SYMPOSIUM REPORT

Phospholipases and fatty acid signalling in exocytosis

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Vesicle fusion is a ubiquitous biological process involved in general membrane trafficking and a variety of specialized events, for example release of neurotransmitters and hormones, sperm acrosome exocytosis, plasma membrane repair and neurite outgrowth. Many vesicle fusion events have long been known to be activated by phospholipases and products of their activity, such as polyunsaturated arachidonic acid. Polyunsaturated fatty acids (PUFAs) have been proposed to have a number of multiple effectors, including ion channels and the cytoskeleton, but the precise mechanism of PUFA action is still unclear. It was recently reported that omega-3 and omega-6 PUFAs can act on syntaxin, a plasma membrane protein directly involved in vesicle fusion. In this review, we will discuss the role of this new mode of PUFA action in exocytosis.

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Exocytosis is a cellular event where cargo-laden vesicles fuse with the plasma membrane. Vesicle fusion is involved in various biological processes including cell growth, plasma membrane repair, axonal branching, recycling of plasma membrane transporters, and release of soluble signalling molecules such as neurotransmitters or hormones into the extracellular space. It is well established that phospholipases A_2 , C and D, which metabolize membrane phospholipids, are somehow involved in vesicle fusion (Vitale *et al.* 2001; Brown *et al.* 2003; Rossetto *et al.* 2006). One class of phospholipid components, long-chain polyunsaturated fatty acids (PUFAs), has emerged as being particularly important for exocytosis. However, the mechanisms of action of PUFAs in the regulation of vesicle fusion are not well understood. They have been proposed to modulate ion channel function and to perturb cytoskeleton (Honore *et al.* 1994; Lesage *et al.* 2000; Mignen & Shuttleworth, 2000; Neco *et al.* 2003).

Recently, PUFAs were shown to act on syntaxin, a plasma membrane protein directly involved in fusion of vesicles with the plasma membrane (Rickman & Davletov, 2005; Darios & Davletov, 2006; Connell *et al.* 2007). Syntaxin belongs to the soluble NSF-attachment receptor (SNARE) protein family responsible for intracellular membrane fusion throughout the cell. A prototypical set of fusion proteins involved in neurotransmitter release consists of the plasma membrane syntaxin 1 together with

SNAP-25 (synaptosome-associated protein of 25 kDa) and vesicular protein synaptobrevin (Rizo & Sudhof, 2002). The three proteins form a slightly twisted four-helical bundle between two approaching membranes (Sutton*et al.* 1998), probably initiating the fusion event. Since the action of PUFAs on ion channels and cytoskeleton has been discussed elsewhere (Nakamura *et al.* 2001; Neco *et al.* 2003), we will focus here on the role of PUFA-releasing enzymes and fatty acid signalling to promote activation of SNARE proteins in vesicle fusion.

PUFAs and neuronal function

The membrane bilayers in which SNARE proteins reside are composed of several phospholipid species as well as sphingolipids and cholesterol. A typical phospholipid structure and the sites of phospholipase action are shown in Fig. 1*A*. Phospholipids often contain PUFAs such as arachidonic or docosahexaenoic acids in the *sn*-2 position, whereas the *sn*-1 position is normally occupied by saturated fatty acids.

Two types of PUFAs, omega-6 and omega-3, are essential for vertebrates as they cannot be formed *de novo* and need to be ingested. Omega-6 arachidonic and omega-3 docosahexaenoic acid are major building blocks of cellular membranes (Svennerholm, 1968). PUFAs possess favourable biophysical properties such as flexibility and solubility, which promote membrane fluidity. In addition, when released through phospholipase action, they directly take part in regulation of many cellular processes and are also converted into eicosanoids, including prostaglandins

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(Brash, 2001). Of note, some of these metabolites were reported to activate vesicle fusion, whereas others inhibit it (Bazan *et al.* 2002).

The importance of PUFAs for neuronal function is well known (Wainwright, 2002). Mutations in PUFA-related enzymes cause mental retardation in humans (Meloni*et al.* 2002), and diets deficient in essential PUFAs are associated with deficits in infant brain function (Wainwright, 2002). In addition, mutations in an enzyme involved in PUFA production cause neuronal impairment in the *C. elegans* model organism, which can be rescued by external application of arachidonic or docosahexaenoic acid (Lesa *et al.* 2003). There was some uncertainty regarding the effect of arachidonic acid on catecholamine secretion

Figure 1. Alternative pathways leading to the release of mobile polyunsaturated fatty acids from the lipid bilayer into the cytosol

A, a typical phospholipid consists of saturated palmitic acid (C16), or more commonly stearic acid (C18), attached to the *sn*-1 position of the glycerol backbone, while unsaturated omega-6 arachidonic acid (C20 : 4) or omega-3 docosahexaenoic acid (C22 : 6) occupy its *sn*-2 position. The glycerol backbone also carries a hydrophilic headgroup shown here as a phosphorylated inositol; other common headgroups are choline, ethanolamine and serine. Phospholipase sites of action are shown by grey arrows. Note the presence of four and six unsaturated double bonds in arachidonic acid and docosahexaenoic acid, respectively. *B*, hydrolysis of the phospholipid molecule by phospholipase A₂ (PLA₂) produces lysophospholipid which is preferentially retained in the membrane and more soluble arachidonic acid (red hairpin) which, following diffusion into the cytosol, is either metabolized into different eicosanoid products or incorporated back in the *sn*-2 position of phospholipids (not shown). Phospholipase C (PLC) acts on the lipid bilayer to release soluble headgroup (red circles) leaving diacylglycerol (DAG) in the membrane. Further action of a DAG lipase results in the release of PUFAs into the cytosol. Monoacylated glycerol remains in the membrane due to straight saturated stearic or palmitic fatty acids. Phospholipase D (PLD) releases the headgroup (preferentially choline) of the phospholipid molecule producing hydrophobic phosphatidic acid which remains in membrane. Phosphatidic acid can in turn be metabolized into DAG by phospatidic acid (PA) hydrolase.

(Frye & Holz, 1984; Morgan & Burgoyne, 1990) but a recent study demonstrated arachidonic acid-induced up-regulation of secretion in both permeabilized and intact cell models (Latham *et al.* 2007). These data together suggest that PUFAs or their metabolites are essential for exocytosis. Interestingly, PUFA-rich diets affect expression of only few genes; amongst them is the syntaxin-binding protein Munc18, suggesting a possible link with SNARE proteins (Barcelo-Coblijn *et al.* 2003).

PUFA-releasing enzymes and exocytosis

Phospholipase A_2 s (PL A_2 s) are a group of enzymes which catalyse the breakdown of phospholipids. They release fatty acids from the *sn*-2 position of the phospholipid molecule (Fig. 1*A*) and are postulated to be involved in vesicle fusion in many biological systems, including neurotransmitter release, insulin secretion, sperm acrosome exocytosis and neurite outgrowth (Freeman *et al.* 1990; Wolf *et al.* 1991; Nakamura, 1993; Roldan & Fragio, 1993; Tsukada *et al.* 1994; Almeida *et al.* 1999; Brown *et al.* 2003; Juhl *et al.* 2003). PLA₂-mediated phospholipid hydrolysis generates free unsaturated fatty acids and lysophospholipids (Fig. 1*B*). Unsaturated fatty acids have intrinsically high rates of dissociation from membranes and, in contrast to lysophospholipids, can diffuse into the cytosol to intracellular sites of action without any transporter molecules (Tanford, 1973; Balsinde *et al.* 2002). It has been shown that PLA_2 acts to 'prime' fusion machinery on the plasma membrane, suggesting up-regulation of SNAREs or SNARE-associated molecules (Karli *et al.* 1990). Interestingly, this priming of vesicle fusion is achieved through the production of arachidonic acid and not lysophospholipid (Karli *et al.* 1990).

Mounting evidence also implicates phospholipase C (PLC) in exocytosis (Hammond *et al.* 2006). PLC releases the soluble headgroup from membrane phospholipids, leaving diacylglycerol (DAG) anchored in the membrane by its two hydrophobic carbon chains (Fig. 1*B*). The PLC inhibitor U73122 potently blocks synaptic vesicle exocytosis in many systems, even in the case of the most powerful secretagogue known – latrotoxin from the black widow spider (Davletov *et al.* 1998). The presence of DAG is necessary for activation of protein kinase C and Munc13, both of which are important positive regulators of vesicle fusion (Rhee *et al.* 2002). DAG can be further metabolized by DAG lipase, releasing arachidonic acid from the phospholipid *sn*-2 position and leaving monoacylated glycerol in the lipid bilayer (Brash, 2001). Interestingly, a DAG lipase inhibitor can block both neurite outgrowth and insulin release (Meiri *et al.* 1998; Guenifi *et al.* 2001). Recently it has been shown that a DAG lipase-like protein is obligatory for neurotransmitter release in the*Drosophila* genetic model (Huang *et al.* 2006). Genetic tests revealed a synergistic interaction between syntaxin and this DAG lipase-like protein, suggesting that the latter molecule could be directly involved in activation of SNARE function in synaptic vesicle fusion (Huang *et al.* 2006).

Finally, phospholipase D (PLD) has also been involved in the induction of vesicle fusion, at a late step of exocytosis (Vitale *et al.* 2001). PLD hydrolyses phosphatidylcholine into the choline headgroup and phosphatidic acid, which can in turn be hydrolysed into DAG by phosphatidic acid phosphohydrolase (Fig. 1*B*). PLD is required for the formation of certain SNARE complexes in yeast (Coluccio *et al.* 2004). It appears that in the case of PLD, early metabolite – phosphatidic acid – may play a role in exocytosis (Metz & Dunlop, 1990; Humeau *et al.* 2001).

Action of PUFAs on syntaxin

Snake neurotoxins that disrupt synaptic function contain PLA2s (Rossetto *et al.* 2006) and their secretogogue action appears to be blocked by application of a SNARE-cleaving botulinum toxin suggesting that PLA_2 s act upstream of SNARE assembly (Tse *et al.* 1982; Prasarnpun *et al.* 2004). We recently reported immediate effects of arachidonic acid on the SNARE fusion machinery (Rickman & Davletov, 2005; Darios & Davletov, 2006; Connell *et al.* 2007).

Syntaxin present in a 'closed conformation' is a target for PUFAs. They can induce a conformational change, leading to an 'open' conformation of the syntaxin and allowing interaction with SNAP-25. Upon this interaction, SNAP-25 acquires its coiled α-helical structure. A subsequent association of the heterodimer with vesicular synaptobrevin leads to formation of the ternary SNARE complex which promotes vesicle fusion.

Addition of arachidonic acid to synaptic membranes or membrane treatment with PLA_2s was sufficient to potentiate SNARE complex formation (Rickman & Davletov, 2005; Darios & Davletov, 2006). It is notable that prostaglandins are ineffective in syntaxin activation (Connell*et al.* 2007), suggesting that it is PUFAs themselves that promote vesicle fusion.

Analysis of individual SNARE components revealed that it is syntaxin that is sensitive to the presence of two major PUFAs enriched in brain, arachidonic and docosahexaenoic acid (Darios & Davletov, 2006; Connell *et al.* 2007). A strong correlation was observed between changes in the structural properties of syntaxin and neuronal growth providing mechanistic insights into PUFA action in brain development (Darios & Davletov, 2006). Although this was demonstrated for syntaxin 3, which mediates vesicle fusion during neuronal growth, it is now clear that syntaxin 1, the major brain isoform, is also sensitive to arachidonic acid (Connell *et al.* 2007; Latham *et al.* 2007) suggesting a conserved mechanism of regulation of the SNARE fusion machinery.

Specificity of PUFA actions

Polyunsaturated fatty acids exhibit high specificity in up-regulation of syntaxin for entry into the SNARE complex, with saturated fatty acids being ineffective (Rickman & Davletov, 2005; Darios & Davletov, 2006; Connell *et al.* 2007). Since saturated fatty acids form micelles at lower concentrations than PUFAs (Tanford, 1973), micelle formation per se is not likely to underlie syntaxin activation. Furthermore, the ability of PUFAs to form micelles is inhibited in the presence of syntaxin (Connell *et al.* 2007). Half-maximal effective concentration for arachidonic acid activation of syntaxin was reported to be around 50 μ M (Rickman & Davletov, 2005). Is such concentration achievable within the cellular environment? A high percentage of phospholipid molecules carry esterified PUFA meaning that, inside the tightly packed lipid bilayer, PUFA concentration will be in a molar range. Local phospholipase action on phospholipid membranes is likely to result in a transiently high concentration of arachidonic acid at the membrane where syntaxin resides, before arachidonic acid diffuses deeper into the cytosol (Brash, 2001). Direct measurements in endocrine insulin-secreting cells demonstrated that, upon stimulation of exocytosis, global arachidonic acid concentration can reach $50-75 \mu$ M (Wolf *et al.* 1991). Furthermore, a very high turnover of arachidonic acid has been demonstrated in isolated nerve terminals and neuronal growth cones, which are enriched in PLA_2 (Negre-Aminou & Pfenninger, 1993). The small volume of these specialized neuronal compartments, containing the SNARE fusion machinery, suggests that syntaxins can encounter transiently high PUFA concentrations that would be sufficient to stimulate SNARE interactions.

Putative mode of action of PUFAs on the syntaxin molecule

How does arachidonic acid promote SNARE interactions? The syntaxin structure is conserved throughout evolution (Munson *et al.* 2000; Rizo & Sudhof, 2002) and consists of a membrane anchor, SNARE motif and an autonomously folded N-terminal domain (Fig. 2). This latter N-terminal domain folds back over the helical SNARE motif thereby inhibiting SNARE complex assembly (Rizo & Sudhof, 2002), either partially (e.g. syntaxins 1 and 2) or completely (e.g. syntaxins 3 and 4) (Connell *et al.* 2007). An essential cytosolic protein, Munc18, is the major binding partner of syntaxins in brain, and the syntaxin 1–Munc18 complex has been crystallised. In this structure, Munc18 acts as a 'clamp' to hold syntaxin 1 in a closed, inactive conformation (Misura *et al.* 2000). Remarkably, arachidonic acid is able to activate syntaxins even in tight association with Munc18, allowing syntaxin engagement of its SNARE partners without abolition of native syntaxin–Munc18 association (Rickman & Davletov, 2005; Connell *et al.* 2007). It is possible that flexible unsaturated fatty acids can penetrate into hydrophobic grooves between the amphipathic syntaxin helices, producing a conformational change and promoting SNARE complex formation. A similar mode of interaction has been suggested for several fatty acid-binding proteins (Hamilton, 2004). Structural studies, e.g. a nuclear magnetic resonance approach, are now required to provide an atomic level description of the dynamic interaction between arachidonic acid and syntaxin.

Conclusion

Although phospholipases and polyunsaturated fatty acids have been known for several decades to play a crucial role in exocytosis, we are still lacking a complete picture of their mode of action. Together with modulation of the ion channel and cytoskeleton functions, PUFAs can act on proteins that are pivotal in vesicle fusion – syntaxins. However, phospholipases not only release PUFAs, but are responsible for the production of other lipid molecules (lysophopholipids, DAG, acylglyerol, phosphatidic acid, see Fig. 1) which may also take part in vesicle fusion (Rhee *et al.* 2002; Rigoni *et al.* 2005; Wang *et al.* 2006). The recent discovery that a ceramidase and a protein closely resembling a diacylglycerol lipase (DAG lipase) are essential for vesicle fusion in *Drosophila* (Rohrbough *et al.* 2004; Huang *et al.* 2006) further highlights the fact that various lipids play critical roles in exocytosis. Furthermore, multiple phospholipase isoforms are present in the body, varying both in distribution and regulatory mechanisms. It still remains to be determined exactly which phospholipase isoforms assist the specific vesicle fusion events, underlying the need for more targeted genetic and pharmacological investigations. Although syntaxins have emerged as likely mediators of PUFA action in exocytosis, we cannot rule out that some of the newly identified proteins essential for vesicle docking, priming, fusion and membrane retrieval may also be sensitive to the products of local lipid metabolism.

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