

REVIEW

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# New insights into the immune functions of podocytes: the role of complement

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## Abstract

Podocytes are differentiated epithelial cells which play an essential role to ensure a normal function of the glomerular filtration barrier (GFB). In addition to their adhesive properties in maintaining the integrity of the filtration barrier, they have other functions, such as synthesis of components of the glomerular basement membrane (GBM), production of vascular endothelial growth factor (VEGF), release of inflammatory proteins, and expression of complement components. They also participate in the glomerular crosstalk through multiple signalling pathways, including endothelin-1, VEGF, transforming growth factor  $\beta$  (TGF $\beta$ ), bone morphogenetic protein 7 (BMP-7), latent transforming growth factor  $\beta$ -binding protein 1 (LTBP1), and extracellular vesicles.

Growing literature suggests that podocytes share many properties of innate and adaptive immunity, supporting a multifunctional role ensuring a healthy glomerulus. As consequence, the “immune podocyte” dysfunction is thought to be involved in the pathogenesis of several glomerular diseases, referred to as “podocytopathies.” Multiple factors like mechanical, oxidative, and/or immunologic stressors can induce cell injury. The complement system, as part of both innate and adaptive immunity, can also define podocyte damage by several mechanisms, such as reactive oxygen species (ROS) generation, cytokine production, and endoplasmic reticulum stress, ultimately affecting the integrity of the cytoskeleton, with subsequent podocyte detachment from the GBM and onset of proteinuria.

Interestingly, podocytes are found to be both source and target of complement-mediated injury. Podocytes express complement proteins which contribute to local complement activation. At the same time, they rely on several protective mechanisms to escape this damage. Podocytes express complement factor H (CFH), one of the main regulators of the complement cascade, as well as membrane-bound complement regulators like CD46 or membrane cofactor protein (MCP), CD55 or decay-accelerating factor (DAF), and CD59 or defensin. Further mechanisms, like autophagy or actin-based endocytosis, are also involved to ensure podocyte homeostasis and protection against injury.

This review will provide an overview of the immune functions of podocytes and their response to immune-mediated injury, focusing on the pathogenic link between complement and podocyte damage.

**Keywords** Podocyte, Complement, Immune system

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### Background

Podocytes are highly specialized epithelial cells of the glomerulus and represent a major component of the GFB [1]. They have a complex architecture including a large cell body facing the urinary space and an interdigitating network of extensions (primary and secondary processes) terminating as (tertiary) foot processes on the GBM [2].

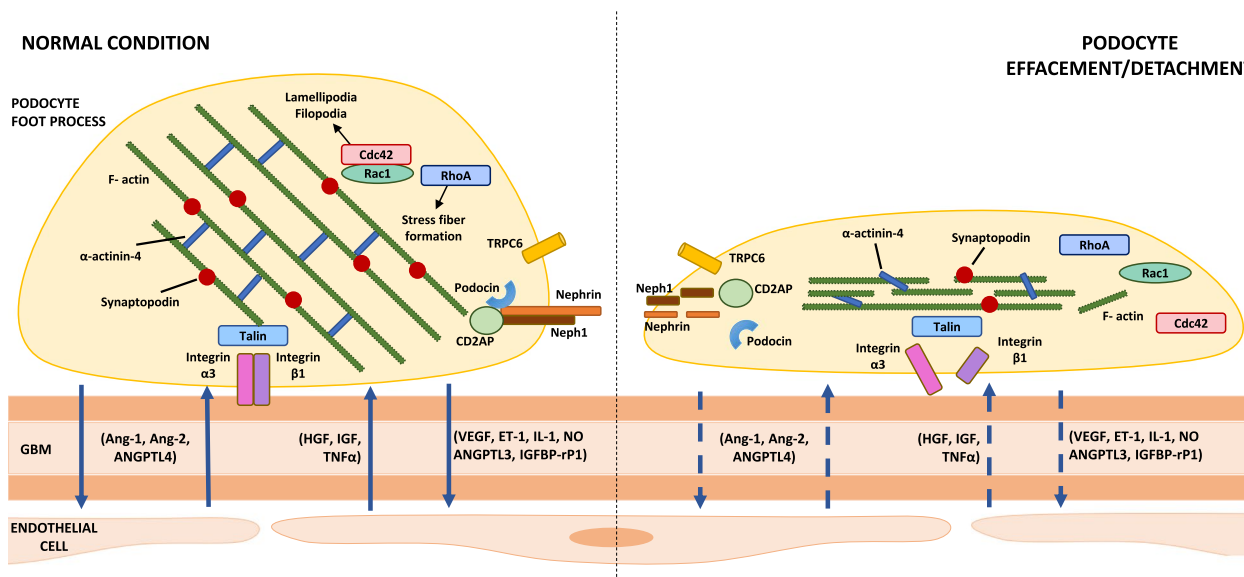
Normal podocyte function is guaranteed by a sophisticated actin cytoskeleton, mainly localized within the foot processes [3]. Podocytes are characterized by a highly complex architecture regulated by multiple proteins, grouped into two main podocyte structures: the slit diaphragm (SD) and focal adhesions (FA). The SD is a unique highly specialized cell–cell junction between two podocyte foot processes (Fig. 1), including key proteins like nephrin, podocin, or synaptopodin [4, 5]. The SD represents not only a size-selective barrier to prevent filtration of large macromolecules but also a signalling platform with critical functions, such as regulation of the actin cytoskeleton and initiation of signalling pathways to modulate the plasticity of foot processes [6]. FA are complex structures which are able to connect the actin cytoskeleton of foot processes to the GBM, thanks to two main molecular components: integrins and GTPases.

Besides contributing to the GFB, podocytes play important functions such as synthesis and repair of the GBM (together with endothelial cells), production of

VEGF, and platelet-derived growth factor (PDGF) [6–9]. Moreover, growing literature suggests that podocytes have many functions of the innate and adaptive immune systems [10–13]. They express cytokine and chemokine receptors to respond to a variety of soluble mediators. They are also able to synthesize inflammatory mediators, such as interleukin-1 (IL-1), which may contribute to local inflammation. Evidence in literature suggests a possible role in the adaptive immune system too, as antigen-presenting cells (APC) to initiate specific T-cell responses, like dendritic cells or macrophages [14, 15].

Furthermore, podocytes express several complement components, such as complement receptor type 1 (CR1) and type 2 (CR2) and complement regulators like CD46, CD55, or CD59, and they can produce complement proteins locally, including complement component 3 (C3) and CFH [16–18]. Nevertheless, the role of complement components expressed or secreted by podocytes in regulation of the local complement reaction is not fully understood.

Podocyte injury is involved in the pathophysiology of several glomerular diseases, like immune-complex glomerulonephritis, minimal change disease (MCD), focal segmental glomerulosclerosis (FSGS), and collapsing glomerulopathy [19, 20], and evidence from the literature suggests that the complement system could be primary or secondary involved in the podocyte damage [21–23].



**Fig. 1** Main components of the slit diaphragm and podocyte-endothelial cell cross talk in healthy versus damaged podocytes. Podocyte slit diaphragm, glomerular basement membrane (GBM), and endothelial cells are the main components of the glomerular filtration barrier. Podocyte effacement/detachment, secondary to mechanical, oxidative, and/or immunologic triggers, is characterized by loss of slit diaphragm integrity, disruption of actin cytoskeleton and focal adhesions, and interruption of the physiological podocyte-endothelial cell cross talk (dashed arrows). Abbreviations: GBM, glomerular basement membrane; Ang, angiotensin; ANGPTL, angiotensin-like protein; IGF, insulin-like growth factor; IGFBP-rP1, insulin-like growth factor-binding protein-related protein 1; ET-1, endothelin-1; HGF, hepatocyte growth factor; IL-1, interleukin-1; NO, nitric oxide; TNF-a, tumor necrosis factor-a; VEGF, vascular endothelial growth factor

### The immune podocyte: innate and adaptive functions

Increasing evidence suggests that podocytes play a role in the innate immune response because of their expression of Toll-like receptors (TLRs), especially TLR4, a subtype able to recognize bacterial lipopolysaccharide (LPS). Those receptors are upregulated in animal models of cryoglobulinemic membranoproliferative glomerulonephritis, and they could mediate glomerular damage by modulating expression of chemokines [12].

TLRs are located on the cell surface or intracellularly and can be expressed by different types of cells, such as dendritic cells, macrophages and monocytes, fibroblasts, B and T cells, and endothelial and epithelial cells. They play an essential role by recognizing pathogen-associated molecular patterns; in particular, cell surface TLRs can mainly recognize microbial membrane components such as LPS, lipids, and proteins, while intracellular TLRs mainly recognize nucleic acids from bacteria and viruses [24]. In addition, TLRs can be activated by endogenous ligands released during stress or tissue injury, such as heat shock proteins, mRNA, and necrotic debris [25]. Cultured human podocytes constitutively express cell surface TLRs (i.e., TLR1, 2, 3, 4, 5, 6, and 10) [26], suggesting a possible role in the defense against microbial agents; however, *de novo* expression of intracellular TLRs subtype has also been reported in podocytes of patients with glomerular disease. In particular, puromycin aminonucleoside (PAN), commonly used to induce a nonimmune podocyte injury *in vitro*, can upregulate TLR9 intracellular expression and activate NF- $\kappa$ B and p38 MAPK in human immortalized podocytes, utilizing endogenous mtDNA as TLR9 ligand to facilitate podocyte apoptosis [27]. This would suggest a bivalent role of TLRs in podocytes, both as major players in response to foreign pathogens and mediators of podocyte damage.

Moreover, podocytes can express MHC class I and II genes [28, 29], as well as B7-1 (or CD80, involved in T-cell activation) [15, 30] and FcRn (IgG and albumin transport receptor, used by podocytes to internalize IgG from the GBM) [31, 32]. In particular, MHC class II expression on podocytes is required for the development of immune-mediated renal injury, as MHC II presentation by podocytes is necessary to induce the CD4+ T-cell-driven glomerular disease [14]. It is reported that these cells can act as antigen-presenting cells (APC), as they can express several macrophagic-associated markers [33, 34], and they are able to process antigens to initiate specific T-cell responses [15], supporting their multifunctional role in the immunological pathogenesis of glomerular diseases.

Furthermore, expression of functional chemokine receptors (CCR4, CCR8, CCR9, CCR10, CXCR1, CXCR3,

CXCR4, and CXCR5) has been demonstrated in cultured human podocytes [35, 36]. Chemokines are small chemoattractant cytokines released by innate immune cells (i.e., neutrophils, eosinophils, macrophages, dendritic cells, natural killer cells), as well as endothelial and epithelial cells. They play a central role in inflammation and immune cell recruitment by guiding circulating leukocytes to inflammation or damage site [37, 38]. They also promote cell growth and tumor angiogenesis and are able to modulate apoptosis by binding G-protein-coupled receptors (GPCRs) on the surface of immune cells. Chemokine receptors are expressed in leukocytes, as well as non-hemopoietic cells, such as endothelial and epithelial cells [39].

CXCR1, CXCR3, and CXCR5 chemokine receptors have been identified in podocytes from kidney biopsies of patients with primary membranous nephropathy (PMN), while they were not expressed in healthy kidneys. Huber et al. suggested that podocyte CXCRs activation may contribute to GFB disruption and onset of proteinuria in PMN through hyperactivation of NADPH oxidases and oxygen radicals production [36].

Podocytes are involved in the inflammatory response of several human glomerulopathies, as suggested by their ability to produce pro-inflammatory cytokines like IL-1 $\alpha$  and IL-1 $\beta$  [40, 41]. It has been reported that they can express inflammasome components, like NOD-like receptor (NLR) family proteins, which contribute to inflammatory response in the local kidney in primary glomerular diseases like lupus nephritis (LN) [42].

Podocytes are also known to secrete and/or express several complement proteins and regulators, suggesting local activation of the complement cascade. Expression of complement genes, including C1q, C1r, C2, C3, C3a receptor (C3aR), C5a receptor (C5aR), C7, CR1, and CR2, has been detected in cultured podocytes under normal physiological conditions, with increased local synthesis of complement proteins following podocyte injury [16, 17]. On the other side, complement regulators have been identified too, both membrane-bound (CD46, CD55, CD59) and soluble (CFI and CFH) forms. In particular, podocytes can express CFH locally to clear subendothelial immune complex deposits [43]. The fact that podocytes are able to produce complement components, including regulators, might have a relevant impact on podocytopathies where the complement system plays a pathogenic role. The balance between local complement activation and regulation is important to maintain the glomerular environment, as podocytes could become both target and source of injury, contributing to local complement activation and amplifying their own damage [44, 45].

**Table 1** Summary of (potential and recognized) podocyte immune functions

Molecules	Expression	Immune function and possible implications	References
CD80 (or B7-1)/CD86 Class I/II MHC	Cultured human podocytes can express antigen-presenting cell molecules	Activation of specific T-cell immune responses in renal diseases	[14, 15, 28–30, 33, 34]
Chemokine receptors (CCR and CXCR)	CCR4, CCR8, CCR9, CCR10, CXCR1, CXCR3, CXCR4, and CXCR5 are expressed in cultured human podocytes CXCR1, CXCR3, and CXCR5 have been identified on podocytes from kidney biopsies of PMN patients	Possible pathogenic role in acute and chronic glomerular inflammation NADPH-oxidases hyperactivation and ROS production — possible contribution to glomerular filtration barrier damage, and onset of proteinuria	[12, 35–39]
Complement system components	Cultured human podocytes can produce and express complement components, including regulators	Possible local activation of the complement cascade	[16, 17, 43]
FcRn	Both in vitro and in vivo podocytes express FcRn	IgG clearance from the glomerular basement membrane, albumin recycling	[31, 32]
Cytokines/growth factors/inflammation components	Both in vitro and in vivo podocytes produce cytokines and growth factors (i.e., TNF- $\alpha$ , IL-1 $\alpha$ and $\beta$ , IL-6, IL-8, VEGF). They can also express inflammasome components (NOD-like receptor family proteins)	Possible role in the local inflammatory response in glomerular diseases	[40–42]
Toll-like receptors (TLRs)	Constitutive expression of cell surface TLRs has been identified on cultured human podocytes De novo expression of intracellular TLRs has been detected in podocytes of patients with glomerular disease (upregulation of intracellular TLR9 with activation of NF- $\kappa$ B/p38 MAPK)	Possible role in the defense against microbial agents Possible role in immune response and glomeruli inflammation	[12, 13, 24–26]

**Abbreviations:** PMN primary membranous nephropathy, FcRn neonatal Fc receptor, VEGF vascular endothelial growth factor, NOD nucleotide-binding and oligomerization domain, NF- $\kappa$ B nuclear factor- $\kappa$ B, MAPK mitogen-activated protein kinase

A summary of the main immune functions of podocytes are summarized in Table 1.

### Podocyte and complement system

The complement system, classically described as part of the innate immune system, represents indeed a functional bridge between innate and adaptive immunity. It consists of more than 30 plasma or membrane-anchored proteins and regulators which play a role in inflammation, opsonization and lysis of pathogens, clearance of apoptotic cells, and enhancement of both innate and adaptive immunity [46–48]. It can be activated by three different pathways, the classical, the lectin, and the alternative pathway [49, 50], which are tightly regulated by several complement components, like the membrane-bound proteins CD46, CD55, and CD59 and the soluble CFH, to prevent uncontrolled complement hyperactivation [51]. All three pathways induce a proteolytic cascade leading to a shared terminal pathway with subsequent membrane attack complex (MAC) assembly in the cell plasma membrane. Once inserted in the lipid bilayer, MAC forms a stable pore with ~10 nm diameter generating several intracellular signals, which have been

characterized by both in vivo and in vitro models as summarized in Table 2 [52].

### Sublytic effects of complement activation on podocytes

Mechanical, oxidative, and immunologic stress can cause podocyte damage and subsequently affect the integrity of glomerular barrier. Complement activation with sublytic MAC formation on podocytes is an example of immunologic stress, which can trigger downstream pathways including protein kinases, lipid metabolism, cytokine production, ROS generation, growth factor signal transduction, endoplasmic reticulum stress, and the ubiquitin–proteasome system, eventually leading to disruption of the podocyte actin cytoskeleton and subsequent cell detachment [53].

More in details, evidence suggests that sublytic amount of MAC on the podocyte surface can induce calcium influx through the membrane pore, as well as calcium release from the intracellular storages, eventually leading to increased intracellular calcium which can activate multiple pathways, such as protein kinase signalling, and in particular protein kinases C (PKC) responsible for membrane vesiculation and internalization of MAC

**Table 2** Signalling pathways activated by MAC (adapted from Takano et al. (2013). *Seminars in Nephrology*. Reference [52])

Pathway	Effects of terminal pathway activation
Intracellular calcium	Calcium influx through MAC and calcium release from intracellular storage sites
Protein kinases	Activation of protein kinase C (PKC), receptor tyrosine kinase (RTK), Ras-ERK, JNK, p38, and ASK1 (HN)
Phospholipases	Activation of phospholipase C (PLC)- $\gamma$ 1, cPLA2, and iPLA2- $\gamma$ (phosphorylation), and AA release
Prostanoids	Upregulation of cyclooxygenase (COX)-2 (cultured podocytes and HN glomeruli) and COX-1 (HN glomeruli), production of prostanoids
ROS	Superoxide production via NADPH oxidase and lipid peroxidation (HN) ROS production via xanthine oxidase pathway (HN) Generation of hydrogen peroxide by cytochrome P450 family of heme protein monooxygenases (cultured podocytes)
Growth factors	Upregulation of platelet-derived growth factor B-chain, HB-EGF (HN), and Ret (HN and cultured mouse podocytes) Increase of p21 and p27 CDK inhibitors and decrease of CDK2 activity Decrease of p57 and increase of Cdc2, cyclins B1, B2, and D1 and phosphorylated histone-3
Transcription factors	Activation of NF- $\kappa$ B (cultured podocytes and in vivo)
DNA damage	Production of interleukin-8 and monocyte chemoattractant protein-1 Increase of p21, p53, GADD45, and checkpoint kinase-1 and kinase-2 (cultured podocytes and HN)
Endocytosis	Endocytosis (podocyte)
Ectocytosis	Ectocytosis in membrane vesicles (urinary space)
ER stress	Damage of ER membrane and unfolded protein response induction Upregulation of ER chaperones, PERK stimulation, eukaryotic translation initiation factor-2 $\alpha$ subunit phosphorylation, and reduction of protein synthesis
Ubiquitin–proteasome system	Polyubiquitination of glomerular proteins (HN) Upregulation of ubiquitin proteasome system (cultured podocytes)
Podocyte cytoskeleton	Disassembly of F-actin filaments and focal adhesion complexes Increase of RhoA and decrease of Rac1 and Cdc42 activities (cultured podocytes) Foot process effacement by induction of active RhoA in podocytes (in mice) TRPC6 upregulation (cultured podocytes)
Slit diaphragm	Decrease of nephrin mRNA and protein (HN) Dissociation of nephrin from actin cytoskeleton and loss of slit diaphragm integrity Alteration of podocin location and nephrin dissociation from podocin
Cell cycle	Increased DNA synthesis without cell proliferation (podocyte)
Anti-apoptosis	PI3K/Akt activation, Bad phosphorylation, and dissociation of the Bad/Bcl-XL complex Upregulation of caspase-8 inhibitor and cFLIPL and downregulation of FasL
Pro-apoptosis	DNA damage via apoptosis regulating proteins (podocytes)

**Abbreviations:** MAC membrane attack complex, PKC protein kinase C, RTK receptor tyrosine kinase, Ras-ERK Ras-extracellular signal regulated kinase, JNK c-Jun N-terminal kinase, ASK1 apoptosis signal-regulating kinase-1, HN Heymann nephritis, cPLA2 cytosolic phospholipase A2, iPLA2- $\gamma$  independent PLA2- $\gamma$ , AA arachidonic acid, NADPH nicotinamide-adenine dinucleotide phosphate, ROS reactive oxygen species, HB-EGF heparin-binding epidermal growth factor-like factor, Ret glial cell-derived neurotrophic factor receptor tyrosine kinase, CDK cyclin-dependent kinase, NF- $\kappa$ B nuclear factor- $\kappa$ B, GADD45 growth-arrest DNA damage-45, ER endoplasmic reticulum, PERK protein kinase R-like ER kinase, TRPC6 transient receptor potential channel 6, PI3K phosphatidylinositol 3-kinase, Akt protein kinase B, Bad BCL2-associated agonist of cell death, Bcl-XL B-cell lymphoma-extra large, cFLIPL cellular FLICE-inhibitory protein long form, FasL Fas ligand

channels [52, 54–57], as suggested by reduction of MAC endocytosis by inhibiting PKC pathway [58].

It is well known that Ca<sup>2+</sup> signalling in healthy podocytes is mainly mediated by angiotensin II and TRPC5 and 6 (nonselective cationic channels, downstream of angiotensin II signalling) [59]; interestingly, TRPC6 can play a dual role, as it has been shown that acute activation of this channel is able to protect podocytes from complement-mediated injury, while gain-of-function mutations/chronic hyperactivation can affect the SD and/or foot processes morphology leading to glomerular diseases, such as FSGS [60].

It has also been described that sublytic MAC can induce transactivation of receptor tyrosine kinases at the plasma membrane of cultured podocytes, resulting in

activation of the Ras-extracellular signal-regulated kinase (ERK) pathway and phospholipase C- $\gamma$ 1. Transactivated receptor tyrosine kinases could play as scaffold for proteins assembly and/or activation, inducing activation of downstream pathways, either dependently or independently the increased cytosolic calcium levels [54, 61].

Other pathways activated by MAC formation on podocyte surface involve arachidonic acid (AA) release by cytosolic phospholipase A2- $\alpha$  (cPLA2), inducing AA metabolism to prostanoids, as described by Cybulsky et al. [62]. Eicosanoids can play a role in complement-mediated podocyte injury, as supported by experimental models of membranous nephropathy. Despite the exact mechanisms of glomerular damage are still unclear, cytotoxic consequences of cPLA2 activation could include

release of free fatty acids and lysophospholipids, as well as ions influx, which could ultimately affect the energy machinery [63].

ROS production has also been described in podocytes exposed to sublytic amounts of MAC; both cultured and in vivo podocytes express components of the NADPH oxidase, a complex enzyme able to reduce molecular oxygen to the superoxide anion, which is further metabolized to other ROS [52]. Lipid peroxidation and changes in the podocyte membrane composition, as well as in the GBM components, have been reported as consequence of ROS production. Moreover, inhibition of ROS and/or lipid peroxidation resulted in reduced proteinuria in animal models of membranous nephropathy, suggesting their pathogenic role in glomerular damage [64].

Endoplasmic reticulum (ER) stress with accumulation of misfolded proteins and subsequent increase of the ubiquitin–proteasome system has been reported as additional response to complement-mediated injury, as possible protective response of podocytes to ongoing complement attack [65].

Sublytic MAC deposition on podocytes can also induce DNA damage, both in vitro and in vivo models, as demonstrated by Pippin et al. [66]. The authors also described that sublytic MAC-induced podocyte injury was associated with an increase in specific cell cycle-related genes, including p53, p21, growth-arrest DNA damage-45, and checkpoint kinase-1 and 2, leading to cell cycle arrest and podocyte growth suppression. This could explain why podocyte proliferation is limited following immune-mediated injury.

#### **Consequences of complement activation on podocyte energy metabolism**

The effects of complement activation on podocyte energy machinery are not fully understood. Brinkkoetter et al. demonstrated that podocyte metabolism is somewhat different from other type of cells, as it primarily relies on anaerobic glycolysis and the transformation of glucose to lactate [67]. More in details, the authors showed a significantly lower mitochondrial density per cell area, compared to other type of renal cells (i.e., renal tubular cells). Also, glomeruli stained for mitochondrial enzyme superoxide dismutase 2 (SOD2) and the glycolytic enzyme pyruvate kinase M2 (PKM2) confirmed the perinuclear localization of mitochondria (and their almost complete absence in secondary and tertiary processes), while PKM2 was ubiquitous, suggesting podocyte processes as a large compartment of anaerobic glycolysis. They also used Tfam (mitochondrial transcription factor A) knockout mice to demonstrate that loss of mitochondrial transcription and lack of the oxidative phosphorylation machinery do not induce podocyte disease. In addition,

transient knockdown of Tfam in human podocytes significantly reduced mitochondrial respiration, while anaerobic glycolysis was significantly increased allowing a normal podocyte function.

It has been demonstrated that sublytic complement-mediated injury induces reduction of intracellular ATP, in addition to reversible disruption of actin stress fibers and focal adhesions, mainly due to dephosphorylation (instead of degradation) of focal contact proteins, as described by Topham et al. using an in vitro model of rat podocytes [68]; however, the precise mechanisms need to be clarified. Also, complement activation on podocytes can cause nephrin dissociation from the actin cytoskeleton with disruption of the slit diaphragm, GFB damage, and subsequent onset of proteinuria, as suggested by the Heymann nephritis (HN) model [54, 61].

#### **Complement-mediated injury and podocyte response**

Podocytes rely on several adaptive mechanisms to mitigate complement-mediated injury. Autophagy, a highly conserved mechanism of lysosome-mediated degradation of damaged organelles or nonfunctional proteins, is enhanced after sublytic complement damage in mouse podocytes, whereas its inhibition amplifies complement-mediated cell injury [69]. Liu et al. investigated the role of autophagy in PMN, comparing podocytes from PMN patients to cultured mouse podocytes exposed to sublytic complement activity, and they found impaired autophagy in podocytes from PMN patients, characterized by intracellular accumulation of p62 (marker of impaired autophagy) and increase in autophagic vacuoles [70].

Podocyte-derived VEGF has also a bivalent function, as it is described that its overexpression can cause a collapsing glomerulopathy, while its inhibition is associated with GFB disruption, proteinuria, and possible development of thrombotic microangiopathy as well [71]. The putative mechanism is that, in normal conditions, VEGF signalling can regulate complement activity on podocytes and protect them from complement-mediated injury by increasing local CFH production, while its inhibition would provoke reduced levels of CFH, and podocytes would become more vulnerable to the injury.

More recently, new interesting mechanisms have been described to protect podocytes from injury, as reported by Medica et al. using a co-culture model of glomerular endothelial cells and podocytes. In particular, they demonstrated that extracellular vesicles derived from endothelial progenitor cells and involved in intercellular crosstalk (by transferring of proteins, lipids, and genetic material) are able to protect both glomerular endothelial cells and podocytes from complement (C5a)- and cytokine-mediated injury [72]. In particular, they showed that pre-stimulation of endothelial cells with extracellular

vesicles prevented podocyte apoptosis and GFB disruption, and this protective effect could be mainly secondary to RNA transfer from the extracellular vesicles to damaged endothelial cells and podocytes.

Despite a tight surveillance of the complement system, including the activity of soluble and membrane-bound regulators, together with the protective mechanisms previously described to escape the injury, unrestricted complement activation can exceed those regulatory mechanisms, causing host tissue injury, as reported in various diseases including glomerulonephritis [73], hemolytic uremic syndrome (HUS) [74], sepsis [75], systemic lupus erythematosus [76], rheumatoid arthritis [77], organ transplant rejection [78], and age-related macular degeneration [79].

### Summary and conclusions

Podocytes play a critical role to ensure the glomerular homeostasis. Over the years, growing literature highlighted the multiple and complex biological functions of these pericytes-like epithelial cells, which are much more than a supporting component of the GFB [1, 80–82].

Several authors described them as “immune podocytes,” to underline their properties as both innate and adaptive immune cells [10, 13, 15]. Understanding their complex biology is essential to unravel the pathogenic mechanisms of several glomerular diseases, where podocyte injury represents a common denominator.

The role of the complement system in podocyte injury has also been evaluated in a multitude of kidney disorders, such as membranous nephropathy, lupus nephritis, HUS, FSGS, and several more [45, 83–90]. The effects of complement activation on podocytes can vary based on the disease pathophysiology, as well as based on the initial trigger, which could induce lytic versus sub-lytic effects. Interestingly, podocytes have developed several protective mechanisms to escape the complement attack, such as autophagy, internalization mechanisms like endocytosis, and expression of complement regulators, and the balance between injury and defense mechanisms can ultimately determine the destiny of the podocyte cell [65, 69, 91].

Future studies, both *in vitro* and *in vivo*, are needed to better understand the role of complement activation in podocytopathies and the rationale for the use of anti-complement therapies in conditions where the complement system appears as main driver of the disease.

### Abbreviations

APC	Antigen-presenting cells
ATP	Adenosine triphosphate
BMP7	Bone morphogenetic protein 7
C1q	Complement component 1q
C1r	Complement component 1r

C1s	Complement component 1 s
C2	Complement component 2
C3	Complement component 3
C3a	Complement component 3a
C3aR	Complement component 3a receptor
C4	Complement component 4
C5a	Complement component 5a
C5aR	Complement component 5a receptor
C6	Complement component 6
C7	Complement component 7
CCR4	CC chemokine receptor 4
CCR8	CC chemokine receptor 8
CCR9	CC chemokine receptor 9
CCR10	CC chemokine receptor 10
CD80	Cluster of differentiation 80
CFH	Complement factor H
CFI	Complement factor I
cPLA2	Cytosolic phospholipase A2
CR1	Complement receptor type 1
CR2	Complement receptor type 2
CXCR1	CXC chemokine receptor 1
CXCR3	CXC chemokine receptor 3
CXCR4	CXC chemokine receptor 4
CXCR5	CXC chemokine receptor 5
DAF	Decay-accelerating factor
ERK	Ras-extracellular signal-regulated kinase
FA	Focal adhesions
FcRn	Neonatal Fc receptor
FSGS	Focal segmental glomerular sclerosis
GBM	Glomerular basement membrane
GFB	Glomerular filtration barrier
GPCR	G-protein-coupled receptor
HN	Heymann nephritis
HUS	Hemolytic uremic syndrome
IL-1 $\alpha$	Interleukin-1 $\alpha$
IL-1 $\beta$	Interleukin-1 $\beta$
LN	Lupus nephritis
LTBP1	Latent transforming growth factor $\beta$ -binding protein 1
MAC	Membrane attack complex
MAPK	Mitogen-activated protein kinase
MCD	Minimal change disease
MCP	Membrane cofactor protein
MHC	Major histocompatibility complex
NADPH	Nicotinamide adenine dinucleotide phosphate
NF- $\kappa$ B	Nuclear factor kappa-light-chain-enhancer of activated B cells
NLR	NOD-like receptor
PAN	Puromycin aminonucleoside
PDGF	Platelet-derived growth factor
PKC	Protein kinase C
PMN	Primary membranous nephropathy
PKM2	Pyruvate kinase M2
ROS	Reactive oxygen species
SD	Slit diaphragm
SOD2	Superoxide dismutase 2
Tfam	Mitochondrial transcription factor A
TGF $\beta$	Transforming growth factor $\beta$
TLRs	Toll-like receptors
TRPC5	Transient receptor potential canonical 5
TRPC6	Transient receptor potential canonical 6
VEGF	Vascular endothelial growth factor

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