Introduction

Extracellular vesicles (EVs) refer to all endogenously produced membrane-bound vesicles that are released from cells into the extracellular space [1]. EVs contain various molecules, such as DNA, messenger RNA (mRNA), proteins, lipids, and microRNAs (miRNAs). Different subtypes of EVs have been described including exosomes, microvesicles or microparticles, apoptotic bodies, ectosomes, and oncosomes based on their origin, size, contents, and biogenesis (Table 1). In the narrow sense, EVs usually refer to exosomes or microvesicles. In this review, we will use the compre-
hensive term ‘EVs’ to focus on exosomes and microvesicles because of overlapping characteristics among the different subtypes. EVs were initially discovered over 30 years ago, but it was thought that their physiological role was limited to the excretion of intracellular or membrane components of cells. However, recent advancements have shown that EVs are involved in multiple biological processes such as immune modulation, hemostasis, and tissue proliferation/regeneration, which can affect the development and regeneration of organs [1–3]. EVs are also widely found in pathologic conditions. Their presence in any body fluid expands the diagnostic role of EVs as biomarkers in various types of diseases. EVs also contribute to disease progression by affecting the level of inflammation, thrombosis, and tumor-ogenesis [4,5]. On the other hand, EVs have received a lot of medical attention as a potential therapeutic vehicle based on their capacity of shuttling proteins and genetic materials. In this review, we will focus on the role of EVs in renal physiology and disease processes as well as their potential applicability as diagnostics and therapeutics in multiple renal diseases.

**Physiologic role of extracellular vesicles in kidneys**

In physiologic conditions, the major biological functions of EVs are to get rid of unwanted substances from host cells. EVs can also transfer important biological information to their recipient cells by delivering genetic material, proteins, lipids, and receptors. The conveyance of RNAs or miRNAs can have a substantial effect on the recipient cells by re-programming their genetic characteristics. Below, we will discuss in detail the biological function of EVs in renal physiology (Fig. 1A [6]).

**Homeostasis and cell survival**

One of the most important biological roles of EVs is in the removal of intracellular toxic materials to extracellular spaces. As cells go through multiple biological processes, they build up damaged organelles and cellular waste, which can induce cellular stress leading to cell death or inflammation. Eukaryotic cells have developed a self-defense mechanism for the removal of intracellular waste—the secretion of EVs into the extracellular space [7]. In this way, cells can effectively eliminate potentially harmful chromosomal DNA fragments through exosomes. The inhibition of this process can induce an innate immune response, DNA damage, and apoptosis in normal human cells. Kidney cells are relatively susceptible to biological stress. Under hypoxic conditions, renal tubular cells proactively augment the amount of exosome secretion and alter the composition of exosomes to meet specific biological needs [8]. EV uptake into recipient cells is also a very important biological process in intercellular communication. The most common mechanism of cellular uptake is endocytosis whereby the EVs are engulfed by recipient cells [9]. The balance between EV release and uptake depends on the physiologic condition of the parent/recipient cells, the type of parent/recipient cells, and the recognition of ligands/receptors on the EVs and recipient cells [10].

**Inflammation/immunomodulation**

EVs are a miniature version of their parent cells, so their immunologic function significantly depends on their origin and the microenvironment their parent cells are exposed to. For example, EVs from dendritic cells can function as
antigen-presenting vesicles through the direct presentation of peptide-major histocompatibility complex (MHC) complexes to T cells [11]. These peptide-MHC complexes can be transferred to other recipient cells, indirectly presenting antigens, and subsequently leading to immune cell stimulation. On the other hand, EVs secreted by tumor cells can mediate an immuno-suppressive response by switching monocytes and T cells to tolerogenic subtypes, inhibiting the differentiation of monocytes, or inducing the apoptosis of bystander T cells, so-called activation-induced cell death [12]. The physiological state of the parent cells significantly affects the content of the EVs. For example, exosomes derived from hypoxic tumor cells could induce higher levels of differentiation and the activation of myeloid-derived suppressor cells compared to exosomes derived from cells under normoxic conditions [13].

**Antimicrobial effect**

The urinary tract system is constantly exposed to microorganisms from the exterior environment, putting it at high risk of urinary tract infection. However, most of the urinary tract, except the urethra, generally remains sterile. This resistance to infection is mediated by several factors. Traditionally, anatomical barriers, such as the glycoprotein plaque and a layer of hydrated mucus, as well as immunologic barriers from various resident immune cells and the epithelial cell lining, were considered to be major host defense mechanisms [14]. Recently, EVs were found to serve a significant role against infection within the urinary tract. A proteomic study showed that enriched innate immune proteins including calprotectin and lysozyme C in urinary EVs could mediate an antimicrobial effect [15]. Even though autophagy is a well-developed cellular defense mechanism to clear foreign pathogens, some
bacteria have evolved defense mechanisms to overcome autophagy by neutralizing lysosomal pH. A study by Miao et al. [16] shows that bladder epithelial cells can overcome this phenomenon through exosomes—by expelling exosome-encased bacteria from the intracellular space.

Kidney regeneration/repair

Mesenchymal stem cells (MSCs) are well-known for their regenerative and reparative potential in different organs including the kidneys [17]. However, the reparative potential of MSCs is mediated through EVs rather than the differentiation potential of the MSCs themselves. MSC-derived EVs contain different types of antioxidants and growth factors to stimulate the differentiation of resident progenitor cells. Delivery of genetic materials like DNA, miRNAs, or mRNAs through EVs can also introduce the reparative potential of parental MSCs to recipient cells. In a study by Ranghino et al. [18], glomerular MSC-derived EVs mediated tubular epithelial cell regeneration and alleviated ischemic acute kidney injury (AKI) by transferring mRNAs and miRNAs with potential proregenerative effects.

Hemostasis and platelet aggregation

An ample number of publications have reported the coagulant properties of plasma EVs in humans [19,20]. EVs originating from platelets, endothelial cells, or leukocytes were found to be involved in the course of hemostasis. These EVs can initiate the coagulation process by exposing phosphatidylserine (PS) on their surfaces to allow binding sites for coagulation factors [21]. Another mechanism is through the expression of tissue factors on their surfaces, initiating the extrinsic pathway of coagulation and leading to thrombin burst and platelet clot formation [22]. A study by Yu et al. [23] showed that patients with diabetic kidney disease had a significantly higher level of PS-expressing microvesicles compared to control groups. They also found that higher levels of PS-positive microvesicles were associated with hypercoagulable status as well as worse renal function. A similar finding was shown in patients with immunoglobulin A nephropathy [24].

Recently, several studies have revisited the procoagulation property of EVs because of technical improvements in blood collection, plasma preparation, and EV isolation [25]. Interestingly, plasma EVs isolated from healthy individuals promote fibrinolysis more dominantly than coagulation. This discrepancy suggests an ambivalent role of EVs in hemostasis processes, which depends on the microenvironment. In addition, proper EV sample preparation is important for a better understanding of their role in the human body.

Vessel integrity and revascularization

Endothelial cell-derived EVs help to maintain vessel integrity and promote vascular endothelial cell survival by removing cell-death signals and releasing excess complements from vascular endothelial cells [26]. In damaged vessels, platelet-derived EVs induce the adhesion of endothelial cells to the extracellular matrix, which helps the regeneration of endothelium and attenuates vascular permeability [27].

Tumor-derived EVs are well-known for their angiogenic properties, but EVs can also induce angiogenesis under physiologic conditions. Endothelial cell-derived EVs modulate angiogenesis by promoting endothelial cell invasion and capillary-like structure formation [28]. Recently, the transfer of certain miRNAs, including miR-126 or miR-124, using endothelial cell-derived EVs and recipient endothelial cells has been found to have a significant role in angiogenesis [29,30]. MSC-induced EVs also induce angiogenesis by transferring proteins like phosphorylated signal transducer and activator of transcription-3 or nuclear factor-kB pathway-associated proteins, leading to the transcription of proangiogenic proteins [31,32]. MSC-derived EVs can also transfer several growth factors including epidermal growth factor and vascular endothelial growth factor to endothelial cells promoting angiogenesis.

More detailed information about the angiogenic mechanism of EVs from multiple cellular origins can be found in a review article by Todorova et al. [2].

Electrolyte and water balance

Sodium/water reabsorption is one of the major functions of renal tubular cells. Two-thirds of filtrated sodium is reabsorbed in proximal tubular epithelial cells and fine-tuning for sodium reabsorption occurs in the distal tubule and collecting ducts. Therefore, interactive communication between proximal and distal/collecting tubular cells is important for proper maintenance of electrolyte balance and volume control. A study by Jella et al. [33] showed that exosomes from proximal tubular cells can regulate epithelial sodium channel
activity in the distal tubule and collecting ducts through the exosomal delivery of glyceraldehyde 3-phosphate dehydrogenase. Upon exposure to vasopressin, cortical collecting duct cells increase the production of aquaporin water channels within themselves as well as within their exosomes [34]. These exosomes could transfer functional aquaporin channels to other tubular cells, increasing their water transport capacity. By mediating inter and intracellular communication of renal epithelial cells among the different nephron segments, exosomes provide a significant role in electrolyte and fluid balance in the human body.

**Extracellular vesicles as mediators of kidney disease**

In pathologic conditions, the production of EVs significantly increases, contributing to the initiation and propagation of disease through immunomodulation, thrombogenesis, and oncogenesis. The detailed role of renal EVs in different disease courses are discussed below (Fig. 1B).

**Glomerular disease**

In advanced glomerular disease, tubular damage and fibrosis are the leading mechanisms of renal dysfunction. A recent study by Jeon et al. [35] showed that miRNAs in EVs released by injured podocytes can induce tubular epithelial cell apoptosis, suggesting possible glomerulotubular crosstalk. This study also showed that the miRNA profile in EVs from injured podocytes are different from those of non-injured podocytes, and miRNA mimics of injured podocytes could reproduce a proapoptotic effect on renal tubular epithelial cells. In diabetic nephropathy, exposure to high glucose levels can induce EV production from podocytes. Tubular epithelial cells uptake these podocyte-derived EVs and generate tubular fibrosis through p38 phosphorylation [36].

**Acute kidney injury**

A recent miRNA sequencing study showed that urinary exosomal miRNAs have characteristic profiles and functional properties that depend on the AKI stage [37]. Several exo-miRNAs (miR-16, miR-24, and miR-200c) were increased during the early injury state, and the expression of their target mRNAs was significantly affected in the renal medulla. During the early recovery phase, exo-miRNAs that share Zeb1/2 as a common target were significantly upregulated, implying their role in transforming growth factor (TGF)-β1-associated renal fibrosis. In ischemia-reperfusion injury, hypoxic damage induces the increased production of EVs in tubular epithelial cells that contains TGF-β1 mRNA [8]. Exosomal delivery of these TGF-β1 mRNAs into fibroblasts ultimately leads to fibroblast activation and proliferation. On the other hand, Ranghino et al. [18] recently showed that MSCs within the glomeruli can contribute to postischemic renal recovery primarily through the release of EVs.

In sepsis-related AKI, platelet-derived and neutrophil-derived EVs demonstrate proinflammatory and procoagulant effects, which can prevent the growth of bacteria and its dissemination [38]. However, these EVs can systematically induce oxidative stress and disseminate inflammatory responses, leading to poorer clinical outcomes.

**Tubulointerstitial inflammation**

Tubulointerstitial inflammation is a common characteristic of both acute and chronic kidney disease. Lv et al. [39] found a significant increase in tubular epithelial cell-derived exosomal miRNA-19b-3p in both a lipopolysaccharide-induced AKI model and an adriamycin-induced chronic proteinuric kidney disease model. Tubular epithelial cell-derived exosomal miRNA-19b-3p promoted M1 macrophage activation and tubulointerstitial inflammation. This group also showed that exosomal delivery of CCL2 mRNA from tubular epithelial cells to macrophages could mediate albumin-induced tubulointerstitial inflammation [40]. Hypoxia is another well-known stimulator of tubulointerstitial inflammation in the kidney. Hypoxic stimulation induces the expression of hypoxia-inducible factor-1α in tubular epithelial cells. This stimulus leads to exosomal miRNA-23a transfer from tubular epithelial cells to macrophages, which triggers macrophage activation and promotes tubulointerstitial inflammation [41]. Thus, modulation of macrophage activation by blocking exosomal miRNA or mRNA delivery from tubular epithelial cells can be a novel therapeutic approach.

**Fibrosis**

Fibrosis in glomeruli and tubules is an ultimate pathologic
process of chronic kidney diseases and end-stage renal disease, and the pathophysiological role of EVs in fibrosis has been explored in several studies [5,8,42,43]. Various stimuli on renal cells, such as hypoxia, oxidative stress, or high glucose, can lead to EV secretion or changes in the EV composition [8,42]. EVs secreted by damaged kidney cells can be transferred to other normal kidney cells, changing their phenotype and activating fibroblasts. In hypoxic conditions, damaged tubular epithelial cells transfer TGF-β1 mRNA to fibroblasts through exosome secretion, ultimately leading to the proliferation/activation of fibroblasts and progression of renal tubular fibrosis [8]. An in vitro study by Wu et al. [42] showed that EVs secreted from high glucose-treated glomerular endothelial cells can trigger podocyte epithelial-mesenchymal transition by transferring TGF-β1 mRNA into glomerular mesangial cells. This process induces cellular proliferation and overproduction of extracellular matrix within glomeruli, which is a key pathological mechanism of diabetic nephropathy. Several miRNAs in EVs, such as miR-21, miR-192, and miR-34a, also affect the biological processes of tubular fibrosis [43,44] and glomerular fibrosis [45], respectively.

Nephrolithiasis

Given the significant role of EVs as mediators of various renal diseases, several studies have speculated that EVs have a role in stone formation. For instance, an in vitro study by He et al. [46] showed that exposure of renal tubular cells to different concentrations of oxalates could affect the size and composition of the exosomes from tubular cells. This result suggests that the biological role of exosomes may depend on the level of oxalate exposure in a given microenvironment. Other in vitro studies by Singhto et al. [47,48] showed that exosomes derived from human macrophages exposed to calcium oxalate crystals could induce the migration/activation of monocytes/T-cell/neutrophils and enhance the production of proinflammatory cytokine interleukin-8. These exosomes had a greater capacity to bind calcium oxalate crystals and subsequently enhanced crystal invasion through the extracellular matrix, suggesting the possible role of macrophage-derived exosomes in the crystal invasion of the renal interstitium.

Renal cancer

EVs mediate intercellular communication between tumor cells. EVs from cancer cells contain proangiogenic mRNAs and various miRNAs with tumor invasive properties [49]. Clear renal cell carcinoma-derived EVs can negatively affect adjacent immune cells by the activation of the TGF-β/SMAD signaling pathway [50]. EVs are also known to mediate chemoresistance in renal cancer cells through the delivery of long noncoding/competing endogenous RNA, ultimately leading to higher expression of prometastatic receptors [51]. Further information about the role of EVs in urological malignancies is well described in a review article by Linxweiler and Junker [4].

Extracellular vesicles as biomarkers in various kidney diseases

Most of the EV studies in kidney diseases have been focused on biomarker discoveries. EVs can be found in multiple types of biological fluids including serum, urine, breast milk, cerebrospinal fluid, saliva, ascites, and bronchoalveolar lavage fluid. Under physiologic conditions, almost all urinary EVs originate from kidney cells because circulating EVs do not generally go through the glomerular basement membrane. Therefore, the organ-specific origin of urinary EVs comprises the major advantage of using urinary EVs as biomarkers in kidney diseases. Also, urinary EVs can be obtained noninvasively and repeatedly, providing important information regarding diagnosis, prognosis, and response to treatment.

Even though circulating EVs can originate from different organs, serum, and plasma, they can still provide significant information about the extent and prognosis of renal disease because the clinical outcome is determined not only from the kidney damage itself but also from the dysfunction of distant organs including the lungs, heart, brain, liver, and immune system [52]. Easier clinical accessibility is also a huge benefit of using circulating EVs as biomarkers in kidney disease.

Recent advancements in ‘omic’ studies have expedited the discovery of novel EV biomarkers enabling the detection of pathologic processes even earlier than traditional biomarkers. Salih et al. [53] compared urinary EVs from autosomal dominant polycystic kidney disease (ADPKD) patients and
healthy controls using quantitative proteomics. Among the proteins with higher abundancy, five proteins including complements and plakins were selected for further analysis based on their significance in pathway analysis. Especially, a higher level of complement in urinary EVs was found in younger ADPKD patients with preserved kidney function, raising its potential as a biomarker. Due to the vast amount of information from various studies and limited space, the role of EVs as biomarkers is summarized in Table 2 [53–78].

### Extracellular vesicles as therapeutics in various kidney diseases

#### Extracellular vesicles as a cargo of therapeutic material

The role of EVs as biocarriers has received a lot of attention

### Table 2. Extracellular vesicles as biomarkers in renal disorders

| Disease                  | Source/method                          | Main finding                                                                                                                                                                                                 | Reference |
|--------------------------|----------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------|
| AKI                      | Human urine/Western blotting           | Increased level of Na⁺/H⁺ exchanger type-3 in urinary exosome was found in patients with acute tubular necrosis but not in other etiologies of AKI.                                                             | [54]      |
|                          | Human or rat urine/proteomics          | Urinary exosomal Fetuin-A increased with AKI in both humans and rats. Exosomal Fetuin-A increase was detectable in the urine even before the development of histologic injury. | [55]      |
|                          | Human or rat urine/Western blotting    | Increased urinary exosomal ATF3 in AKI and WT-1 in podocyte injury.                                                                                                                                         | [56]      |
|                          | Human urine/Western blotting           | Increased urinary exosomal ATF3 in sepsis-induced AKI patients.                                                                                                                                             | [57]      |
| DN                       | Human urine/RT-qPCR                    | In DN patients, urinary exosomal let-7c-5p was upregulated, and miR-29c-5p and miR-15b-5p were down-regulated. These three miRNAs could predict the development of DN.                                         | [58]      |
|                          | Human urine/RT-qPCR                    | Increased level of uromodulin mRNA in urinary EVs derived from patients with type 2 DM nephropathy compared to type 2 DM without nephropathy. Urinary EV uromodulin mRNA level correlates with renal function decline in type 2 DM patients. | [59]      |
|                          | Human urine/microarray                 | Type 2 DM patient with microalbuminuria had increased levels of let-7i-3p, miR-24-3p and miR-27b-3p, and decreased levels of miR-15b-5p. The level of miR-30a-5p in urinary EVs was significantly correlated to macroalbuminuria in type 2 DM. | [60]      |
| IgA nephropathy           | Human urine/RT-qPCR                    | Total amount of urinary exosomes and exosomal CCL2 mRNA are upregulated in IgA nephropathy patients, which reflects active renal histologic injury and renal function deterioration.                          | [61]      |
|                          | Human serum/RT-qPCR                    | IgA nephropathy patients with lower serum miR-182 had more severe tubular atrophy, interstitial inflammation, and fibrotic tendency; IgA nephropathy patients with higher serum miR-192 had a slower disease progression.     | [62]      |
|                          | Human plasma/RNA-sequencing            | Exosomal IncRNA-G21551 was significantly down-regulated in IgA nephropathy patients compared to healthy first-degree relatives.                                                                        | [63]      |
| Nephrotic syndrome       | Human urine/RT-qPCR                    | Urinary exosomal miR-193a is significantly increased in primary FSGS patients compared to those in MCD patients.                                                                                          | [64]      |
|                          | Human urine/RT-qPCR                    | Urinary exosomal miR-194-5p, miR-146b-5p, miR-378a-3p, miR-23b-3p and miR-30a-5p are significantly increased in nephrotic syndrome and markedly reduced during remission. Concentration of miR-194-5p and miR-23b-3p were correlated with the degree of proteinuria. | [65]      |
| Renal transplantation    | Human urine/Western blotting           | Patients with delayed graft function after kidney transplantation had increased levels of NGAL in urinary exosomes.                                                                                     | [66]      |
|                          | Human plasma/RT-qPCR                   | Increased transcript levels of glycoprotein 130, SH2D1B, TNF-β, and CCL4 in plasma exosomes derived from patients with antibody-mediated rejection.                                                            | [67]      |
|                          | Human urine/integrated kidney exosome analysis | High level of CD3 positive urinary EVs in patients with acute cellular rejection.                                                                                                                                   | [68]      |
|                          | Human urine/proteomics                 | Higher levels of tetraspanin-1 and hemopexin protein are found in urinary exosomes derived from patients with T cell-mediated rejection.                                                               | [69]      |

(Continued to next page)
Table 2. Continued

| Disease | Source/method | Main finding | Reference |
|---------|---------------|--------------|-----------|
| PKD     | Human urine/proteomics | Higher levels of complement C3 and C9 are found in urinary EVs derived from ADPKD patients. | [53] |
|         | Human and rat urine/Western blotting | Increased urinary exosomal AGS3 expression in PKD patients/rats. | [70] |
| LN      | Human urine/RT-PCR | Urinary exosomal miR-29C correlates with the degree of renal chronicity but not with renal function in patients with LN. | [71] |
|         | Human urine/RT-PCR | Decreased level of urinary exosomal let-7a and miR-21 in patients with active LN compared to inactive LN. | [72] |
|         | Human urine/miRNA sequencing | Urinary exosomes from LN patients complicated with cellular crescent have a unique miRNA expression profile including miR-3135b, miR-654-5p, and miR-146a-5p. | [73] |
| Renal fibrosis and CKD model | Human urine/RT-qPCR | Down-regulation of urinary exosomal miR-29C level with worsening renal function and progression of tubulointerstitial fibrosis in CKD patients. | [74] |
|         | Human and mouse urine/RT-qPCR | Upregulation of urinary exosomal miR-21 in CKD patients as well as in nephrotoxic serum-treated mice. | [75] |
|         | Human urine/RNA-sequencing | Urinary exosomal miRNA-181a significantly decreased in CKD patients compared to healthy controls. | [76] |
| RCC     | Human plasma/RT-qPCR | Increased hsa-miR-301a-3p and decreased hsa-miR-1293 levels were found in plasma EVs in patients with metastatic clear cell RCC compared to those with localized disease. | [77] |
|         | Human plasma/RT-qPCR | Higher levels of plasma EV-derived TIMP-1 mRNA were found in patients with metastatic RCC or huge tumor burden. Also, patients with a high level of EV-derived TIMP-1 mRNA had poorer survival compared to those with a low level. | [78] |

ADPKD, autosomal dominant polycystic kidney disease; AGS3, activator of G-protein signaling 3; AKI, acute kidney injury; ATF3, activating transcription factor 3; CKD, chronic kidney disease; DN, diabetic nephropathy; DM, diabetes mellitus; EV, extracellular vesicle; FSGS, focal segmental glomerulosclerosis; IgA, immunoglobulin A; LN, lupus nephritis; IncRNA, long noncoding RNA; MCD, minimal change disease; miRNA and miR, microRNA; mRNA, messenger RNA; NGAL, neutrophil gelatinase-associated lipocalin; PKD, polycystic kidney disease; RCC, renal cell carcinoma; RT-qPCR, reverse transcription-polymerase chain reaction; RT-PCR, reverse transcription-polymerase chain reaction; SH2D1B, SH2 domain-containing protein 1B; TIMP, tissue inhibitor of metalloproteinase; TNF-α, tumor necrosis factor alpha; WT-1, Wilms tumor 1.

Recently, EVs carry several biological benefits as a vector over other methods, including stability, small size, and low immunogenicity. Specific cell surface molecules on EVs increase their targeting capacity with fewer off-target effects. Nanosized EVs with certain surface molecules can even penetrate natural barriers such as the blood-brain barrier [79]. Also, EVs use the native mechanisms of recipient cells in the course of internalization and intracellular trafficking [80]. Delivery of biomolecular cargos such as mRNA, miRNA, and proteins can modify the genetic profile and biological response of recipient cells, leading to the modulation of disease processes [81].

EV loading of traditional drugs with anti-inflammatory properties has been studied for therapeutic applicability [82,83]. Dexamethasone, for example, is widely used to suppress inflammation in various renal diseases, but its clinical use is limited due to significant adverse effects. By incubating macrophages with dexamethasone, macrophage-derived dexamethasone containing EVs were successfully produced and delivered to the inflamed kidneys of mice, which showed significant suppression of renal inflammation and fibrosis [82]. Notably, dexamethasone containing EVs substantially decreased the adverse effects from systemic steroid treatment such as hyperglycemia and suppression of the hypothalamic-pituitary-adrenal axis.

During the past decade, there has been an explosive advancement in EV research. Interdisciplinary cooperation between medicine and biomedical engineering has harnessed EVs as potent drug delivery tools through the application of nanotechnology. Several studies have successfully loaded genetic material into EVs and shown effects on modifying disease courses. Yim et al. [84] recently developed a novel technique for the effective intracellular transfer of soluble proteins through exosomes by using optogenetic engineering.
This technology was adopted in an experimental sepsis model, showing the protective effect of exosomal super-repressor inhibitor of kappa B (IκB) delivery in a murine septic AKI model [85]. Recently, our group has also demonstrated the beneficial effect of exosomal super-repressor IκB delivery in an ischemic AKI model (under review).

**Extracellular vesicles for kidney repair**

Stem cell therapy has great benefits in multiple experimental kidney injury models, not through their differentiation potential but the paracrine effects of their EVs [17]. EVs secreted from MSCs or endothelial progenitor cells have been shown to play a significant role in the amelioration of tubular cell apoptosis/fibrosis and proliferation of tubular and endothelial cells in various AKI models [86,87]. This beneficial effect was also found in experimental chronic kidney disease models, ameliorating epithelial-to-mesenchymal transition and preventing inflammatory cell infiltration [88,89]. MSCs that were engineered to overexpress miRNA-let7c could selectively deliver miRNA-let7c to damaged kidney cells using exosomes, which led to protection from renal injury and decreased fibrosis in a unilateral ureteral obstruction model [90]. Nagaishi et al. [91] showed that MSC-derived renal trophic factors including exosomes could alleviate albuminuria and immune cell infiltration into the kidneys in a diabetic nephropathy model. MSC-derived exosomes showed an antiapoptotic effect and protected tight junction structure in tubular epithelial cells. Intravenous injections of urinary stem cell-derived exosomes could also reduce urinary microalbumin excretion and prevent podocyte/tubular epithelial cell apoptosis in diabetic rats [92].

Apart from MSCs, EVs from other origins have shown a renoprotective effect in animal kidney injury models. Exosomes derived from sera of animals after remote ischemic preconditioning could attenuate sepsis-induced renal damage, and this was mediated by the upregulation of exosomal miRNA-21 [93]. Dominguez et al. [94] showed that intravenous delivery of EVs from rat renal tubular cells could markedly reduce renal damage in recipients after ischemic insult. This group also showed that human renal exosomes can prevent ischemic renal injury in nude rats [95].

However, the clinical evidence of the renoprotective effect of EV therapy in humans is relatively scarce. Several problems limit the therapeutic application of EVs in humans; the low yield of EVs from cultured cells and premature EV isolation/ purification with a risk of contamination from other molecules with similar density [96,97]. Although several studies have shown promising advances in EV isolation and purification, which is reviewed elsewhere [98], further efforts are needed to improve isolation efficiency and purity. Also, large-scale production is required for the therapeutic application of EVs in humans. Yang et al. [99] recently used a cellular nanoporation method to produce large-scale exosomes containing therapeutic mRNAs and targeting peptides. By using this strategy, they could produce a significantly higher number of exosomes and exosomal mRNA transcripts compared with bulk electroporation and other exosome production strategies. Lastly, advanced techniques in the fine modulation of EV content using transfection or cell engineering is needed to deliver specific transcripts or protein composition into target cells. This will lead to a better understanding of the biological role of certain EVs in specific renal disease models and will broaden the therapeutic applicability of EVs in various renal diseases.

**Extracellular vesicles as potential therapeutic targets**

As we have mentioned, EVs can mediate the propagation of renal damage in septic, inflammatory, or thrombogenic conditions. Therefore, temporarily blocking the release and uptake of EVs can possibly alleviate tissue damage. Various pharmacologic agents including antiplatelet agents, statins, calcium channel blockers, and abciximab were found to negatively affect the process of EV release and uptake, which is reviewed elsewhere [1]. However, whether blocking the release and uptake of EVs can directly affect the disease course has not been fully investigated. A study by Mossberg et al. [100] showed that C1-inhibitor can significantly reduce the release of chemotactic kinin B1-receptor-positive endothelial microvesicles, suggesting it has therapeutic potential in inflammatory diseases. Further studies are required to understand the clinical applicability of controlling EV interaction with recipient cells in systemic inflammatory conditions.

**Conclusions and perspectives**

EVs are involved in cell-to-cell communications within kidneys, mediating podocyte-tubulointerstitial/proximal-distal tubular/glomerulus-tubular crosstalk. EVs can either pro-
Extracellular vesicles (EVs) have been implicated in physiological roles in renal repair and immunomodulation or mediate pathological processes inducing thrombosis and inflammation. Recent studies have significantly improved our understanding of the roles of EVs in kidney physiology and pathology. However, most of the results are limited to descriptive transcriptomic/proteomic analysis and lack consistency as well as reliability. Stringent validation through clinical trials and large-scale cohort studies is necessary before entering into clinical application. Further optimization of EV isolation techniques and meticulous manipulation of the genetic materials or protein compositions of EVs are also necessary for clinical consideration. These efforts will extend our knowledge on the role of EVs in the development/progression of kidney diseases and will broaden the clinical applicability of EVs as novel diagnostics and therapeutics of renal diseases.

**Conflicts of interest**

Tae-Hyun Yoo is a Scientific Advisory Board member at ILIAS Biologics Inc. and the Editor-in-Chief of *Kidney Research and Clinical Practice*. Chulhee Choi is the founder and a shareholder of ILIAS Biologics Inc. The authors have no additional financial interests.

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**Authors’ contributions**

Conceptualization, Data curation, Investigation, Visualization: SAL

Formal analysis, Project administration: CC, THY

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Alvarez S, Suazo C, Boltansky A, et al. Urinary exosomes as a source of kidney dysfunction biomarker in renal transplanta-
plastic disease (Alzheimer’s disease, Parkinson’s disease) and cancer. These disorders are associated with the formation of amyloid plaques and neurofibrillary tangles, which lead to neuronal dysfunction and cell death. The identification of biomarkers that can predict these disorders is crucial for early intervention and better patient outcomes.

Recent studies have also highlighted the potential of urinary exosomes as diagnostic tools for cancer. For example, exosomal miRNAs have been reported to be upregulated in urine samples from patients with prostate cancer, bladder cancer, and renal cell carcinoma. These miRNAs can serve as novel biomarkers for cancer diagnosis and monitoring.

In conclusion, urinary exosomes represent a promising source of biomarkers for various diseases, including renal disease and cancer. Further research is needed to fully understand the key exosomal components and their roles in disease progression. The potential of urinary exosomes as non-invasive diagnostic tools offers hope for early diagnosis and personalized treatment strategies.

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