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Targeting and Therapeutic Peptide-Based Strategies for Polycystic Kidney Disease

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Abstract

Polycystic kidney disease (PKD) is characterized by progressive cyst growth and is a leading cause of renal failure worldwide. Currently, there are limited therapeutic options available to PKD patients, and only one drug, tolvaptan, has been FDA-approved to slow cyst progression. Similar to other small molecule drugs, however, tolvaptan is costly, only moderately effective, and causes adverse events leading to high patient dropout rates. Peptides may mitigate many drawbacks of small molecule drugs, as they can be highly tissue-specific, biocompatible, and economically scaled-up. Peptides can function as targeting ligands that direct therapies to diseased renal tissue, or be potent as therapeutic agents themselves. This review discusses various aberrant signaling pathways in PKD and renal receptors that can be potential targets of peptide-mediated strategies. Additionally, peptides utilized in other kidney applications, but may prove useful in the context of PKD, are highlighted. Insights into novel peptide-based solutions that have potential to improve clinical management of PKD are provided.

Keywords

Polycystic kidney disease; Targeted drug delivery; Peptide; Chronic kidney disease

1. Introduction

The kidneys are organs responsible for filtering wastes from the blood, as well as maintaining a balance of blood plasma solutes [1]. One condition affecting proper kidney

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function is polycystic kidney disease (PKD), which is the most commonly inherited kidney disease, affecting more than 12 million patients worldwide [2, 3]. PKD is characterized by continuous growth of fluid-filled cysts originating from the nephrons, which are the fundamental filtration units of the kidney. Cyst growth leads to massive enlargement and distortion of renal architecture, ultimately reducing filtration function and resulting in end-stage renal disease (ESRD), at which point patients seek dialysis or kidney transplantation [4, 5]. The majority of clinical treatment options focus on symptomatic management, and do not hinder cyst growth. These include a protein restricted diet, blood pressure control, and inhibiting the renin angiotensin aldosterone system (RAS) [6] with a RAS blocker [7]. However, blood pressure management and protein restricted diet ultimately show minimal benefits in delaying the age of ESRD onset due to cyst progression [8].

Recent efforts have focused on the use of drugs already approved by the Food and Drug Administration (FDA) for other diseases that have also shown efficacy in delaying cystogenesis, including pioglitazone, sirolimus, and metformin (Table 1) [9–12]. Despite initial enthusiasm, these drugs have been largely disappointing due to poor tolerability, off-target organ side effects, and/or limited therapeutic efficacy in patients [13]. Similarly, tolvaptan, the only FDA approved drug for autosomal dominant polycystic kidney disease (ADPKD), is difficult to tolerate due to side effects such as nausea and polyuria, and has been found to cause hepatotoxicity in certain patients [14]. Although tolvaptan slows cyst growth and delays the onset of renal failure by an average of ~5 years after 18 years of treatment, like other small molecules investigated for PKD, tolvaptan does not cure the underlying genetic defect of PKD [15]. Currently, no clinical trials exist for therapies that reverse cyst expansion, but emerging anti-micro RNA (anti-miR) [16, 17], clustered regularly interspaced short palindromic repeats (CRISPR) technologies [18], or other gene therapy solutions may provide promise for PKD reversal upon further investigation.

A major bottleneck in the clinical translation of these small molecule therapeutics however is their nonspecific and untargeted nature. Small molecules are classically categorized as organic, lipophilic compounds that have low molecular weight, typically less than 900 g/mol [19, 20]. These characteristics allow them to diffuse throughout the body, chemically interacting with target as well as off-target tissue. Peptides, generally defined as short chains of <50 amino acids, are one class of biologics which can address these issues, [21]. Due to their biologically based origin, they can be tuned to target discrete tissues or pathways, increasing therapeutic efficacy, as well as degrade into biocompatible amino acid constituents [22, 23].

However, peptides are not without their own set of drawbacks as well. Biologics such as peptides have been found to cause immunogenicity and an anti-drug antibody (ADA) response [24, 25], while also exhibiting unfavorable pharmacokinetic properties when compared to small molecules. Furthermore, *in vivo*, peptides encounter proteolytic enzymes and can be rapidly excreted from the body [26], exhibiting plasma half-life as low as a few minutes. Hence, modifications for extending half-life and increasing stability of peptides have been developed, including pegylation [27, 28], terminal modification [29], antibody conjugation (peptibodies) [30], and inducing cyclic and secondary structures [31]. Moreover, peptides can be incorporated into nanomaterials to improve their physiochemical properties

which is currently an active field of research [32–34]. Nanomedicine strategies are just beginning to emerge specifically for PKD, including recent efforts developing an oral nanocapsule strategy to improve the bioavailability of metformin [35].

Nonetheless, applications of peptides towards PKD offers an exciting, untapped area of research and medicine to advance the standard of care for PKD patients [36, 37]. Peptides can function as kidney-targeting ligands to direct small molecules to renal cysts [34, 38]. Additionally, therapeutic peptides can directly reduce cyst growth and exhibit promise for economical large-scale synthesis [39, 40] and long shelf-life [41]. This review outlines the pathogenesis of PKD and offers insights on aberrant pathways and specific cellular targets that may be exploited in targeting diseased, renal cells. Kidney-targeting and therapeutic peptide-based strategies that have shown success in other indications and have potential to be applied in PKD are also discussed.

2. Polycystic kidney disease

2.1. Kidney physiology

The kidneys function to maintain homeostasis in the body, regulating fluid, pH, and ionic balance, as well as removing waste metabolites [42]. The main filtration unit of the kidney is the nephron, which receives unfiltered whole blood from the afferent arteriole and feeds into the glomerulus (Figure 1). The glomerulus is a mass of capillaries functioning as a molecular sieve, where solutes are drawn out of the blood and passed to downstream renal tubule cells for reabsorption of salts, ions, or water. Solute not reabsorbed back into the blood, which include urea, metabolic waste products, and excess salts are excreted out of the body through collecting ducts which eventually lead to the bladder. The glomerulus is an assembly of four different cell types: the glomerular endothelial cells and podocytes that form a selectively permeable nanoscale barrier for solutes, as well as mesangial cells and parietal epithelial cells that help support the internal and external structure of the glomerulus, respectively [43, 44]. The tubular epithelial cells (Figure 2D) following the glomerulus, which make up the proximal tubule, Loop of Henle, distal tubule, and collecting ducts, are affected by cyst growth in PKD. Hence, the ability to reabsorb solutes from urinary filtrate is compromised.

2.2. Autosomal dominant polycystic kidney disease (ADPKD)

ADPKD is a chronic kidney disease mainly affecting the tubule cells, and is also the most commonly inherited cause of ESRD in adults [45]. The prevalence of ADPKD is estimated between 1:400 to 1:1000 people, and manifests in the form of fluid filled cysts, originating from diseased renal tubular cells that enlarge and destroy healthy nephron architecture and function [46]. ADPKD is diagnosed mainly by past family history, or due to presentation of common symptoms such as hypertension and flank pain, and is typically confirmed by renal ultrasound imaging [47]. Genetic testing is very costly in comparison to ultrasound, and still fail to identify mutations in about 10% of patients, and is therefore not used as a frontline method of diagnosis [48, 49]. What makes new cases of PKD especially tricky to diagnose is that the decrease in kidney function occurs during late stages of cyst formation, when the patient is middle aged. Highly efficient compensatory hyperfiltration by surviving non-cystic

nephrons maintain the glomerular filtration rate (GFR) within a normal range, despite continuous cyst growth and loss of functioning parenchyma [50]. Therefore, even within the same families with the same type of mutation, the age of ESRD and cyst localization can vary greatly [51].

Specifically, ADPKD involves mutations in two genes, PKD1 and PKD2, which encode for polycystin-1 (PC1) and polycystin-2 (PC2), respectively [52]. PKD1 mutations account for ~78% of patients, and PKD2 for ~13%, and no mutation is detected in ~9% of cases [53]. Patients with a mutation in PKD1 will reach ESRD by approximately 53 years of age, while PKD2 mutation patients have a much milder phenotype of kidney cysts compared to PKD1, and have a delayed onset of ESRD of ~20 years [54]. The gene products, PC1 and PC2, are essential proteins that are involved in maintaining the differentiated phenotype of renal tubular epithelial cells. Reduction in protein expression below a critical level in either one of these proteins results in loss of planar polarity, increased proliferation and apoptosis, and the development of a secretory phenotype (Figure 2) [46]. PC1 is a transmembrane protein with domains extending into the cytoplasm [52], while PC2 is a transmembrane, calcium-responsive cation channel [55]. In their native state, PC1 and PC2 interact with each other through their C-terminal tails in a 1:3 ratio of PC1 to PC2 to form the polycystin complex (PC) [56]. A number of intracellular localizations of PCs have been studied, but PC complexes residing on the primary cilia of renal epithelial cells are mainly responsible for PKD pathogenesis [57, 58]. Primary cilia are thin extensions of the plasma membrane and in the context of renal epithelial cells, primary cilia play a role in sensing mechanical and chemical extracellular signals and drive intracellular calcium signaling. The aberrant pathways in ADPKD, such as those stemming from primary cilia dysfunction, will be described in detail in section 3 along with peptide targeting and therapeutic solutions focusing on those pathways.

2.3. ARPKD

A related disease to ADPKD is autosomal recessive polycystic kidney disease (ARPKD), which also produces cyst-like structures in the kidneys as well as the liver. This hepatorenal disease is less frequent than ADPKD, and occurs in 1:20,000 people [59], but exhibits a more aggressive progression of cyst growth, with cysts arising mainly from collecting duct cells. The majority of ARPKD patients show disease phenotype as newborn or young children, and neonatal mortality is 25%–35%. Affected fetuses show massively enlarged kidneys, pulmonary hypoplasia, as well as contracted limbs with club feet [60]. In stark contrast to ADPKD, approximately 50% of ARPKD patients that survive the neonatal period progress to ESRD by 10 years of age [61].

In ARPKD, the genes with mutations are the Polycystic Kidney and Hepatic Disease 1 (PKHD1) gene and the DAZ Interacting Zinc Finger Protein 1 (DZIP1L) [62]. DZIP1L encodes a ciliary-transition-zone protein [63], while PKHD1 encodes for the protein fibrocystin, but the specific function and cellular localization of these proteins are not well understood. Histological assessment of ARPKD patients suggests that fibrocystin regulates cell-to-cell adhesion as well as proliferation [60]. Because disease-specific therapies and patient data for ARPKD is currently lacking, our review focuses on the more prevalent

ADPKD. However, many of the downstream pathways and therapeutic targets are shared between ADPKD and ARPKD, and may be opportunities for future ARPKD studies [64].

3. PKD targets for peptide-based intervention

Peptides offer many benefits over other biologics such as proteins, antibodies, and hormones used for therapeutic and targeting applications [65]. Specifically, peptides are relevant for targeting the kidney as they are small enough to access kidney tubule cells through glomerular filtration, usually limited to substances <15 nm in diameter and below 50 kDa [66–68]. Peptides also have the potential to be therapeutically potent, as they can show specificity to certain receptors and therefore reduce undesired off-target effects [69]. Limitations of peptides have included poor metabolic stability and rapid clearance compared to larger biologics, but many of these drawbacks can be improved by the addition of protecting groups onto the N- and C-terminus [70], exchanging L-amino acids with synthetic D-amino acids [31], and cyclization [71], which can minimize enzymatic degradation and unwanted protein adsorption.

Ease of large-scale chemical synthesis of synthetic or naturally occurring peptides is an attractive property for eventual clinical translation [72, 73]. A common method for peptide synthesis is solid phase peptide synthesis (SPPS), where peptides can be constructed stepwise on a solid resin support [74]. Typically, sequences >15 amino acids in length encounter diminishing yields in SPPS, as aggregation and additive coupling errors decrease purity as chain length increases [75]. However, peptide side chain groups can be kept chemically inert through the incorporation of protecting groups to prevent unwanted side reactions that reduce synthesis purity. These side chains can be later deprotected to add imaging agents, small molecule drugs, and allow for specific secondary structure [76–78]. Other synthesis methods include recombinant bacterial expression systems [79] and chemoenzymatic synthesis [80], where peptide bonds are catalyzed by hydrolase. As such, peptide strategies have begun to emerge to target PKD pathways. In this section, we describe the aberrant pathways in PKD and highlight potential targeting and therapeutic peptides applicable to PKD treatment.

3.1. Cellular signaling pathways as PKD targets

Over the last decade, many cellular signaling and metabolic pathways misregulated in PKD have been identified as drivers of cystogenesis [46]. In ADPKD, low levels of the PC on the primary cilia and endoplasmic reticulum (ER) reduce the amount of intracellular calcium available through the PC2 calcium channel. This triggers g-protein coupled activation of calcium-inhibitable adenylyl cyclase 6 (AC-VI) and inhibition of calcium/calmodulin-dependent phosphodiesterase (PDE), causing enhanced accumulation of cyclic AMP (cAMP) and activation of protein kinase A (PKA) [81]. cAMP/PKA signaling regulates a network of downstream pathways that affect cell cycle progression, cell proliferation, energy metabolism, and cell death (Figure 3) [82]. For example, the upregulation of the mitogen-activated protein kinases (MAPK) pathway, and phosphoinositide 3-kinases, protein kinase B, mammalian target of rapamycin (PI3K/AKT/mTOR) pathways through the inhibition of tuberous sclerosis complex proteins (TSC1 and TSC2) drive aerobic glycolysis and cell

cycle progression. The MAPK pathway also drives caspase-dependent cell death [83], even though abnormally increased proliferation in renal tubule cells is the hallmark of PKD [84]. The increased levels of apoptosis can destroy healthy nephron filtration function, while heightened proliferation of cyst lining cells drives cyst progression and expansion [85]. Downregulation of the AMP-activated protein kinase (AMPK) pathway stimulates ion transport through the cystic fibrosis transmembrane conductance regulator (CFTR) and fluid secretion, driving cyst formation [86]. Apart from the PC1 and PC2 transmembrane proteins, a series of cell surface receptors can contribute to increased cAMP and proliferation, including vasopressin receptor (V2R) and Wnt receptors depicted in Figure 3, and discussed in section 3.1.1 and 3.1.3., respectively. In this section, these pathways are reviewed and relevant peptide sequences showing therapeutic promise in PKD (Table 2) are presented.

3.1.1. cAMP and calcium dependent pathway—In PKD, an increase in cAMP levels stimulates cyst progression, which is led by a decrease in intracellular calcium and increased V2R signaling [85, 87–89]. cAMP is nucleotide second messenger, important for multiple cellular processes such as cell differentiation, DNA synthesis, and cell proliferation [90, 91]. cAMP stimulates cyst growth by promoting epithelial cell proliferation via B-Raf/MEK/ERK pathway activation [91, 92] and by transepithelial fluid secretion, which regulates chloride secretion through CFTR [93, 94]. Intracellular cAMP levels are regulated by adenylyl cyclases (AC), where AC catalyzes cAMP formation cAMP from ATP, and PDEs degrade cAMP to AMP. A study by Igarashi *et al.* studied the role of PC complexes in regulating cAMP in the primary cilia, and showed that the loss of primary cilia results in AC activation and increased cAMP levels, driving cyst growth in PKD [95].

One peptide that has shown efficacy to reduce pathways downstream of cAMP in a polycystic kidney disease rat (PCK) is the B-type natriuretic peptide (BNP) (SPKMVQSGSGCFGRKMDRISSSSGLGCKVLRH), a naturally occurring peptide responsible for blood pressure reduction [96]. Delivery *in vivo* stimulated cyclic GMP (cGMP) production and enhanced intracellular calcium levels, counteracting the effects of increased cAMP in PKD. While the crosstalk between cAMP and cGMP is not well understood in PKD, BNP led to decreased fibrosis, hypertension and vasopressin, thus offsetting PKD pathogenesis [97]. At the endpoints of this study, kidney histology revealed a 40% reduced cystic index in BNP-treated rats compared to PCK controls, and total kidney weight per body weight (TKW/BW) ratio was reduced by 1.9%, suggesting BNP could be a novel treatment for PKD.

Building on this study, a V2R antagonist peptide was used to treat PKD by reducing intracellular cAMP [98]. Ciolek et al. used mambaquaretin-1 (RPSFCNLPVKPGPC-NGFFSAFYYSQKTNKCHSFTYGGCKGNANRFSTLEKCRRTC VG), which was originally identified and purified from the venom of the green mamba snake and found to selectively inhibit major signaling pathways in PKD including both cAMP production and MAPK activity. When mambaquaretin-1 was synthesized by SPSS and administered daily for 99 days to CD1-*pcy/pcy* mice, a juvenile model of PKD, cystic index was reduced by 28% compared to controls, and the average number of cysts was decreased by 33% without causing toxicity (Figure 4). However, increased urine outflow was observed in a dose-

dependent manner with reduced urine osmolality, suggesting an aquaretic effect similar to inhibiting V2R by antagonists such as tolvaptan.

Somatostatin is another peptide hormone that can inhibit downstream intracellular cAMP levels. It is naturally secreted in a broad range of mammalian tissues, including the pancreas, hypothalamus, and from portions of the central nervous system [99]. As somatostatin receptors are highly expressed in the kidney and liver, somatostatin analogs have the potential to slow cystogenesis in both organs in ADPKD patients. In pioneering work, Epstein et al. demonstrated the efficacy of long-acting somatostatin (octreotide long-acting release (LAR)) for ADPKD patients [100]. In this randomized, placebo-controlled clinical trial, patients were intramuscularly treated every 28 days for 6 months with 40 mg of octreotide-LAR, a synthetic analog of somatostatin encapsulated into microspheres for long-acting release, followed by estimated glomerular filtration rate (eGFR) measurements and kidney volume evaluation via computed tomography (CT). Twelve out of 13 patients tolerated somatostatin well and showed decreased kidney volume of 71 ± 107 mL ($P < 0.05$) compared to those receiving the placebo (162 ± 114 mL; $P < 0.010$), as well as decreased cystic volume ($3.0 \pm 6.5\%$ vs. $5.6 \pm 5.8\%$). However, no significant changes in GFR was found, prompting further clinical trials.

In the follow up ALADIN trial (A Long-Acting somatostatin on Disease progression in Nephropathy due to ADPKD), 79 patients with $\text{eGFR} > 40$ mL/min/1.73m² (healthy eGFR is approximately 90 mL/min/1.73m²) were randomized to receive subcutaneous injections of octreotide-LAR or the placebo every 4 weeks for 3 years [101]. After 1 year, the increase in the total kidney volume (TKV) was reduced by 2/3rd with octreotide-LAR compared to the placebo. Over the 3 years of the ALADIN trial, the placebo group doubled in TKV compared to the octreotide group, but was not statistically significant due to a wide range of patient responses. Hence, the effects of octreotide treatment attenuated over time and a larger study may be needed to fully assess its therapeutic benefits. A follow up study, ALADIN 2, is currently recruiting additional patients in Italy. ALADIN 2 is similar to ALADIN except that patients with severely hindered eGFR of 15–40 mL/min/1.73 m² that may also benefit from treatment are being recruited. These clinical trials of peptides to reduce cAMP levels in PKD present promising new avenues for inhibiting cyst growth.

3.1.2. mTOR pathway—The mTOR signaling pathway serves a critical role in regulating several biological processes including cell metabolism, growth, proliferation, and survival. Dysregulation of mTOR complexes, mTORC1 or mTORC2, or overactivation of mTORC1 has been implicated in PKD development [11]. mTORC1 is inhibited by the tuberous sclerosis complex proteins TSC1 and TSC2, and thus, loss-of-function mutations of either TSC1 or TSC2 lead to kidney cyst formation [102, 103]. PC1 plays a vital role in stabilizing the TSC1/TSC2 complex and increased mTOR signaling has been described in multiple PKD models [104, 105], suggesting that PC1 is linked to mTOR expression. mTORC1 activation is linked to the S6 kinase (S6K)-dependent pathway which stimulates lipid synthesis and enhanced glycolysis [106]. mTOR and S6 kinase, a downstream effector of mTOR, has been found to localize to cyst-lining epithelial cells from PKD1 knockout mice, linking loss of PC1 function to heightened mTOR activity [107]. Prominent mTORC1 inhibitors currently in PKD clinical trials include rapamycin itself, analogues such as

sirolimus and everolimus [108], as well as metformin, a known AMP-activated protein kinase (AMPK) activator that also downregulates S6K phosphorylation [109, 110]. Currently, no mTOR peptide inhibitors have been tested for PKD, but in 2017, the first polypeptide named small regulatory polypeptide of amino acid response (SPAR) (MGAKAPRGPKVAQWAMETAVIGV-VVVLFFVVTVAITCVLCCFSCDSRAQDPQGGPGRSFTVATFRQEASLFTGPVRHAQPVPS AQDFWTFM) was found to suppress mTORC1 activity by analyzing and transcribing cytosolic RNA sequences previously thought to be non-coding [111, 112].

Therapeutic peptides demonstrate immense potential in the treatment of PKD as they can offer potent mTOR inhibition and anti-proliferative effects. Li et al. reported a Smac-mimetic peptide to induce TNF- α pathway-dependent cystic renal epithelial cell death [113]. Smac-mimetics are synthetic, cell membrane permeable peptides consisting of the 4 N-terminal amino acids of Smac, a protein naturally found in mitochondria that antagonizes cellular inhibitor of apoptosis protein 1 (cIAP1) and cellular inhibitor of apoptosis protein 2 cIAP2, which is related to downstream effects of mTOR-activated overproliferation [114]. In a PKD1 knockout murine model, upon intraperitoneal administration of Smac-mimetic peptides in lactating mothers on day 3 and 5 after birth of their litters, slowed cyst expansion and cell death of cystic epithelial cells was specifically observed in the kidney tissue of their pups, with no effects on neighboring healthy renal tubule cells. The Smac-mimetic peptide promoted degradation of cIAP1 in renal epithelial cells, and in combination with TNF- α , activated receptor interacting serine/threonine kinase 1 (RIPK1) dependent cell death complex, and degradation of the caspase-8 inhibitor, Fas-associated death domain-like interleukin-1-beta-converting enzyme-inhibitory protein long form (FLIP), in cystic renal epithelial cells.

Xueying Lin and colleagues established a zebrafish ADPKD model by transcription activator-like effector nucleases (TALEN) mediated gene editing to study the effect of autophagy in ADPKD which is directly controlled by the mTOR energy sensing and proliferative pathways. Beclin 1 has been reported to induce autophagy by binding to the autophagy inhibitor, GAPR-1, and was modified with a HIV transactivator of transcription 1 (Tat-1) protein domain to promote cell membrane permeability. The authors demonstrated that by activating autophagy with a specific autophagy inducer via a partial sequence of Beclin 1 (267–284), cystogenesis is reduced and renal function can be restored [115]. As a combinatorial strategy for ADPKD therapeutics, treatment with autophagy activators and mTOR-dependent rapamycin may be a promising strategy to attenuate cyst formation and restore kidney function.

3.1.3. Wnt signaling pathway—Both known wingless integration 1 (Wnt) signaling pathways are involved in formation of kidney cysts; activation of the canonical, β -catenin-dependent pathway drives renal cell nuclear changes (Figure 3) towards an undifferentiated, proliferative state, while activation of the noncanonical pathway alters planar cell polarity [116–118], compared to normal renal tubular cells that divide parallel to the tubular lumen axis. In PKD, alterations in planar cell polarity due to Wnt signaling leads to aberrant orientation of cell division and contributes to cysts sprouting from the main tubule lumen [119–121]. Although canonical Wnt signaling is essential for normal kidney maturation

during development, pathological activation of the canonical Wnt pathway in PKD destabilizes the PC complex and results in severe cystogenesis [122]. Loss of ciliary PC complex function has also been correlated with a substantial increase in canonical Wnt pathway activation [123]. While no studies have investigated peptides in the context of Wnt signaling in PKD, many Wnt peptide inhibitors have been investigated in the context of cancer and may provide efficacy in PKD. For example, stapled peptides investigated by Dietrich et al. (RRWPRXILDHXHVRVWR), were found to inhibit β -catenin dependent Wnt signaling [124]. Incorporation of cationic and amphipathic properties found in classes of cell penetrating peptides (CPPs) motivated the sequence design of these synthetic peptides, and allowed improved passage through the cell membrane and access to cytosolic β -catenin. The macrocyclic peptide (RKLYERFWWCG) has been found to inhibit Wnt signaling through direct interaction with Wnt surface protein [125]. Both may be effective peptide treatments for PKD.

3.1.4. Notch signaling pathway—Notch signaling has also been reported to be a central pathway in PKD development, as it regulates cellular processes such as proliferation [126]. Specifically, Notch3 is a receptor that has been found to be upregulated in the epithelial lining of cysts in both PKD murine models as well as human ADPKD-derived cysts [127]. Furthermore, a study using 3D cultures of primary human ADPKD cells demonstrated inhibition of Notch3 via gamma secretase resulted in reduced cyst proliferation *in vitro* [127]. Thus, targeting Notch3 signaling may result in an effective therapeutic strategy to arrest cyst growth in PKD.

Like Wnt signaling, no peptides have been developed to inhibit Notch signaling specifically in the context of PKD, but peptide-based strategies have been demonstrated against the Notch signaling pathway in other contexts. For instance, a library of 5–15 amino acid length, synthetic peptides were developed by Lin et al. to target Notch receptors 1–4, of which Notch3 is most relevant to PKD. Out of a library of 155 peptides, 15 short synthetic peptide sequences bound to Notch3 receptors, the key receptor in PKD, and induced apoptosis in HCC2429 lung carcinoma cells and HeLa cells [128]. These peptides exhibited a secondary structure containing a core with a β -pleated sheet, three disulfide bonds, and a series of loops [129], which allows for efficient and selective binding to Notch3 [130]. These peptides may be used as a targeting ligand in the context of PKD, and also provide additional benefits by inducing apoptosis in overproliferative cells in cyst epithelia.

3.2. Receptor targets in PKD

Receptor-mediated targeting presents an opportunity to achieve enhanced therapeutic efficiency and reduced off-target effects compared to current therapeutics in PKD [131, 132]. Receptors including megalin and cubilin, folate, and vasopressin V2R receptors are overexpressed in PKD and presents a strategy for binding to target cells [34, 133–136]. The binding to these target receptors may also cause downstream effects, including triggering endocytosis and associated cellular pathways, some of which may be therapeutically beneficial. This section will highlight the receptor targets which may be exploited for delivering PKD peptide and small molecule prodrug therapies (Table 3) and discuss the design insights and origins of these candidates.

3.2.1. Megalin and cubilin receptors—For PKD therapy, ligands with a high affinity for megalin and cubilin receptors provide an opportunity for kidney-targeted delivery of drugs and biologics. These endocytic receptors are expressed on the apical surface of proximal tubule epithelial cells and mediate renal-tubular reabsorption of proteins and several filtered ligands [137]. Cubilin is an intrinsic factor-cobalamin receptor (460 kDa) and interacts with megalin through an alpha-helical amino terminus [137]. Megalin is a low-density lipoprotein (~600 kDa) with a large extracellular domain which allows endocytic trafficking of ligands [138]. Of note, megalin is a key contributor to the reabsorption of albumin [139], and albumin peptide fragments and synthetic albumin-like-peptides can allow for enhanced megalin binding and cell internalization [140, 141]. Therefore, megalin-binding peptides may serve as front runners for selective renal targeting and enhancing uptake of peptide-conjugated payloads.

Binding to megalin as well as cubilin has been shown to regulate the endocytic trafficking of several ligands including lipoproteins, carrier proteins, hormones, and drugs in renal epithelial cells [142]. While relatively few kidney-targeting peptides are available, the synthetic peptide (KKEEE)₃K [143], which was previously found to target the kidneys through the megalin receptor, has been explored in multiple, recent studies [143–145]. In addition to (KKEEE)₃K, related peptide sequences have been produced via Fmoc SPSS, including (KKEE)₅K, (KKQQQ)₃K [145], and (KKEEE)₂K [146], to elucidate effects of peptide size, charge, and payload in relation to kidney targeting. Generally, these studies found acidic, negatively-charged peptides are not as efficiently reabsorbed by megalin and are more readily excreted, demonstrated by highly negative peptides, D₈ and (DSS)₄. Slightly positively-charged sequences investigated in these series of experiments, ranging between 0 to +10.0 mV zeta potential [147], was found to have higher kidney accumulation compared to negatively-charged counterparts. The underlying cause may be due to the affinity of more positively-charged peptides to the negatively-charged surface of renal proximal tubule cells, facilitating further interaction with the megalin receptor [148]. However, using highly-charged peptides like lysine (K) and glutamic acid (E) may confer undesirable properties, including aggregation and immune responses which remain to be fully elucidated. Like nanoparticles, and small molecules to a lesser extent, highly-charged peptides may adsorb serum proteins and molecules of the complement immune system, causing toxicity [149]. Nonetheless, the sequence of specific amino acids still plays a role, as when comparing equally-charged peptide sequences with amino acid substitutions, renal uptake can be nullified. For example, substitution of lysine (K) with arginine (R) to synthesize (RREEE)₃R preserves the number of positively-charged amino acids, but showed a 45% reduction in kidney uptake as well as faster urinary clearance [145].

Taking into account these design considerations, Lenhart et al. later demonstrated the potential of the (KKEEE)₃K peptide to target diseased kidneys in a juvenile cystic kidney (JCK) mouse model. Renal localization of dye-labeled (KKEEE)₃K, as well as a shorter, related sequence (KKEEE)₂K was investigated. Although the shorter, less negatively-charged sequence was hypothesized to improve renal accumulation [145], after 4 hours post-intravenous administration, >90% of the fluorescence signal was found in the kidneys of wild type and JCK mice for both peptides (Figure 5) [150]. In contrast, the free dye

accumulated equally in the liver and kidney. Both peptide-dye conjugates co-localized with megalin and aquaporin 1 (AQ1), markers of proximal tubule cells. Interestingly, slightly better renal retention was observed for the longer (KKEEE)₃K sequence, although not statistically significant. This may suggest the zwitterionic nature of peptides may affect renal retention. Additionally, (KKEEE)₃K has been incorporated into micelle nanoparticles to take advantage of the increased half-life and multifunctionally offered by nanocarriers [32, 151], and was found to have higher kidney accumulation compared to untargeted micelles in wild type mice [144, 147].

3.2.2. Folate receptor—The folate receptor is highly expressed on the apical brush border of renal proximal tubules, and functions to reabsorb filtered folate, a vitamin crucial for cell division and metabolism [152–159]. Very recently, folate-targeting approaches have received attention as a kidney-specific carrier for PKD therapy as folate receptors are highly expressed on renal cysts *in vivo* [160, 161]. Weimbs et al. investigated a folate-conjugated rapamycin (FC-rapa) and showed inhibition of mTOR activity in the kidneys of a Pkd1flox/_: Nestin-Cre murine model at a dose of 0.6 $\mu\text{mol/kg/day}$ [161]. Both FC-rapa and free rapamycin treatment groups were found to reduce the progression of PKD to a similar degree; however, FC-rapa demonstrated significantly less hepatotoxicity [161]. Although folate receptors are expressed in several tissues, expression is orders of magnitude higher in the kidneys [162], and hence, FC-rapa was found to preferentially accumulate in both healthy and polycystic kidneys and showed the highest mTOR inhibition in cystic cells. Notably, because suprapharmacological levels of folate have been shown to cause adverse effects such as ferroptosis (iron dependent cell death) [163], dose must be taken into account during the development of renal-targeting solutions via the folate receptor.

While folate is not a peptide, many peptides have been developed to efficiently bind the folate receptor. The peptide sequence MHTAPGWGWRLS, identified through phage display, exhibited an equilibrium dissociation constant (K_d) of 300 nM [164]. Through computational docking experiments, the peptide was found to bind to the surface of the receptor instead of docking into the folate binding pocket. While this K_d was weaker than that of folate itself ($K_d \approx 1$ nM), one way to improve peptide binding affinity includes altering the stereochemistry of the peptide, as reported by Zhou et al. [165]. In addition to folate-targeting peptides, conjugating peptides to folate can improve the binding affinity of folate beyond normal levels. For instance, the binding affinity of peptide-conjugated folate, GF-folate-IQ, to the folate receptor was found to be 0.18 nM [166]. This strategy may offer both high binding to the folate receptor and a direct conjugation strategy for many therapeutics relevant to PKD.

3.2.3. Vasopressin V2 receptor—Vasopressin V2 receptor antagonists have emerged as a potential therapeutic strategy against disease progression in PKD [167]. These receptors are usually localized to the basolateral side of collecting duct cells in the kidneys, and its activation initiates key homeostatic effects of vasopressin such as water reabsorption from the filtrate. Vasopressin binding to its V2 receptor activates an intracellular cascade, which in turn activates adenylyl cyclase and induces cAMP generation as mentioned above. This results in increased gene expression and stimulates the aquaporin water channel AQP-2,

which ultimately allows reabsorption of water from urine, as well as a PKA-dependent positive feedback expression of the V2R. Consequently, urine volume decreases while urine osmolality increases. Failure of V2 receptor binding causes inability to concentrate urine, thus results in concentrated fluid in the renal cysts.

Given the crucial role of vasopressin in PKD pathogenesis, studies have found that treatment with V2 receptor antagonists can reduce cystic burden in orthologous animal models of human PKD by reducing renal cAMP levels [87, 168]. These studies eventually motivated development of the V2 receptor antagonist, tolvaptan, which received FDA approval in 2018 [169]. In the first seminal clinical trial, the Tolvaptan Efficacy and Safety in Management of Autosomal Dominant Polycystic Kidney Disease and Its Outcomes (TEMPO) 3:4, tolvaptan, as compared with placebo, slowed the increase in total kidney volume and the decline in kidney function.

Despite these outcomes, this first generation, small molecule drug also has undesirable drawbacks. Patients treated with tolvaptan exhibit adverse events related to aquaresis (excretion of dilute urine) and liver toxicity, which led to a higher discontinuation rate (23% dropout rate vs. 14% in the placebo group). Deranged liver function tests detected in clinical trials were found to be potentially life-threatening, and additional, major side effects include dry mouth, polyuria, hypernatremia, nausea/vomiting, and abdominal pain/cramps [170]. Notably, expected therapeutic outcomes have also been lackluster, as perfect adherence to tolvaptan treatment for 18 years is predicted to delay the time to ESRD by only 4.9 years [15]. The oral bioavailability of tolvaptan is expected to be 42% [171], and these drawbacks of off-target side effects and limited therapeutic efficacy calls for opportunities in precision medicine using strategies such as peptide targeting and therapeutics.

As mentioned, among aberrant signaling pathways in PKD, cAMP overexpression driven by upregulated V2 receptor and vasopressin production are highly prevalent [172]. As disease progresses, the V2 receptor is also overexpressed in a positive feedback loop due to stimulation of downstream cAMP pathways [173]. Therefore, a highly promising strategy to target the diseased cells in PKD would be to direct therapeutic payloads to cells overexpressing V2R. Jung et al. utilized the peptide vasopressin analogue desmopressin (dDAVP) conjugated with nine D-arginines (dDAVP-9r), YFQNCpRGGGGGRRRRRRRRR, utilized as a peptide delivery platform for siRNA delivery against aquaporins to collecting duct cells [174]. *In vitro*, Madin-Darby Canine Kidney (MDCK) cells and pig epithelial-like kidney cells (LLC-PK1) exhibited an order of magnitude higher binding when incubated with FITC-siRNA/dDAVP-9r compared to FITC-siRNA alone or FITC-siRNA/9r. As expected, no substantial differences in cellular uptake between FITC-siRNA/9r and FITC-siRNA/dDAVP-9r were found in monkey kidney (Cos-7) cells, which are negative for V2 receptors, indicating that targeting the V2 receptor was essential for cellular uptake of the payload siRNA [175].

3.2.4. Epidermal growth factor receptor—One of the abnormalities in PKD pathogenesis is the mis-localization of epidermal growth factor receptors (EGFR) in cyst lining epithelium, resulting in increased activity of EGFR in cystic tissues [176]. Hence, although the EGFR pathway is highly studied in the context of cancer, this pathway can also

be used as a ligand target in PKD [177]. EGFR is a transmembrane protein and plays a central role in regulating cell proliferation, cell survival, and plays a crucial role in embryonic kidney development [178]. Seven highly effective EGFR binding ligands have been identified to trigger such canonical downstream effects: EGF, transforming growth factor- α (TGF- α), heparin-binding epidermal growth factor (HB-EGF), amphiregulin (AREG), betacellulin, epiregulin, and epigen [179]. Binding of each of these to EGFR triggers a wide variety of downstream effects in renal cells, including changes to inflammation, cell cycle state, and fibrosis [180], and hence, peptide ligand concentrations must be carefully considered to limit such adverse effects. Furthermore, studies assessing immunogenicity of peptides will be crucial before clinical translation.

Richards *et al.* showed that inducing expression of a modified, inactive form of EGFR could block the overexpression in EGFR-specific tyrosine kinase activity that accompanies renal cyst growth, leading to an improvement in kidney function and a significant decrease in collecting duct cyst burden [181]. Hence, these results suggest that changes in EGFR expression may drive cyst formation in the collecting ducts, and that therapeutics targeting the tyrosine kinase activity of EGFR may be therapeutic in PKD. Ahsan et al. synthesized the peptide Disruptin (SVDNPHVC), which was found to dimerize with EGFR causing its degradation and downregulation [182]. Although applied to throat, neck, and lung squamous cell cancers, it is important to note that intraperitoneal injection of Disruptin into mice bearing EGFR-driven human tumor xenografts did not affect adjacent healthy tissue, and only caused apoptosis in overproliferative, cancer cells. Hence, in the context of PKD, this peptide strategy may target cystic regions overexpressing EGFR without disturbing healthy nephrons that show baseline levels of EGFR.

3.2.5. Galectin-3—Galectin-3 is a β -galactoside-binding lectin expressed in the ureteric bud and collecting duct of the nephron during embryonic development [183]. It is also overexpressed in human ARPKD and in cyst epithelia, and is correlated to disease severity in ADPKD [184]. Galectin-3 has been hypothesized to be primarily localized to the cytoplasm; however, there is evidence that it localizes to the nucleus, cell surface, and is secreted in serum and urine [183]. Therefore, cell surface galectin-3 is a potential target for peptide ligands, while serum and urine galectin-3 may be a potential biomarker for PKD [185]. Phage display was previously performed to identify three highly specific, short peptide sequences for binding to galectin-3: G3-A9 (PQNSKIPGPTFLDPH), G3-A9 (SMEPALPDWWWKMFK) and G3-C12 (ANTPCGPYTHDCPVKR) [186]. High expression of galectin-3 in BT549 mammary gland cells corresponded to high peptide binding efficiency, compared to the negative controls, prostate carcinoma cells (PC-3 M) and human breast (MDA-MB-435), that have lower galectin-3 expression, demonstrating target specificity. Although initially investigated for renal carcinoma, these highly specific peptides ($K_d \approx 17\text{--}80$ nM) have potential to target cystic epithelia expressing galectin-3.

3.3 Tubule-targeting peptides

Some kidney targeting peptides may not target PKD specific phenotypes, but can target specific segments of the nephron and may be used as a strategy to prevent ESRD in younger patients in which kidney architecture is still relatively non-cystic [187, 188]. Bidwell et al.

utilized an elastin-like polypeptide (ELP) conjugated to a seven amino acid length sequence, CLPVASC named the kidney targeting peptide (KTP), and demonstrated kidney accumulation that is 5-fold higher than an untargeted ELP and 5-fold higher compared to other peripheral organs in swine [189]. Intrarenal distribution showed association of this KTP ELP around the glomerulus and surrounding proximal and distal tubules. This is the shortest peptide sequence to date to produce such accumulation increases above controls, which may be beneficial for conjugating active pharmaceutical compounds while not hindering their therapeutic efficacy.

Odermatt et al. reported a method of screening phage-display libraries *ex vivo* on intact kidney tubules using microdissection, vs. traditional monolayer-plated cells [190]. The linear peptide ligand, ELRGDMAAL, was found to selectively bind cortical collecting ducts (CCD) compared to other parts of the nephron such as the distal convoluted tubules, glomerulus, proximal convoluted tubules (PCT). Peptide sequences containing this motif exhibited 16-fold higher binding to CCD compared with PCT in wild type rats upon intravenous injection. Using this same microdissection screening method, Odermatt et al. found that peptides with a motif K(X₃)TNHP showed twice as much binding to PCT compared to CCD. Tunable uptake within different nephron cell types may be useful in applications for ARPKD drug delivery, where cysts arise in both the tubule cells and collecting ducts.

Recently, Jung et al. took a machine learning approach to predict the tissue-specific targeting ability of peptides, saving time and cost over traditional phage display [191, 192]. In this approach, prior to *in vitro* studies, candidate peptide sequences can be given a prediction score representing their affinity to various organs. Two sequences demonstrated especially high prediction scores for the kidney, PKNGSDP and DSHKDLK, based on support vector machine (SVM) and artificial neural network (ANN) methods. The authors conclude that these sequences recognize cellular markers present on glomerular endothelial cells. Although, no *in vivo* studies have been performed to verify kidney localization in PKD mice models, these nephron-segment-specific peptides present an opportunity for future prodrug, nanoparticle, and peptide targeting studies.

4. Conclusion

This review focuses on cellular and molecular signaling pathways involved in the PKD progression, which can be exploited for designing effective renal-targeted, peptide-based therapeutics. The only currently available treatment, tolvaptan, experiences many of the drawbacks of typical small-molecule drugs, such as poor delivery, low bioavailability, and/or lack of specific targeting. Hence, targeting using peptide-based ligands and therapeutics specific to PKD cellular pathways, such as cAMP dysfunction and renal cell overproliferation, may be used to overcome these challenges. With future improvements in peptide-based technologies paired with the growing understanding of PKD molecular pathogenesis, the development of PKD-specific peptides will likely emerge as a powerful treatment strategy.

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Abbreviations

ALADIN	a long-acting somatostatin on disease progression in nephropathy due to PKD
AC	adenylyl cyclase
AMPK AMP	activated protein kinase
AREG	amphiregulin
ADA	anti-drug antibody
anti-miR	anti-micro RNA
AQ1	aquaporin 1
ANN	artificial neural network methods
ADPKD	autosomal dominant polycystic kidney disease
ARPKD	autosomal recessive polycystic kidney disease
BNP B	type natriuretic peptide
AC-VI	calcium-inhibitable adenylyl cyclase 6
cIAP1	cellular inhibitor of apoptosis protein 1
CRISPR	clustered regularly interspaced short palindromic repeats
CCD	cortical collecting ducts
cAMP	cyclic adenosine monophosphate
cGMP	cyclic GMP
CFTR	cystic fibrosis transmembrane conductance regulator
DZIP1L	DAZ interacting zinc finger protein 1
ELP	elastin-like polypeptide
ER	endoplasmic reticulum
ESRD	end-stage renal disease
ELISA	enzyme-linked immunosorbent assay
EGFR	epidermal growth factor receptors

eGFR	estimated glomerular filtration rate
ERK	extracellular signal-regulated kinase
FLIP	Fas-associated death domain-like interleukin-1-beta-converting enzyme-inhibitory protein long form
FDA	Food and Drug Administration
GFR	glomerular filtration rate
GSK3β	glycogen synthase kinase 3 β
HB-EGF	heparin-binding epidermal growth factor
KTP	kidney targeting peptide
LAR	long-acting release
mTOR	mammalian target of rapamycin
MAPK	mitogen-activated protein kinases pathway
PDE	phosphodiesterase
PI3K/AKT/mTOR	phosphoinositide 3-kinases, protein kinase B, mammalian target of rapamycin
PKHD1	polycystic kidney and hepatic disease 1
PKD	polycystic kidney disease
PC	polycystin complex
PC1	polycystin-1
PC2	polycystin-2
PKA	protein kinase A
PCT	proximal convoluted tubules
PCK	rat rat model of polycystic kidney disease
Ras	rat sarcoma
RIPK1	receptor interacting serine/threonine kinase 1
RAS	renin angiotensin aldosterone system
S6K	S6 kinase
Src	sarcoma
SPAR	small regulatory polypeptide of amino acid response
SPPS	solid phase peptide synthesis

SVM	support vector machine
SPR	surface plasmon resonance
TEMPO	tolvaptan efficacy and safety in management of autosomal dominant polycystic kidney disease and its outcomes
TKV	total kidney volume
TKW/BW	total kidney weight per body weight
Tat-1	transactivator of transcription 1
TALEN	transcription activator-like effector nucleases
TGF-α	transforming growth factor- α
TSC1	tuberous sclerosis complex protein 1
TSC2	tuberous sclerosis complex protein 2
TNFα	tumor necrosis factor α
V2R	vasopressin receptor
VEGF	vessel endothelial growth factor
B-Raf	v-raf murine sarcoma viral oncogene homolog B
Wnt	wingless integration 1

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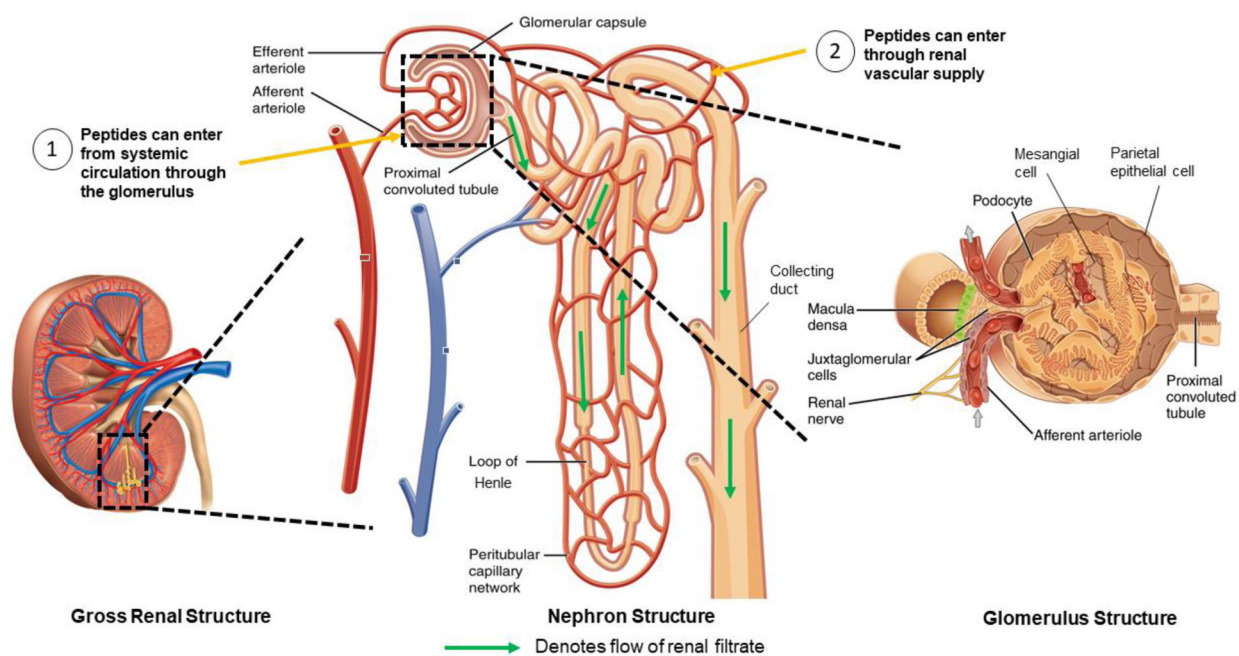


Figure 1. Structure and cellular makeup of the nephron. Adapted and reprinted from ref. [199].

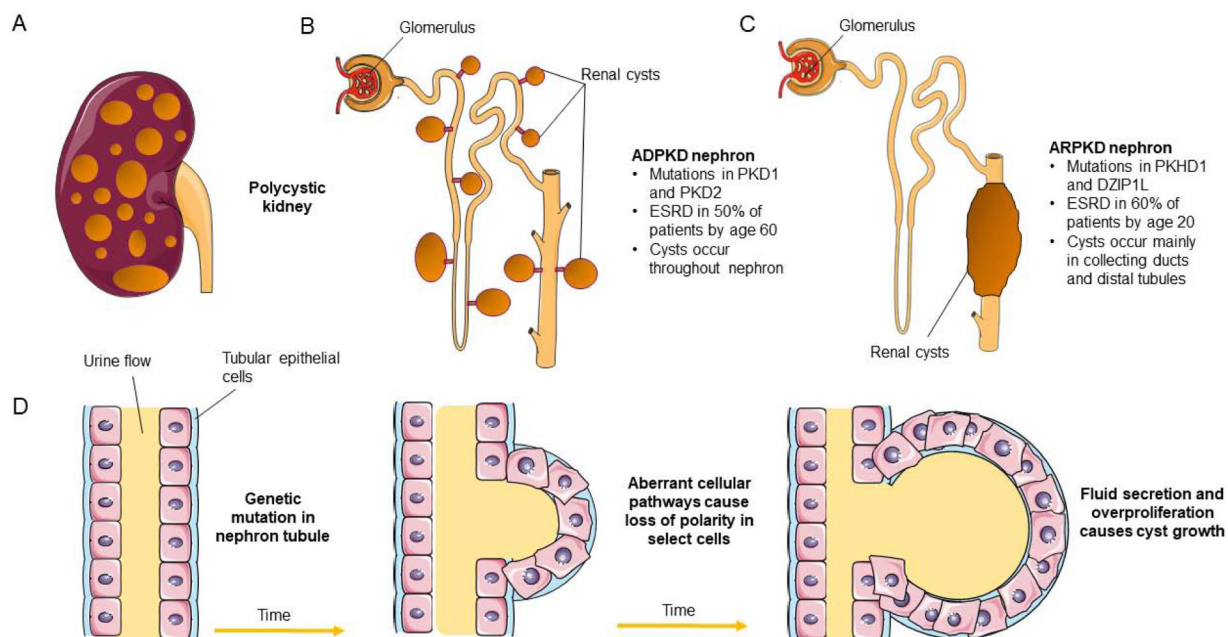
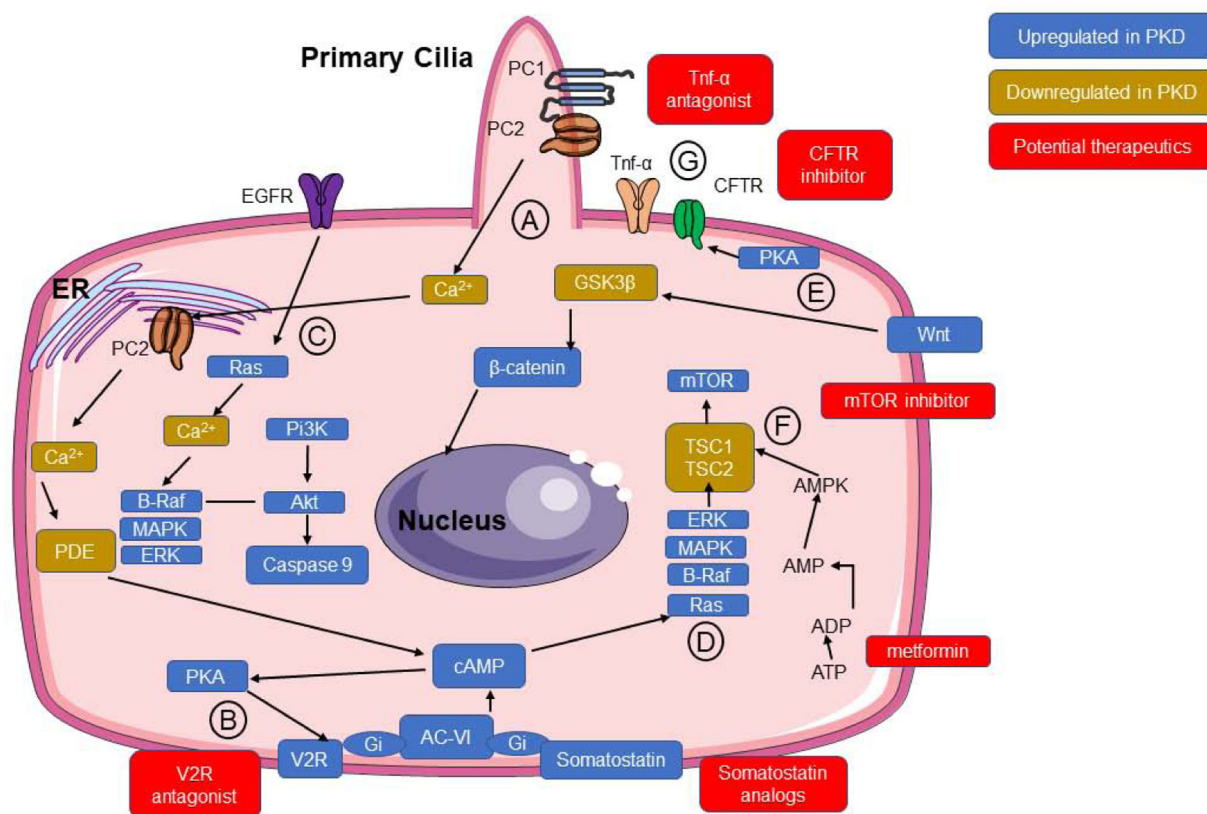


Figure 2.

A) Polycystic kidney disease (PKD) is the most common form of inherited kidney disease and is characterized by cyst growth in kidney parenchyma. B) Autosomal dominant PKD (ADPKD) is caused by mutations in the genes PKD1 and PKD2, affecting their respective protein products polycystin 1 (PC1) and polycystin 2 (PC2). C) Autosomal recessive PKD (ARPKD) is caused by mutations in polycystic kidney and hepatic disease 1 (*PKHD1*) and *DZIP1L* which encodes fibrocystin and DAZ-interacting protein 1-like protein, respectively. D) In kidney tubules or collecting ducts, germline mutations in a few renal cells initiates cystogenesis. As time progresses, loss of planar polarity causes sprouting of cysts from the tubule axis. Fluid secretion and overproliferation continues to drive cyst enlargement until neighboring healthy parenchyma is compressed and filtration function is decreased.



Cellular pathways involved in ADPKD pathogenesis within a renal epithelial cell. A) PC1 or PC2 loss of function on the primary cilia and ER impairs the calcium transient response, resulting in intracellular cAMP accumulation by preventing PDE degradation of cAMP. B) V2R activation is a major source of cAMP production in the cell through g-protein coupled activation of AC-VI. C) Ras dependent EGFR activation also lowers intercellular calcium. D) The response to abnormally increased cAMP levels leads to an activation of B-Raf/MAP2K1/ERK and inhibition of TSC1 and TSC2, driving the cell towards aerobic glycolysis and cell cycle progression. E) Other alterations include loss of planar cell polarity through Wnt dysregulation, and downstream β -catenin-dependent nuclear changes can drive cell de-differentiation into a proliferative phenotype. F) Master regulators of cellular energy metabolism, including the AMPK and mTOR pathways are mis-regulated, further enhancing proliferation. G) Altered CFTR behavior generates a secretory cell phenotype, in contrast to the normal absorptive phenotype, while TNF- α is an upregulated inflammatory mediator of the PKD phenotype. Proteins and receptors upregulated in PKD are highlighted in blue, and those downregulated are highlighted in yellow. Potential therapeutic targets have been listed in red next to the relevant receptor or misregulated protein.

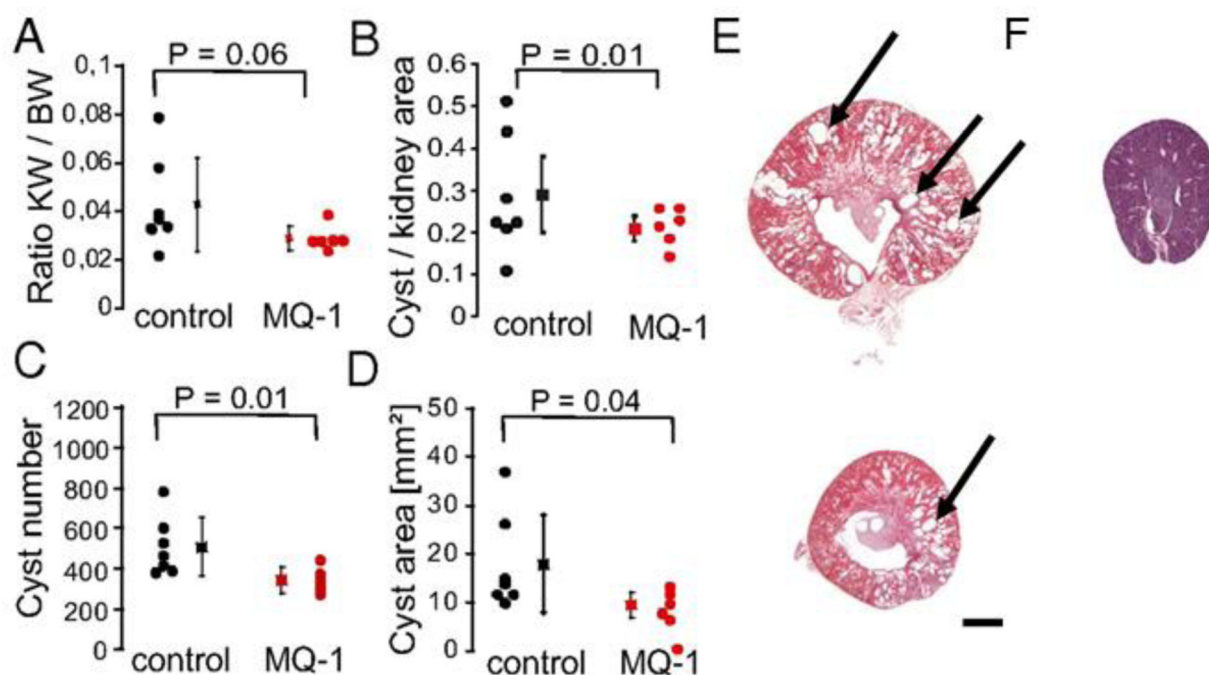


Figure 4.

Therapeutic effect of mambaquaretin-1 (MQ-1) in pcy mice. A) Ratio of kidney weight to body weight (KW/BW), B) ratio of cyst area to kidney area, C) cyst number, D) cyst area, and E) representative H&E-stained sections from kidneys of top: nontreated pcy mice and bottom: treated with mambaquaretin-1. $n = 6-7$. Cysts appear as voids throughout healthy tissue, and large individual cysts have been marked with arrows. F) Representative H&E-stained section from a wild type mouse. Scale bar, 2 mm. Adapted and reprinted with permission from United States National Academy of Sciences [98].

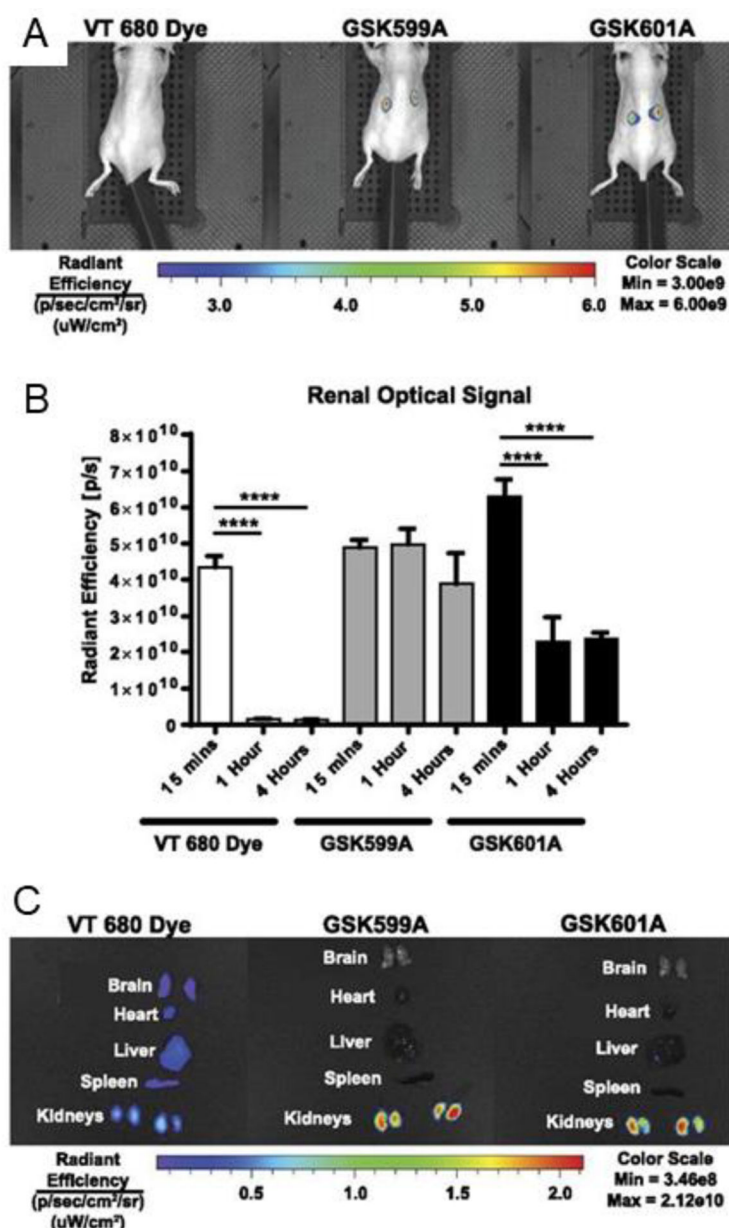


Figure 5.

Biodistribution of kidney targeting peptides. A) *In vivo* and *ex vivo* biodistribution of control VivoTag 680 dye, (KKEEE)₂K (GSK599A), and (KKEEE)₃K (GSK601A) 4 hours post-injection to JCK mice. B) Renal signal at 15 minutes, 1 hour, and 4 hours post-injection. n = 4–6 (****P < 0.0001). C) *Ex vivo* organ optical signals in groups treated with VivoTag 680 dye vs. those treated with GSK599A and GSK601. Reprinted and modified with permission from ASPET [194].

Table 1.

Clinical trials for PKD.

Interventions	Phases	Company/Sponsor	ClinicalTrials.gov Identifiers
KD019 (Tesevatinib)	Phase 1/2	Kadmon Corporation, LLC	NCT01559363 , NCT03203642 , NCT02616055
Rapamune	Phase 1/2	The Cleveland Clinic	NCT00286156
Bosutinib	Phase 2	Pfizer	NCT01233869
Lixivaptan	Phase 2	Palladio Biosciences	NCT03487913
Metformin	Phase 2	University of Maryland, Baltimore, University of Southern California, United States Department of Defense, University of Colorado, Denver	NCT02903511 , NCT02656017
Pasireotide LAR	Phase 2	Mayo Clinic	NCT01670110
Pioglitazone	Phase 2	Indiana University	NCT02697617
Pravastatin	Phase 2	University of Southern California, University of Southern California, National Institute of Diabetes and Digestive and Kidney Diseases	NCT03273413 , NCT04284657 , NCT00456365
Tolvaptan MR vs Tolvaptan IR	Phase 2	Otsuka	NCT01210560 , NCT01451827
Bardoxolone methyl capsules	Phase 2/3	Reata Pharmaceuticals	NCT03918447 , NCT03366337
Rapamycin	Phase 2/3	Yale university	NCT00920309
Venglustat GZ402671	Phase 2/3	Genzyme, a Sanofi Company	NCT03523728
Lisinopril, Telmisartan	Phase 3	National Institute of Diabetes and Digestive and Kidney Diseases	NCT01885559 , NCT00283686
Spirolactone	Phase 3	University of Colorado, Denver	NCT01853553
Tolvaptan	Phase 3	Otsuka	NCT01214421 , NCT02160145 , NCT00428948 , NCT02251275 , NCT00413777
Curcumin	Phase 4	University of Colorado, Denver	NCT02494141

Table 2.

Therapeutic peptides for polycystic kidney disease.

Peptide	Type	Major Findings	Reference
Somatostatin (FCDWKTCT)	Therapeutic	Clinical trial with octreotide-LAR, a somatostatin analogue; patients had lower kidney volume but no change in eGFR compared to controls	[100]
Smac-mimetic (4 N-terminal amino acids of Smac)	Therapeutic	Pkd1-floxed mouse model treated with Smac mimetic demonstrated cystic index reduction and kidney weight reduction	[113]
Tat-Beclin-1 peptide (amino acids 267–284 of Beclin-1)	Therapeutic	Beclin-1 peptide treatment to PKD zebrafish reduced cyst burden through mTOR pathway	[115]
B-type natriuretic peptide (SPKMVQGSGC-FGRKMDRISSS-SGLGCKVLRH)	Therapeutic	Upregulated BNP production in the PKD rat model reduced cyst progression	[193]
Mamabaquaretin-1 (RPSFCNLPVKPGPCNG-FFSAFYYSQKTNKCHS-FTYGGCKGNANRFST-LEKCRRTC VG)	Therapeutic	CD1-psy/psy mice, a juvenile model of PKD, showed less cysts after treatment with the peptide	[98]

Table 3.

Targeting peptides for polycystic kidney disease.

Peptide	Targets	Major Findings	Reference
(KKEEE) ₃ K family	Megalin receptor	In polycystic JCK mice, (KKEEE) ₃ K peptide found in cortex of kidneys	[194]
Desmopressin-conjugated D-arginine (dDAVP-9r)	Upregulated V2R	(dDAVP-9r) delivered siRNA payload to V2R in the kidneys	[174]
Disruptin (SVDNPHVC)	Upregulated EGFR	Disruptin is specific to upregulated EGFR cancer phenotypes	[195]
Library of 5–15 AA length synthetic peptides	Upregulated Notch3 receptors	Identified peptide sequences with specific binding to Notch receptors 1–4 to inhibit tumor growth	[128]
PKNGSDP and DSHKDLK	Glomerular endothelium	Machine learning prediction of sequences from phage display input data	[191]
AVP Peptide (CYFQNCPRG)	Proximal Tubule Cell	Specific renal delivery of sugar-modified low-molecular-weight peptides to proximal tubule cells	[196]
G3-C12 (ANTPCGPYTHDCPVKR)	Proximal Tubule Cell	IV injection of G3-C12 peptide accumulated in proximal renal tubule cells	[197]
CLPVASC	Proximal Tubule Cell	CLPVASC showed high kidney specificity to human podocytes, proximal tubule epithelial, and glomerular microvascular endothelial cells	[198]