



REVIEW

Phosphatidylserine, a death knell[#]

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Received 30.8.00; revised 13.11.00; accepted 27.11.00

Edited by M Piacentini

Abstract

Virtually every cell in the body restricts phosphatidylserine (PS) to the inner leaflet of the plasma membrane by energy-dependent transport from the outer to the inner leaflet of the bilayer. Apoptotic cells of all types rapidly randomize the asymmetric distribution, bringing PS to the surface where it serves as a signal for phagocytosis. A myriad of phagocyte receptors have been implicated in the recognition of apoptotic cells, among them a PS receptor, yet few ligands other than PS have been identified on the apoptotic cell surface. Since apoptosis and the associated exposure of PS on the cell surface is probably over 600 million years old, it is not surprising that evolution has appropriated aspects of this process for specialized purposes such as blood coagulation, membrane fusion and erythrocyte differentiation. Failure to efficiently remove apoptotic cells may contribute to inflammatory responses and autoimmune diseases resulting from chronic, inappropriate exposure of PS. *Cell Death and Differentiation* (2001) 8, 551–563.

Keywords: apoptosis; phosphatidylserine; membrane asymmetry; phagocytosis; macrophages; aminophospholipid translocase

Abbreviations: PS, phosphatidylserine; GlcNAc, N-acetylglucosamine; aPLA, anti-phospholipid antibodies

Introduction

The lipid bilayer of the plasma membrane is often considered as simply an inert sea (or a pair of parallel lines) serving as a matrix for suspending membrane proteins. This view is not dramatically dislocated by the fact that individual phospholipid

molecules are sometimes withdrawn from the matrix and hydrolyzed to produce signaling molecules, including prostaglandins, diacylglycerol and ceramides. In recent years, however, a growing body of evidence has suggested that physical and chemical properties of the bilayer itself, such as the thickness of the hydrophobic core¹ or local lateral domains of specialized lipid composition^{2–4} may play significant roles in the assembly and organization of cellular membranes. In addition to these structural contributions to membrane function, the past few years have also seen the revelation that a phospholipid itself, and not a derived product, acts on the extracytosolic, external face of the plasma membrane to regulate intercellular interactions.

Appreciation of this new role for phospholipids was galvanized by the demonstration that phosphatidylserine (PS) appears on the surface of apoptotic lymphocytes and contributes to their phagocytosis by activated macrophages.⁵ The functional importance of the phagocytosis of apoptotic cells would be difficult to overstate. Although *in vitro* assays of apoptosis often concentrate on autodigestion events such as degradation of nuclear DNA or of specific proteins, apoptosis *in vivo* is primarily a process of cell removal. Indeed, the bulk of autodigestion occurs after apoptotic cells have been safely segregated and are already being digested within a phagolysosome.^{6,7} Autodigestion alone, without phagocytosis, can leave corpses lying about.⁸ In organisms such as mammals, where morphological, neural and immunological development depends heavily on the elimination of unwanted cells, and where tissue inflammation is a risk of uncontrolled cell lysis, the mechanisms of apoptotic cell recognition and removal are of critical physiological importance. Moreover, the apoptotic machinery is phylogenetically very old, and may be the *raison d'être* for the presence and disposition of PS in the plasma membrane bilayer.

PS signals clearance by the reticuloendothelial system

The discovery of the importance of PS for removal of apoptotic cells provided a plausible functional foundation for an older literature implicating PS as a potent signal in cell-cell interactions. The power of PS to act as a signaling molecule was clearly demonstrated over a decade ago in studies with artificial lipid vesicles, or liposomes, which had been formulated to persist in the mammalian circulation.⁹ When such vesicles were doped with as little as 3 mole % PS, they were rapidly removed from the circulation by the liver and spleen. The effect was specific to PS—other anionic phospholipids would not substitute—indicating that PS alone and by itself was a signal for clearance by the reticuloendothelial system. These *in vivo* studies complemented prior and subsequent *in vitro* demonstrations that all types of

[#]American Heritage Dictionary: 'a signal of disaster or destruction'; Webster's New World Dictionary: 'an omen of death'.

macrophages preferentially phagocytose liposomes containing PS.^{10–15}

The obvious question provoked by these studies is why blood cells are not similarly cleared from the circulation, given that their plasma membranes contain PS. For quite some time, it has been generally agreed that normal platelets, erythrocytes, and probably lymphocytes, neutrophils, and monocytes as well, are not cleared because PS is not exposed their cell surface, but rather is restricted to the inner leaflet of the plasma membrane bilayer.^{16–18} This transbilayer asymmetry in lipid composition between the inner and outer leaflet is crucial. If erythrocytes present PS on their surface, they are cleared from the circulation^{19,20} just as are PS-containing liposomes. Again, this behavior *in vivo* is seconded by studies *in vitro* which show that macrophages preferentially phagocytose erythrocytes with PS on their surface.^{21,22} The critical role of the exposed PS in provoking phagocytosis is demonstrated by the ability of PS vesicles to specifically and completely inhibit preferential uptake of erythrocytes with PS on their surface.^{23,24} Indirect evidence suggests that this mechanism contributes to the removal of senescent erythrocytes from the circulation.²⁵

These studies with erythrocytes were complemented by evidence that lymphocytes which present PS on their surface interact with macrophages, whereas normal lymphocytes with PS restricted to the inner leaflet, such as those drawn from the circulation, do not interact.^{26,27} The realization some years later that the interacting lymphocytes were apoptotic^{5,28} extended to lymphocytes the concept that exposure of PS has evolved as a mechanism for eliminating effete blood cells.²⁹ Over this same period, increased understanding of the dynamics of PS movements across the bilayer provided insight into the molecular mechanisms which underlie and regulate exposure of PS on the apoptotic cell surface.

Mechanisms for maintenance and loss of PS asymmetry

Like many other basic principles of membrane structure and function, insights into the mechanisms by which PS becomes exposed on the cell surface come from studies of erythrocytes. PS is not the only phospholipid which is asymmetrically distributed across the bilayer, although it is the only major phospholipid which is completely sequestered away from the outer monolayer. It has been known for 30 years that in the plasma membrane of the erythrocyte the choline phospholipids, phosphatidylcholine (PC) and sphingomyelin, are found predominantly in the outer leaflet of the bilayer, whereas phosphatidylethanolamine, which along with PS are the aminophospholipids, is found predominantly in the inner leaflet.³⁰ However, it wasn't until the 1980s that the mechanism for establishing and maintaining this asymmetry began to emerge. In 1984, Seigneret and Devaux demonstrated that a spin-labeled analog of PS introduced into the outer leaflet of the plasma membrane of erythrocytes was rapidly and quantitatively transported to the inner leaflet in an ATP-dependent manner; an analog of phosphatidylethanolamine was transported with lower efficiency, and an analog of

PC not at all.³¹ This transport activity was termed the aminophospholipid translocase, and was also demonstrated in the membrane of chromaffin granules isolated from adrenal cells³² and in synaptic vesicles from the electric eel.³³

The synaptic vesicle, chromaffin granule, and erythrocyte membranes all contain a PS-stimulated ATPase whose activity *in vitro*, like the translocase activity *in situ*, is sensitive to vanadate and N-ethylmaleimide.^{34–36} Cloning of the ATPase from chromaffin granules revealed it to be a P-type ATPase³⁷ which identifies a new subfamily of P-type ATPases distinct from the light (Na^+ , K^+ , Ca^{2+}) and heavy (Cu^{2+} , Cd^{2+}) metal ion transporter subfamilies with multiple members in yeast, slime mold, nematodes, flies, mammals, protozoa and plants.^{38,39} Although the substrates of other subfamily members are not known, mutations in one cause hereditary cholestasis in humans.⁴⁰

The ancient pedigree of the translocase argues that lipid asymmetry is an old and general feature of eukaryotic cells. While often widely assumed to be the case, until recently arguments for this view rested heavily on the demonstration of aminophospholipid translocase activity in a range of mammalian cell types.¹⁷ However, this conclusion has now been more directly and more generally substantiated using the PS-specific probe annexin V. Van den Eijnde and coworkers⁴¹ injected biotinylated annexin intracardially into viable mouse embryos which were subsequently fixed, sectioned and stained with horse radish peroxidase-avidin to identify those cells in which PS was exposed *in vivo*. Non-apoptotic cells of all types were generally unlabeled, indicating that their PS was not exposed on the outer leaflet of the plasma membrane and thus neatly demonstrating the ubiquity of PS asymmetry. Similar experiments in chick and *Drosophila* embryos confirm that most normal cells in these animals also sequester PS,⁴² arguing that plasma membrane lipid asymmetry was a property of the cells of the common ancestor of vertebrates and invertebrates, which lived sometime in the Precambrian, over 600 million years ago. Notably, two instances of annexin binding by non-apoptotic cells were observed: myoblasts fusing to form myotubes and myocardioblasts to form cardiac muscle, and megakaryoblasts and megakaryocytes from which thrombocytes were forming. These intriguing findings are discussed in more detail below.

While the translocase acts to establish and maintain lipid asymmetry, a separate membrane protein acts to disrupt this asymmetry. When cytosolic levels of Ca^{2+} are elevated in erythrocytes or platelets, PS appears on the cell surface within minutes.^{43,44} Although Ca^{2+} inactivates the translocase,⁴⁵ this event alone is insufficient to bring PS rapidly to the cell surface because of the slow rate of passive diffusion of phospholipids across the bilayer.⁴⁶ However, Ca^{2+} also activates a non-specific, bi-directional lipid flippase, distinct from the translocase and termed the scramblase.^{44,47} This activity rapidly randomizes the transbilayer distribution of all the plasma membrane phospholipids to produce a lipid-symmetric membrane, bringing PS to the cell surface. Scramblase activity from erythrocytes⁴⁸ and platelets⁴⁹ has been reconstituted with low efficiency into artificial lipid vesicles and a 37 kD protein with one membrane-spanning domain has been

cloned.⁵⁰ However, this protein is normal in Scott syndrome patients whose blood cells are defective in scramblase activity,^{51,52} suggesting that other molecules are required for scramblase activity and/or activation. A similar activity has also been demonstrated in T cell hybridomas⁵³ and primary thymocytes,⁵⁴ where a distinct lag is seen between elevation of cytosolic Ca^{2+} and scramblase activity (P Williamson *et al.*, submitted), suggesting that the activation mechanism may be complex.

PS exposure in apoptosis

Several lines of evidence indicate that exposure of PS on the cell surface is a general feature of apoptosis. The first evidence was provided by biochemical detection of PS on the surface of apoptotic but not non-apoptotic lymphocytes⁵ likely from a loss of overall membrane asymmetry.⁵⁵ The advent of annexin binding as a simple assay for PS revealed the exposure of PS on apoptotic cells as disparate as a panel of primary and cultured hematopoietic cells,⁵⁶ smooth muscle cells,⁵⁷ vascular endothelial cells,⁵⁸ spermatogenic cells,⁵⁹ neurons⁶⁰ and even cells at developing cranial sutures.⁶¹ This piecemeal approach has been complemented by the more global observations of van den Eijnde,^{62–67} where annexin binding was studied after *in vivo* injection. Those studies showed that cells derived from all three germ layers were labeled by annexin at sites where apoptosis occurs during embryonic morphogenesis, including neurons, bronchi, somites and degenerating tailgut and interdigital tissue. Similar conclusions were drawn from observations of mammalian, avian and insect embryos, placing the evolutionary origin of PS exposure during apoptosis, like the origin of PS asymmetry itself, to a time over 600 million years ago. Moreover, these results offer the hope that the mechanisms of recognition and elimination of apoptotic cells, like the mechanisms of apoptotic autodigestion, will show a common basis in a wide variety of different organisms.

At least for lymphocytes, coordinate regulation of the translocase and the scramblase controls the exposure of PS on the apoptotic cell surface.^{53,68} Activation of the scramblase rapidly brings PS to the outer leaflet, driven by the standing concentration gradient across the bilayer created by the translocase, and PS remains on the cell surface so long as the translocase is inactivated. As expected, exposure of PS on apoptotic lymphocytes, detected by annexin binding, is observed at the time when activation of the scramblase and inactivation of the translocase occur, as detected by transbilayer movements of phospholipid analogs,⁶⁹ with the extent of translocase inactivation suggested to determine the degree of PS exposure.⁶⁸

The mechanisms which regulate the activities of the translocase and scramblase are not known. Since artificially elevating cytosolic Ca^{2+} concentrations both inactivates the translocase and activates the scramblase, Ca^{2+} could serve as the physiologic signal regulating PS distribution, either alone or through intermediaries. However, cytosolic Ca^{2+} chelators do not prevent PS redistribution in apoptotic cells,⁷⁰ even though external application of EGTA does.^{69,70} More recently, it has been shown that the

genetic abnormality in Scott syndrome, which blocks scramblase activation by elevated cytosolic Ca^{2+} , has no effect on scramblase activation during apoptosis (P Williamson *et al.*, submitted), strongly suggesting that scramblase activation in apoptotic cells does not result from changes in cytosolic Ca^{2+} levels. Activation of the scramblase is not restricted to apoptotic cells, however, and occurs upon activation of lymphocytes, monocytes and neutrophils.¹⁶ In both apoptotic and activated cells, the scramblase is activated upon activation of protein kinase $\text{C}\delta$, although scramblase activation in activated cells is only transient and the translocase does not appear to be affected.⁷¹ Finally, studies with inhibitors suggest that scramblase activation requires polyamines,⁷² but why that should be the case is not clear.

Another obvious possible mechanism for regulating PS exposure is that the translocase and scramblase might be targets of proteolytic caspases activated during the autodigestion or execution phase of apoptosis. Indeed, redistribution of PS is sensitive in some instances to the wide spectrum caspase inhibitor zVAD-fmk.⁷³ However, it has also been shown that exposure of PS and subsequent phagocytosis of apoptotic cells can be inhibited while caspase activation remains normal, as measured by PARP cleavage and DNA degradation.⁷⁴ Moreover, in instances when caspase activation is not required for the induction of apoptosis, PS exposure downstream is not sensitive to zVAD-fmk.⁶⁹ Similarly, induction of apoptosis in sperm and avian erythroid cells results in DNA degradation, as measured by TUNEL-positive nuclei, PS exposure, and eventual cell lysis, but these processes do not appear to be caspase-dependent.⁷⁵ In addition, at least in lymphocytes, application of EGTA to apoptotic cells can restore translocase activity and extinguish scramblase activity.⁶⁹ These findings argue that the regulation of the translocase and scramblase can be reversible, making it unlikely that they are direct targets of caspases, and suggesting that PS exposure is one of several features of apoptosis which are independent of zVAD-sensitive caspases. Since protein kinase $\text{C}\delta$ is cleaved and activated in a caspase 3-dependent manner in apoptotic cells,⁷¹ zVAD-insensitivity of PS exposure suggests that mechanisms independent of protein kinase $\text{C}\delta$ activation are available for activating the scramblase in apoptotic cells. However the translocase and scramblase are regulated, redistribution of PS appears to be downstream of participation by bcl-2,⁵⁶ and analysis of APAF knockout mice suggests that PS exposure may be downstream of this ced-4 homolog as well.⁷⁶ The fact that PS exposure is caspase-independent suggests that APAF may play a role in the regulation of pathways that are not connected to the activation of the zVAD-sensitive caspase-9.⁷⁷

PS as a signal for phagocytosis of apoptotic cells

The universality of PS exposure on apoptotic cells argues for an important role for loss of lipid asymmetry in apoptosis, but provides no real clues by itself as to what that role might be. Nor does it necessarily follow that PS exposure is the property

of functional consequence, rather than just an indicator of loss of asymmetry, since a decrease in lipid order⁷⁸ and packing,⁷⁹ an increase in fluidity⁸⁰ and an increase in hydrophobicity of the outer leaflet²² also accompany loss of asymmetry. However, there is now ample evidence that loss of asymmetry targets apoptotic cells for phagocytosis and that PS is a key element in this process.

As is the case for erythrocytes which have lost their asymmetric distribution of phospholipids, PS, either in the form of liposomes or just its water-soluble headgroup (phosphoserine), can stereospecifically inhibit phagocytosis of apoptotic cells.⁵ These data demonstrate (1) that PS is not just an indicator of loss of lipid asymmetry in apoptotic cells, but rather an active participant in phagocytosis, and (2) that a receptor protein is involved in the recognition of the PS headgroup. For some time, appreciation of the generality of the central role of PS in the phagocytosis of apoptotic cells was limited by the fact that only recognition by macrophages, such as elicited peritoneal macrophages, is inhibited by PS vesicles, while phagocytosis by unactivated macrophages, such as bone marrow or monocyte-derived macrophages, is not sensitive to these vesicles.⁸¹ This difference led to the supposition that macrophages possess either a PS-dependent or PS-independent ($\alpha_v\beta_3$ integrin-independent) recognition system. However, lipid-symmetric erythrocytes (which display PS on their surface and whose phagocytosis is inhibited by PS vesicles) inhibit uptake of apoptotic cells by macrophages which use the $\alpha_v\beta_3$ integrin-dependent mechanism.²⁴ Furthermore, annexin blocks phagocytosis of apoptotic thymocytes by both types of macrophages, although it has no effect on or even enhances phagocytosis resulting from Fc receptor recognition.⁸² These results imply that PS recognition is an integral part of the removal of apoptotic cells by all macrophages.

In the literature, macrophages which utilize the $\alpha_v\beta_3$ integrin system for apoptotic cell recognition have been described as 'unactivated' macrophages and those which are inhibitable by PS vesicles as 'activated' macrophages. This usage is unfortunate because the term 'activation' also refers to macrophages which have become cytotoxic. Until a more appropriate terminology is agreed upon and adopted, the terms activated and unactivated will continue to be used. They are used in the remainder of this review with the understanding that they refer only to the alternate systems used by macrophages to recognize apoptotic cells.

Remarkably, it has recently been shown that exposure of PS is not just required on the apoptotic cell. Over a decade ago, studies with fluorescent membrane probe MC540 suggested that macrophages constitutively express lipid-symmetric domains in the plasma membrane which preferentially cycle into endocytic vesicles.⁸³ Recently, *in vivo* annexin labeling revealed that the macrophage membrane near or surrounding the engulfed apoptotic cell exposes PS like the apoptotic cell itself.⁶⁷ This PS exposed on the phagocyte surface is required in the phagocytosis process, since treating macrophages with annexin is as effective as treating apoptotic lymphocytes in blocking the engulfment process.⁸⁴ The engulfment process thus seems to require interactions between PS-exposing membranes

on both the engulfing and target cells, interactions that are not required for engulfment of other, non-PS-exposing particles by the same phagocytes.

If PS recognition is important, something, and presumably a protein, must do the recognizing. Recently, Fadok and coworkers⁸⁵ have identified a protein molecule required for phagocytosis of apoptotic cells by activated macrophages, which are those sensitive to PS vesicles. This molecule is expressed only at low levels on unactivated macrophages, and its expression on transfected cells renders phagocytosis by those cells sensitive to PS vesicles. Together, these results imply that the molecule is intimately involved in the PS-dependent mechanism of phagocytosis by activated macrophages, and the authors suggest that it may be a receptor for PS. If this exciting possibility is true, the molecule cannot be the only such receptor molecule, since it is apparently not required for the PS-dependent recognition of apoptotic cells by unactivated macrophages, and is presumably not generally present on the target cell surface, which may also possess molecules which recognize the PS on the macrophage surface. Whether a family of these molecules may play similar roles on different cells is an interesting question.

Surface receptors and the engulfment of apoptotic cells

Although PS exposure is central to the recognition of apoptotic lymphocytes by all types of macrophages, the preceding discussion has already hinted that membrane proteins are also required. A great deal of energy has been expended in identifying the proteins that participate in the phagocytosis of apoptotic cells, an effort which has resulted in an embarrassment of riches (Table 1). Several principles are useful in clarifying the profusion and confusion represented by the (growing) size of this table. One is that the involvement of

Table 1 Proteins implicated in the recognition of apoptotic cells

Protein	Cell	Evidence	Reference
Integrins			
$\alpha_v\beta_3$	Phagocyte	Biochem	81
$\alpha_v\beta_5$	"	"	86
$\alpha_7\beta_1$	"	"	162
Scavenger receptors			
CD36	Phagocyte	Biochem	88
OxLDL	"	"	163
SRBI	"	"	89
SR-A	"	"	164
Croquemort	"	Genetic	165
Asialoglycoprotein receptor	Phagocyte	Biochem	166
Lectin-like receptor	Phagocyte	Biochem	167
CD14	Phagocyte	Biochem	95
ICAM3	Target	Biochem	87
ABC1	Both	Gen/Biochem	113
<i>ced-7</i>	Both	Genetic	108
<i>ced-5</i>	Phagocyte	Genetic	168
<i>ced-6</i>	Phagocyte	Genetic	103
Complement proteins	-	Biochem	169-171
Thrombospondin	-	Biochem	88
β_2 -GPI	-	Biochem	160
Gas-6	-	Biochem	172

many of these proteins is not general, but rather is restricted to particular target/phagocyte pairs. Thus, recognition involving the lectin-like receptor is specific to activated macrophages, the $\alpha_v\beta_3$ integrin to unactivated macrophages, the $\alpha_v\beta_5$ integrin to naive dendritic cells. While such proteins may play roles in more than one cell system, they are clearly not general; whether they are functional substitutes for each other is not known, although in some cases, such as the integrins, this conclusion is tempting.

A second principle is that engulfment must involve cooperation among receptors (Table 2). For example, the phagocytosis of simple lipid-symmetric erythrocytes or apoptotic lymphocytes by activated macrophages involves a PS receptor, because it is partially inhibited by PS vesicles, and also a lectin-like receptor, because it is also partially inhibited by N-acetylglucosamine (GlcNAc).²⁴ The inhibition by PS vesicles and GlcNAc are not additive, suggesting that the two receptors must cooperate. Similarly, engulfment of apoptotic lymphocytes by unactivated macrophages is partially inhibited by RGDS, which is not additive with the inhibition provided by lipid-symmetric erythrocytes. These results suggest that multiple signals combine on the apoptotic cell surface to produce a complex ligand recognized by a complex of receptors.²⁴

With these principles in mind, several patterns can be observed in Table 1. The first is the repeated identification of integrins, including the $\alpha_v\beta_3$ ⁸¹ or $\alpha_v\beta_5$ ⁸⁶ receptors, or integrin ligands such as ICAM-3, the only protein on the apoptotic cell surface identified as involved in recognition.⁸⁷ As yet an integrin has not been identified in the recognition of apoptotic cells by activated macrophages, and the general integrin-inhibitory peptide RGDS has no effect on recognition by these cells, but not all integrin binding is sensitive to the inhibitor, leaving open the question of whether the requirement for integrin involvement is general.

A second regularity is the repeated identification of members of the scavenger receptor family, including CD36 in conjunction with thrombospondin,⁸⁸ SR-BI,⁸⁹ and the product of the *croquemort* gene in *Drosophila*.⁹⁰ Since these proteins recognize polyanions, including phospholipid micelles and bilayers containing anionic phospholipids such as PS,⁹¹ they might be capable of recognizing PS on the apoptotic cell surface and contribute to phagocytosis to a varying degree depending on the nature of the target cell

and phagocyte. However, they cannot account for instances where PS has been shown to be the only anionic lipid which inhibits phagocytosis of apoptotic cells nor explain the stereospecificity of PS inhibition. These considerations and others, including indirect evidence regarding CD36 in particular, argue that the PS receptor is not a scavenger receptor.⁹² However, whether these receptors contribute to nonspecific recognition of the anionic surface of a PS-exposing apoptotic cell membrane remains an open question. Moreover, while removal of apoptotic cells is not measurably reduced in mice in which scavenger receptor genes have been (singly) disabled by knockout mutations,⁹³ apoptotic cell removal in *Drosophila* is specifically sensitive to mutation of the *croquemort* gene,⁹⁰ suggesting that these receptors play some central and important part in the process. CD14 is a macrophage-specific receptor which binds lipopolysaccharide (LPS), triggering inflammatory responses and septic shock.⁹⁴ It was identified by expression cloning as the target of the monoclonal antibody 61D3,⁹⁵ which blocks phagocytosis of lymphocytes by both activated and unactivated macrophages.^{24,96} Only anti-CD14 antibodies which inhibit binding of LPS can block phagocytosis of apoptotic lymphocytes, suggesting that the LPS and apoptotic cell binding sites are closely associated or identical.⁹⁵ Moreover, phagocytosis of lipid-symmetric erythrocytes is CD14-dependent.^{24,97} Since the only property which lipid-symmetric erythrocytes are known to share with apoptotic thymocytes is exposure of PS on their cell surface, these observations suggest that CD14 may recognize PS. However, there is no evidence in support of this possibility other than a report that human monocyte THP-1 cells transfected with CD14 display a twofold increase in low affinity binding of PS vesicles in the presence of LPS-binding protein.⁹⁸

Activation of protein receptors for apoptotic cells

Work from Raff's laboratory^{99,100} has shown that all the components necessary for initiation and completion of the apoptotic process pre-exist in the cytoplasm of mammalian cells: synthesis of new proteins is not required for cells to enter and complete the process of apoptosis. Although the cell surface changes which lead to phagocytosis occur early in apoptosis, there is still plenty of opportunity within that timeframe for the modification, removal, and reassignment of cell surface molecules in the service of engulfment. Nevertheless, the fact that the process does not require novel gene expression, combined with the large variety of cell types and environments in which recognition and phagocytosis take place, raises the possibility that the interaction between the apoptotic cell and phagocyte may not depend on dedicated receptor/ligand interactions, but rather may make opportunistic use of the particular cell surface recognition molecules that are available on the target cell and its neighbor before apoptosis begins. This possibility would explain the familiarity of many of the molecules which have been suggested as participants. It would also predict that the proteins involved would differ depending on the specific cells which are

Table 2 Receptors implicated by inhibitors in recognition of apoptotic lymphocytes by macrophages

Inhibitor	Macrophage inhibited		Implied receptor
	Unactivated	Activated	
PS vesicles	No	Yes	PS receptor I
LS erythrocytes	Yes	No	PS receptor II (?)
RGDS, $\alpha_v\beta_3$ Ab	Yes	No	$\alpha_v\beta_3$ integrin
GlcNAc	No	Yes	Lectin-like receptor
61D3 Mab	Yes	Yes	CD14 (LPS receptor)
Annexin V	Yes	Yes	PS receptors on apoptotic cell and/or macrophage
CD36 Ab	Yes	Yes	CD36 (scavenger receptor)

interacting as target and phagocyte. Exactly such differences have been demonstrated when the same target is presented to different kinds of engulfing macrophages²⁴ and when different targets are presented to the same macrophage (Y Ren *et al.*, submitted).

In this view, the nature of the specific ligands and receptors involved becomes a problem which is specific to the particular cells under study. If this view is correct, the current list of cell surface receptors involved in recognition may actually be limited by the small number of target/phagocyte cell pairs which have been examined (largely macrophages and apoptotic hematopoietic cells), and this list may be considerably lengthened as other combinations are studied. The more general question is then the mechanism by which these interactions are usurped by apoptosis for the purpose of recognition and phagocytosis of apoptotic cells, processes not provoked when these receptors and ligands normally interact. Accumulating evidence suggests that the critical activating process on the apoptotic target cell surface is the transbilayer rearrangement of membrane lipids.

When the scramblase is activated in erythrocytes by raising cytosolic Ca^{2+} concentrations, the resulting lipid-symmetric erythrocytes are recognized and phagocytosed by macrophages by a process which involves CD14²⁴ and, in the case of activated macrophages, lectin-like receptors,⁹⁷ just as in the case of apoptotic targets. Yet the absence of a nucleus, protein synthesis apparatus, or internal membrane system in erythrocytes precludes the appearance of new proteins on the cell surface. In this model system, integrin receptors do not seem to play a role, but when lipid asymmetry is similarly abolished in more complex lymphocytes by raising cytosolic Ca^{2+} concentrations, the cells immediately become targets for recognition by the same CD14 receptor, lectin-like receptor, and integrin-based mechanisms that are involved in recognition of authentic apoptotic cells.⁸⁴ On the other hand, apoptotic cells in which PS exposure is blocked or defective are no longer recognized by any of the protein-mediated mechanisms.¹⁰¹ Together, these results argue that loss of phospholipid asymmetry and/or the appearance of PS on the apoptotic cell surface activate the recognition process, including those features which are dependent on specific membrane protein receptors and ligands other than PS. As mentioned above, alterations in membrane physical properties accompany transbilayer lipid rearrangement; in the apoptotic lymphocyte membrane, any of these changes may promote the generation of recognition ligands through the lateral rearrangement and association of existing membrane proteins, formation of PS-protein complexes or the exposure of new epitopes on proteins.¹⁰²

Other proteins implicated in phagocytosis

The list of proteins implicated in phagocytosis of apoptotic cells is not restricted to cell surface molecules. In particular, genes identified as required for removal of apoptotic cells in *C. elegans* encode several proteins which are not cell surface receptors. One of these, the product of the *ced-6* gene,¹⁰³ is a cytoplasmic protein which has been proposed to act as an

adaptor in a signal transduction pathway. The nature of this pathway is currently obscure, but it is clear that a signaling pathway must be activated during removal of apoptotic cells. This pathway suppresses the normal inflammatory responses of macrophages to phagocytic stimuli^{95,104} through autocrine/paracrine mechanisms involving TGF-beta, PGE2 and PAF.¹⁰⁵ When in the phagocytic process this pathway is provoked, whether it involves *ced-6* homologs, and whether its induction involves a receptor/ligand interaction separate from those involved in binding and recognition of the apoptotic cell, are all currently unknown.

Two other proteins, the products of the *ced-5* and *ced-10* loci, are required for removal of apoptotic cells in *C. elegans*. These proteins are members of the crkII/dock180 signaling pathway;¹⁰⁶ in mammalian cells, this pathway is linked to motility, and closely related pathways play important roles in controlling the cytoskeletal rearrangement required for engulfment.¹⁰⁷ These findings confirm the conclusion drawn from the specificity of annexin inhibition of phagocytosis and the specificity of anti-phlogistic effects that engulfment of apoptotic cells is mechanistically distinct from the phagocytosis of other targets. Like those studies, information on how the process differs is only sketchy at this time, and it is not yet clear how the unique events on the cell surface are linked to the unique anti-phlogistic pathway or the unique pathway regulating the cytoskeletal machinery.

Another molecule identified from the collection of *C. elegans* engulfment mutants is the product of the *ced-7* gene,¹⁰⁸ a member of a very large family of membrane ATPases called ABC proteins.¹⁰⁹ Other members of this family include the regulatory subunit of the SUR K^+ transporter,¹¹⁰ the CFTR protein mutated in cystic fibrosis,¹¹¹ and the *mdr* and *MRP* proteins whose over-expression confers non-specific drug resistance on cells.¹¹² A human homolog of the *ced-7* gene, called ABC1, was isolated from macrophages, and a functional homology between the worm and human proteins established by demonstration that intracellular injection of antibodies against ABC1 into macrophages blocked engulfment of apoptotic cells.¹¹³ A knockout mutation of this gene in mice also resulted in a block to the rapid engulfment of apoptotic cells by macrophages.¹¹⁴ Remarkably, in *C. elegans*, the product of the *ced-7* gene was equally required in both the target and the phagocytic cell,¹⁰⁸ and pharmacological inhibition of ABC1 indicates that the same is true in mammals.¹¹⁵ The similarity of this dual requirement to that for PS exposure seems to be no mere coincidence—the same pharmacological inhibitors that suppress engulfment also block Ca^{2+} -induced loss of lipid asymmetry and PS exposure in both targets and macrophages,¹¹⁵ and PS exposure is also abnormal in lymphocytes and erythrocytes from ABC1 knockout mice.¹¹⁴

In a surprising turn of events, the phenotype of the human knockout of the ABC1 gene is also known since the discovery that mutations in this gene are responsible for Tangier disease,^{116–121} a genetic deficit of interest because patients display a defect in reverse cholesterol transport from cells to circulating lipoproteins. This phenotype is not peculiar to humans—the ABC1 knockout

mouse displays a similar defect.¹¹⁴ These findings imply that ABC1 plays an important role in the process of cholesterol efflux. This process is not well understood, so that the participation of ABC1 does not cast any light on how ABC1 might affect this process. The hydrophobic character of cholesterol makes it unlikely that any molecule is required for its transbilayer transport; a more likely role for ABC1 may be that it contributes to the phospholipid efflux which occurs concomitantly with cholesterol efflux,¹²² but experimental evidence will be needed to settle this issue. What is intriguing is the connection that is revealed between recognition of apoptotic cells and lipid transport processes; the same intriguing connection can be drawn from the involvement of lipoprotein-binding scavenger receptors in apoptotic cell phagocytosis.

Professional vs non-professional phagocytes

It should be kept in mind in all these considerations that cell removal requires at least two distinct mechanistic steps. The first is recognition of the apoptotic cell, i.e., the formation of cell-cell interactions specific to the apoptotic nature of the target cell. The second is the actual engulfment process, which must depend on the identification of the first interaction as apoptosis-specific. The large majority of studies on the mechanisms of phagocytosis of apoptotic cells have employed professional phagocytes (macrophages), where these two processes are closely linked to one another in time, with phagocytosis following recognition within a few minutes.¹²³ Under these circumstances, specific phagocyte/target pairs which have not yet completed engulfment are rare²⁴ and difficult to quantify separately. But these mechanisms are in fact separable. Resident peritoneal macrophages specifically bind but do not engulf PS-exposing erythrocytes unless activated¹²⁴ or unless serum is present.¹²⁵ Even more clearly, engulfment by nonprofessional phagocytes such as BHK cells begins with a rapid recognition step, but phagocytosis is delayed for several hours.¹²³ While the characteristics of these individual steps are not yet known, the fact that annexin inhibits the phagocytosis of apoptotic smooth muscle cells by their non-professional normal neighbors⁵⁷ implies that the involvement of PS in phagocytosis is not specific to professional phagocytes.

Overall, it might be expected that the recognition and binding step is primarily a matter of specific receptor-ligand interactions, while the engulfment step is a matter of membrane and cytoskeletal dynamics and the two are presumably linked by signaling events of some kind. But certainly some of the mechanisms uncovered by studies of macrophages are specific to these specialized cells and not generally applicable to non-professional phagocytes. For example, although CD14 appears to be absolutely required by professional phagocytes of all stripes to phagocytose apoptotic lymphoid cells, this requirement cannot extend to non-professional phagocytes since expression of CD14 is restricted to cells of the monocyte/macrophage lineage. While some unknown protein more ubiquitously expressed may play a role in non-professional cells similar to the role of CD14 in professional cells, CD14 must represent a

specialization of professional phagocytes for removal of apoptotic cells, particularly in the immune system. The same logic applies to any of the other surface molecules implicated in phagocytosis—to the extent that they are restricted to professional phagocytes, there must be a functional homolog in other types of cells if the process in which they participate is general to phagocytosis of apoptotic cells. On the other hand, since apoptotic cells of all types express PS on their surface, and presumably undergo the physical membrane changes accompanying loss of asymmetry, the ligands on apoptotic cells recognized by non-professional phagocytes may be generated by the same mechanisms as the ligands recognized by professional phagocytes, though, as discussed above, the actual ligand/receptor pairs may vary depending on the cell types. A key to unraveling the roles of the many molecules which participate in apoptotic cell clearance without provoking inflammation may lie in distinguishing the several steps in the process and identifying the molecules required for each of them.

Other consequences of PS exposure

As mentioned above, apoptosis, and the exposure of PS associated with it, are probably over 600 million years old. Given that every cell in the body has the ability to sequester PS to the inner leaflet or to present it on the cell surface, it is not surprising that over time this process has been appropriated by evolution for specialized purposes. As a result, there are several physiological processes in which PS exposure plays a role, which in turn provide a source for pathological consequences if disrupted by disease. A consideration of these specialized instances may provide insights into the ancestral mechanisms which regulate PS exposure and recognition.

The best studied of these specialized systems is the platelet. When the platelet is activated by thrombin and collagen acting through the α IIb β 1 and gpVI receptors, the translocase is inactivated and a vigorous scramblase is activated,¹²⁶ bringing PS extremely rapidly (seconds) to the platelet surface.¹²⁷ The exposed PS becomes the site of assembly of the tenase and prothrombinase complexes of the proteolytic cascade leading to the thrombin cleavage of fibrinogen and formation of the fibrin matrix of the clot.¹²⁸ Both the tenase and prothrombinase complexes consist of a catalytic subunit and a cofactor; while each of these can stereospecifically bind PS by itself,^{129–131} the enzymatic activity of the complex is much higher than that of either protein alone.¹³² The PS-binding region of the catalytic subunits (factors IXa and Xa) consist of characteristic Gla domains which contain several γ -carboxyglutamic acid residues, a vitamin-K dependent amino acid modification.¹³³ The cofactor subunits (factors Va and VIIIa) do not contain such residues, but are even more specific than the catalytic subunits in their recognition of PS.¹³⁴ This cooperative 'recognition' mechanism may represent a strategy for obtaining highly specific recognition of a ligand such as PS which is structurally simple, but repetitively expressed. It is also superficially reminiscent of the proposal that PS-dependent recognition of apoptotic cells

requires a complex, multicomponent recognition system. More concretely, these proteins contain domains which recognize PS, and their PS-binding domains appear in other proteins which recognize PS in the more primitive process of clearing apoptotic cells. Although macrophages may be involved in resolution of the clot through recognition and phagocytosis of activated platelets with PS on their surface, such a process is secondary to the more important, crucial role of PS in clot formation.

Besides these possible correlates of PS recognition on apoptotic cells, platelets might also illuminate the signaling mechanisms which regulate PS exposure. As in apoptotic cells, PS exposure in platelets requires coordinate regulation of the translocase and scramblase. At present, regulation of these processes is not well understood in either system, but a natural mutation which prevents activation of the scramblase is manifest in Scott syndrome.^{136,137} Since the translocase is regulated normally in the cells of Scott syndrome patients,¹³⁷ it and the scramblase appear to be regulated independently of one another. The scramblase in normal platelets is activated by a Ca^{2+} -mediated signaling pathway which fails to activate the scramblase in Scott platelets. Artificially elevating Ca^{2+} using ionophore similarly fails to activate the scramblase in Scott platelets, erythrocytes and lymphocytes.^{138,139} But activation of the scramblase in apoptotic Scott lymphocytes proceeds normally, implying that the physiological Ca^{2+} signaling pathway utilized by activated platelets is not the pathway used by apoptotic lymphocytes (P Williamson *et al.*, submitted).

As mentioned above, the studies of Van den Eijnde⁶² quite surprisingly reveal that PS exposure occurs on budding megakaryocytes of mouse embryos, as well as on fusing myoblasts, confirming an old report that PS is exposed on myoblasts fusing in culture.¹⁴⁰ The functional significance of PS exposure in these two cases is not clear, although extensive membrane fusion is a critical feature of both myotube formation and thrombocyte release from megakaryocytes, and PS has long been implicated in the fusion process.⁶³ Similarly, it is reasonable to suppose that the mechanism of PS exposure in these two cell types is derived in some way from the more ancestral mechanism in apoptotic cells. Most importantly, however, the presence and persistence of these PS-exposing cells in the normal embryo is evidence that PS exposure is not sufficient by itself to trigger phagocytosis. Whether these cells escape phagocytosis because they lack some other important component required for recognition, because they somehow mask the exposed PS on their surface, or because they occupy some privileged site within the embryo devoid of competent phagocytes will be an interesting subject for further investigation.

Erythropoiesis provides an instructive instance in which localized exposure of PS on the apoptotic cell surface may result in a specialized phagocytic event. In mammals, the development of the nucleated erythroblast comes to a climax upon enucleation, during which the cytoplasm of the polychromatophilic erythroblast becomes the incipient reticulocyte, as the nucleus extrudes from the erythroblast surrounded by only a thin cytoplasmic remnant and is

engulfed by a macrophage even before segregation of the two bodies is complete.^{141,142} When observed in isolation, the process of enucleation was described by Bessis: 'With time lapse photography ..., the cell appears to push pseudopodia frenetically in all directions and after a number of apparent convulsions, the nucleus is extruded',¹⁴³ a description which is highly reminiscent of the morphological description of apoptosis by Parnaik.¹²³ Moreover, the nuclear condensation and margination that occurs during this process are quite similar to the morphology of apoptotic cells. The portion of the plasma membrane which surrounds the lobe of the cell containing the extruding nucleus binds MC540,^{144,145} a fluorescent dye which marks lipid-symmetric membranes⁷⁹ and stains apoptotic cells,^{5,55} suggesting that recognition and engulfment of this portion of the erythroblast occurs by the same PS-dependent mechanisms by which apoptotic cells are recognized and phagocytosed. Remarkably, the reticulocyte lobe of the cell does not bind MC540, and is not phagocytosed, but rather after partition is complete is released into the circulation. How this portion of the membrane escapes losing lipid asymmetry is not known, but it may depend on lateral rearrangement and localization, within the plane of the membrane, of the critical proteins which control lipid asymmetry. In humans, the erythrocytes which mature from the incipient reticulocyte contain very low levels of scramblase compared to platelets and lymphocytes,^{47,146} and pig erythrocytes lack scramblase activity altogether,¹⁴⁷ while translocase levels in both species of erythrocytes are respectable. How lateral rearrangement and segregation of the scramblase to the plasma membrane surrounding the extruding nucleus might occur is certainly not clear, but may be related to the compartmentalization of the erythrocyte spectrin-based cytoskeleton to the reticulocyte portion of the enucleating cell.¹⁴⁸

Given the potent responses to PS exposed on the surface of platelets and apoptotic cells, it is not surprising that inappropriate exposure of PS can have pathological consequences. Abnormal PS exposure has been associated with sickle cell disease for many years. One source of this exposure is the release of small vesicles from sickle erythrocytes;^{149,150} these vesicles become lipid symmetric,¹⁵¹ perhaps because of ATP depletion arising from their high surface/volume ratios or perhaps during the fusion process by which they are released. In addition, it has become clear that PS exposure also occurs in a small fraction of circulating sickle cells^{152,153} which may contribute to the anemia as well as the hypercoagulability¹⁵⁴ associated with the disease. Because human erythrocytes contain both translocase and scramblase activities, the exposure of PS must require translocase inactivation and may also involve scramblase activation. One possible mechanism for PS exposure in sickle erythrocytes might be the release of Ca^{2+} from internal vesicles contained within sickle erythrocytes¹⁵⁵ which originate from the plasma membrane and thus concentrate Ca^{2+} inside themselves by the action of the plasma membrane Ca^{2+} -ATPase.¹⁵⁶ Sporadic lysis and release of high concentrations of Ca^{2+} from the vesicles might account for the

appearance of PS on only a small fraction of circulating cells.

One intriguing finding which may be related to the exposure of PS on sickle erythrocytes is that sickle cell disease patients tend to develop antiphospholipid antibodies (aPLA).¹⁵⁴ These autoimmune antibodies are better known from studies of patients with aPLA syndrome and lupus patients, where it has been established that the antibodies are actually low affinity antibodies directed against plasma proteins, including β -2 glycoprotein I, thrombin, and proteins S and C.¹⁵⁷ The antibodies were originally thought to react with phospholipids because they bind to ELISA plates coated with phospholipids. It is now clear that this behavior results from the ability of the plates to bind and concentrate the protein 'cofactors' which are the actual target of the antibodies.¹⁵⁸ The appearance of the 'aPLA' in patients with sickle cell disease, where the pathological appearance of PS is well documented, lends weight to the proposal that the antibodies may arise generally from the abnormal persistence of PS-exposing cell surfaces. A more general source for such surfaces might be from delay or failure in the system of apoptotic cell removal.¹⁵⁹ The resulting persistence of PS-exposing apoptotic cells could permit accumulation of cofactors on the cell surface, thus generating the provoking autoantigens. An important ramification may be the effect on recognition of apoptotic cells coated with these proteins. Because of its ability to bind PS, the major cofactor, β -2 glycoprotein I, has been proposed as a marker for apoptotic cells.¹⁶⁰ The presence of β -2 glycoprotein I does not enhance the efficiency of recognition of apoptotic cells by macrophages,¹⁶¹ arguing against such a role for the protein in normal recognition of apoptotic cells. However, the aPLA which recognize and bind to β -2 glycoprotein on the cell surface do enhance the phagocytosis of apoptotic cells through Fc-mediated phagocytosis;¹⁶¹ under such circumstances, phagocytosis of apoptotic cells results in a proinflammatory response by the phagocytic cells. Thus, aPLA may convert the removal of apoptotic cells from a benign, physiological to a pathological process.

Together, these results point out how failure to remove apoptotic cells in a timely fashion may have pathological consequences in addition to the local inflammation which is usually invoked as the physiological rationale for rapid removal of apoptotic cells. Whether such a failure is a contributor to autoimmune diseases such as lupus, and if so, where the breakdown in removal occurs, will be important areas for studies exploring the clinical implications resulting when removal of apoptotic cells is compromised.

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