

An Indole-2-Carboxamide Derivative, LG4, Alleviates Diabetic Kidney Disease Through Inhibiting MAPK-Mediated Inflammatory Responses

Jianchang Qian^{1,*}Sihui Yin^{1,*}Lin Ye¹Zhe Wang²Sheng Shu^{1,3}Zhenxin Mou^{1,3}Mingjiang Xu¹

Nipon

Chattipakorn⁴Zhiguo Liu¹Guang Liang^{1,3,5}

¹Chemical Biology Research Center, School of Pharmaceutical Sciences, Wenzhou Medical University, Wenzhou, Zhejiang, People's Republic of China;

²Department of Pharmacy, The Second Affiliated Hospital of Wenzhou Medical University, Wenzhou, Zhejiang, People's Republic of China; ³School of Pharmacy, Hangzhou Medical College, Hangzhou, Zhejiang, People's Republic of China;

⁴Cardiac Electrophysiology Research and Training Center, Faculty of Medicine, Chiang Mai University, Chiang Mai, Thailand; ⁵Wenzhou Institute, University of Chinese Academy of Sciences, Wenzhou, Zhejiang, People's Republic of China

*These authors contributed equally to this work

Correspondence: Guang Liang; Zhiguo Liu
Chemical Biology Research Center, School of Pharmaceutical Sciences, Wenzhou Medical University, Wenzhou, 325035, People's Republic of China
Tel/Fax +86-577-86699396
Email wzmclianguang@163.com; lzgcnu@163.com

Aim: Elevated inflammatory signaling has been shown to play an important role in diabetic kidney disease (DKD). We previously developed a new anti-inflammatory compound LG4. In the present study, we have tested the hypothesis that LG4 could prevent DKD by suppressing inflammation and identified the underlying mechanism.

Methods: Streptozotocin-induced type 1 diabetic mice were used to develop DKD and evaluate the effects of LG4 against DKD. To identify the potential targets of LG4, biotin-linked LG4 was synthesized and subjected to proteome microarray screening. The cellular mechanism of LG4 was investigated in HG-challenged SV40MES13 cells.

Results: Although LG4 treatment had no effect on the body weight and blood glucose levels, it remarkably reversed the hyperglycemia-induced pathological changes and fibrosis in the kidneys of T1DM mice. Importantly, hyperglycemia-induced renal inflammation evidenced by NF- κ B activation and TNF α and IL-6 overexpression was greatly ameliorated with LG4 treatment. Proteome microarray screening revealed that JNK and ERK were the direct binding proteins of LG4. LG4 significantly reduced HG-induced JNK and ERK phosphorylation and subsequent NF- κ B activation in vivo and in vitro. In addition, LG4 did not show further anti-inflammatory effect in HG-challenged mesangial cells with the presence of JNK or ERK inhibitor.

Conclusion: LG4 showed renoprotective activity through inhibiting ERK/JNK-mediated inflammation in diabetic mice, indicating that LG4 may be a therapeutic agent for DKD.

Keywords: indole-2-carboxamide derivative, diabetic kidney disease, inflammation, MAPK, NF- κ B

Introduction

Although a great improvement has been made in blood glucose management, the number of patients with diabetic kidney disease (DKD) is still growing due to the increasing of diabetic population.¹ Microalbuminuria is a key feature in the initial stage of DKD (30–300 mg/day), and macroalbuminuria (>300 mg/day) and overt nephropathy occur as the disease progressed.^{2–4} Meanwhile, glomerular dysfunction and morphological changes that lead to subsequent development of glomerulosclerosis and nephron dropout will be seen with or without tubular atrophy, tubulointerstitial fibrosis and renal arteriolar hyalinosis.^{5–8} Although the progression of DKD could be slowed with tight glycemic control and antihypertensive/lipid-lowering therapies, these

interventions do not prevent the progress of DKD to end stage of renal disease (ESRD) or renal failure.⁹

Chronic inflammation in the kidney plays a key role in the development and progression of DKD.^{10,11} Hyperglycemia induces glomerular hyperfiltration, mechanical stress, and glycocalyx dysfunction that result in the overexpression of proinflammatory cytokines such as tumor necrosis factor α (TNF- α), interleukin-6 (IL-6) and interleukin-1 β (IL-1 β) through activating mitogen-activated protein kinases (MAPKs) and NF- κ B signaling pathways in the kidney.^{12,13} Stress-activated protein kinases, especially p38 mitogen activated protein kinase and the c-Jun amino terminal kinase signaling, are the validated upstream transducer that modulate activation of NF- κ B in hyperglycemia.¹⁴ These inflammatory responses are also causal factors of glomerulosclerosis and tubulointerstitial fibrosis by inducing extracellular matrix deposition and differentiation/proliferation of myofibroblast. The underlying signaling pathways include NF- κ B, JAK/STAT, TGF- β /Smad, etc.^{15,16} Therefore, targeting MAPK/NF- κ B transduction could be a promising strategy to block hyperglycemia-induced inflammation for DKD treatment.

Indole is an important scaffold in the field of medicinal chemistry. Many indole derivatives possess significant pharmacological activities such as anti-inflammatory, analgesic and antipyretic effects.^{17–20} For instance, indomethacin is one of the indole acetic acid derivative and a non-steroidal anti-inflammatory drug (NSAID), it however is known to cause ulcers for the users, and its safety profile has been improved by chemical modifications.²¹ We had designed and synthesized a series of indole derivatives and evaluated their anti-inflammatory property by measuring the production of TNF- α and IL-6 in activated macrophages and LPS-induced sepsis in mice. Among them, an indole-2-carboxamide derivative, LG4, exhibited significant anti-inflammatory effects in vitro and in vivo.²² In this study, we explored the potential effect of LG4 on chronic inflammation and explored the underlying molecular mechanisms in a mouse DKD model and mesangial cells.

Materials and Methods

Chemicals and Reagents

Compound LG4 was synthesized as described previously.²² The chemical synthesis of biotin-labeled LG4 was described in the [Supplementary File](#). UO126 and SP600125 were purchased from MedChemExpress (Monmouth Junction, NJ). The compounds used in the study were dissolved in dimethyl

sulfoxide (DMSO) at 20 mM as stock solutions. Other reagents used in this study were purchased from Sigma-Aldrich (Burlington, MA, USA).

Cell Culture

A mouse glomerular mesangial cell line, SV40MES13, was obtained from Shanghai Institute of Biochemistry and Cell Biology (Shanghai, China) and cultured in low-glucose (5.5 mM) DMEM (Gibco, Eggenstein, Germany) containing 5% fetal bovine serum (FBS), 100 U/mL penicillin, and 100 mg/mL streptomycin at 37°C with 5% CO₂. When cells are treated with high-concentration glucose (HG), DMEM with 33 mM glucose was used.

Animal Model

Male C57BL/6 mice were obtained from Animal Center of Wenzhou Medical University and housed at a constant room temperature with a 12:12 h light-dark cycle, chow diet and water *ad libitum*. Animal study protocols were approved by Wenzhou Medical University Animal Policy and Welfare Committee in accordance with the NIH guidelines (Guide for the care and use of laboratory animals). Type 1 diabetic (T1DM) mouse model was prepared as previously described.²³ Briefly, diabetes was induced in the mice at 8 weeks of age by intraperitoneal injection of 50 mg/kg streptozotocin (STZ, 10 mL/kg in citrate buffer, pH 4.5) for five consecutive days, while the control animals were received citrate buffer in the same schedule. One week later after first STZ dosing, blood glucose was measured through tail vein using a glucometer. Mice with fasting blood glucose >300 mg/dL were considered as T1DM.

Eight weeks after setting up the T1DM, randomly selected diabetic animals were divided into three groups, ie, T1DM+vehicle, T1DM+LG4 (5 mg/kg), and T1DM+LG4 (10 mg/kg); n = 5 mice per group. LG4 (5 and 10 mg/kg) was administered by oral gavage once every two days for 10 weeks. The T1DM+vehicle and age-matched control group (n = 5) was given with vehicle (0.5% CMC-Na solution) in the same schedule. Blood glucose and body weight were recorded every two weeks. After 10 weeks of treatment, the mice were sacrificed under sodium pentobarbital anesthesia (i.p. injection of 0.2 mL sodium pentobarbital at 100 mg/mL), and blood samples and kidney tissues were collected and processed for further analyses.

Real-Time Quantitative PCR

Cells or kidney tissues were homogenized in TRIZOL (Invitrogen, Carlsbad, CA) for total RNA extraction. Reverse transcription and quantitative PCR (RT-qPCR) were performed using M-MLV Platinum RT-qPCR Kit (Invitrogen, Carlsbad, CA). Primers for genes including *Tnf- α* , *Il-6* and *β -actin* were synthesized by Invitrogen (Invitrogen, Shanghai, China), [supplementary table s1](#). The relative amount of target genes was normalized to the amount of *β -actin*.

Immunoblotting Assay

Cell and tissue lysates were separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to a polyvinylidene fluoride membrane (Bio-Rad Laboratory, Hercules, CA). The membrane was blocked for 2 hours at room temperature, and then incubated with primary antibodies overnight, followed with 2-h incubation with secondary antibody that conjugated with horseradish peroxidase. Finally, the specific bands were visualized using enhanced chemiluminescence reagents (Bio-Rad, Hercules, CA, USA). Densitometric quantification for blots was performed using ImageJ software.

The primary antibodies against BCL-2 (Cat. sc-7382, 1:200) and GAPDH (Cat. sc-47724, 1:200) were purchased from Santa Cruz Technology (Santa Cruz, CA, USA). Anti-TGF- β (Cat. ab92486, 1:1000), anti-Collagen IV antibodies (Cat. ab6586, 1:1000) and BAX (Cat. ab32503, 1:1000) were obtained from Abcam (Cambridge, MA, USA). Rabbit mAbs against p-p44/42 (Erk1/2)_Thr202/Tyr204 (Cat. 4370S, 1:1000), p-SAPK/JNK_Thr183/Tyr185 (Cat. 4668, 1:1000), P-IKK (Cat. 2697T, 1:1000) and I κ B α (Cat. 4812S, 1:1000) were purchased from Cell Signaling (Beverly, MA, USA). These antibodies were all diluted with primary antibody dilution buffer (Beyotime, Cat. P0023).

Histological Analyses

Kidney was fixed in 4% paraformaldehyde solution and paraffin embedded. Tissues were cut into 5 μ m sections and stained with Hematoxylin and Eosin (H&E), Masson's trichrome or anti-TNF- α antibody (Cat. ab6671, Abcam), respectively. Subsequently, the slides were imaged and captured under light microscope (200 \times amplification; Nikon, Tokyo, Japan). The ImageJ software (NIH) was applied to quantify the level of positive staining area. At least 6 non-overlapping fields in the view were scored and

the value determined was related to total tissue area in the field.

Glomerulosclerosis was evaluated using Masson's trichrome staining images. Glomerular sclerosis was assigned score from 0 to 4 based on sclerotic area. Grade 0 means normal; grade 1, area up to 25%; grade 2, area from 25% to 50%; grade 3, area from 50% to 75%; grade 4, area from 75% to 100%. The Glomerular sclerosis score was determined using formula: $(1 \times A_1 + 2 \times A_2 + 3 \times A_3 + 4 \times A_4) / (A_0 + A_1 + A_2 + A_3 + A_4)$, where A_x is the number of glomeruli in each grade.

Assessment of Kidney Function

The levels of urine albumin creatinine ratio (UACR), serum creatinine (SCR), and blood urea nitrogen (BUN) were detected using commercial kits (Nanjing, Jiancheng, Jiangsu, China).

Immunofluorescence Staining

The paraffin sections (5 μ m thickness) from kidney were dewaxed and dehydrated. After antigen retrieval and blocking with 5% BSA, the sections were incubated with anti-collagen IV antibody overnight at 4 $^{\circ}$ C, followed by incubation with a FITC-conjugated secondary antibody. After counterstaining with DAPI for 5 min, the slides were viewed by a fluorescent microscope (200 \times amplification; Nikon, Tokyo, Japan). The semiquantitation was performed on ImageJ to determine relative fluorescence intensity.

HuProtTM Human Proteome Chip

HuProtTM chip was blocked (5% BSA in 0.1% Tween 20, 1 \times TBST, pH=7.4) for 2 hours at room temperature with gentle agitation, and then washed with TBST (pH 7.5) for 5 minutes and repeated for three times. The chips were then incubated with D-biotin or biotin-labeled LG4 (10 μ M) in blocking buffer for 1 hour at room temperature. Rinse the chip for 3 times with TBST, and then incubate with 1% Cy5-Streptavidin solution for 20 minutes at room temperature protected from light. The microarrays were spun dried for 2 minutes and scanned with a GenePix 4000B (Axon Instruments, USA). The GenePix Pro 6.0 was used for data analysis.

Statistical Analysis

All experiments were randomized and blinded. In all in vitro experiments, data represented 3 independent experiments and expressed as means \pm SEM. Statistical analysis was performed with GraphPad Prism 6.0 software (San Diego, CA, USA). We used one-way ANOVA followed by

Dunnett's post hoc test when comparing more than two groups of data and one-way ANOVA, non-parametric Kruskal–Wallis test, followed by Dunn's post hoc test when comparing multiple independent groups. P values of < 0.05 were considered to be statistically significant. Post-tests were run only if F achieved $P < 0.05$ and there was no significant variance inhomogeneity.

Results

LG4 Treatment Improved Renal Function in T1DM Mice

The chemical structure of LG4 is shown in [Figure 1A](#). To evaluate the effect of LG4 on DKD, we treated STZ-induced T1DM mice with LG4 (at 5 and 10 mg/kg/2 days) for 10 weeks. As shown in [Figure 1B](#), the blood glucose levels of T1DM mice were higher than control group, and there was no difference in blood glucose between vehicle and LG4 treatment groups. During the administration period, mouse body weight was stable, and LG4 treatment showed no effect on the body weight of the T1DM mice ([Figure 1C](#)). Interestingly, BUN, UACR, and SCR, which are important indicators in assessing DKD, were significantly increased in the vehicle-treated group compared to the control group, and they were significantly decreased with LG4 treatment for 10 weeks in the T1DM mice ([Figure 1D–F](#)).

LG4 Attenuates Renal Fibrosis in T1DM Mice

To further evaluate the effect of LG4 on kidney morphological changes of the T1DM mice, H&E and Masson trichrome staining were conducted using the kidney tissues. Firstly, H&E staining showed classic DKD morphological changes in the kidney of the T1DM mice, including mesangial expansion, and LG4 treatment significantly improved these morphological changes ([Figure 2A](#)). Secondly, Masson trichrome staining and anti-collagen IV (COL-4) staining showed obvious accumulation of collagen in the glomerulus and tubulointerstitium of the T1DM mice, and LG4 treatment greatly mitigated these collagen accumulation in the kidney ([Figure 2A–C](#)). Meanwhile, glomerulosclerosis was assessed and scored according to Masson trichrome staining images. As shown in [Figure 2D](#), the glomerular sclerosis index of T1DM group was up to 2.6 which means moderate sclerosis compared with control group. However, treatment with LG4 significantly downgraded the glomerulosclerosis.

Furthermore, Western blot showed significant increases in COL-4 and TGF- β expression in the kidneys of the vehicle-treated T1DM mice, and they were ameliorated with LG4 administration ([Figure 2E and F](#)).

LG4 Treatment Inhibits Renal Inflammation in T1DM Mice

Chronic inflammation is one of the causing reasons for renal fibrosis in DKD. We then tested the effect of LG4 on pro-inflammatory cytokine expression in the kidney. Firstly, immunohistochemical staining showed increased TNF- α expression in the kidney of T1DM mice compared to control group ([Figure 3A and B](#)), and it was significantly blunted with LG4 treatment. Similarly, LG4 treatment significantly prevented the increases of transcription levels of *Tnf- α* and *Il-6* in the kidney of T1DM mice ([Figure 3C and D](#)). Furthermore, the protein level of p-I κ B α was enhanced in vehicle-treated T1DM mice, which was obviously suppressed with LG4 administration. The changes of I κ B α and p-I κ B α indicated LG4 inhibited diabetes-induced activation of NF- κ B in diabetic mouse kidneys ([Figure 3E and F](#)).

LG4 Directly Binds to Proteins of MAPKs Pathways

To explore the underlying mechanism of the renoprotection of LG4 in DKD, HuProtTM human proteasome microarray was used to screen potential targets of LG4. This methodology has been used to finding the molecular target of natural product in our previous study.²⁴ Firstly, the biotin-labeled LG4 (LG4-biotin) ([Figure 4A](#)) was synthesized and structurally characterized as described in the [supplementary file \(Supplementary Figure S1–7\)](#). Next, the anti-inflammatory activity of LG4-biotin was confirmed in LPS-challenged mouse primary peritoneal macrophage. LG4-biotin and LG4 showed equal effect on suppressing LPS-induced IL-6 and TNF- α expression, indicating the biotin conjugation did not affect the bioactive structure of LG4 ([Supplementary Figure S8](#)). Then, the LG4-biotin and D-biotin were binding to the proteasome microarray, [Figure 4B](#). Compared to the D-biotin control, a total of 406 proteins were identified with SNR >2 ($p < 0.05$). Pathway enrichment of the differential binding proteins are shown in [Figure 4C](#). Among them, MAPKs were the top 2 signaling pathway that had direct binding activity to LG4. As shown in [Figure 4D](#), LG4 could bind to MAPK10 (JNK3), MAPK9 (JNK2),

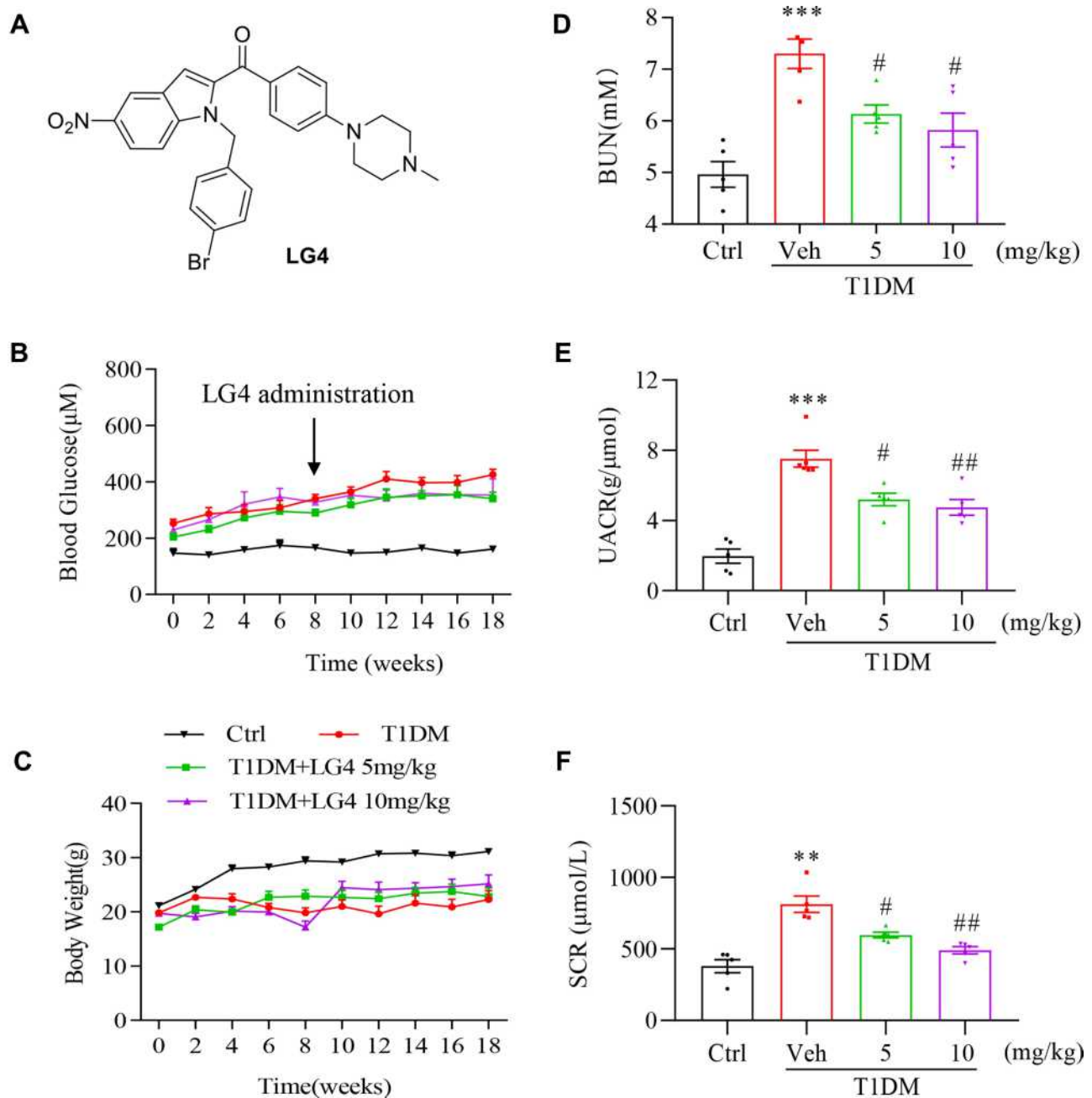


Figure 1 LG4 treatment improved renal function in T1DM mice. **(A)** Chemical structure of LG4. **(B and C)** Type 1 diabetes mellitus mice (T1DM) were orally administrated with vehicle (Veh) or LG4 (5, 10 mg/kg, every other day) from week 8 to week 18. Blood glucose and body weight were measured every other week. **(D–F)** Blood Urea Nitrogen (BUN), Urine Albumin Creatinine Ratio (UACR), and serum creatinine (SCR) of each group were determined at 18th week. ** $p < 0.01$; *** $p < 0.001$ vs Ctrl; # $p < 0.05$, ## $p < 0.01$ vs T1DM, $n = 5$.

MAPK8 (JNK1) and MAPK1 (ERK) that are key signal cascades of NF- κ B pathway in response to external stimuli and production of pro-inflammatory cytokines.^{25,26}

LG4 Inhibits JNK and ERK Signaling in vitro and in vivo

To determine the effect of LG4 on modulating MAPKs activation, high glucose (HG)-treated SV40MES13 cells

were pretreated with LG4 at 2.5 and 5 μM . HG significantly increased JNK and ERK phosphorylation, and these were blunted with LG4 pretreatment at 2.5 and 5 μM (Figure 5A and B). Similarly, phosphorylated JNK and ERK levels were significantly higher in the kidney of the vehicle-treated group than the control group, and LG4 treatment significantly suppressed this phosphorylation in the T1DM mice (Figure 5C and D).

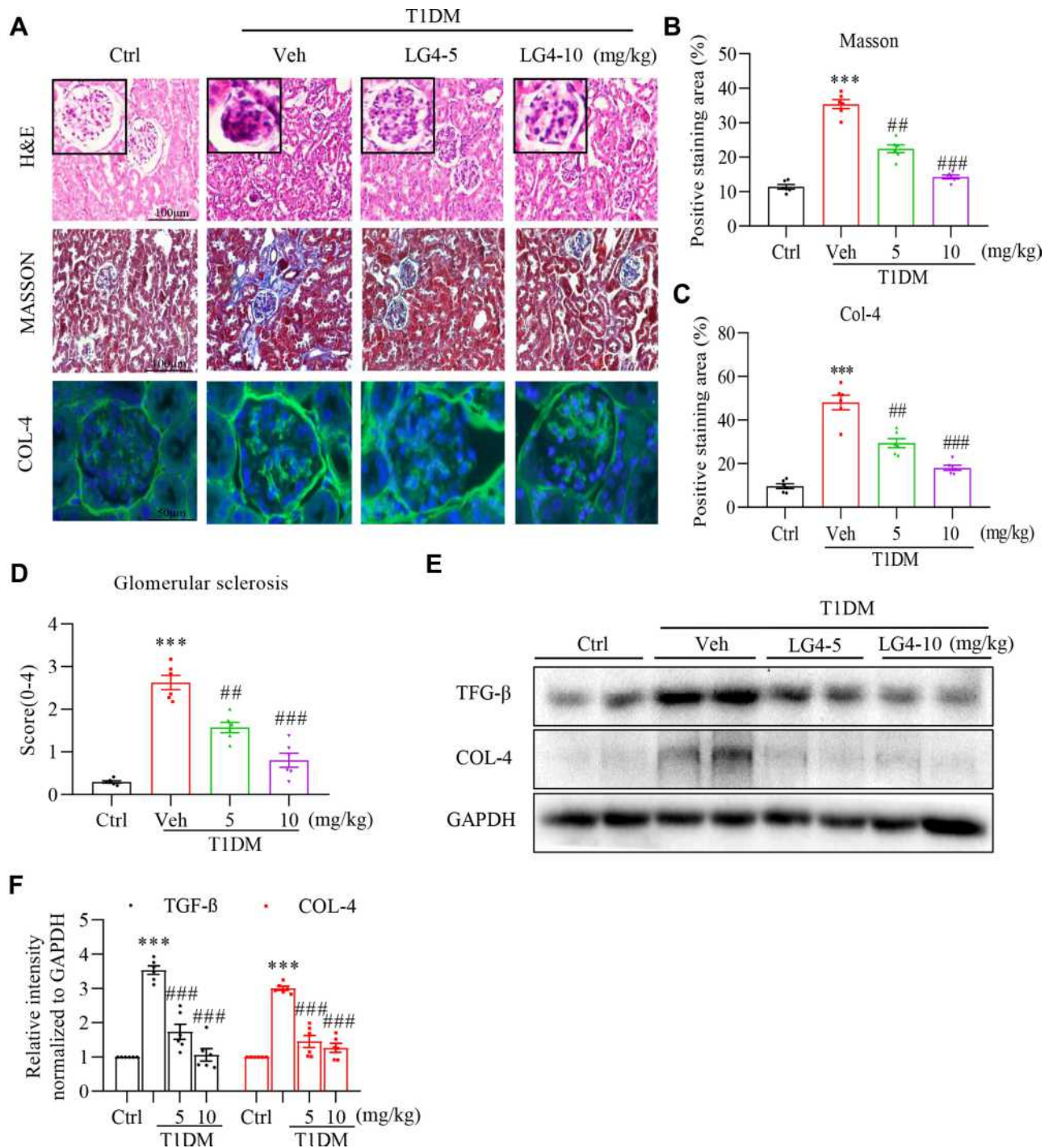


Figure 2 LG4 attenuates renal fibrosis in T1DM mice. (A) Kidney tissue sections were from the control, T1DM, and T1DM+LG4 mice. Representative images of H&E staining (200× magnification), Masson's staining (200× magnification), and anti-COL-4 immunofluorescence staining (green, 400× magnification) were shown. (B and C) Quantitative date of the Masson's (expressed as percentage of positive area) and COL-4 staining (expressed as relative fluorescence intensity). (D) Glomerular sclerosis score (0–4) was determined and plotted. (E and F) Western blot showed COL-4 and TGF-β levels in the kidney tissues, and relative protein levels to GAPDH were shown. ***p < 0.001 vs Ctrl; ##p < 0.01, ###p < 0.001 vs T1DM, n = 5.

LG4 Inhibits HG-Induced Inflammation via Targeting JNK and ERK Pathways

Next, we tested the role of LG4 in HG-induced inflammation in vitro. Firstly, HG induced IKK and IκBα

phosphorylation and IκBα degradation in SV40MES13 cells, while these changes were prevented by LG4 pretreatment at 2.5 and 5 μM (Figure 6A), indicating that LG4 inhibited HG-induced NF-κB activation. Accordingly,

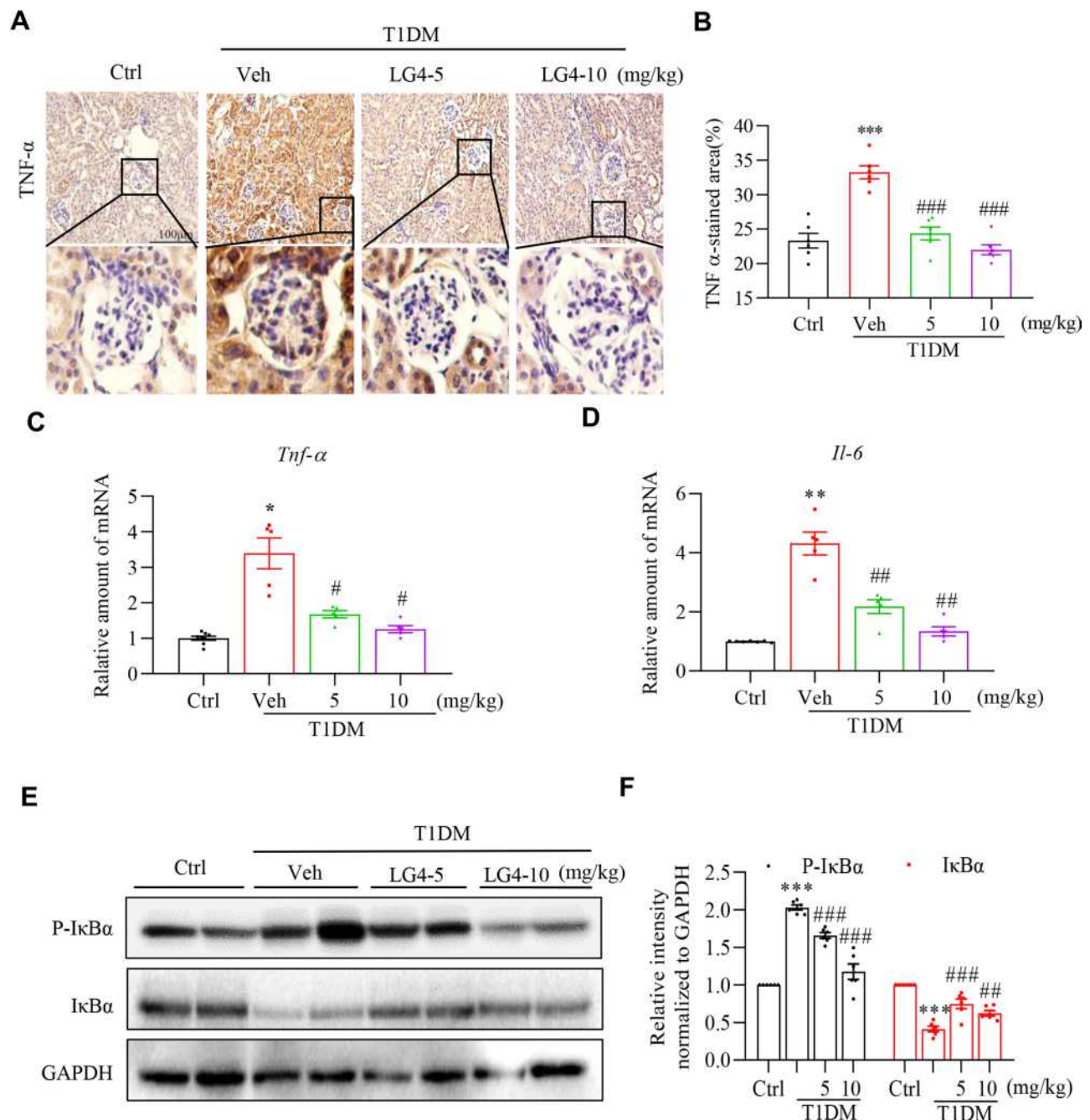


Figure 3 LG4 treatment inhibits renal inflammation in T1DM mice. (A and B) Representative images and quantitation of immunohistochemistry staining against TNF- α in the kidneys (200 \times). A typical view of glomerulus was selected and showed. (C and D) Real-time qPCR analyzed gene expression levels of *Tnf- α* and *Il-6* in the kidneys. (E and F) Western blot assay detected p-I κ B α and I κ B α level in the kidneys. Relative densitometric quantification to GAPDH was shown. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, T1DM vs control (Ctrl); # $p < 0.05$, ## $p < 0.01$, ### $p < 0.001$, T1DM + LG4 vs Veh, $n = 5$.

LG4 pretreatment significantly inhibited HG-upregulated *Il-6* and *Tnf- α* gene transcription in SV40MES13 cells (Figure 6B). As expected, JNK and ERK inhibition with SP600125 (10 μ M) and U0126 (10 μ M), respectively, significantly reduced HG-induced *Il-6* and *Tnf- α* expression in

SV40MES13 cells (Figure 6C and D), and LG4 had no further inhibitory effect on HG-induced *Il-6* and *Tnf- α* expression at the presence of SP600125 or U0126. These data validated that the anti-inflammatory effect of LG4 might be through inhibiting MAPKs signaling.

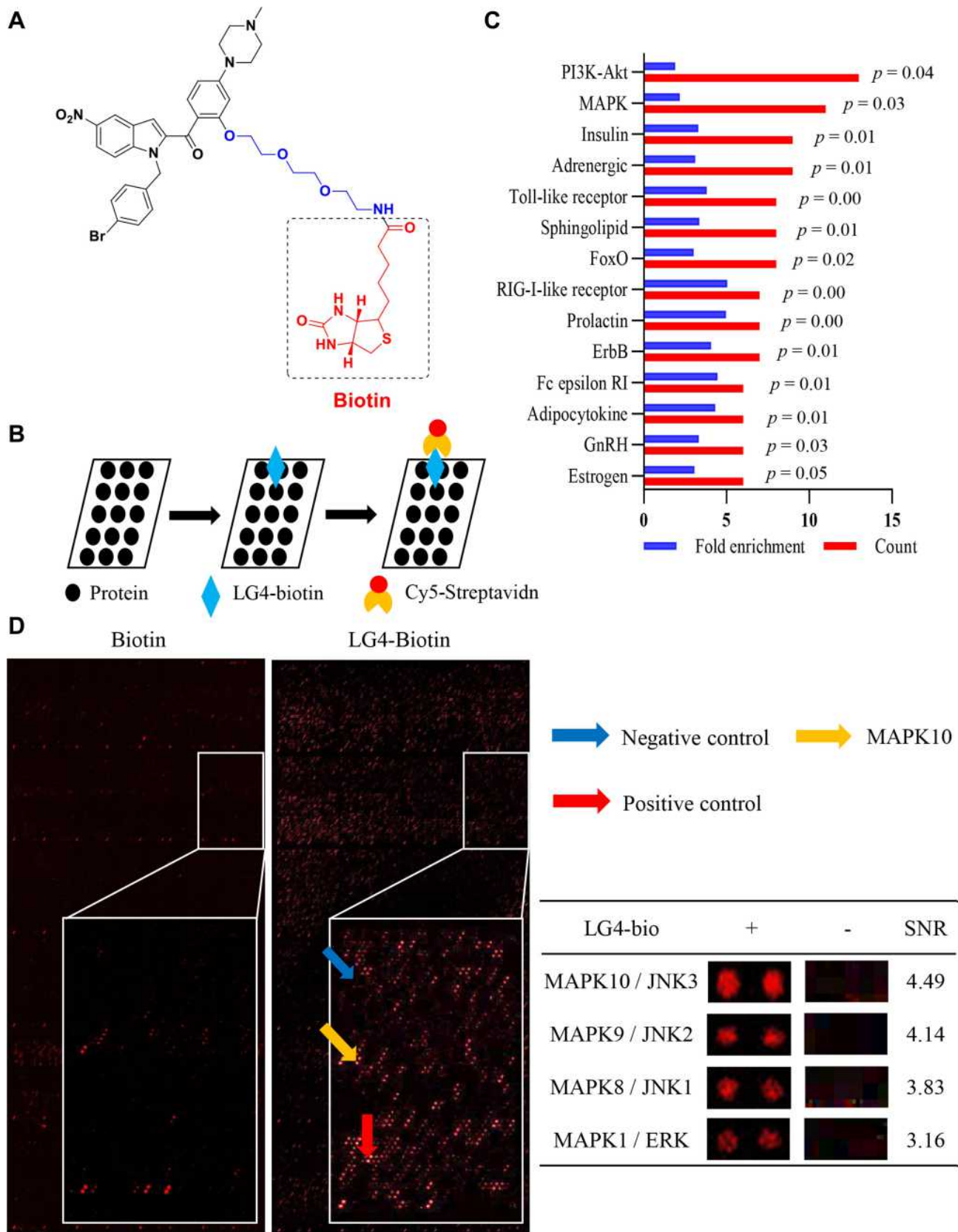


Figure 4 LG4 directly binding to proteins of MAPK pathways. **(A)** Chemical structure of biotin-labelled LG4. **(B)** Schematic showing the main steps for biotin or LG4-biotin using HuProt™ microarray. **(C)** Pathway enrichment of the differential binding proteins (fold enrichment ≥ 2 and p value < 0.05) were listed. **(D)** Representative image of the microarray showing positive (red arrow), negative (blue arrow) and MAPK10 (yellow arrow) spots. The table showed the magnified images of LG4-biotin binding to MAPKs spots and signal-to-noise ratios (SNR).

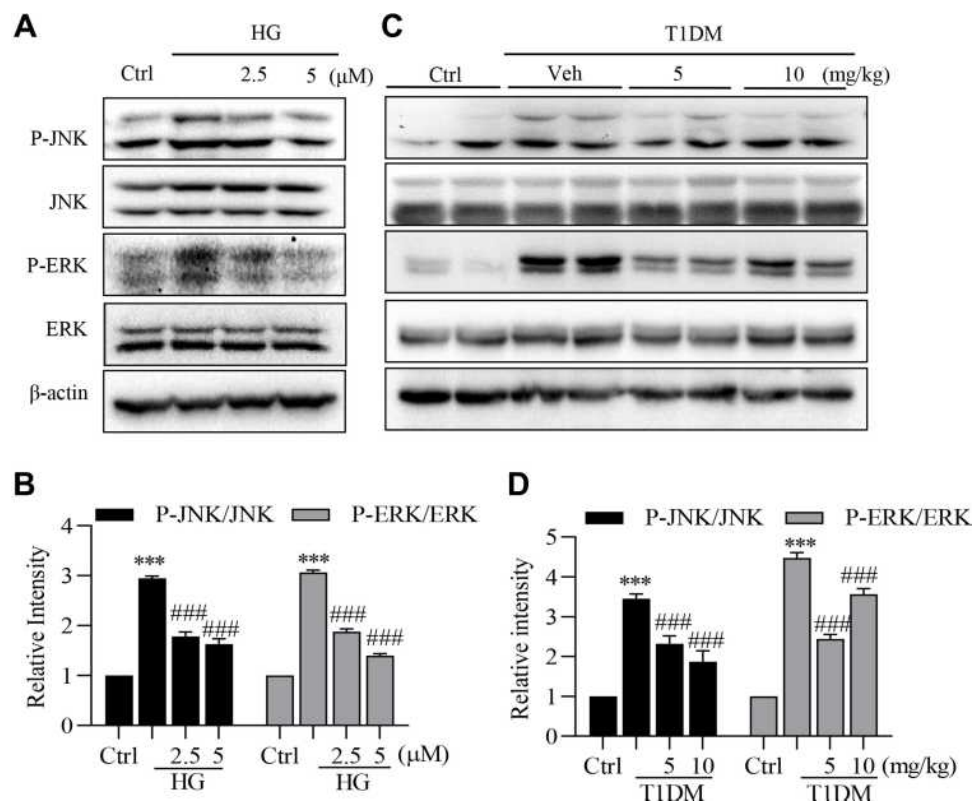


Figure 5 LG4 inhibits the activation of JNK and ERK in vivo and in vitro. **(A and B)** SV40MES13 cells were pretreated with LG4 (2.5 and 5 μ M) for 1 h and incubated with HG (33 mM) for 2 h. Cells were harvested and processed for phosphorylated JNK and ERK detection by Western blot. GAPDH, total JNK and ERK were used as loading controls. Corresponding densitometric quantifications were shown. **(C and D)** Western blots detected p-JNK and p-ERK levels in the kidneys. Corresponding densitometric quantifications were shown. *** $p < 0.001$ vs Ctrl; ### $p < 0.001$ vs HG or Veh.

Discussion

Inflammation plays a key role in the development and progression of DKD.^{27,28} Inflammatory responses are causal factors of glomerulosclerosis and tubulointerstitial fibrosis by inducing extracellular matrix deposition and differentiation/proliferation of myofibroblast. Therefore, blocking inflammation in kidney could be a strategy for the treatment of DKD. Hyperglycemia induces glomerular hyperfiltration, mechanical stress, and glycocalyx dysfunction that result in the overexpression of proinflammatory cytokines such as TNF- α , IL-6, and IL-1 β through activating MAPKs and NF- κ B signaling pathways in the kidney.^{12,13} From the prospective of molecular mechanism, MAPKs have been validated as upstream transducer of NF- κ B activation in hyperglycemia.¹⁴ As a key transcription factor mediating inflammatory responses, NF- κ B has been demonstrated to regulate the expression of a variety of inflammatory cytokines, chemokines, and profibrotic factors.^{29,30} Of note, NF- κ B, JAK/STAT, and TGF β /Smad signaling pathways have been demonstrated

involving in the fibrotic process.^{15,16} Therefore, MAPKs/NF- κ B axis could be the potential therapeutic target for DKD.

Indole is one of the important structural motifs in medicinal chemistry and has shown considerable pharmacological activities against inflammatory conditions, hyperlipidemia, and tumor growth.^{31–33} LG4, a novel indole compound developed in our laboratory, has been shown to exert anti-inflammatory effects both in vitro and in vivo.³⁴ In this study, we found that LG4 treatment remarkably reversed the function, histological abnormalities, fibrosis, and inflammation in the kidneys of T1DM mice without affecting their blood glucose and body weight. The effect of LG4 is highly related to its anti-inflammation property. However, a limitation of this study is the absence of in vivo pharmacokinetic data of LG4. The combination of pharmacokinetic (PK) and pharmacodynamic (PD) study will help to determine the potency of LG4 as a candidate for drug development. Our preliminary in vitro PK study demonstrated that LG4 was probably not

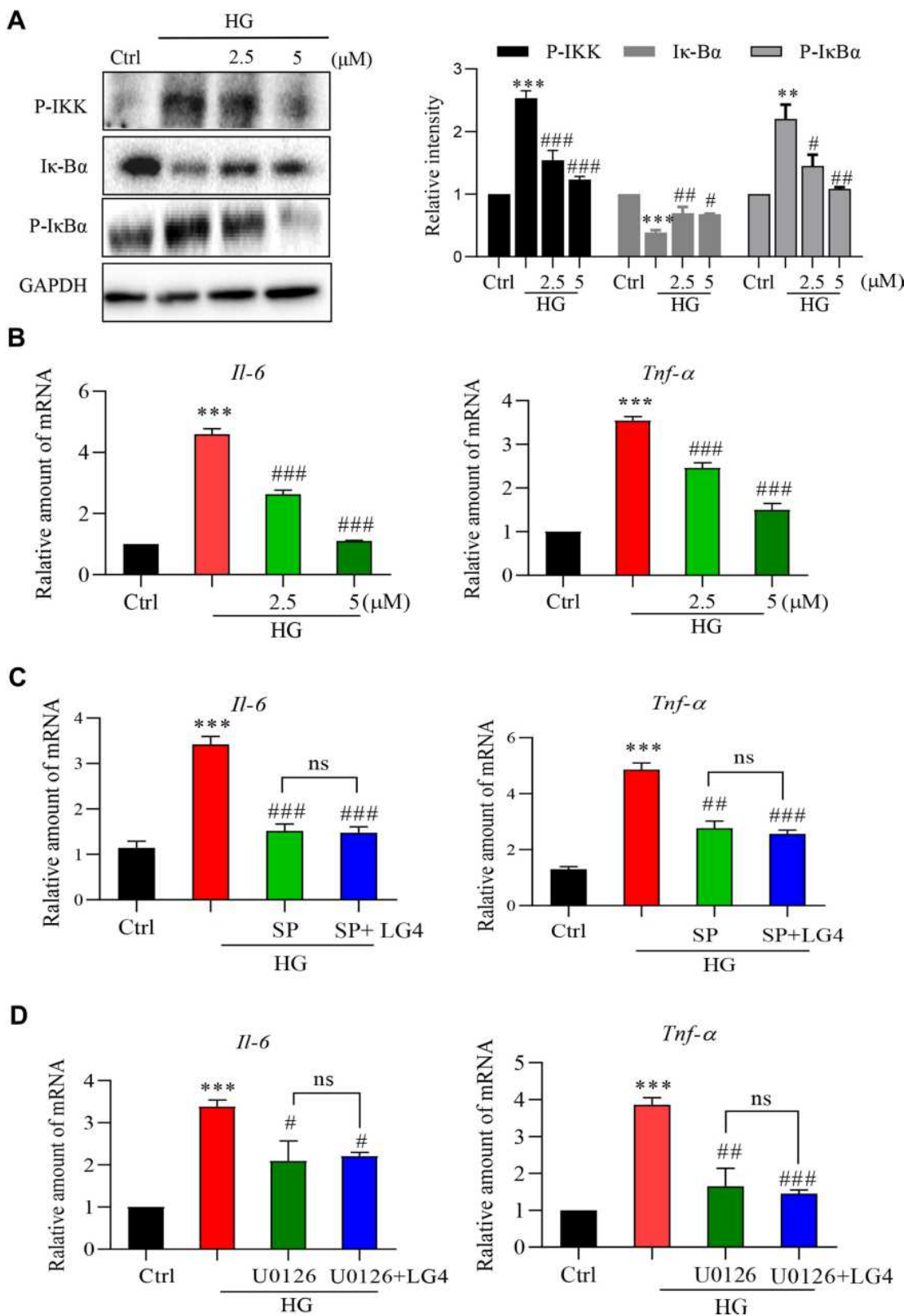


Figure 6 LG4 inhibits high glucose-induced inflammation via targeting JNK and ERK pathways. **(A)** SV40MES13 cells were pretreated with LG4 (2.5 and 5μM) for 1 h and incubated with HG (33 mM) for 2 h. Cells were harvested for p-IKK and Iκ-Bα analysis with Western blot. Corresponding densitometric quantifications were shown. **(B)** SV40MES13 cells were pretreated with LG4 (2.5 and 5 μM) for 1 h and incubated with HG (33 mM) for 8 h. Gene expression levels of *Tnfα* and *Il6* were analyzed with real-time qPCR. **(C and D)** SV40MES13 cells were pretreated with LG4 (5 μM) with/without SP600125 (10 μM) or U0126 (10 μM) for 1 h, followed by incubation with 33 mmol/L HG for 8 h. Gene expression levels of *Tnfα* and *Il6* were examined by RT-qPCR. Data were normalized by β-Actin level. ***p* < 0.01; ****p* < 0.001 vs Ctrl. #*p* < 0.05, ###*p* < 0.01, ####*p* < 0.001 vs HG, ns: no significant difference.

metabolized by cytochrome P450, indicating that the first pass elimination effect is weak ([Supplementary Figure S9](#)). Our data show that LG4 is a candidate compound with potential clinical utility in prevention of DKD. Further in vivo study is necessary for the pre-clinical study of LG4.

Based on the proteasome microarray assay, LG4 showed significant interactions with proteins of multiple signaling pathways including PI3K-AKT, MAPK, toll-like receptor, etc. There may be multiple signaling pathways mediating the therapeutic effect of LG4. We focused on MAPK pathways in this study, because MAPKs are functionally connected kinases that regulate key cellular process involved in kidney disease such as all survival, death, differentiation and proliferation, and MAPKs also have been proved to mediate inflammation in the kidney injury.³⁵⁻³⁷ In this study, phosphorylation of MAPKs (including JNK and ERK1/2) was increased in HG-challenged SV40MES13 cells and diabetic mouse kidneys. Importantly, LG4 could directly bind to MAPKs (JNK and ERK) and inhibit HG/diabetes-induced MAPKs activation. Because LG4 did not show further anti-inflammatory effect in HG-challenged mesangial cells with the presence of JNK and ERK inhibitors, LG4 might exhibit anti-inflammatory effect through inhibiting MAPKs and NF- κ B signaling pathways in kidney of T1DM mice.

However, a limitation of this study is that only a mouse model of STZ-induced type 1 DKD was used. DKD occurs in many patients with type 2 diabetes and a progression of DKD may be better reflected in type 2 diabetic mice.^{38,39} The renal protective effect of LG4 on DKD of type 2 diabetic mice needs to be further explored.

Conclusion

In conclusion, this study demonstrated a therapeutic role of LG4 in DKD through inhibiting renal inflammation and fibrosis in T1DM mice. Mechanistically, LG4 may exhibit anti-inflammatory and anti-fibrotic effects through directly binding to MAPKs and subsequent MAPK/NF- κ B signaling inhibition.

Abbreviations

BAX, BCL2-Associated X; BCL2, B-cell lymphoma-2; BSA, Bovine serum albumin; BUN, blood urea nitrogen; CMC-Na, carboxymethyl cellulose sodium; COL-4, collagen IV; DAPI, 4',6-Diamidino-2-Phenylindole, Dihydrochloride; DKD, diabetic kidney disease; DMEM, Dulbecco's Modified Eagle Medium; DMSO, dimethyl

sulfoxide; ERK, extracellular regulated protein kinases; ESRD, end stage of renal disease; FBS, fetal bovine serum; FITC, Fluorescein Isothiocyanate; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; H&E, Hematoxylin and Eosin; HG, high glucose; IKK, inhibitor of nuclear factor kappa-B kinase; IL-1 β , interleukin-1 β ; IL-6, interleukin-6; I κ B α , inhibitor of NF- κ B; JAK, Janus Kinase; JNK, c-Jun N-terminal kinase; MAPK, activating mitogen-activated protein kinases; NF- κ B, nuclear factor-kappa; NSAID, non-steroidal anti-inflammatory drug; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis; STAT, signal transducer and activator of transcription; STZ, streptozotocin; T1DM, type 1 diabetic mellitus; TGF- β , transforming growth factor- β ; TNF- α , tumor necrosis factor α ; UACR, urine albumin creatinine ratio.

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Disclosure

The authors declare that they have no conflicts of interest.

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