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Lysophosphatidylcholine as a ligand for immunoregulation

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Abstract

Despite the recognized effects of lysophosphatidylcholine upon cells of the immune system and its association with inflammatory processes, its mechanism of action has remained poorly characterized. Our recent identification of the first lysophosphatidylcholine receptor as an immunoregulatory G protein-coupled receptor named G2A whose genetic ablation results in the development of inflammatory autoimmune disease has, therefore, provided a new perspective on the role of this lysophospholipid as a modulator of immune responses. This commentary discusses the biological properties of lysophosphatidylcholine as an immunoregulatory ligand for cells of the innate and adaptive arms of the immune system. Although we focus primarily on ligand interactions with G2A, we also discuss the issue of possible functional redundancy with other receptors with recently established ligand specificities towards phosphorylcholine-containing lysolipids including lysophosphatidylcholine. © 2002 Published by Elsevier Science Inc.

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1. Lysophospholipids

Bioactive lysophospholipids are a group of naturally occurring lipids regulating a wide variety of cellular activities including proliferation, smooth muscle contraction, wound healing, and tumor cell invasiveness [1]. Formation of lysophospholipids is enhanced under inflammatory conditions, in ischemic tissue, and during oxidation of LDL. Lysophosphatidic acid and sphingosine-1-phosphate are the best characterized lysophospholipids, and eight receptors (Endothelial Differentiation Gene receptors; EDG) have been identified thus far with varying ligand specificities towards them [2]. Although most EDG receptors are broadly expressed and mediate the diverse effects of lysophosphatidic acid and sphingosine-1-phosphate in multiple tissues, others such as EDG 6 are expressed predominantly in lymphocytes and accessory immune cells [3], suggesting

that their primary roles may be immunoregulatory. However, for certain lysophospholipids, biological actions have been established primarily within the immune system. For example, LPC is an important inflammatory mediator with recognized effects in multiple immune cell types and pathophysiological processes. Most notably, as a major component of oxLDL, LPC plays a major etiological role in atherosclerosis [4]. Oxidative modification of phospholipids is associated with the acute phase response and chronic inflammation, and its by-products are increasingly recognized as key factors in the initiation and progression of chronic inflammatory autoimmune diseases [5–7]. Among these, LPC is implicated in the pathogenesis of the autoimmune disease SLE [8,9].

2. Biological actions of LPC

Molecular species of LPC, distinguished by the lengths and saturation of their acyl chains, are produced from cell membrane-derived phosphatidylcholine (PC) as a result of hydrolysis by PLA₂ [10]. PLA₂ activation is a rate-limiting event in the production of several proinflammatory lipids (Fig. 1) and is associated with cellular responses to multi-

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Abbreviations: APCs, antigen presenting cells; GPCR, G protein-coupled receptor; LDL, low-density lipoprotein; LPC, lysophosphatidylcholine; oxLDL, oxidized low-density lipoprotein; PLA₂, phospholipase A₂; SLE, systemic lupus erythematosus; and SPC, sphingosylphosphorylcholine.

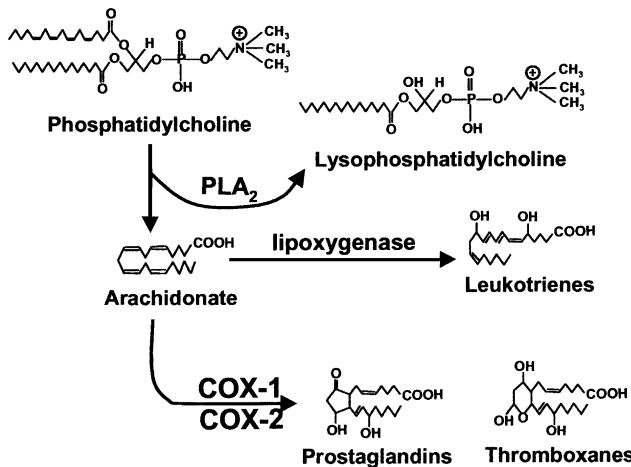


Fig. 1. Phospholipase A₂ (PLA₂), a rate-limiting enzyme in the generation of multiple lipid inflammatory mediators. A large proportion of cellular membrane phosphatidylcholine contains an arachidonoyl acyl moiety at the *sn*-2 position, the target of PLA₂ hydrolysis. Consequently, in addition to lysophosphatidylcholine, arachidonic acid is a major product of PLA₂ action. Arachidonic acid is a key intermediate for the production of various other lipids via the actions of lipoxygenase and cyclooxygenase (COX) enzymes. COX-2 is a transcriptionally regulated isoform of COX induced by multiple mitogenic and stress stimuli, and the target of highly efficacious inhibitory drugs used in the treatment of chronic inflammatory conditions.

ple stimuli, including stress and inflammatory signals. Among the different forms of PLA₂, secretory forms (sPLA₂) are released from macrophages and other cell types at sites of inflammation and tissue injury [11]. LPC is

also derived from PC-containing lipoproteins as a result of oxidative processes via activation of endogenous PLA₂ [12]. Early studies demonstrating the ability of high concentrations of LPC (50 μ M) to modulate plasma membrane integrity via micelle formation suggested that its biological role may be primarily structural and cytolytic. However, specific effects of low concentrations of LPC at the level of several physiological and pathological processes through an uncharacterized GPCR(s) have since been widely reported. Extracellular production of LPC by sPLA₂ promotes inflammatory effects including up-regulation of endothelial cell adhesion molecules and growth factors [13,14], chemotaxis of monocytes and T lymphocytes [15,16], and stimulation of macrophage activation [17]. Although some effects of LPC were believed to be mediated by platelet-activating factor (PAF) receptors in certain cell types [18], the identification of a specific high-affinity LPC receptor remained elusive. We have recently identified two LPC receptors, one of which plays an important immunoregulatory role and exhibits higher affinity binding to this lysophospholipid [19–21].

3. LPC receptors

We identified G2A, a G protein-coupled receptor, as a transcriptional target of the leukemogenic tyrosine kinase BCR-ABL in murine B lymphoid progenitor cells [22].

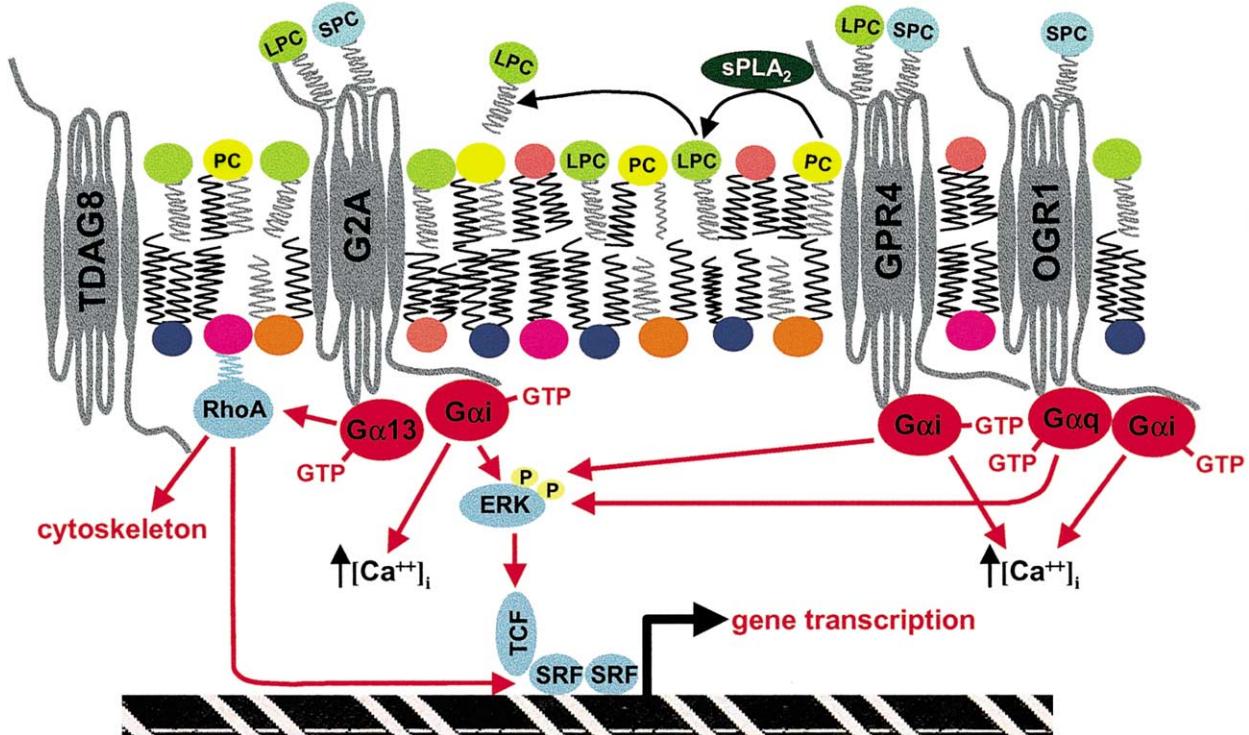


Fig. 2. Overlapping ligand specificities of G2A and its relatives for lysophosphatidylcholine (LPC) and/or sphingosylphosphorylcholine (SPC). G2A elicits RhoA-dependent actin cytoskeleton rearrangement and transcriptional activation of serum response factor (SRF) via G α 13 heterotrimeric G proteins. G2A, GPR4, and OGR1 mediate ERK MAP kinase activation and intracellular calcium increase (\uparrow [Ca⁺⁺]_i) via G α i and/or G α q heterotrimeric G proteins [19,21,29].

While G2A is expressed predominantly in hematopoietic cells including T lymphocytes, it is also transcriptionally induced in response to stress stimuli in B lymphoid cells. G2A couples to G α 13 heterotrimeric G proteins to activate RhoA, leading to the assembly of actin stress fibers and transcriptional activation of serum response factor [23,24]. Although studies of biological and signaling events downstream of G2A suggested that transcriptional induction of this GPCR may function at the level of the actin cytoskeleton to regulate lymphocyte activation/proliferation and/or cellular migration, more recently, genetic ablation of G2A function in mice has revealed a role for G2A in the homeostatic regulation of lymphocyte pools and the maintenance of immunological tolerance. G2A-deficient mice develop a late-onset systemic autoimmune syndrome exhibiting many features in common with the human autoimmune disease SLE, and their T lymphocytes exhibit hyperproliferative responses to antigen receptor stimulation [25].

Based upon protein sequence homology, G2A is closely related to three GPCRs (GPCR 4, GPR 4; ovarian cancer related GPCR 1, OGR 1; and T cell death associated GPCR 8, TDAG 8) [26–28] with overlapping expression in cells of the innate (macrophages and dendritic cells) and adaptive immune systems (Kabarowski JHS, Lim R and Witte ON, unpublished observations). We have identified LPC as a ligand for G2A and established G2A, GPR 4, and OGR 1 as representative members of a receptor family with ligand specificities towards the phosphorylcholine-containing lysophospholipids LPC and SPC [19,21,29]. These receptors mediate intracellular calcium flux and ERK MAP kinase activation in response to their ligands via G α i or G α q heterotrimeric G proteins (Fig. 2). Although no direct binding studies have been reported, TDAG 8 is activated by the glycosphingolipid psychosine (D-galactosyl- β -1,1'-sphingosine) and several related lipids [30]. TDAG 8 was originally identified by virtue of its transcriptional induction during thymocyte apoptosis and is also induced in dexamethasone-treated T cells [28]. It is not clear how, if at all, this receptor–ligand pair could modulate physiological apoptotic responses during T lymphocyte development or immune function. However, we do not rule out the possibility that, as for its relatives, TDAG 8 may exhibit broader ligand specificity than that established thus far.

4. Biological forms of LPC

LPC can exist in several physiological forms, including free, micellar, LDL, bound to hydrophobic serum proteins such as albumin, consumed within immune complexes, and incorporated into plasma membranes. While application of LPC elicits short-term responses via G2A and GPR4 [19,21], it is likely that long-term exposure of cells to exogenously applied LPC results in its conversion to a form with altered receptor binding properties. Indeed,

physiological concentrations of LPC in body fluids are high (up to 100 μ M) and therefore probably exist predominantly in an “inactive” form. A large proportion of this LPC is present in association with hydrophobic proteins such as albumin and within lipoprotein complexes that are incapable of eliciting LPC-dependent biological responses [31] including inhibition of T cell proliferation and ERK MAP kinase activation via G2A (Kabarowski JHS and Witte ON, unpublished observations). It is likely, therefore, that the actions of LPC are spatially and temporally constrained, possibly occurring in an autocrine/paracrine fashion. For example, activation of receptors by LPC produced from plasma membranes of neighboring cells may represent one mode of spatially restricted paracrine action. Temporal restriction of LPC action is also likely imposed by the presence of extracellular hydrolytic enzymes capable of converting this lysophospholipid [32]. These are all important considerations with respect to modulation of immunological processes within the highly organized microenvironments of secondary lymphoid organs and at sites of pathogen encounter by macrophage and dendritic cells. Nevertheless, long-range effects of LPC are recognized, but may be restricted to its transport function as a preferred carrier of certain essential fatty acids to specific tissues [33,34].

5. LPC, T lymphocytes, and autoimmunity

Increased levels of antibodies against LPC in patients with SLE [8,9,35] and the development of systemic autoimmune disease in G2A-deficient animals [25] suggest a pathophysiological connection. How the pathology of this disease could relate to this receptor/ligand pair is likely to be complex considering the multiple susceptibility factors involved in SLE [36]. Further complexity is introduced by the broad cellular involvement in SLE and the presence of related LPC receptors in multiple immune cell types. Nevertheless, several possibilities can be proposed. For example, several studies suggest that endogenously produced LPC may influence T cell responses and that receptor-mediated signals are involved. For example, PLA₂-mediated LPC production is stimulated in T cells following antigen receptor (TCR) cross-linking [37] and may modulate long-term responses of activated T lymphocytes in an autocrine/paracrine fashion. Consistent with this hypothesis, a primary defect in young disease-free G2A-deficient mice is a hyperproliferative response of T lymphocytes to antigen receptor stimulation. While this suggests that loss of G2A-dependent growth inhibitory responses of T lymphocytes to LPC may underlie the age-related autoimmune syndrome developing in G2A-deficient mice, pathophysiological involvement for other immune cell types is likely considering the broad cellular involvement in systemic autoimmune disease. Expression

of G2A in macrophages and dendritic cells expands the repertoire of immunological processes in which G2A may function to include innate immune responses and antigen presentation.

6. LPC as a regulator of innate immunity

The innate immune system is the first line of host defense against infectious challenge and subsequently controls the induction of adaptive immune responses via antigen presentation and cytokine production. Dendritic cells and macrophages are the major APCs of the immune system. APCs present antigenic peptides in the context of major histocompatibility complex to naïve T lymphocytes in the initiation of T-dependent immune responses. Monocyte derived and resident dendritic cells take up antigen in peripheral tissues and migrate to T cell areas of draining lymph nodes where interaction of co-stimulatory molecules on the surface of the APC with their cognate T cell expressed molecules results in activation of antigen-specific T lymphocytes [38]. The induction of these co-stimulatory molecules (B7.1,CD80; B7.2,CD86), as well as other key immunoregulatory receptors and cytokines, requires a “danger signal” identifying an infectious non-self encounter triggered by recognition of “pathogen-associated molecular patterns” such as lipopolysaccharide, bacterial lipoproteins, and bacterial DNA via pattern recognition receptors [39]. Any defect in these processes is likely to deregulate T lymphocyte activation/proliferation and peripheral immunological tolerance.

Therefore, in addition to a direct role for G2A in the modulation of T lymphocyte responses, innate immune responses of macrophages and dendritic cells may be regulated via this receptor in response to local fluctuations in lysolipids encountered during infection and inflammation. It is possible that LPC produced at sites of inflammation and from disintegrating apoptotic/necrotic cell membranes may be a novel “molecular pattern” recognized by G2A to modulate innate immune processes and the initiation and/or resolution of inflammatory responses. For example, G2A may relay signals regulating APC migration between tissue and lymph in response to LPC produced at inflammatory sites through its effects on actin cytoskeleton reorganization [23]. On the other hand, APC/T lymphocyte interactions within lymph nodes may be modulated by autocrine/paracrine production of LPC through G2A to influence the threshold for T cell activation.

In addition to the elimination of pathogens, macrophage recruitment to inflammatory sites is critical for the efficient clearance of necrotic cell debris and apoptotic cells. Recognition and uptake of apoptotic cells by macrophages and newly generated bone marrow derived immature dendritic cells are also important for maintaining peripheral tolerance to self-antigens [40]. By “sampling” and subsequently presenting self-antigens derived from apoptotic

cells, these monocytic derived cells constitute a critical homeostatic mechanism contributing to the suppression of autoreactivity. In addition to alterations in surface expression of phospholipids, their increased oxidation is also a characteristic feature of apoptotic cells, which express oxLDL-specific ligands on their surfaces increasingly recognized as targets of autoantibodies in SLE [41–43]. Indeed, impaired apoptotic clearance can promote the development of autoimmunity, and such a defect can be induced by antibodies against LPC [44,45]. Although increased sPLA₂-mediated LPC production in the outer plasma membrane leaflet of injured and apoptotic cells has been suggested to potentiate binding of the acute phase protein C-reactive protein (CRP) leading to complement activation via the classical pathway and their consequent phagocytosis [46], whether increased levels of cellular LPC are a physiological determinant of apoptotic cell recognition via G2A and/or GPR4 receptors is presently unknown.

7. LPC as an etiological factor in atherosclerosis

The key role of LPC as a major antigenic component of oxLDL has implicated this bioactive lipid in atherosclerosis, the primary cause of heart disease and stroke. However, its role is poorly understood. Although monocyte/macrophage recruitment to the arterial endothelium and uptake of proinflammatory oxLDL and apoptotic cells likely serve a protective function, their progressive accumulation in hypercholesterolemic subjects is believed to initiate the development of atherosclerotic lesions. Later stages of lesion development are associated with T lymphocyte infiltration, their immune activation and cytokine production (TH1/TH2), and humoral responses to antigens, most significantly oxLDL epitopes [4]. Indeed, all aspects of immune function discussed thus far as potential targets of LPC action play some role in atherogenesis. Most notably, macrophage responses to oxLDL such as migration and activation are likely determined to a significant extent by its LPC content, and may therefore be mediated to some degree via G2A and/or GPR4. Monocyte recruitment to lesion-prone arterial sites is also regulated by the expression of surface adhesion molecules and chemokines by endothelial cells in response to inflammatory mediators. Among these, intercellular adhesion molecule-1 (ICAM-1) and monocyte chemotactic protein-1 (MCP-1) expression has been shown to be induced in endothelial cells by exposure to LPC [47,48]. Following their recruitment and migration into the subendothelial space, monocytes differentiate into macrophages and take up oxLDL via scavenger receptors, leading to the formation of cholesterol ester loaded “foam cells.” Here again, the LPC component of oxLDL is a key mediator of this process via up-regulation of scavenger receptor expression on the surface of macrophages [49]. LPC may even propagate the oxidative

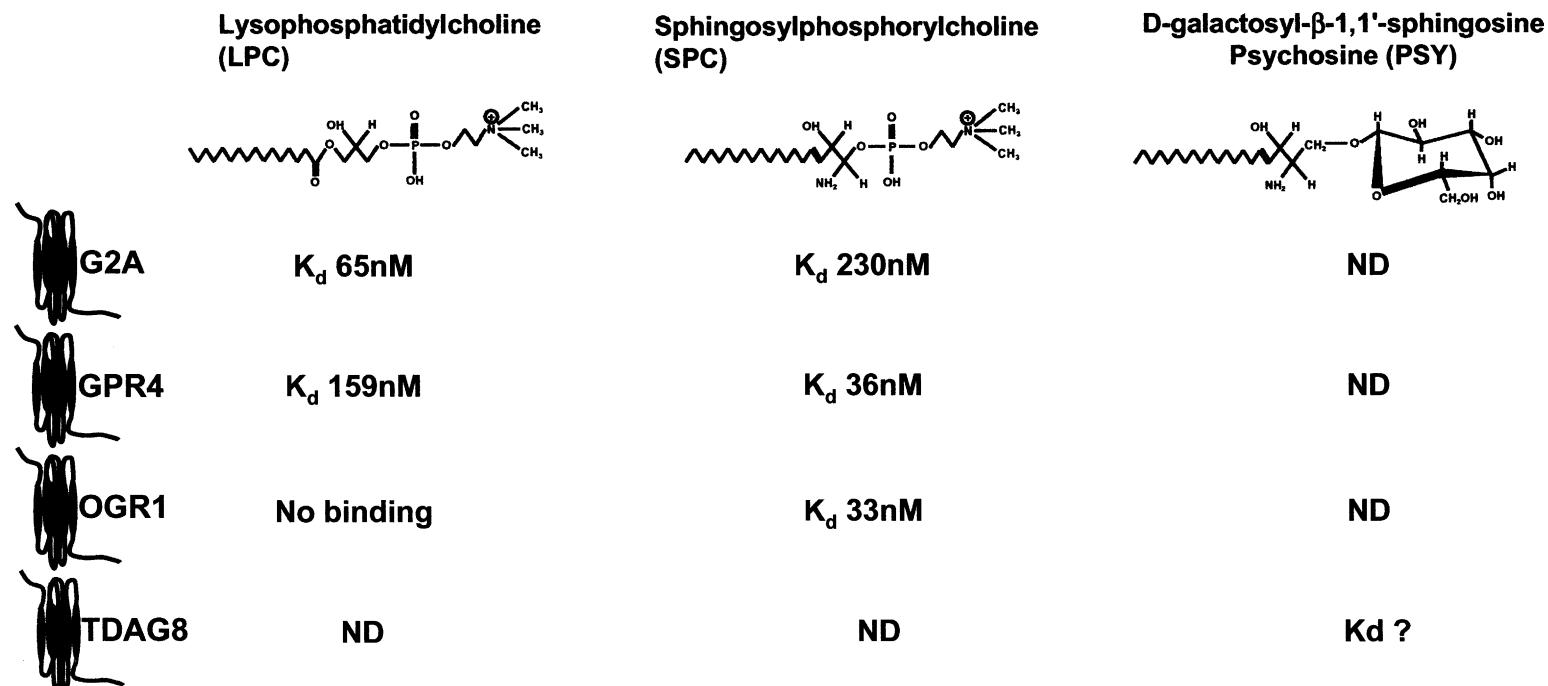


Fig. 3. Ligand binding specificities and affinities of G2A, GPR4, OGR1, and TDAG8. Binding studies were performed on membrane preparations from HEK293 cells overexpressing the indicated receptors [19,21,29]. (ND: not determined; K_d ? no direct binding studies reported.)

processes that lead to the accumulation of oxLDL through its action on endothelial NADH/NADPH oxidase and endothelial/macrophage nitric oxide synthase mediated production of reactive oxygen species [50,51]. Which, if any, of these LPC-dependent cellular events are mediated through G2A and GPR4 are questions currently being addressed in our laboratory. To directly test the pathophysiological role of these receptors in atherogenesis, we are examining the susceptibility of receptor-deficient mice harboring a targeted deletion of the LDL receptor to the development of atherosclerotic lesions.

8. LPC receptors as pharmacological targets

The suitability of GPCRs as targets of drug-based therapeutic intervention promises that further studies of G2A function may uncover clinical benefits of modulating G2A activity in the treatment of chronic inflammatory diseases. The identification of LPC as a ligand for G2A is a major step towards achieving this goal as high throughput screening of chemically modified variants of this lysophospholipid may uncover molecules with agonistic/antagonistic properties towards G2A function. However, receptor/ligand relationships within GPCR families exhibit significant promiscuity, with many receptors recognizing more than one ligand and vice versa. We have found this to be the case for members of the “G2A receptor sub-family,” which exhibit both distinct and overlapping expression patterns together with varying ligand binding affinities (Fig. 3). This suggests that these receptors may perform both redundant as well as unique biological roles, which have important implications for the design of pharmacological agents with the potential to selectively agonize or antagonize one or more of them without suppressing or potentiating immunological processes in a global and deleterious manner. In addition, sequence homology comparisons suggest that one or more orphan GPCRs may prove to be additional members of this receptor sub-family (Ferl GZ, Kabarowski JHS, Radu CG, Roy MP and Witte ON, unpublished observations). Nevertheless, the fact that minor alterations of acyl chain length abolish LPC binding to G2A [19] suggests that other structural modifications may uncover ligand mimetics with higher receptor binding affinities sufficient to effectively compete LPC binding.

An alternative approach to modulate receptor function *in vivo* may be through specific targeting with monoclonal antibodies recognizing extracellular epitopes of the ligand binding pocket formed by the packed α -helical transmembrane domains. Irrespective of methodology, however, an understanding of the mechanisms by which loss of G2A function disrupts immunological tolerance to precipitate autoimmunity and the key cellular and biochemical processes impacted by LPC ligation of its receptors in alternative cell types is a prerequisite for the successful employment of any modulatory agent, as well as the design

of animal model systems for their preliminary evaluation. We hope our future studies as well as those conducted in other laboratories will lead to opportunities for such clinical applications.

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