

Voltage-Dependent Calcium Channels at the Plasma Membrane, but Not Vesicular Channels, Couple Exocytosis to Endocytosis

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SUMMARY

Although calcium influx triggers endocytosis at many synapses and non-neuronal secretory cells, the identity of the calcium channel is unclear. The plasma membrane voltage-dependent calcium channel (VDCC) is a candidate, and it was recently proposed that exocytosis transiently inserts vesicular calcium channels at the plasma membrane, thus triggering endocytosis and coupling it to exocytosis, a mechanism suggested to be conserved from sea urchin to human. Here, we report that the vesicular membrane, when inserted into the plasma membrane upon exocytosis, does not generate a calcium current or calcium increase at a mammalian nerve terminal. Instead, VDCCs at the plasma membrane, including the P/Q-type, provide the calcium influx to trigger rapid and slow endocytosis and, thus, couple endocytosis to exocytosis. These findings call for reconsideration of the vesicular calcium channel hypothesis. They are likely to apply to many synapses and non-neuronal cells in which VDCCs control exocytosis, and exocytosis is coupled to endocytosis.

INTRODUCTION

Endocytosis is coupled to exocytosis, allowing recycling of vesicles and maintaining exocytosis (Royle and Lagnado, 2003; Schweizer and Ryan, 2006). Many studies suggest that calcium influx regulates endocytosis at synapses and non-neuronal secretory cells (Marks and McMahon, 1998; Cousin and Robinson, 1998; Gad et al., 1998; Sankaranarayanan and Ryan, 2001; Balaji et al., 2008; Artalejo et al., 1995; Neves et al., 2001; Wu et al., 2005; Clayton et al., 2007; but see von Gersdorff and Matthews, 1994; Leitz and Kavalali, 2011). Recent studies of a large nerve terminal, the calyx of Held, suggest that regulation of endocytosis by calcium reflects the trigger of endocytosis (Wu et al., 2009; Hosoi et al., 2009). Although the calcium channel that couples exo- to endocytosis was not identified, the voltage-dependent calcium channel (VDCC) was often implicitly assumed to be the candidate because calcium influx via VDCCs

triggers exocytosis, and exocytosis is coupled to endocytosis. However, a recent study of *Drosophila* synapses proposed that the calcium channel is Flower, a vesicular membrane protein transferred from the vesicle to the plasma membrane via vesicle fusion (Yao et al., 2009). This proposal may have significant impact for four reasons indicated by many reviews (Yao et al., 2009; Brose and Neher, 2009; Shupliakov and Brodin, 2010; Vogel, 2009; Kuo and Trussell, 2009). First, it could be a universal mechanism because Flower is conserved from worm to human (Yao et al., 2009). Second, experiments suggesting that calcium regulates or triggers endocytosis were mostly done by manipulating extracellular calcium or intracellular calcium buffers, which could be accounted for with the Flower hypothesis. Although the calcium current charge was found to correlate with the endocytosis rate (Wu et al., 2009), the exocytosis amount was not well controlled, making it possible for the vesicular calcium channel to account for this correlation. Thus, in principle, Flower could replace the plasma membrane VDCC to trigger endocytosis. Third, the Flower hypothesis explains exo-endocytosis coupling better than plasma membrane VDCCs because (1) Flower channels inserted to the plasma membrane may keep track of the level of exocytosis and thus be responsible for the exo-endocytosis match, and (2) Flower channels could diffuse to the endocytic zone, explaining why endocytic zones may differ from active zones. Fourth, a study in sea urchin eggs found that upon exocytosis, VDCCs are translocated from the vesicle membrane to the plasma membrane to mediate calcium influx required for endocytosis (Smith et al., 2000). These studies led to a more general hypothesis that vesicular calcium channels conserved from sea urchins to nerve terminals couple exo- to endocytosis (Vogel, 2009).

The vesicular channel hypothesis is potentially critical not only in exo-endocytosis coupling but also in many other calcium-dependent exocytosis processes (Brose and Neher, 2009; Kuo and Trussell, 2009). Despite such potential roles, whether fusion-inserted vesicle membrane generates calcium currents or influx at nerve terminals was not tested directly at nerve terminals. Whether plasma membrane VDCCs participate in triggering endocytosis with vesicular calcium channels is also unclear. The present work addressed these issues at the calyx of Held nerve terminal. We found that fusion-inserted vesicle membrane did not generate calcium currents or influx. Instead, the VDCC at nerve terminal membrane coupled exo- to endocytosis.

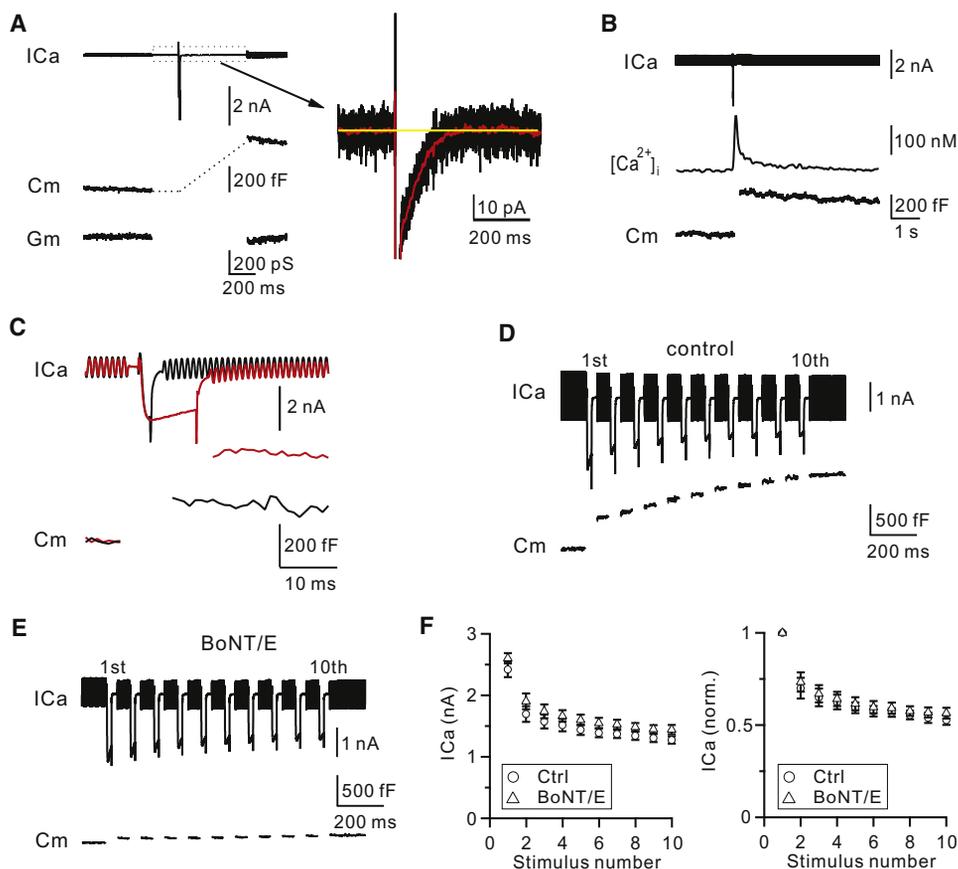


Figure 1. Vesicle Fusion Does Not Generate Calcium Currents at Nerve Terminals

(A) Left view shows sampled I_{Ca} , C_m , and G_m (membrane conductance) induced by a $depol_{10ms}$. I_{Ca} was low-pass filtered at 3 kHz. Right view presents I_{Ca} from the left view plotted in larger vertical and time scales (black). The red trace was the black trace low-pass filtered at 100 Hz. Yellow line is the mean baseline value.

(B) Sampled I_{Ca} , $[Ca^{2+}]_i$, and C_m induced by a $depol_{10ms}$.

(C) Sampled I_{Ca} and C_m induced by 2 (black) and 10 ms (red) depolarization.

(D and E) Sampled I_{Ca} and C_m induced by a $depol_{20ms \times 10}$ with a pipette containing 150 nM boiled BoNT/E (D, control) or BoNT/E (E).

(F) I_{Ca} (mean \pm SE, left) and normalized I_{Ca} (right) induced by each 20 ms pulse during $depol_{20ms \times 10}$ in the presence of 150 nM boiled BoNT/E (circles) or BoNT/E (triangles).

See also Figures S1 and S2A.

RESULTS

Exocytosis Does Not Generate Calcium Currents

Vesicular calcium channels, if present, could be either voltage-independent Flower channels or VDCCs. We tested these two possibilities by measuring calcium currents (I_{Ca}) after and during depolarization, respectively. Depolarization at the calyx was 10 ms from the holding potential of -80 to $+10$ mV (applies unless mentioned otherwise, $depol_{10ms}$), which induced an I_{Ca} of 2.10 ± 0.11 nA and a capacitance jump (ΔC_m) of 427 ± 24 fF ($n = 12$, Figure 1A). At the ΔC_m peak, ~ 400 – 500 ms after $depol_{10ms}$, the steady-state I_{Ca} was undetectable (<3 pA), suggesting that exocytosis did not insert calcium channels to generate steady-state I_{Ca} at -80 mV. Alternatively, a calcium-dependent outward current could be in balance with the exocytosis-generated I_{Ca} . This latter possibility was ruled out because we detected no outward current (<3 pA at 400 – 500 ms after

$depol_{10ms}$) when the ΔC_m was reduced to 83 ± 9 fF ($n = 5$) or $\sim 19\%$ of control by botulinum neurotoxin E (BoNT/E, 150 nM, in pipette, Figure S1A). Similar results were obtained when BAPTA (10 mM) was included with BoNT/E to block potential calcium-dependent currents (Figure S1B, $n = 5$).

Because a vesicle's membrane capacitance is ~ 0.07 fF (Sun et al., 2002; He et al., 2006), the 427 fF ΔC_m induced by $depol_{10ms}$ corresponded to $\sim 6,100$ vesicles. If each vesicle contained exactly 1 calcium channel, $\sim 6,100$ channels would be inserted to the plasma membrane. I_{Ca} traces were originally low-pass filtered at 3 kHz (Figure 1A, black). After we low-pass filtered I_{Ca} traces at 100 Hz (Figure 1A, red), the SD of I_{Ca} traces was 0.83 ± 0.08 pA ($n = 8$). We then set our detection limit at 3 pA, which was greater than three times the SD with $>99.7\%$ confidence for detecting changes every 10 ms. If there were 6,100 channels inserted at the plasma membrane, each channel would generate <0.0005 pA of I_{Ca} , ~ 600 times less than the

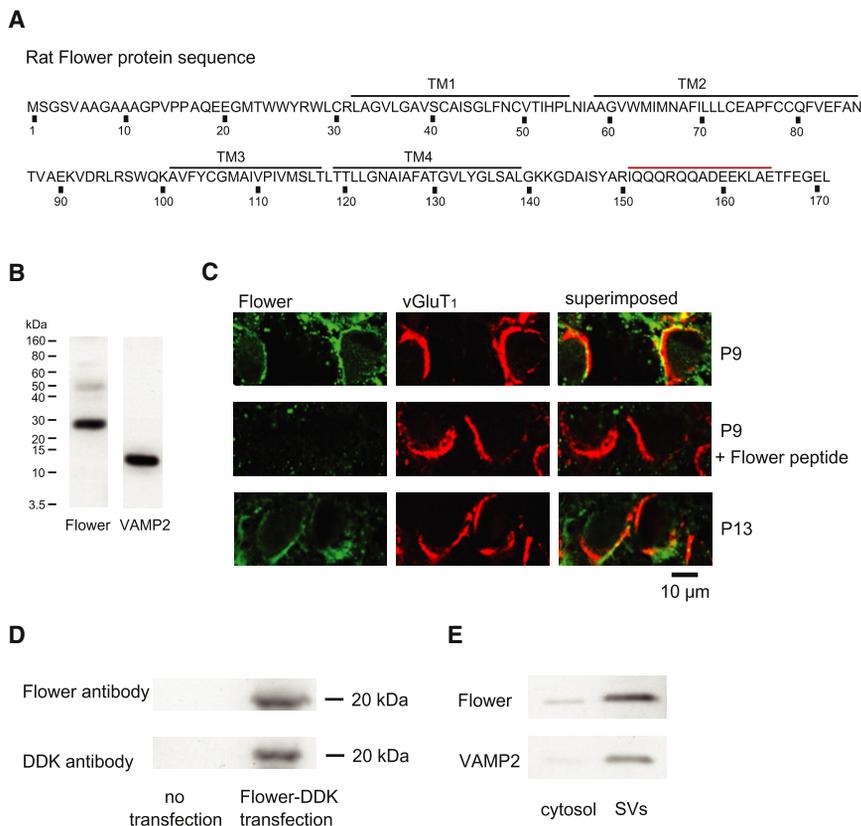


Figure 2. Flower Localization in Rat Brain, Calyces, and Synaptic Vesicles

(A) Rat Flower sequence (NP_001100031.1). Black lines denote the transmembrane domains (TM). Red line denotes the Flower peptide used for Flower antibody generation.

(B) Western blot of Flower and VAMP2 from rat brain.

(C) Confocal images showing the antibody staining against Flower (left) and vGluT₁ (middle, calyx labeling) at P9 (upper) and P13 (lower) calyces (right panel is left and middle panels superimposed). Flower peptide preincubation blocked Flower antibody staining of P9 calyces (middle).

(D) Western blot of DDK-tagged Flower protein from HEK293 cells with or without Flower transfection.

(E) Western blot of Flower and VAMP2 from cytosol and synaptic vesicles (SVs) purified from rat brain.

single-VDCC current (~ 0.3 pA at -80 mV) (Li et al., 2007; Weber et al., 2010). There was no such small single-channel current observed. If calcium channels with a single-channel current similar to VDCCs (0.3 pA) were in a fraction of vesicles, the fraction would be $<0.16\%$ (≈ 3 pA/0.3 pA/6,100). Thus, fused vesicles did not contain voltage-independent calcium channels with any reported conductance.

Consistent with this conclusion, the volume-averaged $[Ca^{2+}]_i$ detected with fura-2 decayed to within 10 nM above baseline at 2–3 s after depol_{10ms}, at which $>75\%$ ΔC_m remained, indicating that fused vesicle membrane could not generate significant $[Ca^{2+}]_i$ increase ($n = 5$, Figure 1B). Similar results (<10 nM at 2–3 s, $n = 4$) were observed when endocytosis was blocked by 0.3 mM GDP β S in the pipette.

Two sets of data suggest that exocytosis did not generate I_{Ca} during depolarization. First, depol_{10ms} induced a ΔC_m 267% \pm 13% ($n = 6$) of that induced by a 2 ms depolarization (Figure 1C). However, I_{Ca} during depol_{10ms} did not increase. I_{Ca} at 9.9 ms during depolarization was 84% \pm 2% ($n = 6$) of that at 2 ms (Figure 1C). This was not due to a balance between calcium-dependent outward currents and exocytosis-inserted I_{Ca} because in the presence of BoNT/E that blocked exocytosis, I_{Ca} at 9.9 ms during depolarization was 85% \pm 2% ($n = 5$, Figure S1A, inset) of that at 2 ms, similar to control. Similar results (86% \pm 1%, $n = 5$) were obtained in the presence of BAPTA (10 mM) and BoNT/E that blocked potential calcium-dependent currents (Figure S1B, inset). Second, in control (150 nM boiled BoNT/E in pipette) during ten pulses of 20 ms depolarization at 10 Hz

(depol_{20ms} \times 10), the total ΔC_m after the first pulse was 503 \pm 33 fF but increased by 938 \pm 55 fF after the tenth pulse ($n = 5$), corresponding to fusion of additional $\sim 13,000$ ($\approx 938/0.07$) vesicles. This additional fusion predicts an increase of 1.3 nA of I_{Ca} if each fused vesicle contains a VDCC with reported ~ 0.1 pA single-channel current at +10 mV (Weber et al., 2010). However, I_{Ca} decreased from 2.42 \pm 0.12 nA at the first pulse to 1.27 \pm 0.06 nA (53% \pm 3%, $n = 5$) at the tenth pulse (Figure 1D), suggesting no vesicular VDCCs. This I_{Ca} decrease was not due to a balance between I_{Ca} inactivation and exocytosis-induced VDCC insertion because a similar I_{Ca} decrease was observed in the presence of BoNT/E (150 nM), which blocked the total ΔC_m induced by depol_{20ms} \times 10 to 179 \pm 27 fF ($\sim 12\%$ of control, $n = 5$, Figures 1E and 1F). Addition of the calcium buffer BAPTA (10 mM) together with BoNT/E (150 nM) partially relieved I_{Ca} decrease (Figures S1C and S1D, $n = 5$), consistent with calcium-dependent I_{Ca} inactivation (Forsythe et al., 1998; Xu and Wu, 2005). We concluded that exocytosis did not generate calcium currents during and after depolarization (Figure 1).

Flower at Nerve Terminals

Results in Figure 1 prompted us to determine whether Flower is expressed in rat brain. An antibody we generated against Flower amino acids 151–165 (Figure 2A) detected a major band in rat (P8) brain (Figure 2B). Immunostaining with this antibody was colocalized with the antibody staining against vesicular glutamate transporter 1 (vGluT₁), which labeled calyces (Figure 2C, upper). Our antibody is specific to Flower because (1) Flower peptide (Flower amino acids 151–165) blocked Flower antibody staining of the calyx (Figure 2C, middle), and (2) Flower antibody specifically recognized DDK-tagged Flower protein overexpressed in HEK293 cells, which contained no endogenous Flower protein (Figure 2D). The molecular weight of Flower overexpressed in

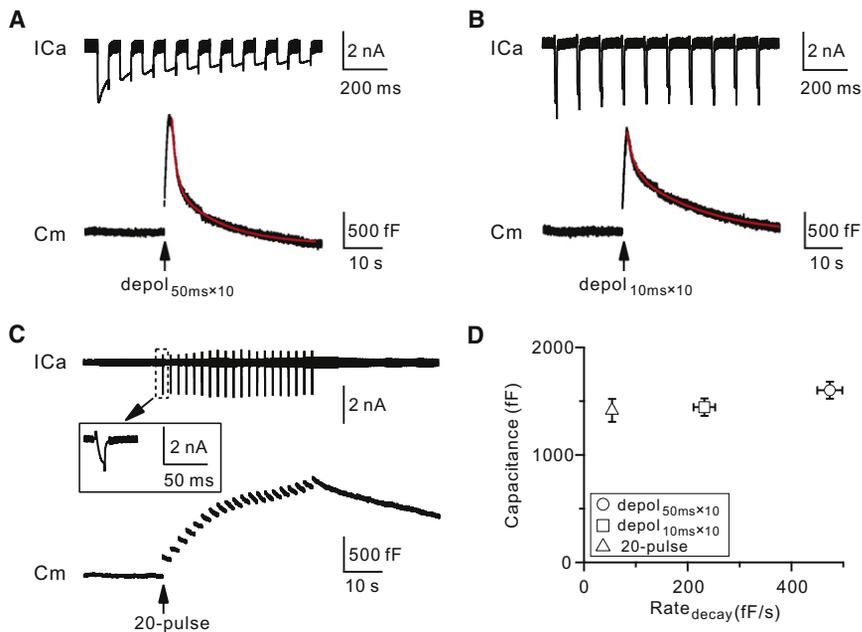


Figure 3. Voltage-Dependent Calcium Currents Determine the Endocytosis Rate

(A–C) Sampled I_{Ca} and Cm induced by depol_{50ms}×10 (A), depol_{10ms}×10 (B), and 20 pulse train (C). Red curves are biexponential fit of the capacitance decay (A: τ , 1.1 and 12 s; B: τ , 1.8 and 18.7 s). The inset shows I_{Ca} induced by the first pulse (–80 to –5 mV) in the 20 pulse train. (D) The Δ Cm (mean \pm SE) is plotted versus the Rate_{decay} (mean \pm SE) induced by depol_{50ms}×10 (circle, n = 12), depol_{10ms}×10 (square, n = 13), and the 20 pulse train (triangle, n = 18). See also Figures S2B–S2D.

HEK293 cells (~20 kDa) was similar to that in *Drosophila* (Yao et al., 2009) but smaller than that in rat brain extracts (~25 kDa, Figures 2B and 2D), likely due to posttranslational modification in rat brain.

Western blot of proteins from cytosol and synaptic vesicles purified from the whole rat brain showed that Flower was preferentially localized in vesicles, similar to vesicle protein VAMP2 (Figure 2E). These results suggest that Flower is expressed in synaptic vesicles of rat brain and calyces, consistent with the report of *Drosophila* (Yao et al., 2009). Although these results suggest expression of Flower at many synapses, we do not know whether Flower is expressed at all or only a fraction of synapses in the brain. Nevertheless, Flower expression at the calyx and synaptic vesicles (Figure 2), together with the lack of exocytosis-generated I_{Ca} (Figure 1), suggests that vesicular Flower insertion to the plasma membrane does not generate I_{Ca}.

VDCC, but Not Exocytosis, Determines Endocytosis Rate

To determine whether the plasma membrane VDCC controls endocytosis, we measured endocytosis after different VDCC-mediated I_{Ca} charges (QICas) while maintaining similar amounts of exocytosis (similar inserted vesicular proteins). Ten pulses of 50 ms depolarization at 10 Hz (depol_{50ms}×10) induced a Δ Cm of 1603 ± 78 fF (n = 12) and an QICa of 453 ± 35 pC (n = 12, Figures 3A and 3D). The capacitance decay, which reflects endocytosis, was biexponential with a τ of 1.2 ± 0.1 s (n = 12; weight, $62\% \pm 4\%$) and 12.5 ± 0.4 s (n = 12), respectively. The initial decay rate (Rate_{decay}) was 474 ± 24 fF/s (n = 12), which reflected mostly the initial rate of the rapid endocytosis component because the rapid component contributed >80% of the Rate_{decay} after this stimulus (Wu et al., 2009).

Ten pulses of 10 ms depolarization at 10 Hz (depol_{10ms}×10) induced a similar Δ Cm (1445 ± 80 fF, n = 13; p = 0.17) as

n = 13; p < 0.01) and Rate_{decay} (231 ± 21 fF/s, n = 13; p < 0.01) than depol_{50ms}×10 (Figures 3B and 3D). The capacitance decay was biexponential with a τ of 2.1 ± 0.2 s (n = 13; weight, $31\% \pm 4\%$) and 18.1 ± 1.0 s (n = 13), respectively (Figure 3B). The τ and the weight of the rapid component were slower and smaller than those induced by depol_{50ms}×10 (p < 0.01). However, the rapid component of endocytosis still contributed to the Rate_{decay} by more than 80% (calculation not shown). Evidently, the Rate_{decay} difference induced by depol_{50ms}×10 and depol_{10ms}×10 is not due to the exocytosis difference (Figures 3A, 3B, and 3D). It must be due to the difference in the VDCC-mediated QICa (~453 versus ~193 pC) because calcium influx triggers and speeds up endocytosis at calyces (Wu et al., 2009; Hosoi et al., 2009).

Twenty pulses of 10 ms depolarization from –80 to –5 mV at 0.5 Hz (20 pulse train) induced a Δ Cm (1414 ± 107 fF, n = 18) similar to those induced by depol_{10ms}×10 or depol_{50ms}×10 (p > 0.1; Figures 3C and 3D), but a much slower endocytosis, reflected as a mono-exponential capacitance decay with a τ >30 s (n = 18) and a much smaller Rate_{decay} (53 ± 4 fF/s, n = 18, Figures 3C and 3D; p < 0.01). This slower Rate_{decay} was clearly not caused by smaller Δ Cm and, thus, not by smaller amount of inserted vesicular proteins (Figure 3D). It was caused by smaller calcium concentration for two reasons. First, each pulse from –80 to –5 mV induced an I_{Ca} (1.4 ± 0.1 nA, n = 18; Figure 3C) much smaller than that by depolarization to +10 mV during depol_{10ms}×10 (2.2 ± 0.1 nA, n = 13; p < 0.01). Second, the pulse was repeated at 0.5 Hz, at which calcium summation was negligible compared to the 10 Hz train (Figure 1B).

P/Q-type Calcium Channels Control Endocytosis

To provide more direct evidence that VDCCs couple exo- to endocytosis, we blocked VDCCs with the P/Q-type VDCC blocker, ω -Agatoxin-IVA. Calyces in P7–P10 rats contain

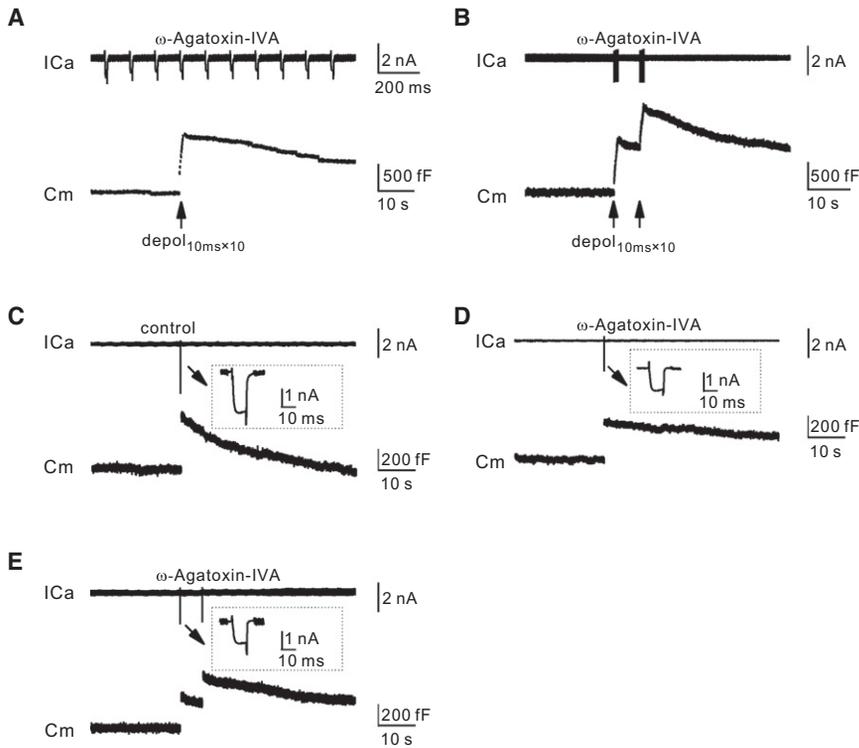


Figure 4. P/Q-type Calcium Channels Are Involved in Rapid and Slow Endocytosis

(A and B) Sampled I_{Ca} and C_m induced by a $depol_{10ms \times 10}$ (A) or a pair of $depol_{10ms \times 10}$ at 5 s interval (B) in the presence of ω -Agatoxin-IVA (200 nM, bath).

(C–E) Sampled I_{Ca} and C_m in the absence (control, C) or the presence of ω -Agatoxin-IVA (200 nM, D and E). The stimulation was a $depol_{10ms}$ in (C) and (D), and a pair of $depol_{10ms}$ in (E).

P/Q-, N-, and R-type VDCCs, among which P/Q-type contributes about two-thirds of I_{Ca} (Wu et al., 1999; Iwasaki et al., 2000). In the presence of ω -Agatoxin-IVA (200 nM, bath, 10–40 min), $depol_{10ms \times 10}$ induced a QICa (72 ± 6 pC, $n = 12$) and a ΔC_m (812 ± 41 fF, $n = 12$; Figure 4A) smaller than control (193 ± 14 pC, 1445 ± 80 fF, $n = 13$; Figure 3B; $p < 0.01$). Endocytosis was much slower because the capacitance decay was too slow to fit exponentially, and the $Rate_{decay}$ (50 ± 4 fF/s, $n = 12$, Figure 4A) was much slower than control (231 ± 21 fF/s, $n = 13$, Figure 3B; $p < 0.01$). The slower $Rate_{decay}$ was not due to the ΔC_m decrease because a pair of $depol_{10ms \times 10}$ values at 5 s interval induced a ΔC_m (1260 ± 36 fF, $n = 10$, Figure 4B) similar to that by a $depol_{10ms \times 10}$ in control ($p > 0.05$; Figure 3B) but induced a $Rate_{decay}$ (61 ± 5 fF/s, $n = 10$, Figure 4B) much slower than that by a $depol_{10ms \times 10}$ in control ($p < 0.01$; Figure 3B). Thus, VDCCs, including P/Q-type, couple exocytosis to rapid endocytosis.

To study slow endocytosis, $depol_{10ms}$ was applied. $depol_{10ms}$ induced a ΔC_m of 427 ± 24 fF ($n = 12$), followed by a mono-exponential decay with a τ of 15.8 ± 0.7 s ($n = 12$) and a $Rate_{decay}$ of 36.7 ± 2.7 fF/s ($n = 12$, Figure 4C). The τ (~ 15.8 s) of this slow endocytosis was much shorter than that after the 20 pulse train, which also induced slow endocytosis with a $\tau > 30$ s ($n = 18$; Figure 3C), but a ΔC_m $\sim 231\%$ higher than that induced by $depol_{10ms}$. Thus, more exocytosis was followed by endocytosis with a slower τ , inconsistent with the vesicular channel hypothesis. The slower τ was likely due to two factors. First, I_{Ca} (1.4 ± 0.1 nA, $n = 18$) induced by each pulse during the 20 pulse train was less than that induced by a $depol_{10ms}$ (2.10 ± 0.11 nA, $n = 12$; $p < 0.01$). Although there were 20 pulses, the low-

frequency (0.5 Hz) train generated negligible (nanomolar range) calcium summation at terminals (Figure 1B), whereas more than a few micromolars of calcium are needed to trigger endocytosis at calyces (Wu et al., 2009; Hosoi et al., 2009). Second, ΔC_m induced by the 20 pulse train was much higher, which may saturate the endocytosis capacity and thus slow down endocytosis (Sankaranarayanan and Ryan, 2000; Wu et al., 2005).

In the presence of ω -Agatoxin-IVA (200 nM), $depol_{10ms}$ induced a QICa (8.8 ± 0.4 pC, $n = 14$, Figure 4D; peak amplitude, 0.91 ± 0.06 nA) $\sim 45\%$ of control (19.5 ± 1.0 pC, $n = 12$, peak amplitude: 2.10 ± 0.11 nA; Figure 4C), and a ΔC_m (270 ± 17 fF, $n = 14$, Figure 4D) about 63% of control (427 ± 24 fF, $n = 12$, Figure 4C). The capacitance decay was too slow to fit with an exponential function (Figure 4D). The $Rate_{decay}$ was 16.3 ± 1.6 fF/s ($n = 14$, Figure 4D), much slower than control (36.7 ± 2.7 fF/s, $n = 12$, Figure 4C; $p < 0.01$). The slower $Rate_{decay}$ and endocytosis time course were not caused by the ΔC_m decrease because in the presence of ω -Agatoxin-IVA, a pair of $depol_{10ms}$ values at 5 s interval induced a ΔC_m (393 ± 27 fF, $n = 9$; Figure 4E) similar to that by a $depol_{10ms}$ in control ($p = 0.31$), but a $Rate_{decay}$ (18.7 ± 1.6 fF/s, $n = 9$; Figure 4E) much slower than that by a $depol_{10ms}$ in control ($p < 0.01$; Figure 4C). Thus, VDCCs, including P/Q-type, couple exocytosis to slow endocytosis.

Results shown thus far were from P7–P10 rats, in which calyces are not mature. In P13–P14 rats in which calyces are mostly mature (Schneppenburger and Forsythe, 2006), we repeated experiments similar to those shown in Figures 1A and 2C (upper) and Figures 3A and 3B and found similar results (Figure S2; Figure 2C, lower). Thus, our observations applied to both immature and mature calyces.

DISCUSSION

We found that fusion did not generate calcium currents or influx at the plasma membrane (Figure 1), that Flower was localized at synaptic vesicles of rat brain and calyces (Figure 2), and that the current charge of VDCCs at the plasma membrane, including P/Q-type, determined the rate of rapid and slow endocytosis (Figures 3 and 4). These results suggest that fused vesicle

membrane and proteins, including Flower, could not generate calcium currents to initiate endocytosis. Instead, plasma membrane VDCCs, including P/Q-type, couple exocytosis to two widely observed forms of endocytosis: rapid and slow endocytosis.

Our findings are different from a study showing that vesicular VDCCs inserted at the plasma membrane via exocytosis are required for endocytosis in sea urchin eggs (Smith et al., 2000). The difference is likely due to the significant differences of the two systems. Calyces are nerve terminals containing small (~50 nm) vesicles with endocytosis lasting for less than tens of seconds (Figures 3 and 4), whereas sea urchin eggs contain very large (~1 μm) vesicles with endocytosis lasting for ~15–30 min.

Our findings are different from the study of *Drosophila* synapses (Yao et al., 2009). Although the synapse specificity could provide an explanation, the Flower hypothesis is not well established as discussed below. At *Drosophila* nerve terminals, vesicular Flower insertion into the plasma membrane via exocytosis increases the $[\text{Ca}^{2+}]_i$ at the submicromolar range after prolonged tetanic stimulation (Yao et al., 2009), whereas endocytosis at nerve terminals is initiated by more than a few micromolars of calcium (Wu et al., 2009; Hosoi et al., 2009; Beutner et al., 2001). Furthermore, the time course of the calcium rise via Flower channels is in the order of 10–60 min (Yao et al., 2009), which was too slow to initiate endocytosis that typically lasts for a few to tens of seconds at many synapses, including *Drosophila* synapses (Poskanzer et al., 2003; Wu et al., 2007). Such slow calcium permeation might explain why we did not detect calcium currents or influx within a few seconds after exocytosis.

Endocytosis quantified with FM1-43 uptake at one time point after prolonged (1 min) high potassium stimulation was reduced by ~41% in Flower mutants (Yao et al., 2009). Compared to a complete block of endocytosis by calcium buffers (Wu et al., 2009; Hosoi et al., 2009; Yamashita et al., 2010), such a small defect suggests that Flower is not essential in initiating endocytosis. Furthermore, whether Flower plays a role under physiological conditions is unclear because endocytosis after brief physiological depolarization was not quantified in Flower mutants (Yao et al., 2009).

Although the Flower hypothesis is not well established, the present work firmly established a critical role of plasma membrane VDCCs, but not vesicular channels, in initiating and speeding up endocytosis at calyces. Whether our results apply to other synapses has not been tested. However, except that the calyx is much larger than conventional boutons, it is similar to other nerve terminals in many aspects, such as the active zone size, the vesicle number per active zone, short-term plasticity, transmitter release properties, calcium currents, and endocytosis time course (Schneggenburger and Forsythe, 2006; Wu et al., 2007; Xu et al., 2007). Our findings are, therefore, likely to apply to other synapses. Considering that VDCCs control exocytosis in all chemical synapses and many non-neuronal secretory cells, we suggest that VDCC-mediated exo-endocytosis coupling is a common mechanism at synapses and many non-neuronal secretory cells. Consistent with this suggestion, calcium influx was found to regulate endocytosis at many synapses and non-neuronal secretory cells (Ceccarelli

and Hurlbut, 1980; Marks and McMahon, 1998; Cousin and Robinson, 1998; Gad et al., 1998; Sankaranarayanan and Ryan, 2001; Balaji et al., 2008; Neves et al., 2001; Clayton et al., 2007; Artalejo et al., 1995; He et al., 2008).

EXPERIMENTAL PROCEDURES

Slice preparation and capacitance recordings were described previously by Sun and Wu (2001) and Sun et al. (2004). Briefly, parasagittal brain stem slices (200 μm thick) containing the medial nucleus of the trapezoid body were prepared from Wistar rats using a vibratome. Unless mentioned, rat age was 7–10 days old. All animal protocols followed the guidelines of the National Institutes of Health, USA. Whole-cell capacitance measurements were made with the EPC-10 amplifier together with the software lock-in amplifier (PULSE, HEKA, Lambrecht, Germany) that implements Lindau-Neher's technique. The frequency of the sinusoidal stimulus was 1,000 Hz, and the peak-to-peak voltage of the sine wave was $\leq 60\text{mV}$. We pharmacologically isolated presynaptic I_{Ca} with a bath solution (~22°C–24°C) containing 105 mM NaCl, 2.0 mM TEA-Cl, 2.5 mM KCl, 1 mM MgCl₂, 2 mM CaCl₂, 25 mM NaHCO₃, 1.25 mM NaH₂PO₄, 25 mM glucose, 0.4 mM ascorbic acid, 3 mM *myo*-inositol, 2 mM sodium pyruvate, 0.001 mM tetrodotoxin (TTX), 0.1 mM 3,4-diaminopyridine (pH 7.4) when bubbled with 95% O₂ and 5% CO₂. When ω -Agatoxin-IVA was applied to the bath, cytochrome *c* (0.1 mg/ml) was also included. The presynaptic pipette (3.5–5 M Ω) solution contained 125 mM Cs-gluconate, 20 mM CsCl, 4 mM MgATP, 10 mM Na₂-phosphocreatine, 0.3 mM GTP, 10 mM HEPES, 0.05 mM BAPTA (pH 7.2) adjusted with CsOH.

The statistical test was a *t* test. Means are presented as \pm SE. Rate_{decay} was measured from 0.3 s to 3 or 6 s after stimulation (Wu et al., 2009). The $[\text{Ca}^{2+}]_i$ was measured with fura-2 (50 μM , replacing BAPTA in the pipette), the Olympus upright epifluorescence microscope (BX51WI, LumplanFI 40 \times , n.a. 0.8), a polychromatic illumination system (T.I.L.L. Photonics, Munich), and a photodiode for fluorescence recording (Xu and Wu, 2005).

Cytosol and synaptic vesicle proteins were prepared by sucrose gradient fractionation (Yao et al., 2009). P8 rat brain was homogenized and centrifuged to yield a crude synaptosomal pellet, which was then resuspended in 100 mM NaCl, 20 mM HEPES (pH 7.4) solution and loaded onto sucrose step gradient (0.2–0.4 M) for centrifugation (20,000 $\times g$, 5 hr). Synaptic vesicles were collected from 0.2 to 0.4 M sucrose interface. The proteins were quantified and subject to SDS-PAGE for western blot assay (Invitrogen).

For western blot, P8 rat brain was homogenized and centrifuged to yield supernatants, which were quantified and loaded to SDS-PAGE for immunoblotting using a custom-made rabbit antibody against Flower (1:50; GenScript) and a mouse antibody against VAMP2 (1:10,000; Synaptic Systems).

For immunohistochemistry, rats were anesthetized with Nembutal, fixed by paraformaldehyde, and infiltrated with 30% sucrose. OCT (Electron Microscopy Sciences)-embedded rat brain was sectioned using a cryostat (Leica CM3050S) at 30 μm thickness. The calyces were recognized by a guinea pig antibody against vGluT₁ (1:5,000; Millipore), and Flower proteins at calyces were identified using the rabbit antibody against Flower (1:50; GenScript). For the peptide-blocking experiment (Figure 2C, middle), Flower antibody was preincubated with the Flower peptide (Figure 2A, red) overnight at 4°C. Dylight-488 donkey anti-rabbit and rhodamine-conjugated donkey anti-guinea pig antibodies (1:100; Jackson ImmunoResearch Laboratories) were used as secondary antibodies. Images were collected by Zeiss LSM510 confocal microscopy (40 \times , N/A 1.3).

SUPPLEMENTAL INFORMATION

Supplemental Information includes two figures and can be found with this article online at doi:10.1016/j.celrep.2012.04.011.

LICENSING INFORMATION

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