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Mechanisms Involved in Calcium-Dependent Exocytosis^a

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Although a great deal is known concerning the membrane events controlling Ca²⁺ influx and metabolism, the events triggered by Ca²⁺ that lead to exocytosis in neurons and other cells are poorly understood. The purpose of this article is to consider some of the processes that control exocytosis, with special emphasis on the bovine adrenal chromaffin cell as a model system. Chromaffin cells are excitable cells and, like sympathetic nerves, are derived from the neural crest. Although chromaffin cells do not contain long processes or nerve terminals, secretion from chromaffin cells is stimulated by Ca²⁺ influx and inhibited by the clostridial neurotoxins (see below), which block exocytosis from the nerve terminals of peripheral and central nervous system neurons. Thus, it is likely that the secretory pathway in chromaffin cells is similar to that in neurons. Chromaffin cell secretory vesicles (chromaffin granules) are large, dense core vesicles that contain proteins, peptides, ATP, and the catechol-amines epinephrine and norepinephrine. Ca²⁺ influx is stimulated in bovine chromaffin cells by activation of the nicotinic receptor/channel complex and voltage-sensitive Ca²⁺ channels.

EXOCYTOSIS AND GTP-BINDING PROTEINS

Exocytosis is the last fusion event in the repeated sequence of vesicle budding and fusion that characterizes the passage of secretory proteins from the endoplasmic reticulum to the cell exterior (for review see ref. 1). Secretory vesicles leaving the *trans* golgi network are targeted to different locations. Some vesicles are targeted to lysosomes. Some go quickly to the plasma membrane where exocytosis occurs in a seemingly unregulated manner (constitutive secretion). Other vesicles fuse with the plasma membrane in a highly regulated manner. In the neuron, regulated exocytosis can occur meters away from the cell body where synthesis of the secretory vesicle membrane and intravesicular proteins occurs. Secretory vesicles involved in synaptic transmission journey to the nerve terminal by way of a microtubule-based transport system. After fusion with the plasma membrane, vesicle membrane recycles by way of endocytosis. In the case of small synaptic vesicles that store nonprotein neurotransmitters (e.g., acetylcholine), the recycled vesicles are reused locally for exocytosis.

In endocrine cells and nerve terminals, the events leading to fusion of the

^aThis work was supported by NIH Grants RO1 DK27959 and PO1 HL18575. Part of the work presented was done when R. W. Holz was an Established Investigator of the American Heart Association.

secretory vesicle with the plasma membrane are highly regulated. In most cases a rise in cytosolic Ca^{2+} from $0.1~\mu M$ to micromolar or higher concentrations is the primary trigger for exocytosis. One of the major challenges in the study of the biochemistry of exocytosis is to distinguish those processes that are intimately involved in regulating exocytosis from the many processes activated by Ca^{2+} . Although there are many possibilities (e.g., protein phosphorylation, cytoskeletal changes, changes in lipid metabolism: see below), at this time we do not know the critical events activated by Ca^{2+} . An alternative approach is to consider whether regulated exocytosis in either endocrine cells or in nerve terminals has elements in common with the multiple fusion events occurring earlier in the biosynthetic pathway of secretory proteins.

The genetic analysis of protein secretion in yeast led to the identification and purification of SEC4p, which is necessary for fusion of post-Golgi vesicles with the plasma membrane, 4.5 and YPT1p, which is necessary for protein transfer from endoplasmic reticulum (ER) to Golgi.6 Both SEC4p and YPT1p are 23 kDa, GTP-binding proteins homologous to the *ras* family of GTP-binding proteins. They share 45% homology with each other. GTP-binding proteins are involved in the transfer and fusion of vesicles formed from donor cisternae (ER or Golgi compartments) to subsequent cisternae in the eukaryotic biosynthetic pathway. A nonhydrolyzable analogue of GTP (GTPγS) prevents transfer of protein from ER to Golgi, 7 and between Golgi compartments, 8 and induces the accumulation of coated vesicles in the Golgi region 9 and the accumulation of tubular-vesicular membranes near the ER. It has been postulated that GTP binding to vesicular protein and subsequent GTP hydrolysis are required for proper targeting and transfer of budding vesicles. 10 An epitope in common with YPT1p is associated with mammalian Golgi.6

Do low molecular weight, GTP-binding proteins generally play an important role in targeting secretory vesicles or controlling fusion in regulated exocytosis? At this time the answer is unclear. Both chromaffin granules, the secretory vesicles in adrenal chromaffin cells,^{2,11} and bovine brain synaptic vesicles^{12,13} contain numerous GTP-binding proteins between 20 and 30 kDa. FIGURE 1A demonstrates the multitude of these proteins that are found on secretory vesicles. One can distinguish at least six proteins on bovine chromaffin granule membranes and on bovine brain synaptic vesicles. (These brain synaptic vesicles display active glutamate uptake. ^{14,15}) GTP-binding proteins from chromaffin granules and synaptic vesicles have identical mobilities in SDS-PAGE. A GTP-binding protein of approximately 45 kDa can also be detected that probably corresponds to either α_0 or α_i , the pertussis toxin substrates for ADP-ribosylation that have been previously demonstrated on chromaffin granules. A quantitative analysis of GTP-binding sites on chromaffin granules (Fig. 1B) indicates that there are approximately 60 GTP-binding sites per chromaffin granule, which accounts for approximately 5% of the total granule membrane protein (see FIG. 1B legend). Nonhydrolyzable guanine nucleotides have a number of effects on secretion in permeabilized secretory cells. GTPyS inhibits Ca²⁺-dependent secretion from electropermeabilized bovine chromaffin cells when added together with Ca²⁺¹⁷ and from digitonin-permeabilized bovine chromaffin cells when a low concentration of GTP_YS is incubated with permeabilized cells prior to stimulation with Ca²⁺ (Fig. 2). These effects are consistent with the effects of GTP_γS to inhibit transfer and fusion of vesicles in earlier steps in the biosynthetic pathway of secretory proteins. The effects, however, depend upon cell type, method of plasma membrane permeabilization and experimental design. For example, nonhydrolyzable guanine nucleotides stimulate Ca2+-dependent secretion in electropermeabilized chicken chromaffin cells¹⁷ and in bovine chromaffin cells permeabilized by α-toxin.¹⁸ Guanine nucleotides and Ca²⁺ are required for secretion from streptolysin-O- or digitoninpermeabilized mast cells in the presence of high concentrations of Cl⁻. ^{19,20} When the

Cl⁻ concentration, however, is reduced¹⁹ or when protein kinase C is activated,²⁰ either GTP_γS or Ca²⁺ alone stimulates submaximal secretion. GTP_γS also stimulates Ca²⁺-independent secretion from digitonin-permeabilized bovine chromaffin cells,^{21,22} and PC12 cells,²³ and from mast cells using whole cell recording with the patch clamp.²⁴⁻²⁶ The multitude of effects of nonhydrolyzable guanine nucleotides on secretion is not surprising inasmuch as guanine nucleotides can activate a host of

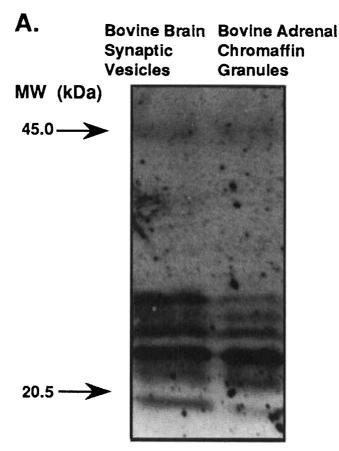


FIGURE 1A. GTP-binding proteins associated with bovine chromaffin granules and bovine brain glutamatergic synaptic vesicles. Chromaffin granules were purified from fresh bovine adrenal medulla. Adrenal glands were homogenized in 0.29 M sucrose, 10 mM Hepes (pH 7.1), 1 mM Na₂EDTA, 1 mM PMSF, and chromaffin granules purified using a discontinuous sucrose gradient according to Smith and Winkler, ⁷³ except that the large granule fraction was layered over a 1.7 M sucrose solution. The gradient was centrifuged at 146,000 × g_{ave} × 70 min in a Beckman 50.2 Ti rotor. The chromaffin granules were collected in the pellet, lysed in 10 mM Hepes (pH 7.1), and washed twice and stored at -70. Synaptic vesicles were purified from bovine brain according to the methods of Ueda and colleagues^{14,15} with an additional gel filtration step using Sephacryl 1000. The vesicle fraction concentrated glutamate in an ATP-dependent manner. The molecular masses of the GTP-binding proteins are 20.1 kDa, 22.4 kDa, 23.0 kDa (dense band), 24.6 kDa, 26.5 kDa, 27.5 kDa, and 45 kDa.

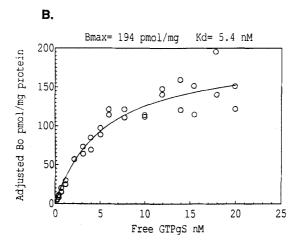


FIGURE 1B. Specific GTPγS binding to chromaffin granule membranes. Purified chromaffin granules (2 μg/mL) were incubated with various concentrations of nonradioactive GTPγS and 0.235 nM [35 S]GTPγS (0.23 μC₁/mL) for 90 min at 30°C. The suspension was passed through a nitrocellulose filter. The filter was washed extensively and the radioactivity determined. The data were analyzed by a nonlinear, least squares algorithm. Assuming a chromaffin granule diameter of 2800 angstroms, 17 μL chromaffin granule volume/mg membrane protein and maximal binding of 194 pmoles/mg membrane protein, one calculates 78 GTPγS binding sites/chromaffin granule. A similar experiment using a different preparation of granules yielded 130 pmoles GTPγS binding/mg membrane protein and 52 GTPγS binding sites/chromaffin granule. Approximately 4–6% of the total chromaffin granule membrane protein is GTP-binding protein, assuming an average molecular mass of 30 kDa for GTP-binding proteins.

enzymes either directly or indirectly (e.g., adenylate cyclase, phospholipase C, phospholipase A_2 , protein kinase C) that could influence the secretory response. Thus, it is unclear which, if any, of the effects of guanine nucleotides on secretion requires GTP-binding proteins on secretory vesicles. Furthermore, it is possible that the GTP-binding proteins associated with secretory vesicles are not directly involved in secretion but with other functions of the vesicles related to biosynthesis of the membrane²⁷ or recycling after exocytosis.

THE ROLE OF ATP IN SECRETION: PRIMING OF SECRETION, PROTEIN PHOSPHORYLATION, AND MAINTENANCE OF POLYPHOSPHOINOSITIDES

Experiments with permeabilized chromaffin cells by Baker and Knight^{28,29} conclusively demonstrated a role for ATP hydrolysis in the intracellular pathway for exocytosis. Synaptic transmission occurs within less than a millisecond after invasion of the terminal by the action potential and the opening of voltage-sensitive Ca²⁺ channels. Because this is insufficient time for a cascade of biochemical reactions,³⁰ the exocytotic machinary must be poised to respond rapidly to the rise in Ca²⁺. In experiments with the patch clamp in which incorporation of the granule membrane into the plasma membrane is measured by increases in plasma-membrane capacitance, depolarization-induced secretion from chromaffin cells occurs within the time

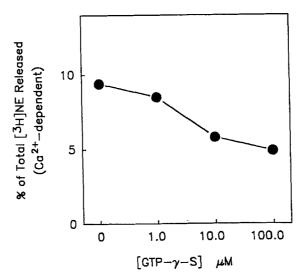


FIGURE 2. GTP γ S inhibits secretion from digitonin-permeabilized chromaffin cells. [3 H]nor-epinephrine-labeled chromaffin cells were permeabilized by incubation in KGEP [139 mM potassium glutamate, 5 mM EGTA, 20 mM PIPES (pH 6.6), 1 mM MgCl $_2$, and 5 mg/mL BSA] containing 2 mM MgATP, 20 μ M digitonin, and various concentrations of GTP γ S. After 4 min, the solution was replaced with digitonin-free KGEP without MgATP and GTP γ S, and with and without 10 μ M free Ca $^{2+}$. Ca $^{2+}$ -dependent secretion was determined after 2 min. Secretion in the absence of Ca $^{2+}$ was 1.6–2.4%. GTP γ S at all concentrations released less than 1% of the catecholamine during the incubation with digitonin. There were 3 wells/group.

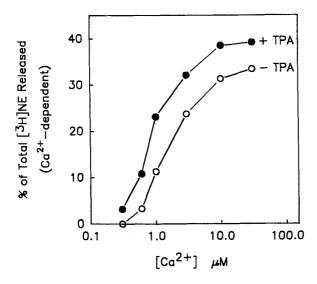
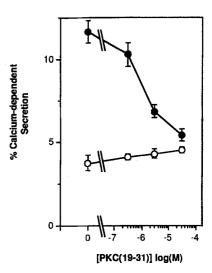


FIGURE 3. The phorbol ester TPA enhances secretion from digitonin-permeabilized chromaffin cells. [3H]norepinephrine-labeled cells were incubated for 30 min in MEM with or without 20 nM TPA. The cells were permeabilized in KGEP containing 2 mM MgATP with indicated [Ca²⁺]. Ca²⁺-dependent secretion was determined after 15 min. Secretion in the absence of Ca2+ was 3.4-3.6%. There were 3 wells/ group.

FIGURE 4. The protein kinase C inhibitor peptide PKC(19-31) inhibits Ca²⁺-dependent secretion in TPA-treated but not in TPA-untreated chromaffin cells. [3H]norepinephrine-labeled cells were incubated for 15 min with (filled circles) or without (unfilled circles) 60 nM TPA in physiological salt solution. Cells were permeabilized for 6 min in Ca2+-free potassium glutamate solution (KGEP) containing 20 µM digitonin in the presence or absence of increasing concentrations of PKC(19-31). The solution was replaced with KGEP with or without 1 µM Ca2+ in the continuing presence or absence of PKC(19-31). Ca²⁺dependent secretion was measured after 5 min. Release in the absence of Ca²⁺ was 3.1-4.4%. There were 3 wells/group.⁴²



resolution of the experiments (several hundred milliseconds).³¹ In experiments in which release of catecholamine is detected biochemically, the maximal rate of secretion occurs within the first 1–2 min and then slows³²; secretion can be reproducibly detected within 10 seconds.³³ We found that the rapid initial rate of catecholamine secretion requires prior incubation with ATP.³² Although the time scale of the experiments is orders of magnitude slower than the events following Ca²⁺ entry in the nerve terminal, the experiments nevertheless demonstrate that ATP acts before Ca²⁺ to prime the cell to respond to Ca²⁺. A similar conclusion can be drawn from experiments with mast cells.³⁴

ATP is required for protein phosphorylation, and there is abundant evidence that protein phosphorylation modulates exocytosis. In the squid giant synapse³⁵ and mammalian synaptosomes³⁶ Ca/calmodulin kinase II enhances secretion, perhaps by

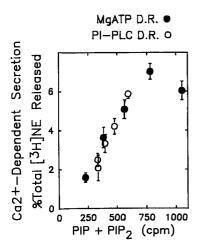
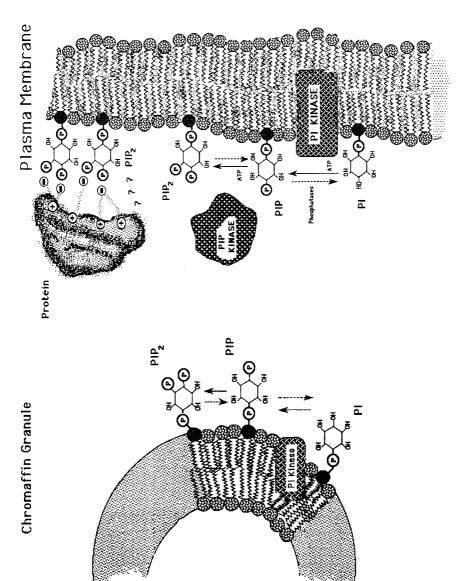


FIGURE 5. The relationship between the levels of the polyphosphoinositides and Ca2+-dependent secretion in digitonin-permeabilized chromaffin cells. The polyphosphoinositides were varied in digitoninpermeabilized cells by varying the concentrations of MgATP or by varying the concentrations of a PtdInsspecific phospholipase C from Bacillus thuringiensis. [3H]Inositol-labeled cells were permeabilized for 8 min in Ca2+-free potassium glutamate solution (KGEP) containing either various amounts of MgATP or 2 mM MgATP and various amounts of bacterial phospholipase C. The levels of [3H]inositollabeled PtdIns, PtdInsP, and PtdInsP2 were determined after an additional 3 min incubation in 10 µM Ca²⁺ in the continuing presence of MgATP. Bacterial phospholpase C was not present in the last incubation. Ca²⁺-dependent secretion from [³H]norepinephrine-labeled cells was determined in parallel experiments at 6 min. For details see Fig. 5 in ref. 45.





phosphorylation of synapsin I. In many secretory cells including chromaffin cells, activation of protein kinase C increases the ability of cells to respond to submicromolar and micromolar Ca²⁺. ³⁷⁻³⁹ FIGURE 3 demonstrates effects of the protein kinase C activator TPA (12-O-tetradecanoylphorbol acetate) on the Ca²⁺ dose-response relation for secretion from permeabilized cells. Because cellular protein kinase C can be activated by Ca²⁺ in the absence of exogenous activators, ^{33,40} an important issue is whether protein kinase C is required for secretion and is responsible for the ATP dependency. We have recently addressed this issue in permeabilized chromaffin cells using a specific inhibitor of protein kinase C. The synthetic peptide PKC(19-31), which is derived from the pseudosubstrate sequence of protein kinase C, is a potent inhibitor of protein kinase C in vitro⁴¹ and in situ within permeabilized cells.⁴² Although it inhibited phorbol ester-induced secretion in the presence of Ca²⁺ almost completely, it had no effect on Ca2+-dependent secretion in the absence of phorbol ester (Fig. 4). Thus, it is unlikely that the ATP-dependency for secretion reflects the necessary involvement of protein kinase C in the secretory pathway. Similarly, it is unlikely that calmodulin-dependent protein kinase or other calmodulin-dependent processes play a necessary role in secretion from permeabilized cells. Relatively high concentrations of the calmodulin antagonists trifluoperazine and calmidazolium have either partial effects in electropermeabilized chromaffin cells^{29,43} or virtually no effects in digitonin-permeabilized chromaffin cells (Holz, unpublished observations). Furthermore, an antical modulin antibody (Holz, unpublished observations) and a calmodulin-binding peptide⁴² have no effect on secretion from digitonin-permeabilized chromaffin cells. The ability of an anticalmodulin antibody to inhibit secretion when injected into intact cells⁴⁴ may reflect a role for calmodulin in the events leading to a rise in cytosolic Ca²⁺.

These data led us to consider other reactions in which ATP may be involved. We found that the maintenance of the polyphosphoinositides by PI and PIP kinases occurred over the same MgATP concentration range as that necessary to maintain exocytosis. 45 We also found that in the presence of MgATP we could manipulate the levels of the polyphosphoinositides by a bacterial phospholipase C that hydrolyzed their precursor, phosphatidylinositol. The polyphosphoinositides declined because of the action of endogenous lipid phosphatases in the absence of continued synthesis. The relationship between Ca²⁺-dependent secretion and the polyphosphoinositides was the same whether the lipids were manipulated by varying the MgATP concentration or by varying the concentration of the bacterial phospholipase C (FIGURES 5 and 6). The inhibitory effects of the exogenous phospholipase C could not be explained by the products of the bacterial phospholipase C reaction (IP₁ and diacylglycerol) or the loss of products from the endogenous phospholipase C (due to the loss of the polyphosphoinositide substrates). The simplest and most compelling interpretation is that the polyphosphoinositides are necessary in the secretory pathway. Thus, ATP

FIGURE 6. Polyphosphoinositide metabolism in chromaffin cells. The polyphosphoinositides are synthesized from phosphatidylinositol by PI and PIP kinases. In resting cells, PIP₂ phosphatase and PIP phosphatase are responsible for degrading the polyphosphoinositides. Phospholipase C activated by receptor-mediated mechanisms or by a rise in cytosolic Ca²⁺⁷⁵ breaks down PIP₂ to IP₃ and diacylglycerol (not shown). Experiments in which the polyphosphoinositides are decreased by ATP or by hydrolysis of PI with a bacterial phospholipase C demonstrate a close correlation between the levels of the polyphosphoinositides and Ca²⁺-dependent secretion. Thus, reducing either substrate for PI kinase reduces the polyphosphoinositides and Ca²⁺-dependent secretion. The requirement for the polyphosphoinositides in secretion could reflect the importance of a protein that binds to and is regulated by these lipids.

is important not only for modulation of secretion by protein kinases, but also for maintenance of the polyphosphoinositides.

Polyphosphoinositides are synthesized both on the plasma membrane⁴⁷ and on the chromaffin granule membrane ⁴⁸⁻⁵⁰ (FIG. 6). The binding of a protein to PIP₂ (and/or PIP) on either membrane may regulate its function in secretion. There are at least two proteins, myosin I and gelsolin, that bind to the polyphosphoinositides and to actin. Myosin I, which causes actin-dependent organelle movement, has recently been demonstrated to bind to PIP₂.⁵¹ The ability of gelsolin to cause Ca²⁺-dependent severing of f-actin is inhibited by polyphosphoinositides⁵² (for review see ⁵³). Because chromaffin granules also interact with f-actin,⁵⁴⁻⁵⁶ it is possible that the PIP₂ or PIP localizes these proteins to secretory sites and controls the interaction of chromaffin granules with the cortical cytoskeleton and the plasma membrane. In addition, calpactin I, which interacts with acidic phospholipids including the polyphosphoinositides,⁵⁷ aggregates and under some conditions allows fusion of chromaffin granules *in vitro*.^{58,59} Indeed, recent reports suggest that calpactin is in intimate contact with chromaffin granules, and the plasma membrane in intact cells,⁶⁰ and can maintain secretion in digitonin-permeabilized chromaffin cells.^{61,62}

CLOSTRIDIAL NEUROTOXINS DIRECTLY INHIBIT THE EXOCYTOTIC PATHWAY

Botulinum and tetanus neurotoxins (clostridial neurotoxins) block neurotransmitter release from various neurons and are thought to interact with critical sites in the exocytotic pathway. 63 Botulinum type D neurotoxin interacts with bovine tissues and inhibits secretion from both intact and electropermeabilized bovine chromaffin cells.⁶⁴ Extracellular application of botulinum types A, B, or E neurotoxin or tetanus neurotoxin has no effect on intact bovine chromaffin cells because of the absence of receptors on the plasma membrane. The introduction of these toxins, however, into cells through a patch-clamp electrode⁶⁵ or by incubation with digitonin- or streptolysin O (SL-O)-permeabilized cells⁶⁶⁻⁶⁹ inhibits secretion. Experiments with permeabilized cells demonstrate that the light chain (50 kDa) of the dichain molecules (150 kDa) is responsible for the inhibition of secretion. ^{67,68,70} The ability of these classical inhibitors of synaptic transmission to inhibit exocytosis from chromaffin cells indicates that there are common elements in the secretory pathway in neurons and chromaffin cells. The toxins decrease the maximal extent of secretion and not the sensitivity to Ca²⁺. 66.67 Interestingly, the toxins do not all act alike. Both tetanus toxin and botulinum B neurotoxins completely inhibit secretion, whereas botulinum A neurotoxin, even at high concentrations, inhibits secretion by no more than 70 percent. Thus, the toxins may interact with different sites in the secretory pathway. At the intact neuromuscular junction, the effects of tetanus and botulinum B neurotoxins can also be distinguished from those of botulinum A neurotoxin. Tetanus and botulinum B neurotoxins reduce evoked miniature end-plate potentials (MEPPS) and desynchronize those MEPPS that are observed.⁷¹ Botulinum A neurotoxin also reduces evoked MEPPS but without desynchronizing the residual MEPPS. Key issues for future investigation are the sites of interaction of the light chains of these toxins within nerve terminals and chromaffin cells and the possibility that they may act through an unknown enzymatic activity. The ADP-ribosylating activity of neurotoxin preparations is likely to be caused by contaminating ADPribosyl transferase activity (C-3 toxin).⁷²

ACKNOWLEDGMENTS

We are grateful to Dr. T. Ueda and Dr. J. Tabb (Mental Health Research Institute and the Department of Pharmacology, University of Michigan) for providing us with purified bovine brain synaptic vesicles. We thank Peter Wick (Department of Pharmacology, University of Michigan Medical School) for preparing FIGURE 6.

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