompanying changes in p or IP size." Here, an experimentally defined phenomenon (LFD) is directly equated with changes in model-derived variables. Such statements should be revised to make clear that δ , p , and IP size are inferred quantities within the model, and that the experimental data are interpreted through this framework rather than directly confirming changes in these parameters. Similarly, overgeneralizing statements such as "Undocking therefore represents the key mechanism controlling short-term depression across stimulation frequencies" should be softened to reflect that this conclusion emerges from the model rather than from direct experimental evidence.

As suggested, we clarify the distinction in the revised version between experimental data and modelling, and we refrain from making definitive statements on underlying cellular mechanisms.

(2) Address the biological interpretation of time-dependent model regimes.

The model relies on distinct parameter regimes applied at different time points, with some transitions effectively suppressed in certain regimes. While this approach captures the data well, its biological interpretation remains unclear. The authors should either (i) expand the discussion to outline plausible biological processes that could give rise to such regime changes (for example, calcium-dependent modulation of transition rates or

activity-dependent changes in vesicle state stability), or (ii) more explicitly frame this aspect of the model as a descriptive abstraction rather than a mechanistic proposal. This further underscores the need to clearly separate the descriptive role of the model from claims about underlying biological mechanisms.

We thank the Reviewer for drawing our attention to this important point. Below 10 ms, rate constants are largely determined by the large-amplitude, fast-decaying Ca^{2+} signal occurring near voltage-dependent Ca^{2+} channels (' Ca^{2+} nanodomain'). After 10 ms, the rate constants depend on the low-amplitude, slowly decaying Ca^{2+} signals averaged over the entire varicosity ('volume-averaged Ca^{2+} '). We explain this better in the revised version (Materials and Methods, p. 21).

(3) Reframe conclusions drawn from calcium-related experiments.

The calcium imaging data demonstrate no detectable changes in the measured presynaptic calcium signals under the tested conditions, but they do not rule out that calcium signals contribute in ways undetectable by the assay. Conclusions should therefore be revised to reflect this limitation, avoiding statements that exclude a role for calcium-dependent mechanisms. Wording such as "we did not detect evidence for..." would be more appropriate than conclusions implying the absence of an effect.

Similarly, while low-frequency depression is still observed at reduced extracellular calcium (1.5 mM Ca^{2+}), the specific mechanistic signature emphasized elsewhere in the manuscript - namely a selectively reduced first response during a high-frequency recovery train - is no longer apparent. These experiments should therefore be discussed as consistent with the proposed framework, but not as providing independent support for a selective reduction in docking-site occupancy. Explicitly acknowledging this limitation would improve clarity and avoid overinterpreting these data.

This has been discussed above ('weaknesses').

(4) Soften interpretations based on non-significant comparisons.

In several places, comparisons that do not reach statistical significance are used to argue for equivalence between conditions (for example, comparisons involving failure versus non-failure trials or different LFD conditions). These conclusions should be revised to emphasize the limits of statistical power and framed as a lack of evidence for a difference rather than evidence of independence.

We have amended this point in the revised version.

Reviewer #2 (Public review):

Summary:

Silva and co-workers exploit their previously established methods of analyzing release events at single parallel fiber to molecular layer interneuron synapses. They observed synaptic depression at low transmission frequencies (< 5 Hz), which rapidly recovers during high-frequency transmission. Analysis of the time course of low-frequency depression revealed an initial rapid and a slow linearly increasing time course. Strikingly, the initial depression occurred even in the absence of preceding release, arguing against vesicle depletion as the underlying mechanism.

Strengths:

The main strength of the study is the careful demonstration of an interesting synaptic phenomenon challenging the classical vesicle-centered interpretation of synaptic depression.

We thank the Reviewer for his positive assessment of our work.

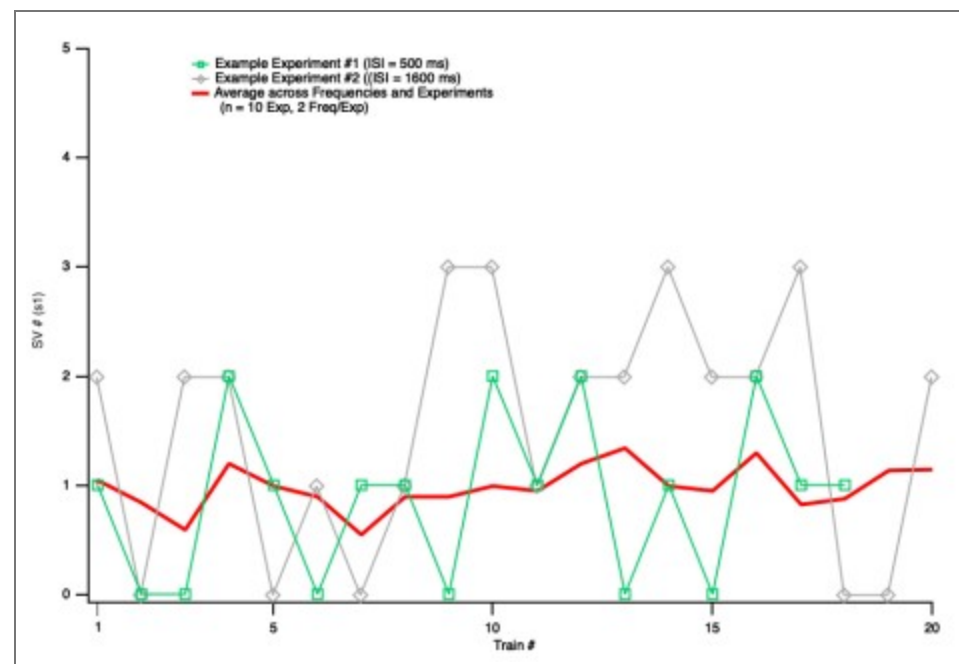
Weaknesses:

No major weaknesses were identified by this reviewer.

The finding of release-independent synaptic depression is important and would have widespread implications. Therefore, some more analyses to increase the confidence in these findings could be performed.

My concern is whether rundown could explain the findings. If the rate of failures in s1 increases and at the same time the amplitude decreases during the experiments, an apparent depression in s2 could arise. The Supplementary Figure 5A addresses run-down, but the figure is not easy to understand, and, as far as I understood, it does not address the question of whether the release-independent depression could be caused by a rundown. To address this, the analysis of Figure 5 could be repeated by investigating the failure rate and amplitude separately or by analyzing the 1st and 2nd half of the recordings separately.

The Reviewer makes a very important point that had escaped our attention. If the responses were declining over the course of an experiment, near the end of the recordings, a high proportion of failures would be associated with a weak response to the second AP. This could distort the relation between initial failures and amount of LFD, perhaps to the point of indicating LFD after failures when there were none. As suggested by the Reviewer, we tested this possibility by examining the stability of the synaptic responses during experiments. We found a mean s_1 value of 0.87 ± 0.13 for the first half of the experiments used in Fig. 5, and of 1.10 ± 0.17 for the second half ($p > 0.05$, $n = 10$). This analysis shows that there was no rundown during these experiments. We show in Author response image 1 a plot of s_1 as a function of train number in these experiments, for two examples as well as for the average across frequencies and across experiments. These plots do not suggest any artefactual correlation between failures, mean s_1 , and rundown.



Author response image 1. Plot of s_1 as a function of train number for the experiments of Fig. 5

In response to a request of Reviewer 2, Author response image 1 illustrates the evolution of s1 values as a function of train number for the experiments used to produce Figure 5. In each experiment, about 20 s1 values were obtained at two ISIs (either 10 ms and 500 ms, or 800 ms and 1600 ms). Author response image 1 shows two examples of s1 values as a function of train number (these values fluctuate widely between 0 and 3), and the average across cells and ISI values. There is no indication of a rundown of s1 values as a function of train number.

Reviewer #3 (Public review):

Summary:

The manuscript builds on the observation that, at some synapses, low-frequency stimulation causes synaptic depression, which can be reversed by subsequent high-frequency stimulation. Such low-frequency depression (LFD) cannot be easily explained by the depletion of a single vesicle pool. Here, Silva and colleagues propose a model of activity-dependent vesicle trafficking to explain LFD at synapses between cerebellar granule cells and molecular layer interneurons.

Strengths:

Overall, LFD is interesting and worthy of examination, and the authors provide new experimental results that are of the high quality expected from this group.

Weaknesses:

The study proposes a novel model of vesicle trafficking that is not explained by known biological mechanisms, and the manuscript does not adequately compare or discuss alternative models.

I have several concerns about how the authors interpret the data. First, the manuscript's primary conceptual advance is the idea that LFD involves vesicle undocking, rather than depletion. However, most experiments were performed under conditions that promote vesicle depletion (3 mM extracellular Ca²⁺). When experiments were repeated in physiological Ca²⁺, there appeared to be little or no LFD (stats are not provided). Second, the RS/DS/DU/undocking model, though not outside the realm of possibility, is not readily explained by known mechanisms and is only loosely supported by experimental findings. Third, when simulating LFD, the authors do not compare alternative models and use inappropriate language to imply that a model fit represents the truth (e.g., "the finding of identical experimental and simulated values confirms that the undocking mechanism accounts for LFD"). Finally, the model is presented in an overly complicated manner. The sheer amount of terms and nomenclature makes the manuscript confusing and difficult to read. Overall, the manuscript would benefit from added experiments and more statistics, a better justification and evaluation of the model, and more nuanced language.

We respectfully disagree with these sweeping criticisms, as described in more detail below.

Major concerns:

(1) Most experiments were performed under conditions that exacerbate depletion

In order to attribute LFD to vesicle undocking rather than depletion, it is important to show LFD under conditions where depletion is minimal. As mentioned above, the authors only report significant LFD in elevated extracellular Ca²⁺. In a small number of experiments performed in more physiological Ca²⁺ (1.5 mM), there is no depression after

a single stimulus, and it is not clear that there was statistically significant depression during a low-frequency train. Several studies cited in support of LFD share this problem:

Abrahamsson et al., (2007) recorded from Schaffer collaterals in 4 mM Ca, 3-4X physiological Ca²⁺.

Doussau et al., (2010) recorded from Aplysia synapses in 3X Ca compared to seawater.

Rudolph et al., (2011) is cited as an example of LFD. However, this study performed experiments at high release probability cerebellar climbing fibers, and reported depression that increased monotonically with stimulation frequency, so it does not resemble the phenomenon studied in this paper. Lin et al., (2022) also largely describe monotonic depression at the calyx.

The Reviewer suggests that LFD may only occur under non-physiological conditions, if the release probability has been increased by artificially elevating the extracellular Ca²⁺.

The implication is that LFD is at best a curiosity with little or no significance for brain signalling. We disagree with this point of view for several reasons.

Concerning the statement ‘In order to attribute LFD to vesicle undocking rather than depletion, it is important to show LFD under conditions where depletion is minimal’: This is the purpose of the analysis shown in Fig. 5.

The statement ‘the authors only report significant LFD in elevated extracellular Ca²⁺’ is inaccurate. Fig. S2C shows a clear LFD in 1.5 mM Ca²⁺, as acknowledged by Reviewer 1 (‘low-frequency depression is still observed at reduced extracellular Ca²⁺’). However, we failed to provide a p-value for the depression in the initial version of the paper (p was 0.004, n = 5 with this data set; paired t-test, one-tail). In the revised version, we document the 1.5 mM results more extensively, including the incorporation of the results of an additional experiment, and an explicit statistical analysis of the data (p = 0.00058, n = 6; paired t-test, one-tail).

Concerning the statement ‘there is no depression after a single stimulus’: We find that the onset kinetics of LFD is slower in 1.5 Ca²⁺ than in 3 Ca²⁺ (respectively 1.8 ISI and 0.51 ISI, Fig. 2C and Fig. S2C). This explains that the PPR is not significantly <1 in 1.5 Ca²⁺ without implying any weakening of the extent of LFD at steady state.

As explained in the manuscript (p. 5), in a previous work, we developed a method to ascribe changes in SV pools, within the RS/DS model, with specific modifications of s1, s2 and s5-s8 during test 100 Hz trains (Tran et al., 2022). This method was developed in 3 mM Ca²⁺ conditions, and for this reason, we performed most experiments for the present work in 3 mM Ca²⁺.

Chiu and Carter (2024) demonstrated LFD in neocortical synapses; they performed their study in 1.2 mM Ca²⁺, not in elevated Ca²⁺.

Rudolph et al., (2011) showed low-frequency depression not only in elevated external Ca²⁺, but also in 0.5 mM Ca²⁺. While Rudolph et al., (2011) did not make an explicit link between their observations and LFD, there is no reason to doubt that these observations are an example of LFD. They showed a biphasic depression when switching the stimulation frequency from 0.05 Hz to 2 Hz. In one of the founding papers of LFD, Doussau et al., (2010) describe a biphasic depression when switching the stimulation frequency from 0.025 Hz to 1 Hz; Fig. 1 of the two papers (Rudolph 2011 and Doussau 2010) are strikingly similar.

Lin et al., (2022) would probably not agree with the statement that the depression at the calyx is ‘largely monotonic’, as they stress the finding of quasi-constant depression between 5 and 50 Hz.

The authors note that their results differ from those of Atluri and Regehr, but do not mention that a possible reason for the difference is the increased release probability in their experiments.

In fact, we clearly listed the difference in external Ca^{2+} as a likely source of the discrepancy by saying ‘This discrepancy presumably stems from differences in experimental conditions (room temperature, stimulation of multiple presynaptic PFs and 2 mM external Ca^{2+} concentration in the previous work, vs. near-physiological temperature, single presynaptic stimulation and 3 mM external Ca^{2+} here)’.

The authors should provide statistics for the data obtained in 1.5 mM Ca, and discuss why LFD is increased in conditions that also elevate vesicle release probability.

See our comments above: the revised version includes the requested statistics. On p. 6 of the manuscript, we do provide an explanation for the apparent lack of LFD at 1.5 Ca^{2+} and 2 Hz, namely a superimposition of LFD with facilitation. At 1.5 Ca^{2+} and 0.5 Hz, our LFD numbers are not weaker than at 3 mM Ca^{2+} and 0.5 Hz of 1 Hz.

Altogether, it is correct that many LFD experiments have been carried out in high release probability synapses and/or under conditions of elevated Ca^{2+} . However, the reasons underlying these choices are diverse (in our case, to build on the previous SV pool analysis developed in Tran et al., 2022 in 3 Ca^{2+} conditions) and do not imply a limitation to the phenomenon. LFD is present in physiological conditions for low-to-moderate release probability synapses (as shown in our work), and altogether, there is no reason to dismiss LFD as non-physiological.

(2) Lack of biological mechanisms supporting the model

The model is presented without compelling biological support. The evidence in support of vesicle undocking comes from experiments by the Watanabe lab, which showed fewer-than expected docked vesicles under EM when cultured synapses were stimulated immediately prior to high-pressure freezing. Kusick et al were careful to note that these vesicles may have been lost to fusion.

The Watanabe lab showed an SV deficit at docking sites at times ranging from about 100 ms to several seconds (Kusick et al., 2020, their Fig. 5E). This corresponds to the ISI values where we see paired-pulse depression. In their Summary, Kusick et al., raise the possibility of SV fusion as an alternative to undocking at the 100 ms time point. But the same issue had previously been considered in Miki et al., 2018 with other techniques (their Fig. 2d), where it was shown that the SV deficit seen in paired-pulse experiments could not be explained by fusion. This leaves undocking as the most likely explanation, at least in our preparation. We have added a new paragraph on p. 14 to clarify this point.

The putative undocking Kusick describes is immediate (< 5 ms after stimulation), and it was not shown to be Ca^{2+} sensitive. This manuscript describes "calcium-dependent undocking" that proceeds from 10 ms - 200 ms. Multiple studies from the Watanabe lab show that a single stimulus lowers the number of docked vesicles, and subsequently, there is a transient redocking of vesicles that can be blocked by EGTA or Syt7 knockout.

This is not an accurate description of the Kusick results or of our results. In the Kusick paper, the SV deficit seen at <5 ms after stimulation is attributed to exocytosis, not to undocking. Clearly, it is Ca^{2+} -dependent. Our manuscript describes potential calcium-dependent undocking not during the time 10 ms- 150 ms, during which our undocking rate is assumed to be calcium-independent, but starting at 150 ms and lasting a few hundred ms thereafter.

I also question the rationale for the authors' model that 2 vesicles are coupled in series to a single release site. Previous papers from this lab cited EM studies from frog and neuromuscular that showed filamentous connections between vesicles (do these synapses show LFD?). Here, the authors primarily cite their previous models to support their arguments. I encourage them to continue searching for ultrastructural evidence for 2-vesicle-docking-units and to cite such studies.

It is important to remember that our sequential two-step model was not based on EM data, but on a series of functional data including variance-mean analysis of summed SV release numbers; covariance analysis among subsequent SV release numbers; analysis of release latencies as a function of stimulus number during an AP train; analysis of SV release numbers under conditions of very high release probability. We note that the phenomenon of Ca²⁺-dependent docking that we proposed based on these observations has been consistent with flash-and-freeze or zap-and-freeze results from several laboratories. Concerning potential filamentous connections between SVs and the AZ plasma membrane at a distance of several 10s of nm, this has been seen not only in frog or mice neuromuscular junctions, but also at brain synapses (ex: Siksou et al., Journal of Neuroscience 2007; Cole et al., Journal of Neuroscience 2016; Fernandez-Busnadiego, Journal of Cell Biology 2010; 2013).

(3) Comparison to other vesicle models

The authors use overly assertive language to suggest that the model proves a mechanism. "Altogether, these results indicate that the slow phase of LFD ... reflects a δ decrease without significant changes in pr, in p or in IP size". Simulating data does not conclusively "indicate" the underlying mechanism, but the authors could state their data can be "explained by a model where..".

Please see our response above to a similar point by Reviewer 1.

However, LFD does not require activity-dependent undocking. Instead, the phenomenon has been explained by high-release probability, paired with an activity-dependent increase in either docking or release probability (Chiu and Carter, 2024; Doussau et al., 2017). Does the new model do a better job of replicating some facet of the data? If multiple models can explain the same data, how can we determine which model is correct? The "Alternative Presynaptic Depression Mechanisms" should be expanded to discuss these issues.

We could not find statements in the Chiu and Carter paper or in the Doussau et al., paper explaining LFD 'by high-release probability, paired with an activity-dependent increase in either docking or release probability'. As far as we can see, Chiu and Carter do not propose any specific mechanism for LFD, beyond saying that depression and facilitation must be separate. Doussau et al., (their Fig. 6) clearly frame their interpretation in a sequential two-step model. As in the preceding Miki et al., paper (which they cite extensively), they assume a rapid (a few ms), Ca-dependent transition between their 'reluctant pool' and their 'fully-releasable pool', respectively homologous to RS and DS. Thus, the Doussau et al., interpretation is close to that presented in our present work, even though significant differences exist. An important difference is that Doussau et al., did not use simple synapses, so that they did not have access to key synaptic parameters such as the number of docking sites or the release probability per docking site. Consequently, the model in Doussau et al., does not have the same level of detail as ours. The revised version explains better the differences and similarity between the models of Doussau et al., and that exposed in our work (new paragraph on p. 14).

Recommendations for the authors:

Reviewing Editor Comments:

Three reviewers have seen the paper. They agree that the evidence for low-frequency depression (LFD) is solid, but they all suggest that the data with 1.5 mM extracellular calcium should be analyzed and discussed in more detail. The finding of release-independent depression is surprising, but further data analysis to investigate whether it could be caused by run-down is required. The reviewers think the study will be more complete if the authors add N's and stats for LFD in 1.5 mM Ca. In addition, the most important change would be adding clarity in the text about the limitations and simplifications of the model. Please edit the text to clearly state that the presented model provides one possible framework to account for the data, and that other models might also be plausible, and clearly describe the limitations and simplifications of the model.

Thank you for your positive assessment of our work. As explained in more detail below, we have added the requested statistics for 1.5 mM Ca data. We have also thoroughly revised our text to separate more clearly facts and interpretation. Finally, we now explain in more detail the limitations of our model.

Reviewer #1 (Recommendations for the authors):

Minor points

(1) Statistical analysis and reporting.

Please justify the use of one-tailed statistical tests and consider whether two-tailed tests would be more appropriate in some cases.

In Materials and Methods (p. 17), we explain that one-tailed statistics were used when the sign of any possible deviation was predicted; otherwise two-tailed statistics were used.

Exact p-values should be reported consistently rather than using threshold notation.

We have corrected this as far as we could. Exact p-values that are still missing will be provided in the version of record.

For analyses using rank-based statistics (e.g., calcium imaging and recovery experiments), please clarify which comparisons were paired and which were unpaired, and ensure that this is clearly indicated in the Methods and figure legends.

We have now clearly indicated whether these statistics were paired or unpaired.

(2) Figure clarity and presentation.

In Figure 1, clarify whether the responses shown in panel C represent measured EPSCs or derived quantities, as this is not immediately clear from the labelling.

Thank you for spotting this, the figure has been corrected.

In Figure 2, please show the variability of control responses used for normalization.

The variability of control responses used for normalization is now clearly explained in Materials and Methods (p. 18).

In Figure 3, the use of similar color schemes for different model states and for data-model comparisons is confusing and should be revised.

Thank you for spotting this; has been corrected.

In Figures 4 and 5, consider reintroducing or clarifying schematic representations of the model states referred to in the text to aid reader comprehension.

Figure 4 had already such representations. We have now modified these representations to make them simpler and clearer. We considered adding some model to Fig. 5 but could not come to any satisfactory option.

(3) Controls and interpretation of pharmacological experiments.

For experiments involving intracellular BAPTA, it would strengthen the interpretation to either include or explicitly discuss positive controls demonstrating that the manipulation was effective under the recording conditions. Alternatively, loosen the interpretation of those data.

We have modified the text as suggested by the Reviewer.

(4) Terminology and consistency.

Ensure consistent use of terminology for vesicle states and model variables throughout the text and figures, and clarify whether different labels (e.g., RS/DS versus alternative notations) refer to equivalent states.

We have revised our manuscript as suggested.

(5) Model fit at short inter-stimulus intervals.

In Figure 3B, the model appears to capture the overall trend of facilitation but shows a noticeable deviation from the data at the shortest inter-stimulus intervals. It would be helpful to clarify whether this reflects limitations of the current parameterization, known simplifications in the model (e.g., assumptions about fast calcium-dependent processes), or variability across recordings. Briefly commenting on this mismatch would help readers interpret the strengths and limits of the model in this regime.

Because of the statistical nature of data (quantal fluctuations) and simulations (Monte Carlo trials), some differences between data and simulations are unavoidable. The low simulation point at 10 ms ISI in Fig. 3B may in addition result from an imperfect transition between the two time domains (time domains 1 and 2) of the simulation. We have added a sentence in Materials and Methods (p. 22) to explain this.

(6) Model parameterization and fitting.

Please clarify how model parameters were estimated, including the fitting procedure, cost function, and any constraints applied. It would be helpful to report confidence intervals or other measures of uncertainty for key fixed parameters, and to indicate how variable these parameters are across cells or recordings. In addition, a brief discussion of how sensitive the main conclusions are to variation in key parameters (such as release probability or transition rates) would help readers assess the robustness and identifiability of the model-derived interpretations.

In addition, it is not entirely clear how model parameters and transitions differ across the concatenated temporal regimes used in the simulations, nor which parameters are held fixed versus altered or effectively suppressed. Summarizing, for each regime, the parameter values (or ranges) and active transitions—either in a table or schematic—would greatly improve transparency and help readers assess parameter identifiability and the robustness of the model-derived conclusions.

For each group of experiments, results under control conditions (8-APs at 100 Hz) were averaged together, leading to the determination of one set of parameters (N , ρ , δ , r , s , p^I). Next the parameters for the second time domain were obtained by trial and error to optimize the fit for Fig. 1 (PPR) data. The parameters were constrained such the PPR recovered to 1

with a 10-second ISI. Unfortunately, the parameter domain was too complex to make an unsupervised fitting procedure practical. We have now added a paragraph on p. 22 to indicate which model parameters were heavily constrained by data and which ones were less constrained. Individual synapses were not simulated. Regarding the different time domains, the probabilities r_b and s_b are introduced for the second time domain and r_f and s_f are changed as shown in Fig. Supp. 3. The kinetic parameters are reset at the beginning of each simulated AP.

(7) Clarify the scope of the model with respect to the slow component of low-frequency depression.

The manuscript identifies a slow component of low-frequency depression that persists over long timescales. While this is an interesting observation, it is not fully clear whether this component is intended to be captured by the current model or represents an additional process outside its scope. Making this distinction explicit would help readers better grasp the scope and limitations of the model.

As stated in our manuscript, modelling the slow component of LFD falls outside the scope of the present work. The set of rate constants in Fig. S3 does not produce any slow component, and we failed to find a combination of rate parameters that would produce a slow component. This is now clearly stated in the Materials and Methods section.

Reviewer #2 (Recommendations for the authors):

I have only the following suggestions to further improve the manuscript.

(1) It is argued that LFD is alleviated when using doublets rather than singlets (Figure 8), but is the steady-state depression in s_1 between singlets and doublets at the end of the experiment statistically different?

It is unclear what should be considered steady state here. Average s_1 values are smaller for singlets than for doublets over the last 25 points (0.58 ± 0.05 and 0.74 ± 0.05 , $p = 0.02$, unpaired t-test).

(2) I do not understand Supplementary Figure 4. Does the value of zero in this plot indicate that the stimulation does not evoke any release anymore? How can this be differentiated from rundown? Can the LFD be reversed by a high-frequency stimulation?

We have rewritten the description of this figure. We have also modified the figure labelling. Finally, we have changed the size and color of the symbol at zero delay to stress that the linear fit is constrained to include this point. Altogether we trust that these changes have clarified the presentation of these results.

(3) The findings are related to Kusick et al., (2020). However, I think that in Kusick et al., (2020) as well as in Watanabe et al., (2013; doi:10.1038/nature12809), all tested time points ranging from a few milliseconds to several seconds after a stimulus show vesicle depletion except the time point of 14 ms in Kusick et al., (2020). I think these findings are, therefore, inconsistent with the much slower observed LFD.

The increase in docked SV# at 14 ms is reported not only in Kusick et al., 2020 but also in Wu et al., 2023 as well as in Ogunmowo et al., 2025. Later, at 100 ms, the number of docked SVs drops again (Kusick et al., 2020, their Fig. 5 d-e, and Ogunmowo et al., 2025), to finally recover on a time scale of seconds (Kusick et al., 2020). These data fit with our observations if LFD is due to undocking, as we propose.

(4) Page 13, 4th paragraph:

Eshra et al., (2021) provide experimental evidence for "a Ca²⁺ -dependent rate of SV entry into the RRP".

The Reviewer is right: Eshra et al., actually reported a weak Ca²⁺-dependence of the replenishment. We have therefore eliminated the Eshra reference in this sentence (note that this paper is mentioned elsewhere in the manuscript).

Reviewer #3 (Recommendations for the authors):

(1) Complicated terminology: I had a hard time understanding the model due to terms like δ , ρ , IP, DS, RS, RS gate, etc. Anything the authors can do to reduce the number of acronyms and simplify their description/depiction of the model will be helpful for readers.

This resembles recommendation (4) of Reviewer 1. As stated in our response above, we have entirely revised the manuscript with these issues in mind. In addition, the abbreviation list at the onset of the manuscript should help the readers.

(2) Figure 5 may represent the strongest evidence in support of an undocking model. This is the smallest figure, and I found it harder to understand. For example, I was initially confused because the blue "fail S1" traces seemed to suggest a very large PPR value. I suggest vastly expanding the size of this figure and the associated results section.

Thank you for your interest in Fig. 5. We have explained the normalization procedure in more detail in the revised version. Also, we have modified the axis label of Fig. 5B to improve clarity. Finally, we have added a paragraph to explain the reason why the RS/DS model accounts for the results of Fig. 5.

(3) Given the similar results but differing mechanistic conclusions of Doussau et al., (2017), that study merits more discussion in the manuscript.

See our response to this point above (main comment (3) by the same Reviewer).

(4) Several studies of LFD synapses have found a role for kinase activity (Silverman-Gavrila et al., 2005; Doussau et al., 2010). This mechanism could be discussed or experimentally tested here.

As mentioned in the Discussion section, the nature of the proteins involved in LFD, or in docking/undocking, remains uncertain. It is not surprising that broad-spectrum kinases and phosphatases affect LFD, as reported in the papers quoted by the Reviewer, but the full implications of these findings will only become clear

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