



The Aryl Hydrocarbon Receptor in Chronic Kidney Disease: Friend or Foe?

Yenan Mo^{1,2†}, Zhaoyu Lu^{1,2†}, Lixin Wang², Chunlan Ji^{1,2}, Chuan Zou² and Xusheng Liu^{2*}

¹ The Second Clinical College of Guangzhou University of Chinese Medicine, Guangzhou, China, ² Department of Nephrology, The Second Affiliated Hospital of Guangzhou University of Chinese Medicine, Guangzhou, China

The aryl hydrocarbon receptor (AhR) is a ligand-activated transcription factor that promotes cell responses to small molecules derived from the diet, microorganisms, metabolism and pollutants. The AhR signal regulates many basic cellular processes, including cell cycle progression, adhesion, migration, apoptosis and cell proliferation. Many studies have shown that AhR is associated with chronic kidney disease (CKD) and its complications. This article reviews the current knowledge about the role of AhR in CKD, showing that AhR mediates CKD complications, including cardiovascular disease, anemia, bone disorders, cognitive dysfunction and malnutrition, and that it influences drug metabolism in individuals with CKD. AhR enhances the intestinal barrier function to reduce the harmful effects of uremic toxins. Therefore, understanding the complex roles of AhR during CKD is important to be able to target this transcription factor safely and effectively for CKD prevention and treatment.

Keywords: aryl hydrocarbon receptor, chronic kidney disease, tryptophan metabolism, uremic toxin, CKD complications, intestine homeostasis

INTRODUCTION

Chronic kidney disease (CKD) has become an emerging global disease during the past decade. The all-age prevalence of CKD increased 29.3% worldwide from 1990 to 2017, involving 697.5 million cases in 2017 (representing a global prevalence of 9.1%) (Bikbov et al., 2020). CKD results in irreversible kidney structural and functional alterations that lead to end-stage renal disease, with patients ultimately requiring dialysis or renal transplantation.

Chronic kidney disease is characterized by the accumulation of a mixture of uremic toxins/solutes that persist even after standard dialysis (Vanholder and Glorieux, 2014; Sirich, 2017). These toxins are divided into three categories: small soluble compounds, middle molecules and protein-bound molecules (Duranton et al., 2012). Tryptophan-derived toxins are of particular interest because they exhibit cardiovascular toxicity and are ligands of the aryl hydrocarbon receptor (AhR) (Heath-Pagliuso et al., 1998; Schroeder et al., 2010). Multiple sensors detect changes in the cellular milieu to adapt to toxic environments and generate responses to molecular environment changes due to diet, symbiotic microbiome and host metabolism factors. One of these sensors is AhR, a ligand-activated transcription factor involved in various cellular processes, such as cell cycle, epithelial barrier function and neurological signaling, as well as responses to antioxidants and xenobiotics (Murray et al., 2014).

Aryl hydrocarbon receptor regulation and its role during CKD has gained interest as a potential source of biological processes that may guide the development of new therapeutic interventions.

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*Correspondence:

Xusheng Liu liuxu801@126.com

[†]These authors have contributed equally to this work

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Mo Y, Lu Z, Wang L, Ji C, Zou C and Liu X (2020) The Aryl Hydrocarbon Receptor in Chronic Kidney Disease: Friend or Foe? Front. Cell Dev. Biol. 8:589752. doi: 10.3389/fcell.2020.589752 In this review, we summarize the current knowledge on the role of AhR in CKD and/or its complications, and we address the potential role of AhR as a therapeutic target.

AHR SIGNALING AND ITS LIGANDS

AhR Signaling Pathway

Aryl hydrocarbon receptor belongs to the periodic-AhR nuclear translocator-single-minded protein (PER-ARNT-SIM [PAS]) superfamily. The PAS domain senses both endogenous factors (such as cellular metabolites) and exogenous factors (such as environmental toxins) (McIntosh et al., 2010).

As shown in **Figure 1**, there are two AhR signaling pathways. One is the xenobiotic-responsive element (XRE)-dependent control of gene expression by AhR, also called canonical AhR signaling. Prior to AhR activation, AhR is inactive as part of a protein complex that consists of two molecules of heat-shock protein 90, one X-associated protein 2 (also known as an AhRinteracting protein), one co-chaperon p23 and the protein kinase Sarcoma (SRC) in the cytoplasm. The AhR chaperone complex stabilizes AhR in the cytoplasm and keeps it in a conformation that has high affinity for its ligands (Pongratz et al., 1992). Once a ligand binds to AhR, the molecule undergoes a conformational change to expose its N-terminal nuclear localization sequence, facilitating translocation of the liganded AhR protein complex into the nucleus. In the nucleus, AhR binds to ARNT (also known as HIF-1 β) through its PAS domain, forming a ligandbound AhR-ARNT dimer. The dimer gets recruited to a DNAspecific sequence (referred to as a DRE or XRE for a dioxinor xenobiotic-responsive element) located within the promoters of target genes, to promote transcription of a wide variety of genes, such as cytochrome P450, family 1, member 1A (Cyp1A1); cytochrome P450, family 1, member 2A; cytochrome P450, family 1, subfamily B; AhR repressor (AhRR) and cyclooxygenase-2 (Denison and Nagy, 2003). AhR-dependent gene transcription ends upon separation of the AhR-ARNT complex from the DRE. The N-terminal nuclear export sequence mediates AhR export from the nucleus into the cytoplasm, where ubiquitin-mediated AhR proteasome degradation occurs (Pollenz, 2002). The other AhR signaling pathway is non-canonical AhR signaling, which controls AhR gene expression through non-XRE DNA-response elements. AhR can also interact with additional transcription factors such as nuclear factor-KB (NF-KB), programmed death ligand 1, signal transducer and activator of transcription (STAT) and nuclear factor-erythroid-2-related factor 2 by binding to them and modulating the expression of their target genes through a non-genomic pathway that does not require ARNT (Kimura et al., 2008; Puga et al., 2009; Wakamatsu et al., 2018; Singh et al., 2019; Wang et al., 2019). Moreover, AhR functions as an E3 ubiquitin ligase, regulating the half-life of other transcription factors. For example, indoxyl sulphate (IS) directly activates AhR, extending the half-life of tissue factor (TF) by inhibiting its ubiquitination in vascular smooth muscle cells; TF is a crucial mediator of injury-related thrombosis (Shivanna et al., 2016). In addition, AhR activation triggers phosphorylation cascades driven by SRC kinase released from the AhR chaperone

complex. For instance, SRC phosphorylates a broad variety of target proteins including indoleamine 2,3-dioxygenase 1 (IDO1) (Bessede et al., 2014), which promotes tryptophan transformation into kynurenine, a ligand of AhR.

AhR Ligands

Aryl hydrocarbon receptor was discovered to be a receptor binding to 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) (also known as dioxin) with high affinity. The receptor can be hyperactivated in a sustained manner, resulting in numerous toxicological outcomes. In addition, many commercial and consumer products, vegetables, fruits, and spices have exhibited AhR activation potential (Jeuken et al., 2003; Zhao et al., 2013) and can facilitate the induction or repression of the expression of a range of genes, thereby regulating a diverse spectrum of biological and toxic effects. Environment pollutants and myriad endogenous molecules such as microbial metabolites (e.g., indolic solutes) and naturally occurring flavonoids (e.g., resveratrol and quercetin) are also AhR ligand candidates (Murray et al., 2014; Gutierrez-Vazquez and Quintana, 2018; Rothhammer and Quintana, 2019; **Table 1**).

For patients with CKD, the most important AhR ligands are uremic toxins, especially those derived from tryptophan metabolism. This essential aromatic amino acid is a biosynthetic precursor of a large number of microbial and host metabolites (Alkhalaf and Ryan, 2015). Tryptophan metabolism follows three major pathways in the gastrointestinal tract: (1) the indolic pathway in intestinal microorganisms that directly transforms tryptophan into several AhR ligand molecules, such as indole-3acid-acetic (IAA), indole-3-aldehyde, indole-3-acetaldehyde, and indole-3-propionic acid (Zelante et al., 2013; Hubbard et al., 2015; Alexeev et al., 2018); (2) the kynurenine pathway via the rate-limiting enzyme IDO1 that produces kynurenine and downstream products such as quinolinic acid and kynurenic acid (Clarke et al., 2012; Cervenka et al., 2017; Kennedy et al., 2017) and (3) the serotonin pathway in enterochromaffin cells that involves tryptophan hydroxylase-1 (Yano et al., 2015). Some tryptophan metabolites such as IS, IAA and kynurenine represent both important uremic toxins and potent AhR ligands. We will summarize the associations between AhR and uremic toxins in CKD below.

ACTIVATED AHR AGGRAVATES RENAL DAMAGE

Patients with CKD are exposed to a variety of uremic toxins, which are agonists of AhR. Early Lu et al. (2006) provided evidence for a positive correlation between renal AhR expression and CKD severity, suggesting that AhR activation has a pathogenic effect in both male and female rats that have undergone nephrectomy. Compared with healthy controls, CKD patients and 5/6 of nephrectomy mice exhibited increased serum AhR-activating potential, which was strongly correlated with IS concentration and estimated glomerular filtration rate, and which upregulated expression of the AhR target genes AhRR and Cyp1A1 (Dou et al., 2018). Kim et al. (2020) reported higher



AhR transactivating (AhRT) activity in patients with CKD than in controls, and they showed that peritoneal dialysis treatment was more effective at reducing this AhRT activity than haemodialysis.

Aryl hydrocarbon receptor signaling regulates many fundamental cellular processes, such as cell cycle progression, apoptosis and cell proliferation by regulating P53, FasR, Bcl-2 and cell cycle kinases (Zaher et al., 1998; Mohammadi et al., 2017). Exposure to adenine and IS increases AhR activity in the periglomerular region as well as in proximal and distal renal tubules, leading to renal fibrosis (Walker et al., 2020). AhR activated by IS induces podocyte injury, progressive glomerular damage and a pro-inflammatory phenotype in vitro and in vivo (Ichii et al., 2014). IS downregulates the expression of the Mas receptor, which is associated with the inhibition of the renin-angiotensin system, via the organic anion transporter 3/AhR/STAT3 pathway to upregulate TGF-β in proximal tubular cells, thereby aggravating CKD (Ng et al., 2014). A possible connection between AhR and renal fibrosis has been further supported by the association between indole solutes and renal

fibrosis (Mutsaers et al., 2015b), which may be mediated by AhR signaling.

In patients with diabetes mellitus, AhRT activity has been positively correlated with the progression of diabetic nephropathy (DN) and the severity of renal insufficiency (Kim et al., 2013). Lee et al. (2016) investigated the role of AhR in the pathophysiological processes of DN using AhR-knockout and pharmacological-inhibitor α -naphthoflavone models, and they found that AhR mediated renal oxidative stress in a diabetic mouse model, resulting in macrophage infiltration, extracellular matrix accumulation and mesangial cell activation.

Translocation of microbial antigens due to gut barrier breakdown has been associated with gut microbiota and the presence of systemic lupus erythematosus (SLE) and lupus nephritis in animal and human studies. Clinical studies have shown reduced gut microbiota diversity in patients with SLE, and gut-derived pathobionts including *Enterococcus gallinarum* have been detected in the liver (Manfredo Vieira et al., 2018). Animal studies have shown that gut-derived *E. gallinarum* can reach

TABLE 1 | exogenous and endogenous AhR ligands.

Ligand		Biochemical pathway
Xenobiotic	2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)	Halogenated aromatic hydrocarbons
	3-methylcholanthrene	
	Benzanthracenes	
	Benzoflavones	
	Benzo[a]pyrene	
	Biphenyls	
	Dibenzofurans	
	Polyaromatic hydrocarbons	
	Leflutamide	Pharmaceuticals
	Tranilast	
	Omeprazole	
	6,2',4',-trimethoxyflavone	Compound from chemical library
	CH-223191	
	GNF351	
	StemRegenin 1 (SR1)	
	Indirubin	Phytochemicals from plants
	Indigo	
Diet	Galangin	Flavonoids
	Quercetin	
	Kaempferol	
	Besveratrol	Polyphenol
	Indole-3-acetonitrile	Dietary metabolite of cruciferous vegetables
	Indole-3-carbinole	
	Indolo[3,2-b]carbazole (ICZ)	
	3.3'-diindolvlmethane (DIM)	
Tryptophan metabolites	3-Methylindole (Skatole)	Microbiota metabolism
	Indole-3-acetaldehvde	
	Indole-3 acetic acid (IAA)	
	Indole-3-aldebyde (IAId)	
	Tryptamine (TA)	
	5-bydroxy-tryptophan (5HTP)	Host metabolism
	Cinnabarinic acid (CA)	
	Kynurenic acid (KA)	
	Kynurenine (Kyn)	
	Xanthurenic acid	
	Indovul sulfate (IS)	Microbiota and bost metabolism
	6-formylindolo[3 2-b]carbazole (EICZ)	Photo-oxidation
Other metabolites	Bilirubin	Haem metabolism
	Biliverdin	Hadminictabolism
		Arachidonic Acid Metabolism
	Prostaglandin PGG2	
	Hudroxyaicosatrianoic acid ([12/B]-HETE])	
	1 4 dibudrow 2 pophtojo opid	Miarabiata matabaliam
		MICrobiola melabolism
	Indirubin	
	Malassezin	
	Irypthantrin	

the liver and induce the development of lupus autoantibodies, partly via activation of the AhR/CYP1A1 pathway to trigger Th17 and T follicular helper cell activation, crucial steps for the production of systemic autoantibodies including anti-dsDNA antibodies (Manfredo Vieira et al., 2018). Therefore, AhR may be a therapeutic target in CKD. In fact, resveratrol, a natural AhR antagonist, inhibits proteinuria, hypoalbuminemia and hyperlipidemia in nephritic rats (Nihei et al., 2001). Atrazine exposure has been shown to cause environmental nephrosis via direct kidney injury, whereas

supplementary lycopenes provide significant protection against atrazine-induced renal injury and environmental nephrosis via their inhibition of the activation of AhR and the expression of cytochrome P450 isoforms (Xia et al., 2018). On the other hand, ARNT, a transcriptional co-activator of AhR, has been implicated in renal fibrosis and is sought as a therapeutic target in CKD; strong experimental support exists for a mechanistic model in which FK506 induces renoprotection (Tampe et al., 2018; Haase, 2019).

AHR ACTIVATION MEDIATES VARIOUS COMPLICATIONS IN INDIVIDUALS WITH CKD

AhR and Cardiovascular Disease

Cardiovascular disease is a major cause of mortality in patients with CKD. AhR is an important receptor for various toxins and can activate pro-thrombotic and pro-inflammatory pathways; thus, there is a close association between the accumulation of uremic toxins and the cardiovascular complications of CKD (Vanholder et al., 2016). In fact, even under nonuremic conditions, AhR activation is associated with increased cardiovascular risks (Sallee et al., 2014). Accordingly, individuals exposed to AhR agonists are prone to increased cardiovascular risks (Humblet et al., 2008). For example, individuals exposed to the TCDD compound in Agent Orange during the Vietnam War have been shown to have associated cardiovascular complications ranging from coronary heart disease to stroke (Lowenstein, 2014).

The accumulation of uremic toxins has effects similar to those of AhR antagonists, making patients with CKD more prone to cardiovascular disease through a series of possible mechanisms. Ito et al. (2016) proved that AhR activated by IS can stimulate the transcriptional activity of activator protein-1 and induce the expression of E-selectin in vascular endothelial cells, thereby increasing the level of vascular inflammation and leukocyte adhesion. Assefa et al. (2019) showed that IS-activated AhR upregulates the activity of SRC to mediate the activity of vascular endothelial cadherin, thereby increasing endothelium permeability and promoting atherosclerosis. In human hepatoma HepG2 cells, IS-induced AhR activation inhibits the expression of fetuin-A, a liver-derived circulating protein that effectively suppresses calcification. Therefore, IS-activated AhR may increase the incidence of vascular calcification and cardiovascular mortality (Ochi et al., 2015). In human umbilical vein endothelial cells, AhR also mediates IS-induced cellular senescence (Koizumi et al., 2014; Eckers et al., 2016) and stimulates the expression of monocyte chemoattractant protein-1, a chemokine recruiting monocytes from the bloodstream into the subendothelial space to participate in the early stages of atherosclerosis (Watanabe et al., 2013). AhR further plays a key role during platelet production (Lindsey and Papoutsakis, 2011; Strassel et al., 2016) and function (Lindsey et al., 2014). AhR agonists enhance platelet aggregation by activating the AhR/p38MAPK pathway in platelets (Pombo et al., 2015). In addition, activated AhR increases the expression and activity of TF to induce a procoagulant state in vascular smooth muscle and endothelial cells (Gondouin et al., 2013; Shivanna et al., 2016); moreover, it enhances the activation of the pro-inflammatory enzyme cyclooxygenase-2 in endothelial cells (Dou et al., 2015). Vascular dysfunction caused by AhR activation may lead to atherosclerotic thrombosis, thus increasing the risk of myocardial infarction, stroke and peripheral artery disease in patients with CKD (Vanholder et al., 2016).

These summarized mechanisms of AhR activation in CKD support efforts to target AhR and limit the detrimental effects of uremic toxins on vascular outcomes. AhR has been considered a therapeutic target for improving the management of the cardiovascular complications of CKD (Shivanna et al., 2016). In Gondouin's study, IS and IAA elevated the production of TF in peripheral blood mononuclear and endothelial cells after AhR activation, generating a "dioxin-like" effect (Gondouin et al., 2013). Addi et al. (2019) confirmed that CH223191, an AhR-specific antagonist, could decrease the expression of TF in a process mediated by IAA in human endothelial cells. Meanwhile, Assefa et al. (2019) showed that resveratrol inhibits the activation of SRC and downregulates the activity and permeability of the vascular endothelium by blocking the IS/AhR pathway, thereby provoking protective effects.

AhR and Anemia

Renal anemia is common in CKD patients and is an important risk factor for CKD progression and mortality (Mohanram et al., 2004; Levin et al., 2006). The prevalent cause of renal anemia is the insufficient production of erythropoietin (EPO) that results from renal interstitial fibrosis (Souma et al., 2015). The transcription of EPO mRNA is mediated by a hypoxia-inducible transcription factor (HIF) formed by HIF-as (including HIF-1a and HIF-2 α) and ARNT. The expression of HIF- α s is strictly regulated by the concentration of cellular oxygen molecules. HIF- α degradation is inhibited under hypoxic conditions, and the resulting nuclear accumulation of HIF-a and its dimerization with ARNT promotes the expression of target genes including EPO. Because both HIF and AhR are members of the PASdomain protein family and contain basic helix loops, AhR competes with HIF-a subunits to combine with ARNT and form heterodimeric transcription factor complexes in the nucleus (Soshilov and Denison, 2008). In individuals with CKD, uremic toxins elevate AhR levels and may provoke the accumulation of AhR-ARNT complexes while inhibiting the formation of HIF-α-ARNT complexes in the nucleus, leading to abnormal intracellular HIF signal transduction and the transition of renal interstitial EPO-producing tissues into a pathological fibrotic state associated with impaired EPO production capacity (Souma et al., 2013, 2016; Mascanfroni et al., 2015).

Asai et al. (2016), first reported that IS suppresses hypoxic HIF activation and inhibits the production of EPO in rats and HepG2 cells, confirming the presence of IS-impaired HIF signaling due to AhR activation in renal anemia. Later, the same researchers found that indole glucuronic acid inhibits the expression of HIF-dependent EPO by activating AhR, whereas the drug antagonist

CH-223191 blocks AhR and abolishes the inhibiting effect of indole glucuronide on HIF (Asai et al., 2018).

Renal anemia has the same signs as iron deficiency anemia, which is caused by impaired iron utilization. Hepcidin, a liversecreted protein, is essential for iron metabolism regulation (Park et al., 2001); it induces the internalization and degradation of a cellular iron exporter called ferroportin that regulates intracellular iron efflux (Nemeth et al., 2004). Therefore, the increased hepcidin levels associated with CKD cause duodenal iron absorption disorder and intracellular iron retention, which aggravate the functional iron deficiency state in CKD. Hamano et al. (2018) showed that the expression of hepcidin is increased by IS in a dose-dependent manner and suppressed by AhR silencing in HepG2 cells.

AhR and Bone Disorders

Chronic kidney disease mineral and bone disorders are generally characterized by changes in bone metabolism. These systemic disorders are common in patients with CKD and are characterized by calcium and phosphorus abnormalities, vitamin D deficiency, vascular calcification, secondary hyperparathyroidism and bone abnormalities (Kidney Disease: Improving Global Outcomes (KDIGO) CKD-MBD Update Work Group, 2017).

Vidal et al. (2015) observed that the amount of tryptophan degraded through the kynurenine pathway increases significantly during the formation of osteoblasts and that osteoblast formation is inhibited when IDO1, the enzyme that stimulates the conversion of tryptophan to kynurenine, is blocked. Data published by Refaey et al. (2017) showed that intragastric or intraperitoneal kynurenine administration impairs osteoblast differentiation and increases osteoclast absorption to accelerate age-related bone loss.

Aryl hydrocarbon receptor is a cytoplasmic receptor of several low-molecular-weight exogenous and endogenous molecules that can control bone homeostasis by regulating osteoclast differentiation through the RANKL/AhR/c-Fos signaling axis (Izawa et al., 2016). Exposure to TCDD (the most widely known exogenous AhR ligand) weakens the mechanical strength of bone, hardens the bone matrix and makes the cortical bone thinner and looser in wild-type mice, but these effects are diminished in AhR-knockout mice (Herlin et al., 2013). In addition, AhRknockout mice exhibit decreased bone resorption and increased bone mass. Kynurenine has been identified as an endogenous AhR agonist (Opitz et al., 2011). Kalaska et al. (2017) observed that the level of peripheral kynurenine is negatively correlated with bone biomechanical variables, bone geometric variables and bone mineral density, and that high peripheral kynurenine levels cause pathological changes in bone microstructure and strength through the AhR pathway in CKD rats that had undergone subtotal nephrectomy.

AhR and Cognitive Dysfunction

The severity of cognitive impairment varies from mild cognitive impairment to severe dementia and is characterized by the loss of independence in daily activities (American Psychiatric Association, 2013). Cognitive impairment may already be present

during the early stages of CKD, and it becomes more severe as CKD progresses (Bugnicourt et al., 2013; Brodski et al., 2019). In a 6-year longitudinal study conducted in the general population, the impact of CKD on the risk of mild cognitive impairment and dementia exceeded the impacts of long-term use of anti-anxiety drugs, stroke or genetic factors (Lipnicki et al., 2017).

Factors such as white matter damage, asymptomatic infarction and microhaemorrhages may be involved in cognitive impairment (Bugnicourt et al., 2013), but vascular injury is the basic factor underlying all these pathological manifestations. AhR is thought to play an essential role in the vascular dysfunctions seen in patients with cardiovascular disease.

Aryl hydrocarbon receptor is expressed in the cerebral cortex, hippocampus, cerebellum and other sites of the brain (Lin et al., 2008), and the receptor is thought to be associated with cognitive dysfunction caused by oxidative stress or excitotoxicity (Kim and Yang, 2005; Williamson et al., 2005). Walker et al. (2020) found that activated AhR in brain microvascular endothelial cells in CKD models induced by adenine and IS treatment (in an uremic environment) plays a role in the biology and pathology of central nervous system endothelium. An increase in the production of reactive oxygen species (ROS) aggravates inflammation in patients with neurodegenerative diseases (Hsieh and Yang, 2013). In Adesso's study, increased expression of AhR in primary astrocytes and mixed glial cells was found after IS treatment; in addition, ROS production was significantly reduced after the addition of an AhR inhibitor, showing that the AhR/ROS pathway is activated by IS in the central nervous system of patients with CKD (Adesso et al., 2018).

AhR and Protein Energy Waste (PEW)

Protein energy waste (PEW) is a common complication with adverse consequences in patients with CKD. PEW is characterized by the gradual loss of protein and energy fuels stored in the body (i.e., body muscle and fat mass) (Ikizler et al., 2013). The loss of body muscle mass is caused by many factors, including insufficient energy and protein intake, protein loss in the urine and chronic inflammation (Carrero et al., 2013).

Enoki et al. (2016) demonstrated that IS activates the AhR/NADPH oxidase/NF- κ B pathway, leading to increased ROS production, which then triggers the production of inflammatory markers and increases the expression of atrogin-1, myostatin, IL-6 and TNF- α to cause muscle atrophy. AhR inhibitors (e.g., CH-223191) and AhR-siRNA significantly inhibit the production of inflammatory cytokines (ROS and atrophy-related genes in C2C12 myoblasts), suggesting that the activation of AhR in cells is associated with muscle atrophy (Enoki et al., 2016).

AhR and Drug Metabolism

Patients with CKD have high rates of complications and comorbidities. Thus, they require the prescription of numerous medications. However, the pharmacokinetics of drugs in these patients are changed by the uremic environment, making drug dosage adjustments challenging and drug-related adverse events frequent. However, the cellular and molecular mechanisms involved remain unclear. There are more than 200 P450 proteins, and AhR signaling regulates three of them (CYP1B1, CYP1A1,



and CYP1A2) (Fujii-Kuriyama and Mimura, 2005). Mutsaers et al. (2015a) showed that uremic toxins reduce the activity of some drug transporters. Santana et al. (2018) proved that IS affects the blood levels of cyclosporine by regulating transport proteins such as P-glycoprotein through AhR activation. Therefore, AhR seems to be associated with the altered drug clearance seen in CKD. The regulation of AhR activity may be a target for improving drug clearance in patients with CKD.

AHR ENHANCES INTESTINAL BARRIER FUNCTION

As discussed above, AhR mediates renal damage and CKD complications and seems to be harmful in CKD patients (Figure 2). However, AhR is thought to play an active role in intestinal homeostasis (Figure 3). AhR was found to be expressed in almost all colonic muscularis layers, whereas it

was absent from the duodenal and jejunal muscular layers. At the same time, only relatively weak signals were detected in ileum terminal neurons, suggesting that AhR expression in intestinal neurons is synchronous with the intestinal microbial load (Obata et al., 2020). AhR influences different aspects of intestinal barrier function. First, AhR is considered important for specific monitoring pathways of the intestinal nervous system that regulate intestinal responses to colonization, the maintenance of intestinal functions and host defenses (Obata et al., 2020). Therefore, pharmacological or dietary interventions that modulate AhR activity in the cellular circuitry to control intestinal peristalsis offer a realistic strategy for the management of conditions associated with gut dysmotility, including the constipation due to slow colonic transit common in patients with CKD.

In addition to the role of AhR in neurogenic motility, AhR-dependent transcriptional procedures are also essential for the barrier function of intestinal epithelial cells (IECs)



(Metidji et al., 2018; Rothhammer and Quintana, 2019). AhR in intestinal stem cells stabilizes intestinal epithelial barrier function and controls the regeneration of IECs. Lack of IECspecific AhR results in the failure to protect against Citrobacter rodentium infection due to reduced quantities of mucusproducing goblet cells (Metidji et al., 2018). AhR activation by dietary ligands may restore barrier homeostasis and protect the stem cell niche. Furthermore, AhR can upregulate epithelial tight junctions. 6-formylindolo[3,2-b]carbazole (FICZ), a highaffinity endogenous ligand of AhR, prevents the decrease of ZO-1, occludin and claudin-1 to modulate intestinal epithelial barrier function in dextran sulfate sodium-induced IECs (Yu et al., 2018). A metabolite of Lactobacillus, indole-3-aldehyde, stimulates lamina propria lymphocytes to secrete AhR/IL-22 pathway products and induces the phosphorylation of STAT3 to accelerate intestinal epithelial proliferation, thereby healing damaged intestinal mucosae (Hou et al., 2018).

Furthermore, the expression of AhR by many adaptive and innate immune cells in the lamina propria, including macrophages, mast cells, eosinophils, dendritic cells (DCs), B cells and T cells, suggests that AhR signaling is essential for the regulation of intestinal mucosal immune responses. DCs are prototypic targets of environmental exposure and endogenous AhR ligands (Quintana et al., 2015). Macrophages and DCs produce high levels of IL-1 β , which promotes neutrophil recruitment and pathogen elimination. AhR is also essential for the function of innate lymphoid cells (ILCs) and Th17/Th22 cells that control symbiosis and pathogens through IL-22-pSTAT3 (Wang et al., 2014; Piccinni et al., 2019) and mediate the production of antimicrobial peptides by IECs (Sonnenberg et al., 2011). Antimicrobial peptides can also participate in colonization resistance against pathogens by shaping the microbiota, which compete with pathogens for nutrients. Genetic AhR ablation enhances gut ILC2 function during anti-helminth immunity, and cell-specific activation of AhR enhances gut ILC3 function for anti-bacterial immunity, suggesting the central role of AhR expression for gut adaptation and ILC2-ILC3 balance (Li et al., 2018).

Since Meijers proposed the existence of a gut-kidney axis (Meijers and Evenepoel, 2011), some studies have discovered that damage to the gut barrier function induced by intestinal dysbiosis in CKD allows the translocation of endotoxin and bacterial metabolites into the circulation, which causes uremic toxicity, inflammation, CKD progression and cardiovascular disease (Meijers et al., 2019). As our understanding of the association between the gut and the kidney continues to grow, various therapies targeting the colonic microenvironment in CKD have been developed, such as regulating gut microbiota, blocking lipopolysaccharides, attenuating inflammation or targeting uremic toxin end-products of microbial fermentation (Ramezani and Raj, 2014). Although few therapies target AhR to improve intestinal barrier function in patients with CKD, the potential exists to uncover many novel avenues to delay CKD progress; for example, baicalein, a flavonoid from Scutellaria baicalensis used in Chinese herbal medicine, induces regulatory T-cell differentiation through AhR and enhances intestinal barrier function through its regulation of tight junctions in a mouse model of food allergies (Bae et al., 2016).

CONCLUDING REMARKS

Initial studies on AhR were focused on its role as a chemical sensor signaling molecule responding to environment pollutants, but the range of subjects researched has gradually expanded to include diseases such as cancer and cardiovascular and kidney disorders. AhR mediates CKD progression and its complications, including cardiovascular disease, anemia, bone disorders, cognitive dysfunction and malnutrition, and it affects the drug metabolism of individuals with CKD. On the other hand, AhR enhances intestinal barrier function and reduces the harmful effects of uremic toxins. Therefore, whether AhR helps patients with CKD to deal with the pathological changes they face or increases their risk for pathologic complications remains unclear. Based on the available data, we could not draw a definite conclusion about a positive role for AhR in intestinal homeostasis, but we know that it mediates renal damage and CKD complications outside the gastrointestinal tract. AhR-mediated detoxification mechanisms stimulate the clearance of endobiotics and xenobiotics by the kidney. AhR may help individuals to survive in changing environments or promote self-recovery during the early stages of disease, but once the normal physiological or protective responses of the AhR pathway are interrupted, the severity of disease increases. Regulating the AhR pathway is without a doubt an attractive treatment strategy for CKD. However, more studies are needed to overcome the interpretation challenges of the current data.

First, AhR ligands have different sources and structures, present diverse receptor affinities and produce different downstream signal transduction effects. Whether a certain type of AhR ligand is beneficial or detrimental under a specific situation needs to be determined. New metabolomics methods are helping researchers to discover, isolate and identify new exogenous and endogenous AhR ligands. An effective AhR ligand may not be able to play a "one size fits all" role throughout the entire CKD treatment process. In fact, even in a single model, an AhR-ligand combination may be temporarily effective in terms of protecting the body, but as the disease progresses, the interaction may turn harmful, or vice versa. In view of

the conflicting observations on whether AhR promotes or inhibits inflammation, identifying and developing specific high-affinity AhR ligands to slow down the development of CKD and its complications is difficult. The ligands need to display targeted absorption, distribution, metabolism and excretion characteristics.

Second, a food-as-medicine approach could be used as a strategy based on bioactive nutrients to target the AhR pathway in patients with CKD. Natural bioactive compounds, including galangin, quercetin, kaempferol, resveratrol and dietary metabolites of cruciferous vegetables, are AhR ligands and potential nutritional therapeutic agents that may modulate the expression of pro-inflammatory factors. Suitable dietderived AhR ligands should be considered during the dietary management of nutrients in patients with CKD, to put the ancient adage of "let food be thy medicine" into practice. Moreover, bioactive nutrients such as flavonoids and polyphenol compounds are widely distributed in natural products such as medicinal plants, tea, fruits and vegetables. In view of the important roles of the above-mentioned compounds in regulating AhR activity, further studies should address the regulation of AhR activity by natural compounds, to reveal their potential as therapeutic agents for CKD and demonstrate their molecular mechanisms.

Third, inducing the organism to produce appropriate AhR ligands is also a potential therapeutic strategy. Both indole-3-pyruvate (IPYA) and IS are derived from gut bacterial metabolism of the essential amino acid tryptophan. IPYA depends on AhR; it protects the intestine against excessive permeability by maintaining the integrity of apical junctional complexes and the associated actin regulatory proteins (Mafra et al., 2020), whereas IS induces intestinal barrier injury by impairing IRF1/DRP1 axis-mediated mitophagy (Huang et al., 2020). Therefore, inhibiting or promoting certain enzyme activities or regulating the intestinal flora for the metabolism of some key substances may induce the production of AhR ligands that are beneficial to the body.

Finally, although AhR is a promising target for clinical applications, most of the knowledge about its physiological and pathological functions in CKD comes from animal models, and the translation of experimental results into clinical applications for patients with CKD is difficult. Therefore, further research is needed to better understand the complex roles AhR plays in individuals with CKD, so as to be able to use it as a safe and effective target for CKD prevention and treatment.

AUTHOR CONTRIBUTIONS

XL: conceptualization. YM, ZL, CJ, and XL: literature search. YM, ZL, and XL: writing, formatting, and figures. LW, CZ, and XL: review and editing. All authors read and approved the final manuscript.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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