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# Editorial: Diabetic kidney disease: routes to drug development, pharmacology and underlying molecular mechanisms

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#### Editorial on the Research Topic

Diabetic kidney disease: routes to drug development, pharmacology and underlying molecular mechanisms

# Introduction

Diabetic kidney disease (DKD) is one of the leading causes of kidney failure, which leads to end-stage renal disease (ESRD) and affects almost one-third of the total diabetic patients worldwide (Cooper and Warren, 2019; Tuttle et al., 2022). DKD is also one of the significant long-term complications, which not only imposes healthcare costs on patients with diabetes but substantially increases the rates of morbidity and mortality. A wide knowledge gap exists between biology and drug development which contributes to the suboptimal treatment options against DKD. An increased understanding of DKD is urgently needed for the development of novel therapeutics, which will help in targeting the changes that occur during the early stages of the disease. Several attempts have also been made to hybridize the process of biomarker discovery and drug development to arrest and reverse the inevitable events, which occur during the progression of DKD. However, currently approved therapeutic regimens such as angiotensin-converting enzyme (ACE) inhibitors, angiotensin II receptor blockers (ARBs), and statins, etc. can simply delay, but do not prevent the decline in kidney function and progression of ESRD (Gentile et al., 2014; Srivastava et al., 2020a; Hartman et al., 2020). Although these therapeutic agents help in reducing albuminuria in patients with DKD, they expose patients to multiple adverse reactions and drug intolerance. Thus, to improve kidney function in diabetic patients, there is an unmet need for new therapeutic strategies.

A variety of pharmacological agents are currently in clinical trials such as sodium glucose transporter-2 (SGLT-2) inhibitors and in preclinical settings including dipeptidase transferase-4 (DPP-4) inhibitors, mineralocorticoid receptor antagonists, N-acetyl-seryl-

aspartyl-lysyl-proline (Ac-SDKP), and endothelin receptor antagonists (ERAs) (Kanasaki et al., 2014; Li et al., 2020b; Srivastava et al., 2020b). These molecules have been shown to exhibit protective effects against increases in inflammatory and fibrotic responses in the kidney and worsening of kidney function in preclinical settings.

In this Research Topic, we provided a deeper insight into the recent advances in the pathophysiologic pathways, which can be modulated to prevent the worsening of diverse phenotypes of DKD. We also addressed the molecular mediators, which can be targeted to attenuate the progression of kidney fibrosis during DKD. Understanding these critical pathways and potential mediators will guide the discovery process of future therapeutic approaches against DKD.

Broadly we categorized this editorial into two major sections.

#### New molecular mechanisms in DKD

In recent years, we have worked on identifying new cellular and molecular mechanisms causative or protective in DKD. Defective central metabolism, i.e., reduced fatty acid oxidation (FAO) and induced abnormal glycolysis, accelerates the mesenchymal switching, and finally fibrogenesis (Kang et al., 2015; Srivastava et al., 2018). Dichloroacetate, 2-deoxyglucose, and C75 by correcting defective metabolism via promoting FAO levels in tubules, help in abolishing the fibrotic phenotype and improving kidney structure and functions (Kang et al., 2015). Sirtuin (SIRT)3 is the key regulator of central metabolism. SIRT3 induction provides protection to both tubules and kidney vasculature. SIRT3 abrogates abnormal glycolysis and improves lipid metabolism by transconversion of pyruvate kinase isozymes M1/M2 (PKM2) dimer to tetramer. SIRT3 also reduces the accumulation of hypoxia-inducible factor (HIF)-1α in the kidney tubules and endothelium (Srivastava et al., 2018; Srivastava et al., 2021b).

In this Research Topic, Li et al. described that Yin and Yang (YY)1 is involved in the upregulation LncRNA-ARAP1-AS2, which by promoting aberrant glycolysis caused fibrosis. The impaired glycolysis was associated with the activation of epidermal growth factor receptor (EGFR)/PKM2/HIF1 $\alpha$  pathway. Their results highlight the role of YY1 in promoting glycolysis during DKD and suggest it as a potential therapeutic strategy for consideration.

Other molecules, i.e., endothelium glucocorticoid receptors (GR) and podocyte glucocorticoid receptors are critical in restoring cellular antifibrotic mechanisms, abolishing the mesenchymal metabolic shift and, associated kidnev inflammation and fibrosis in diabetes (Srivastava et al., 2021c; Srivastava et al., 2021d). Glucocorticoid receptor negatively targets canonical Wnt signaling pathway, inflammation, and improves tubular and endothelial health (Srivastava et al., 2021d). Kidney endothelium-specific deficiency of SIRT3, GR, or fibroblast growth factor receptor (FGFR)1 accelerates the endothelial-tomesenchymal transition (EndMT) by disrupting neighboring cell homeostasis and promoting epithelial-mesenchymal transition (EMT) in tubular cells during diabetes (Li et al., 2020a; Srivastava et al., 2021b; Srivastava et al., 2021d). This EndMTmediated activation of partial EMT is caused due to systemic inflammation, accompanied by elevating levels of promesenchymal and fibrotic cytokines, and reducing levels of antimesenchymal cytokines. Cumulative effects of systemic inflammation, defective central metabolism, and upregulated mesenchymal transcription factors, finally contribute to severe fibrosis in diabetic kidneys. Another article in this Research Topic described the role of nucleotide-binding oligomerization domain (NOD)-like receptors (NLPR3)-mediated pyroptosis in diabetic nephropathy (Wan et al.). This study demonstrated that the canonical NLRP3 inflammasome pathway, which involves the formation of membrane pores, and promotes inflammation, can also induce pyroptosis in the kidneys during DKD. This study suggests that targeting pyroptosis-associated proteins has therapeutic potential in the management of diabetes and related kidney disease.

Another study in this Research Topic by Cen et al. reported the protective effects of tetramethylpyrazine nitrone (TBN) in improving renal anemia, by maintaining iron homeostasis through activating the HIF via 5' adenosine monophosphateactivated protein kinase (AMPK)/mammalian target of rapamycin (mTOR)/4E-binding protein 1 (4E-BP1) pathway. TBN, a nitrone derivative of tetramethylpyrazine also acts as a free radical scavenger against hydroxyl, superoxide, and peroxynitrite. Recently, we reported the therapeutic benefits of repleting the labile iron pool (LIP) of kidney macrophages in improving their antioxidant response and attenuating inflammation, and fibrosis during chronic kidney disease (CKD) (Patino et al., 2023). We also observed that the expression of superoxide dismutase-2 (SOD-2), an antioxidant enzyme decreased in the kidney during CKD, which was associated with an increase in mROS production by kidney macrophages (Bhatia et al., 2022). Therefore, it is critical to further investigate the therapeutic benefits of TBN as reported by Cen et al. against increases in oxidative stress, inflammation, and CKD-associated anemia.

### New molecules against DKD

The discovery of SGLT-2 inhibitors has influential effects in the treatment of diabetes and associated kidney disease. The data from preclinical settings suggest that SGLT-2 inhibitors not only reduced worsening of kidney function and fibrosis, but, improved the overall structural impairments in the kidney of patients with DKD. The data from EMPA-REG OUTCOME clinical trials demonstrated that the use of SGLT-2 inhibitor, empagliflozin is safe and effective in reducing renal-related complications in patients with type 2 diabetes, suggesting the remarkable discovery in the treatment of DKD (Mayer et al., 2019). The SGLT-2 inhibitors provide protection against DKD through promoting the excretion of blood glucose in the urine and reducing intraglomerular pressure. However, the data from the diabetic mouse models have shown that the empagliflozin did not cause any change in blood pressure and blood glucose levels. These data from experimental models suggested that empagliflozin was able to reduce the aberrant glycolysis and induce the FAO in the tubules (Li et al., 2020b). Increased aberrant glycolysis and diminished FAO promote mesenchymal metabolic shift, which contributes to the deterioration of kidney function, structural abnormalities, and kidney fibrosis (Srivastava et al., 2018).

In this Research Topic, Hu et al. performed a systematic review to understand the efficacy and safety of SGLT-2 inhibitors in patients stratified in different groups based on their estimated glomerular filtration rate (eGFR). Based on the search using four databases, they reported that the SGLT2 inhibitors decreased the glycated hemoglobin (HbA1c) levels and body weight, which were in parallel with the baseline eGFR levels. However, SGLT2 inhibitormediated reductions in blood pressure levels were independent of the baseline eGFR in patients. Other reported potential therapeutic mediators, which have also shown to be effective against DKD include DPP-4 inhibitors and incretins analogs. Among different classes of DPP-4 inhibitors, linagliptin has been reported to be most effective in combating diabetes and associated kidney disease (Kanasaki et al., 2014). Linagliptin intake increased the levels of Ac-SDKP tetrapeptide and exhibited renal protection by delaying the progression of kidney disease.

Plant-derived synthetic compounds such as ursolic acid, paeoniflorin, and saponins are found to be effective against DKD. Interestingly ursolic acid improved the renal structural impairments and function through elevating FGFR1 and SIRT3 levels and reducing the expression of DPP-4 (Liu et al.). DPP-4 is known to exhibit profibrotic functions in the tubules. FGFR1 activation is related to the upregulation of microRNAs (miRNAs), such as miRlet-7s and miR-29s, these miRNAs exert antifibrotic effects (Srivastava et al., 2016). In addition, induction of SIRT3 also exhibits protective effects by abolishing the mesenchymal metabolism shift. Ac-SDKP mediates renal protection by binding with FGFR1 and elevating SIRT3 levels (Li et al., 2017; Srivastava et al., 2020b). Therefore, this tetrapeptide has great potential as a future therapeutic option for combating diabetes and related kidney disease (Srivastava et al., 2021a). The author Fang et al. reported that the Niaoduqing molecules mitigate high-glucose-induced podocyte injury by regulating advanced glycation endproducts/receptor for AGE (AGE/RAGE) signaling pathway.

The activation of SIRT3 is also known to attenuate NLRP3-mediated inflammation, apoptosis, and kidney fibrosis during DKD by inducing mitophagic activities (Feng et al., 2018). Mitochondrial-targeted approaches have also shown great efficacy against DKD in experimental models and patients (Bhatia et al., 2020). Mitoquinone (MitoQ), a mitochondrial-targeted coenzyme Q has been reported to mitigate albuminuria, glomerular hypertrophy, mesangial matrix expansion, and kidney fibrosis by restoring the expression of mitophagy regulators, including mitofusin (MFN)-2 and reducing mitochondrial fragmentation (Xiao et al., 2017). We have also observed the myeloid lineage-specific protective functions of MFN2 against increases in kidney macrophage-derived inflammatory and fibrotic responses during CKD (Bhatia et al., 2022). The efficacy of MitoQ was also investigated as part of a Phase IV controlled, doubleblind clinical trial in stage 3-5 patients with CKD (ClinicalTrials.gov, Identifier number: NCT02364648). Szeto-Schiller-31 peptide (H-D-Arg-Dmt-Lys-Phe-NH2) (SS-31), a mitochondrial antioxidant also reduced macrophage recruitment, glomerulosclerosis, podocyte and endothelial cell death in the experimental model of DKD (Szeto et al., 2016).

In this Research Topic, a study by Wang et al. reported that sappanone A (SA), a plant-derived homoisoflavanone, prevented the worsening of kidney function in uninephrectomized and streptozotocin-induced DKD, by restoring the expression of nuclear factor kappa B (NF- $\kappa$ B) inhibitor alpha (I $\kappa$ B $\alpha$ ). Their findings

highlighted the protective effects of SA against increases in proinflammatory (interleukin-1 $\beta$  and tumor necrosis factor- $\alpha$ ) and profibrotic (transforming growth factor- $\beta$ 1 and collagen-IV) responses in the kidney during DKD. Restoration of IkB $\alpha$  is also critical in averting mitochondrial fragmentation and cytochrome (Cyt C)-mediated apoptotic cell death (Laforge et al., 2016). We also observed increases in circulating and urinary levels of Cyt C in experimental and human kidney fibrosis (Bhatia et al., 2022). The increased levels of Cyt C were associated with macrophage-mediated inflammatory and fibrotic events in the kidney (Bhatia et al., 2022). Therefore, the study by Wang et al. is important in confirming, that the SA-dependent protective effects against increases in inflammation and cell death during DKD are exerted via suppression of NF- $\kappa$ B pathway.

Another study by Chen et al. reviewed the therapeutic benefits of Chinese herbal medicines in suppressing oxidative stress, inflammation, and mitochondrial dysfunction. They highlighted that combining these herbal medicines with conventional therapies may further delay the progression of DKD effectively. These herbal medicines have been shown to modulate various pathways, including, i) Glucagon-like Peptide-1 (GLP)-receptor, ii) SGLT2, iii) SIRT1/ AMPK, iv) AGE/RAGE, v) NF-κB, vi) nuclear factor erythroid 2-related factor 2 (Nrf2), vii) NLRP3, viii) Peroxisome proliferator-activated receptor-gamma coactivator (PGC)-1a, and ix) PINK1/Parkin-mediated mitophagy. We also reported that PINK1/Parkin-mediated mitophagy is downregulated in experimental and human kidney fibrosis (Bhatia et al., 2019). The expression of PGC-1a, which is a chief regulator of mitochondrial biogenesis has also been found to be suppressed in the kidney during fibrosis (Bhatia et al., 2022). Fenofibrate has been reported to promote mitochondrial fatty acid β-oxidation and regulate kidney function in patients with DKD by activating PGC-1a (Davis et al., 2011). Considering that kidneys are the second richest organ after the heart in mitochondrial content, it is anticipated and widely reported that mitochondrial dysfunction exaggerates the worsening of kidney diseases including DKD (Forbes and Thorburn, 2018; Bhatia and Choi, 2019; Bhatia et al., 2020). Mitophagy helps in the recycling of damaged mitochondrial components, which can act as mitochondrial damaged-associated molecular patterns (mtDAMPs) and promote inflammatory events in the kidney during DKD (Bhatia and Choi, 2023). These mtDAMPs, which include mitochondrialderived reactive oxygen species (mROS), Cyt C, and cell-free or intracellular naked mitochondrial DNA (mtDNA) can exaggerate inflammation and fibrosis via activating NF-kB by toll-like receptor (TLR)-mediated signaling pathway. The mechanism of action of Chinese herbal medicines against the progression of DKD warrants further investigation.

# Conclusion

Diabetic kidney disease is the chief cause of kidney failure and ESRD, which also increases cardiovascular events and mortality in the suffering patient population globally. It is important to understand the underlying mechanisms of diabetic milieuinduced kidney damage and progression of DKD. In this Research Topic, we have discussed potent therapeutic regimens for the management of DKD, tissue/cell/mitochondrial-specific novel biological mechanistic approaches, functions of regulatory non-coding RNAs, and clinical data sets. We speculate that this Research Topic will provide basic essential information, that could be used in the design of potential therapeutic agents and help in the management of patients with DKD.

# Author contributions

SS and DB wrote and edited the draft. All authors contributed to the article and approved the submitted version.

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## **Conflict of interest**

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