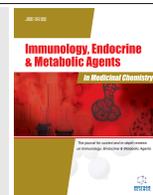

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SCIENCE**

# Anti-Phospholipase A<sub>2</sub> Receptor Autoantibody: A New Biomarker for Primary Membranous Nephropathy

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**Abstract:** Primary membranous nephropathy (also known as idiopathic membranous nephropathy, IMN) is an organ specific autoimmune kidney disease characterized by the development of immune complex deposits in the sub-epithelial spaces, podocyte effacement and glomerular capillary wall thickening in the later stages. Clinical studies have demonstrated that over 70% of patients with IMN possess circulating autoimmune antibodies specifically targeting the phospholipase A<sub>2</sub> receptor (PLA<sub>2</sub>R) on the surface of podocytes. The autoantibodies only bind to the extracellular portion of PLA<sub>2</sub>R under the non-reducing condition, indicating that the epitope in PLA<sub>2</sub>R is conformational requiring specific disulfide bonds to maintain its structure. We recently have successfully located the dominant epitope in PLA<sub>2</sub>R to the extreme N-terminus of the receptor. This finding has opened a new direction for understanding the pathogenesis of anti-PLA<sub>2</sub>R autoantibody induced IMN and offered a strong basis for developing sensitive clinical assays for IMN diagnosis and prognosis, and potentially, new therapeutic approaches for IMN treatment.

**Keywords:** Apheresis column, enzyme-linked immunosorbent assay (ELISA), epitope, idiopathic membranous nephropathy (IMN), immunotolerance therapy, integrin  $\alpha3\beta1$ , kidney, mechanism, pathogenesis, phospholipase A<sub>2</sub> receptor (PLA<sub>2</sub>R).

## INTRODUCTION

Membranous nephropathy (MN) is a common cause of nephrotic syndrome, accounting for ~20-40% of clinical cases in adults over the age of 40 [1-7]. It can be a primary form without identified causes (also known as “idiopathic MN”, IMN), or a secondary form associated with various autoimmune diseases, infections, cancers and exposure to drugs or toxic agents [1, 7]. IMN is an organ specific autoimmune disease. The major antigen responsible for the autoantibody binding in IMN patients was identified to be the phospholipase A<sub>2</sub> receptor (PLA<sub>2</sub>R) [8], an integral transmembrane receptor that binds and removes the secreted phospholipase A<sub>2</sub> enzyme (sPLA<sub>2</sub>) from circulation. Clinical studies have demonstrated that over 70%

of IMN patients possess circulating autoantibodies that specifically target the PLA<sub>2</sub>R expressed on the surface of human podocytes [8] and moreover, the level of autoantibodies in circulation correlates with the severity of proteinuria in patients [9-11]. We recently successfully identified the location of the dominant epitope in PLA<sub>2</sub>R in IMN [12]. This finding has opened a new direction for understanding the pathogenesis of anti-PLA<sub>2</sub>R autoantibody induced IMN and offered a strong basis for developing sensitive clinical assays for IMN diagnosis and prognosis, and potentially, new therapeutic approaches for IMN treatment.

## Pathophysiology of IMN

IMN is an autoimmune kidney disease that accounts for ~80% of all MN cases in adults [1, 13]. It affects 10-12 people per million population. Clinically, about a third of IMN patients undergo spontaneous remission and about 40% progress to the end-stage renal disease in ~10-15 years [14-16]. The common clinical manifestation of IMN is

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edema and proteinuria. Pathologically, IMN is characterized by the development of immune complex deposits in the subepithelial spaces causing a membrane-like thickening of the capillary wall and eventually the glomerular basement membrane (GBM). The name of membranous nephropathy derives from the prominent thickened capillary wall that is visible under light microscope in the later stages of the disease. Currently, the diagnosis of IMN is strictly dependent on immunofluorescence and electron microscopy of the renal biopsies, which detects antibody deposition in the peripheral capillary loop, electron dense deposits in the subepithelial spaces and podocyte foot process effacement, respectively. However, such pathology cannot distinguish between the primary and the secondary forms of MN easily.

The immune deposits in the capillary wall in MN consist of IgG, antigens, and complement components including the membrane attack complex (complement components C5b-9) [17]. IgG4 is usually the dominant IgG subclass in the immune deposits in IMN, although variable amounts of IgG1, IgG2 and IgG3 are also detected. In contrast, IgG1, IgG2 and IgG3 are usually the dominant depositions in the secondary membranous nephropathy [18-21]. The formation of subepithelial immune deposits and complement activation is believed to be the cause of glomerular capillary wall impairment that leads to proteinuria in patients.

Three mechanisms have been proposed for immune complexes formation and deposition in the lesion of MN [4, 22]: 1) immune complexes are formed in circulation and then deposited in the glomerular subepithelial spaces; 2) immune complexes are formed directly *in situ* with local antigens; and 3) immune complexes are formed locally with antigens that are planted in the glomerular subepithelial spaces. These mechanisms have been successfully demonstrated in the animal models such as rabbit chronic serum sickness [23, 24], Heymann nephritis [25] and cationic bovine serum albumin induced rabbit MN [26]. In humans, MN cases with all three of these mechanisms have been demonstrated.

The current understanding of IMN pathogenesis largely derives from an experimental rat model of MN, the Heymann nephritis. In this model, antibodies directly bind to megalin [27, 28], a mem-

brane receptor located on the basal surface of the rat podocytes that forms immune deposits *in situ*, which activates the complement pathway that damages podocyte biology leading to proteinuria [17, 29, 30]. Complement activation and C5b-9 membrane attack complex formation play a key role in causing sublethal podocyte injury and proteinuria in the Heymann nephritis. The sublytic level of C5b-9 complex activates various mediators including phospholipases, protein kinases, cyclooxygenases, transcription factors and cytokines that alter podocyte biology over a period of time. The signals generated by these pathways interfere with podocyte metabolism, structure and function of cytoskeletal proteins, expression and localization of nephrin, turnover of extracellular matrix, and DNA integrity. The assembly of the sublytic C5b-9 complex on podocytes was also shown to up-regulate NADPH oxidoreductase expression and translocation to the cell surfaces. Subsequently, reactive oxygen species are produced and accumulate locally, which leads to oxidation of podocyte membrane lipids, membrane proteins and glomerular basement membrane components [4, 13, 17, 29-31].

Although megalin was identified to be the local podocyte antigen in Heymann nephritis, it is not found in the human podocytes [6, 32]. Two membrane proteins, the neutral endopeptidase (NEP) [33] and PLA<sub>2</sub>R [8] expressed on the basal surface of podocytes have been identified to serve as the local antigens for the neonatal alloimmune and adult IMN respectively. More recently, a new antigen, thrombospondin type-1 domain-containing 7A was also identified to serve as a local antigen in ~ 2.5-5% of adult patients with IMN [34].

### Anti-PLA<sub>2</sub>R Autoantibody and IMN

In 2009, Beck and coworkers made a seminal discovery that, about 70% of patients in the United States with IMN, but not in those with secondary MN or in controls, possess anti-PLA<sub>2</sub>R autoantibodies in circulation that specifically bind to the PLA<sub>2</sub>R expressed in podocytes [8]. The autoantibodies were found to be deposited within the subepithelial immune complexes in patients with IMN, and the eluted autoantibodies from biopsy specimen reacted with PLA<sub>2</sub>R on the Western-blot. The antigen, PLA<sub>2</sub>R was also found in the immune complex deposits at a very high level. Importantly, the autoantibody only recognized the

non-reduced form of PLA<sub>2</sub>R, indicating that the epitope in PLA<sub>2</sub>R is conformational requiring specific disulfide bonds to maintain its structure.

Since this initial discovery, multiple cohort studies have been performed across the world to determine the prevalence of anti-PLA<sub>2</sub>R autoantibodies in IMN patients [9-11, 35-44]. Four techniques have been used to detect the presence of autoantibodies in patients: Western-blotting of PLA<sub>2</sub>R protein under the nonreducing condition (WB), recombinant cell-based indirect immunofluorescence assay (IFA), enzyme-linked immunosorbent assay (ELISA) and immunofluorescence staining of renal biopsies. The available results indicated that the prevalence of anti-PLA<sub>2</sub>R autoantibodies in IMN patients varies significantly in different regions. Specifically, cohort studies showed that the prevalence of anti-PLA<sub>2</sub>R autoantibodies in IMN patients in China [36], Korea [37], Japan [38], Iran [39], Germany [40], UK [9] and Netherlands [41] is 81.7%, 69%, 53%, 75%, 69%, 75% and 74% respectively. Of all the cohort studies, 5 had a sample size over 100 cases, and the average prevalence of anti-PLA<sub>2</sub>R autoantibodies across the world in IMN patients is about 70%. Results from 3 European studies indicated that the level of autoantibodies in circulation correlates with the severity of proteinuria and the clinical outcomes in patients [9-11]; yet results from 2 Asian studies indicated that no significant correlations could be detected [37, 38]. This discrepancy may be caused by different techniques used in the cohort studies. Occasionally, the anti-PLA<sub>2</sub>R antibody was detected in the secondary MN (cancer and hepatitis B) [36] but it is unknown if this is merely a chance occurrence.

The pathogenic role of anti-PLA<sub>2</sub>R autoantibody in IMN was also studied in the kidney transplant recipients. It was reported that IMN could recur one week after kidney transplantation when circulating anti-PLA<sub>2</sub>R antibodies were present at the time of transplantation [45, 46]. It was also reported that some patients with high titers of autoantibodies at the time of kidney transplantation did not have recurrence [45]. Moreover, the recurrence of anti-PLA<sub>2</sub>R autoantibody in IMN patients after kidney transplant [47, 48] was found to be associated strongly with the disease relapse, supporting that the autoantibody serves as the cause of IMN in patients.

Unlike the anti-NEP autoantibody in neonatal alloimmune MN [33], transfer of anti-PLA<sub>2</sub>R patient sera to animals failed to reproduce nephrotic syndrome as that in human. This is likely due to the fact that PLA<sub>2</sub>R is not expressed in the glomerulus of rabbit, rat or mouse [8]. A recent study successfully generated a PLA<sub>2</sub>R knock-in mouse model with human PLA<sub>2</sub>R specifically expressed in the mouse podocytes [49]. Transfusion of human anti-PLA<sub>2</sub>R sera into the PLA<sub>2</sub>R knock-in mice indeed triggered immune complex deposition in the glomerular subepithelial spaces; however, no proteinuria was detected in these animals. This unexpected finding seems to indicate that formation of the immune deposit itself is not sufficient enough to cause proteinuria in patients, and that complement activation or additional mechanisms must be involved, as shown in the Heymann nephritis.

In Heymann nephritis, immune deposits formation and complement activation, as well as a T cell (CD8) involvement are critically required for developing proteinuria in the animals. Depletion of complement components obviated proteinuria in the animal models despite the presence of IgG containing immune deposits in the GBM [50]. The dominant anti-PLA<sub>2</sub>R autoantibody detected in the renal biopsies of IMN patients is IgG4 (other subclasses are also detected at relatively low levels). IgG4 is known to be ineffective to activate the classical complement pathway, suggesting that the mechanism of anti-PLA<sub>2</sub>R autoantibody induced proteinuria in patients may differ significantly from that in the Heymann nephritis. A recent study suggested that aberrant glycosylated IgG4 may play an important role in activating complements via the mannose-binding lectin (MBL) pathway [51], and indeed, MBL components can be found in the immune complex deposits of IMN [52]. In a special clinical case where a patient had recurrent IMN after kidney transplantation, the anti-PLA<sub>2</sub>R autoantibody was identified to be IgG3k, which is capable of activating the classical complement pathway directly [53].

Similar to the type I autoimmune pancreatitis [54], human IMN is an IgG4-related disease with the glomerular immune deposits containing predominantly IgG4-subclass antibodies, indicating the involvement of T helper 2 cells (CD4) in immune response [55, 56]. Clinical studies indeed showed that in IMN patients, the CD4<sup>+</sup>/CD8<sup>+</sup> ratio

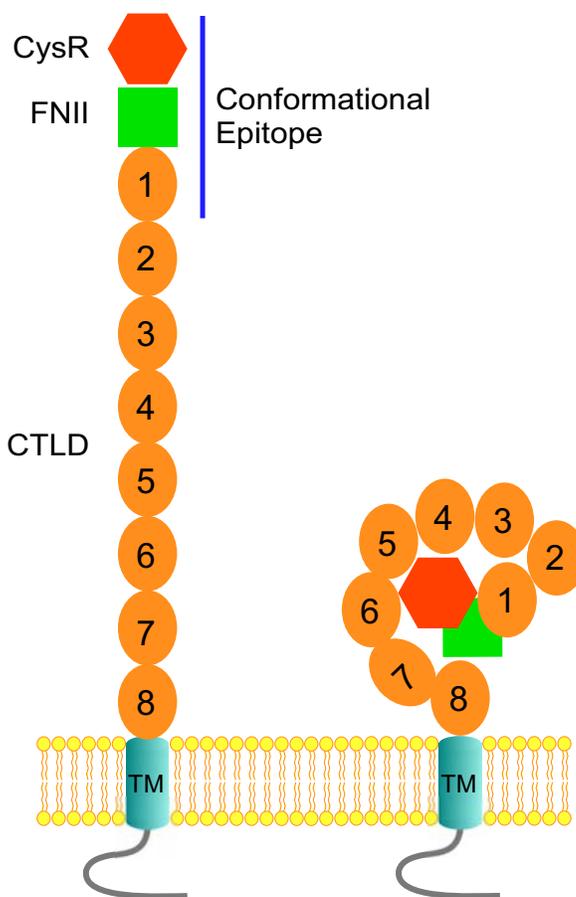
of peripheral T cells and the level of IL-10 and IL-13 mRNA expression in peripheral blood mononuclear cells significantly increased [56]. It is likely that in IMN patients, the antigen primed T helper 2 cells engaged with specific B cells secret IL-10 and IL-13 that activates B cells and promotes IgG4 autoantibody production.

## Structure and Function of PLA<sub>2</sub>R

### Molecular Structure

Two forms of PLA<sub>2</sub>R exist in the body: a membrane associated form (PLA<sub>2</sub>R) and a circulative form (cPLA<sub>2</sub>R) [57, 58]. The membrane associated PLA<sub>2</sub>R is a type I transmembrane glycoprotein (MW ~180 kDa) consisting of a large glycosylated extracellular portion, a single transmembrane helix and a short cytoplasmic tail. The circulating PLA<sub>2</sub>R is a truncated form of the membrane associated PLA<sub>2</sub>R that only contains the extracellular portion due to alternative splicing [57]. In the kidney, both forms of PLA<sub>2</sub>R are present with a ratio of 1.6:1 (PLA<sub>2</sub>R:cPLA<sub>2</sub>R) at the mRNA level [57]. Surprisingly, the anti-PLA<sub>2</sub>R autoantibody only recognizes and binds to the membrane associated but not the circulative PLA<sub>2</sub>R [8], suggesting that the two forms of PLA<sub>2</sub>R may fold differently. The current review focuses on the structure and function of the membrane associated PLA<sub>2</sub>R.

PLA<sub>2</sub>R belongs to the mannose receptor family that consists of four members: the mannose receptor (CD206), Endo180 (also known as the urokinase type plasminogen activator receptor-associated protein uPARAP or CD280), the dendritic cell receptor DEC205 (CD205) and PLA<sub>2</sub>R [59, 60]. All the 4 receptors share a similar three-dimensional protein structure, especially in the extracellular portion, which contains an N-terminal cysteine-rich domain (CysR) followed by a single fibronectin-like type II domain (FnII), and eight to ten C-type lectin-like domains (CTLD), Fig. (1). The protein structure of the individual CysR, FnII and CTLD domain has been solved at the atomic level which showed the critical roles of disulfide bonding in the proper folding of each of the domains [61-63]. Single particle electron microscopy images of the extracellular portion of mannose receptor and Endo180 are also available [64], which indicated that both molecules have two distinctive configurations: a bent conformation with the N-terminal domains folding back toward the middle



**Fig. (1).** The extended (left) and the bent (right) configurations of PLA<sub>2</sub>R in the cell membranes. The immunodominant epitope region for autoantibody recognition is indicated as a solid line.

of the molecule, and an extended conformation with the N-terminal CysR domain pointing outwards from the cell membrane [59]. Transitions between the “bent” and “extended” conformation have been predicted to serve as a general structural mechanism for the mannose receptor family to regulate their ligands binding.

As a member of the mannose receptor family, the extracellular portion of PLA<sub>2</sub>R has also been predicted to have a bent and an extended conformation on the cell surfaces [8] Fig. (1). Whether each of the domains in PLA<sub>2</sub>R is arranged in a similar configuration to that of the mannose receptor in three-dimensional structure is not clear. The difference in autoantibody recognition between the circulating and the membrane associated PLA<sub>2</sub>R seems to indicate that a conformational mechanism is involved in exposing the epitope in PLA<sub>2</sub>R for the autoantibody binding.

## **Biological Function**

Of all the mannose receptor family members, the mannose receptor is known to be involved in the pathogen recognition in the innate immune system; DEC-205 involved in antigen presentation in the dendritic cells; Endo 180/uPARAP involved in regulating extracellular matrix degradation and remodeling. However, the biological function of PLA<sub>2</sub>R has remained largely unclear. PLA<sub>2</sub>R was identified two decades ago based on its high affinity for the venom secreted phospholipase A<sub>2</sub> [65]. Following its cloning, the affinity of PLA<sub>2</sub>R for different sPLA<sub>2</sub> family members was analyzed, and a pancreatic form of sPLA<sub>2</sub> (sPLA<sub>2</sub>-IB) was shown to have the highest affinity [58]. Binding of sPLA<sub>2</sub> to PLA<sub>2</sub>R triggers rapid receptor endocytosis, indicating that PLA<sub>2</sub>R functions as a scavenger receptor for removal of the bound sPLA<sub>2</sub> from circulation [66]. Binding of sPLA<sub>2</sub> to PLA<sub>2</sub>R was also found to activate signaling pathways that triggers cellular responses [67, 68]. Indeed, studies from cultured cells as well as experimental animal models both supported the contention that PLA<sub>2</sub>R is a signaling receptor in cell membranes. Recent reports also suggested that PLA<sub>2</sub>R plays an important role in cell senescence [69] and tumor suppression [70], further supporting the concept that PLA<sub>2</sub>R functions as a signaling molecule in the cell membranes.

PLA<sub>2</sub>R has also been reported to function as an endogenous inhibitor for sPLA<sub>2</sub> in the lung with elevated expression levels during lung infection [71], suggesting its important role in inflammation. However, the biological role of PLA<sub>2</sub>R in the kidney has remained completely unknown despite it being highly expressed in the basal membrane of podocytes [8]. PLA<sub>2</sub>R was predicted to have a protective role for podocyte plasma membranes against being attacked by sPLA<sub>2</sub> enzymes. Intriguingly, *in vitro* biochemical assays showed that although sPLA<sub>2</sub>-IB actively digests phospholipids when mixed with bile, it has minimal activity in digesting membrane phospholipids [72]. sPLA<sub>2</sub>-IB has a molecular weight of 14 kDa and is readily filtered, so it is unlikely to cause damage to the podocyte plasma membranes. Surprisingly, when the PLA<sub>2</sub>R gene was knocked out, mice appeared completely normal with no disease phenotype [73]. These observations seem to suggest that PLA<sub>2</sub>R has a minimal role in podocyte function; however, clinical evidence clearly demonstrated

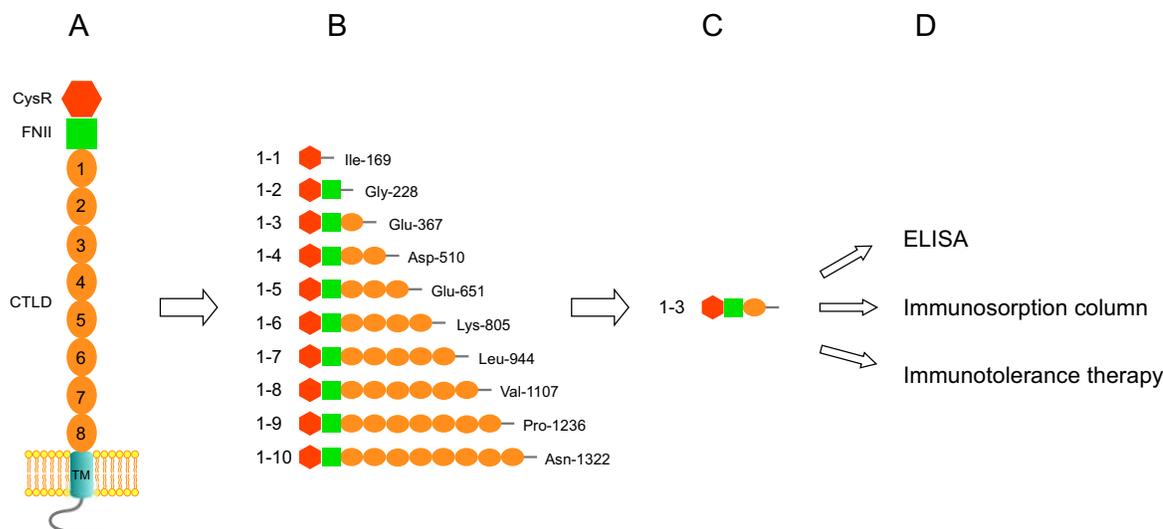
its involvement in mediating podocyte injury in IMN when bound with the anti-PLA<sub>2</sub>R autoantibodies, suggesting that additional physiological roles other than being a functional inhibitor for sPLA<sub>2</sub> in human kidney might be involved.

## **Tissue Distribution**

After its cloning, the distribution of PLA<sub>2</sub>R in human tissue was analyzed by RNA blotting, which showed that PLA<sub>2</sub>R is highly expressed in the pancreas [74], kidney [57, 74], placenta [57, 74], lung [57] and skeletal muscle [57]. Interestingly, PLA<sub>2</sub>R mRNA was only detected in the kidney at a very high level in the human fetus [74]. Lately, PLA<sub>2</sub>R expression was also detected in the human neutrophils [75], gestational tissues [76], the primary human mammary epithelial cells [69], human diploid cells [69] and leukemic cells [77]. Based on the mRNA expression pattern of PLA<sub>2</sub>R in GeneAtlas (provided by Genomics Institute of the Novartis Research Foundation), PLA<sub>2</sub>R is widely expressed throughout the body.

## **Epitope in PLA<sub>2</sub>R for Autoantibody Binding in IMN**

The anti-PLA<sub>2</sub>R autoantibody specifically binds to the extracellular portion of PLA<sub>2</sub>R under the nonreducing condition [8], indicating that the epitope is conformational and requires specific disulfide bonds to maintain its structure. A disulfide bond is formed by sulfhydryl groups of two adjacently located cysteine amino acids in a folded protein. PLA<sub>2</sub>R contains 66 endogenous cysteines throughout the extracellular portion, and disulfide bonds are likely to be present in every one of the 10 domains. The conformational epitope in PLA<sub>2</sub>R could be formed either by a single or by multiple extracellular domains in a folded conformation. Genetic studies have determined that several polymorphisms exist in the extracellular portion of PLA<sub>2</sub>R in IMN patients, especially, M292V and H300D in the CTLD1 domain and G1106S in the linker region between CTLD6 and CTLD7 domain may correlate with the occurrence of IMN in patients with susceptible genetic backgrounds [78, 79]. By using small peptide mapping approach, a number of regions in the PLA<sub>2</sub>R extracellular portion have been predicted to bind to the autoantibodies, suggesting that the epitope in PLA<sub>2</sub>R could potentially be distributed throughout the extracellular portion [80].



**Fig. (2).** Identification of the dominant epitope in PLA<sub>2</sub>R in IMN and its clinical applications. **A.** Topological model of PLA<sub>2</sub>R in the cell membranes. **B.** Truncated PLA<sub>2</sub>R extracellular domains for IMN patient sera probing on the Western-blot. **C.** The identified dominant epitope region. **D.** Clinical applications of the identified epitope in PLA<sub>2</sub>R.

Since PLA<sub>2</sub>R is widely expressed yet IMN is an organ specific disease, we predicted that the PLA<sub>2</sub>R expressed in human podocytes adopt a unique conformation with a specific epitope region highly exposed for the autoantibody binding. To test this hypothesis, we sequentially truncated the extracellular portion of PLA<sub>2</sub>R into 10 protein fragments that contain CysR, CysR and FNII, CysR, FNII and CTLD1-10 domains, Fig. (2). These protein fragments were selectively truncated in the linker regions between each of the domains, to minimize the potential structural disturbance. By probing the truncated PLA<sub>2</sub>R domains expressed in the HEK 293 cells, we successfully determined that the dominant epitope for autoantibody binding is exclusively located in the extreme N-terminus of the receptor, specifically, a region encompassing the CysR-FNII-CTLD1 domain [12]. This epitope was recognized strongly by the sera from various patients that contained anti-PLA<sub>2</sub>R autoantibodies, but not by the sera that were negative of the autoantibodies. Importantly, the isolated epitope region completely blocked the autoantibody binding to the full length PLA<sub>2</sub>R protein on the Western-blot, indicating that the epitope in its native conformation binds strongly to the autoantibody. Our data also demonstrated that all the three domains are required to form an integrated structure for the autoantibody recognition, and neither of the individual domains alone could serve as the epitope on the Western-blot. Therefore, we propose that the epitope in PLA<sub>2</sub>R is

formed by regions from the CysR and the CTLD1 domain with FNII domain serves as a critical structural component to bring the two domains in a close proximity. Indeed, the EM structures of both mannoses receptor and Endo180 showed that the CysR-FNII-CTLD1 domains are compactly folded, indicating that regions in the CysR and CTLD1 domains are close enough to form a conformational epitope [64, 81].

Shortly after our discovery, a similar study on identifying the epitope in PLA<sub>2</sub>R in IMN patients was reported [82]. This study initially determined that the epitope was exclusively located in the CTLD3 domain based on the results using small-angle X-ray scattering, electron microscopy and surface plasmon resonance approaches [83], and in the final published manuscript [82], the epitope was relocated to the CysR domain that corroborated our finding. Specifically, the study determined that a 31 amino acid peptide in the CysR domain forms a major epitope that is recognized by the anti-PLA<sub>2</sub>R autoantibodies. Moreover, the study determined that this epitope could only be maintained under the non-denaturing condition, and that the CTLD3 domain is essentially required for keeping the epitope conformation under the denaturing condition. In our study, the region encompassing the CysR-FNII-CTLD1 domain was well recognized by the autoantibodies from various patients on the Western-blot under the denaturing condition. This discrepancy requires further investigation. Nevertheless, this study clearly sup-

ported our conclusion that the dominant epitope in PLA<sub>2</sub>R in IMN is located at the extreme N-terminus of the receptor.

## **Pathophysiological and Clinical Significance of Epitope Identification**

### ***Pathophysiological Significance***

IMN is an organ specific autoimmune disease. The endogenous antigen, PLA<sub>2</sub>R, is not only expressed in the human kidney, but also highly expressed in the lung, pancreas, placenta, neutrophils, cardiomyocytes and many other tissues. Why the autoantibodies only target the PLA<sub>2</sub>R expressed on the surface of podocytes in the kidney, but not in other organs, has remained completely unclear. Based on the NMR structure of the mannose receptor, PLA<sub>2</sub>R is likely to have an extended and a bent conformation on the cell surface, and the autoantibodies were thought to bind only one of the conformations. Therefore, IMN was predicted to be an autoimmune conformational disease, termed as “conformeropathy” [84], which means the antigen is formed due to a specific conformation of the receptor.

Our finding that the dominant epitope in PLA<sub>2</sub>R is exclusively located at the extreme N-terminus has offered a new level of understanding of the disease pathogenesis and potentially a plausible explanation why IMN is an organ specific autoimmune disease. We predict that the PLA<sub>2</sub>R expressed on the basal surface of the podocytes adopts an extended conformation towards the GBM facing the incoming plasma fluid and as a result is highly accessible to autoantibody binding; yet in other organs, PLA<sub>2</sub>R adopts a bent conformation that precludes the epitope region being recognized by the autoantibodies. In supporting of this hypothesis, we have determined that the extracellular domains of PLA<sub>2</sub>R associate with integrin  $\alpha 3\beta 1$  [3, 85], an adhesion molecule in the podocytes. Integrin  $\alpha 3\beta 1$  is highly expressed in the basal surface of the podocytes, where it mediates podocytes attachment to the GBM. Integrin molecule is known to have a bent conformation in the inactive state, and become extended in the active state when interacting with its ligand [86]. Association of PLA<sub>2</sub>R extracellular domains with the active form integrin  $\alpha 3\beta 1$  is likely to induce the

receptor to adopt an extended conformation that highly exposes the N-terminal dominant epitope for autoantibody binding.

Although the anti-PLA<sub>2</sub>R autoantibody is now considered to be the cause of IMN in patients, how the autoantibody is produced initially in the body has remained unclear. In human glomeruli, PLA<sub>2</sub>R is only expressed in the podocytes [8], and podocytes are epithelial cells separated from circulation by the GBM. Therefore, it is unlikely that exposure of the dominant epitopes in PLA<sub>2</sub>R on the surface of podocytes can serve as the trigger for the autoantibody production since they are inaccessible to the circulating T cells, which help to activate the antigen specific B cells and promote antibody formation and secretion. Identification of the dominant epitope in PLA<sub>2</sub>R has offered an important clue in searching for unknown environmental agents that are potentially to serve as the trigger for the autoantibody production in the body, and subsequently the produced antibodies cross-reacting with the PLA<sub>2</sub>R epitope on the surface of human podocytes through molecular mimicry mechanism.

Comparison of the 31 amino acids in CysR domain against the microbial protein database revealed that a bacterial cell wall enzyme, D-alanyl-D alanine carboxypeptidase, shares a sequence of “LTLENCK” completely with the CysR domain [82], suggesting its potential role to serve as an environmental trigger for IMN. Our study indicated that the CTLD1 domain is critically required for forming the conformational epitope [12], and therefore we compared the amino acid sequence of CTLD1 domain against the NCBI database of non-redundant protein sequences. We found surprisingly that a bacterial subgroup of the C-type lectin-like domain and an invertebrate C-type lectin, CEL-I in the secreted toxin of two ocean organisms Table 1 are about 20-25% homologous to CTLD1. We postulate that exposure to these environmental agents might induce the production of pathogenic anti-PLA<sub>2</sub>R autoantibodies in a genetically susceptible population.

### ***Clinical Significance***

Currently, clinical diagnosis of MN is strictly dependent on renal biopsies that requires signify

Table 1. Human PLA<sub>2</sub>R CTLD-1 domain homologous protein in a group of bacterial and two ocean organisms.

Human PLA <sub>2</sub> R CTLD-1 homologous domain in a subgroup of bacterial C-type lectin-like domain proteins			
Gene ID	Definition	Source organism	Alignment score
gi 78171192	C-type lectin	<i>Chlorobium chlorochromatii</i> CaD3	21.58
gi 33863107	C-type lectin domain-containing protein	<i>Prochlorococcus marinus str.</i> MIT 9313	15.83
gi 86135736	Hypothetical protein MED193_16474	<i>Roseobacter sp.</i> MED193	15.11
gi 87308660	Hypothetical protein WH7805_05191	<i>Synechococcus sp.</i> WH 7805	10.79/17.99
gi 32471540	Hypothetical protein RB1661	<i>Rhodopirellula baltica</i> SH 1	12.23
gi 32472568	Heme/hemopexin utilization protein huxA	<i>Rhodopirellula baltica</i> SH 1	12.23
gi 57506013	TraN protein, homolog	<i>Campylobacter upsaliensis</i> RM3195	12.95
gi 59802590	Cyclin-dependent kinase-activating kinase	<i>Prostheco bacter de jone g e i i</i>	13.67
gi 72003075	C-type lectin	<i>Prochlorococcus marinus str.</i> NATL2A	15.11
gi 78166157	VCBS protei	<i>Chlorobium luteolum</i> DSM 273	15.83
gi 86134618	VCBS	<i>Tenacibaculum sp.</i> MED152	15.83
gi 86749366	VCBS	<i>Rhodopseudomonas palustris</i> HaA2	9.35
gi 87124836	Hypothetical protein RS9917_01402	<i>Synechococcus sp.</i> RS9917	10.79
gi 32477673	Mannan-binding protein MBP (lectin)	<i>Rhodopirellula baltica</i> SH 1	15.83
gi 87310299	Serine/threonine protein kinase	<i>Blastopirellula marina</i> DSM 3645	15.11
gi 67925505	YD repeat	<i>Crocospaera watsonii</i> WH 8501	19.42
Human PLA <sub>2</sub> R CTLD-1 homologous domain in CEL-1 from <i>Cucumaria echinata</i> and Echinoidin from <i>Anthocidaris crassisipina</i>			
Gene ID	Definition	Source organism	Alignment Score
gi 126127	Echinoidin	<i>Heliocidaris crassisipina</i>	25.18
gi 3378108	secreted lectin homolog	<i>Heliocidaris erythrogramma</i>	20.86
gi 17385630	GalNAc-specific lectin	<i>Asterina pectinifera</i>	19.42
gi 21637389	C-type lectin domain protein	<i>Strongylocentrotus purpuratus</i>	20.14
gi 37732133	spEchinoidin	<i>Strongylocentrotus purpuratus</i>	19.42
gi 68357792	PREDICTED: similar to hCG1657150	<i>Danio rerio</i>	12.23
gi 72010149	PREDICTED: similar to secreted lectin homolog; HeEL-1	<i>Strongylocentrotus purpuratus</i>	19.42
gi 72015052	PREDICTED: aggrecan core protein-like	<i>Strongylocentrotus purpuratus</i>	25.18
gi 72085529	PREDICTED: echinoidin-like	<i>Strongylocentrotus purpuratus</i>	22.3
gi 72125841	PREDICTED: aggrecan core protein-like isoform 2	<i>Strongylocentrotus purpuratus</i>	24.46
gi 73959018	PREDICTED: similar to Neurocan core protein precursor (Chondroitin sulfate proteoglycan 3)	<i>Canis familiaris</i>	12.96

cant sample processing time, and the method itself potentially poses a risk of kidney bleeding. The discovery of PLA<sub>2</sub>R as the major antigen in IMN has opened new avenues into the diagnosis and treatment of IMN. Serological tests, such as indirect IFA [35] and ELISA [87, 88], are available on the market with sensitivity for anti-PLA<sub>2</sub>R autoantibodies around 70%. IFA utilizes formalin fixed PLA<sub>2</sub>R expressing HEK 293 cells coated on a coverslip, and ELISA utilizes purified full extracellular portion of PLA<sub>2</sub>R protein, as the target antigens for detecting the presence of autoantibodies in patient sera. It is known that human sera are a mixture containing variety of antibodies; application of such mixture on the fixed human cells will inevitably produce high background. Indeed, IFA could not be used to monitor the autoantibody titer in patient sera and analysis of IFA results is subjective depending on experiences [89]. Identification of the antigenic epitope in PLA<sub>2</sub>R has now made it possible to develop a sensitive and cost-effective epitope specific ELISA assay for detecting the autoantibodies in patient sera. We coated the identified epitope region on an ELISA plate and tested its efficiency for the anti-PLA<sub>2</sub>R autoantibody recognition. Our preliminary data showed that the new assay can detect the presence of autoantibodies up to 1:6,400 dilutions, and moreover, has a ratio of ~10-15 fold of the autoantibody positive vs. negative sera. Our assay has offered a new level of sensitivity and cost effectiveness for detecting the autoantibodies in patient sera, monitoring the disease prognosis, and differentiating the primary and the secondary MN in patients.

Identification of the dominant epitope in PLA<sub>2</sub>R has also offered a strong basis for developing specific treatments for patients with anti-PLA<sub>2</sub>R autoantibody induced IMN. Clinically, IMN patients are treated routinely with high doses of steroids and immunosuppressive agents, which are non-specific and have significant side effects. In some patients, these treatments are not effective. Moreover, the slow decline in circulating autoantibodies after treatment exposes the podocytes to the risk for ongoing injury [90, 91]. If the autoantibodies can be removed selectively from the patient circulation, it may effectively stop the disease progress in the early stage, which would

reverse the abnormalities in kidney function with minimal side-effects [92]. We have successfully built an epitope coated apheresis column and tested its effectiveness for autoantibody removal from the patient sera. Our obtained results have demonstrated that the column has a very high capacity and efficiency to bind and remove the autoantibodies from patient sera, indicating its future potential for IMN treatment. We envision this new treatment will specifically remove the anti-PLA<sub>2</sub>R autoantibodies from circulation without affecting the level of normal antibodies and therefore poses no undesirable immunosuppressive side-effects to the patients.

Alternatively, the identified epitope could be used as a reagent for developing immunotolerance therapies [93, 94] for IMN patients. Immunotolerance is a state of immune unresponsiveness toward a particular antigen due to repeated antigen exposure that results in peripheral T cell tolerance. Immune tolerance therapy intends to reprogram the immune system so that it stops producing autoantibodies. We predict that IMN patients could be exposed to the purified epitope region repeatedly over a period of time until the body tolerates the PLA<sub>2</sub>R epitope without reacting to it. This treatment may potentially stop the anti-PLA<sub>2</sub>R autoantibody production in IMN patients and have long lasting effects.

## CONCLUSION

For the past half century, the pathogenesis of primary membranous nephropathy has been uncertain, and it was referred to as “idiopathic”. With the application of modern technologies in renal disease research, especially protein mass spectrometry, we now have entered into a new era of elucidating the pathogenesis of renal autoimmune diseases. In regard to IMN research, we expect in a near future that a panel of antigens responsible for IMN in patients will be discovered. As a result, we believe that the strict dependence of IMN diagnosis on renal biopsies will one day be fully replaced by the laboratory tests.

## CONFLICT OF INTEREST

The author confirms that this article content has no conflict of interest.

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