Paeoniflorin Inhibits Mesangial Cell Proliferation and Inflammatory Response in Rats With Mesangial Proliferative Glomerulonephritis Through PI3K/AKT/GSK-3β Pathway

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Mesangial proliferative glomerulonephritis (MPGN) is the most common type of chronic kidney disease in China, characterized by mesangial cell proliferation and inflammatory response. Paeoniflorin, an effective composition extracted from Radix Paeoniae Alba, has been used for various kinds of kidney diseases. However, there are no studies reporting the effects of paeoniflorin on MPGN. The present study aims to investigate whether paeoniflorin plays a role in MPGN and confirm the underlying molecular mechanisms. Our results manifested that paeoniflorin strongly restrained 24 h urinary protein and promoted renal function and dyslipidemia in a MPGN rat model. Moreover, paeoniflorin attenuated mesangial cell proliferation and inflammation both in MPGN rats and human mesangial cells (HMCs) treated with lipopolysaccharide (LPS). In detail, paeoniflorin decreased the number of mesangial cells and expressions of proliferation marker Ki67 in MPGN rats. Paeoniflorin also inhibited HMC proliferation and blocked cell cycle progression. In addition, the contents of inflammatory factors and the expressions of macrophage marker iNOS were decreased after paeoniflorin treatment. Furthermore, we found that the protective effect of paeoniflorin was accompanied by a strong inhibition of the phosphatidylinositol 3-kinase (PI3K)/AKT/glycogen synthase kinase (GSK)-3β pathway. Paeoniflorin enhanced the inhibitory effect of PI3K inhibitor LY294002 and suppressed the activated effect of PI3K agonist insulin-like growth factor 1 (IGF-1) on PI3K/AKT/GSK-3β pathway. In conclusion, these results demonstrated that paeoniflorin ameliorates MPGN by inhibiting mesangial cell proliferation and inflammatory response through the PI3K/AKT/GSK-3β pathway.

Keywords: mesangial proliferative glomerulonephritis, paeoniflorin, mesangial cells proliferation, inflammatory response, PI3K/AKT/GSK-3β pathway
INTRODUCTION

Mesangial proliferative glomerulonephritis (MPGN) is defined by pathological injury pattern with increased cell number and extracellular matrix in the mesangial area (Qin et al., 2013; Yang et al., 2018). Abnormally proliferative mesangial cells would release inflammation mediators, which often leads to interstitial fibrosis and glomerulosclerosis, resulting in irreversible progressive glomerulosclerosis, eventually turning into end-stage renal disease (ESRD) (Geng et al., 2016). Proliferation and inflammation of mesangial cells plays a crucial role in the progression of MPGN. However, the underlying mechanism of MPGN is still unknown, and there is no effective therapeutic drug to treat MPGN (Bai et al., 2017). Therefore, it is of vital importance to explore the pathogenesis of MPGN and search for effective drugs.

Phosphoinositide-3-kinase (PI3K) is a significant regulator of various signal transduction pathways, which mainly controls cell growth, apoptosis and metabolism (Zhang et al., 2014; Fruman et al., 2017). The serine/threonine kinase (AKT), a downstream kinase of PI3K, plays an important role in cell death and survival (Li et al., 2018). After receptor stimulation, AKT targets numbers of substrates for phosphorylation. One of the major effectors downstream of AKT is glycogen synthase kinase (GSK)-3β, by initiating phosphorylation at its serine 9 residue (Wang et al., 2014). Research proved that the activation of the PI3K/AKT/GSK-3β signaling pathway is vital for cell proliferation (Wang et al., 2012). Our previous study verified that the AKT/GSK-3β pathway is activated when mesangial cells proliferate, and it could be an efficient path to suppress abnormal proliferation and inflammation response through down-regulating the AKT/ GSK-3β pathway (Wu et al., 2018).

Paeoniflorin is an active ingredient isolated from of Radix Paeonieae Alba, the dried root of Paeonia lactiflora Pallas which has been used in traditional Chinese medicine for many years. Paeoniflorin exhibits numerous pharmacological effects, such as anti-inflammation, immunomodulation, antioxidation and anti-proliferation of cancer cells (Gu et al., 2016; Zhai et al., 2016; Song et al., 2017; Zhang et al., 2018). Recent research and our previous study found that paeoniflorin improved some experimental models of kidney disease, including diabetic nephropathy (Zhang et al., 2017), nephrotic syndrome (Lu et al., 2017), acute renal injury (Wang et al., 2016) and renal fibrosis (Zeng et al., 2013). More importantly, paeoniflorin ameliorated advanced glycation end product (AGE)-induced mesangial cell injury partly by anti-inflammation (Zhang et al., 2013; Chen et al., 2017). In recent research, paeoniflorin played a neuroprotective effect through inhibiting the calpain/AKT/GSK-3β pathway in SH-SY5Y cells (Ma et al., 2018). Meanwhile, some studies have shown that paeoniflorin promoted phosphorylation of AKT and GSK-3β in HepG2 cells and diabetic rats (Ma et al., 2017; Sun et al., 2017). Therefore, it is unclear whether paeoniflorin could protect MPGN, as well as the underlying mechanism between them. The present study aims to investigate the effects of paeoniflorin on MPGN in vivo and in vitro and explore the molecular mechanism related to the PI3K/AKT/GSK-3β pathway.

MATERIALS AND METHODS

Animal Treatment

Fifty male Sprague Dawley rats weighting 180-220 g were provided by Guangzhou University of Chinese Medicine Research Center for Experimental Animal (Certificate NO. SCXK 2013-0020). All the rats were housed in an air-conditioned room at 25 ± 2 °C and 65% humidity with a 12 h light/12 h dark cycle in specific pathogen-free conditions (Environment certificate NO. 2013-0085).

The MPGN rat model was established under an improvement of Hogendoorn’s method (Hogendoorn et al., 1990). Except for 8 rats in the normal group, the other rats underwent left nephrectomy after being anesthetized with 35 mg/kg pentobarbital sodium by intraperitoneal injection. One week later, all the model rats were subcutaneously injected with 3 mg bovine serum albumin (BSA, Sigma-Aldrich, St. Louis, MO, USA) emulsified in 0.1 ml Freund’s complete adjuvant (Sigma-Aldrich, St. Louis, MO, USA) twice in two weeks. At the end of the third week, the rats were intraperitoneally injected with 0.5, 1.0, 1.5 and 3 mg BSA every 1 hour respectively. Afterwards, every rat was injected with BSA in the tail vein from 0.5 mg to 3.0 mg every other day for 11 days with an increment of 0.5 mg; then the increment became 0.5 mg per week until the ninth week. In the interval time between tail vein injection, every rat was given BSA by intraperitoneal injection twice the dose of that with tail vein injection. At the end of the fifth week, every rat was injected with 100 μg lipopolysaccharide (LPS, Sigma-Aldrich, MO, USA) in the tail vein. At the same time, 24 h urinary protein of all rats was detected using Coomassie brilliant blue method (CoWin Biosciences, Beijing, China). Rats with higher urinary protein than the normal rats were randomly divided into four groups (each with 8 rats): model group, prednisone group (5 mg/kg, Guangdong Huanan Pharmaceutical, China), high dose group and low dose group of paeoniflorin (100 mg/kg, 50 mg/kg, Chengdu Herbpurify, China). The dose of paeoniflorin was decided according to our previous study (Lu et al., 2017). Prednisone was regarded as a positive drug, which is a common drug in clinic to treat kinds of chronic kidney diseases. Drugs were orally administered to rats daily from the sixth week to the tenth weeks.

All animal experimental programs were in accordance with the guidelines of the Animal Ethics Committee of Guangzhou University of Chinese Medicine submitting to the guidelines of the European Community and the National Institute of Health of the USA. All efforts were made to minimize suffering during animal experiments.

Measurement of 24 h Urinary Protein and Biochemical Parameters

Metabolic cages were used to collect 24 h urine samples weekly to detect urinary protein from the fifth week. During the urine collection period, rats were fasting and given free access to water. On the last day of animal experiment, animals were euthanized by intraperitoneal injection with pentobarbital sodium. Blood from abdominal aorta was collected and then centrifuged at
3000 rpm for 10 min at 4°C to acquire serum for biochemical analysis. The level of serum creatinine (SCr), blood urea nitrogen (BUN), total cholesterol (TC) and triglycerides (TG) were measured according to the manufacturer’s instructions (Nanjing Jiancheng Bioengineering Institute, China).

Histological Analysis
Kidneys of each rat were divided into two parts; one was fixed in 4% paraformaldehyde for histology staining and immunohistochemistry while the rest were frozen at -80°C for western blot analysis. Kidneys fixed for 48h were dehydrated in ethanol and embedded in paraffin. Renal tissue sections (5-μm thick) were cut and dewaxed by gradient series of ethanol, then stained with hematoxylin-eosin (HE) or periodic acid-Schiff (PAS). The mesangial matrix index (MMI) of PAS stained sections were measured by Image-Pro Plus 6.0 software, representing the ratio of the mesangial matrix area and glomerulus area. Twenty glomeruli were assessed in each rat kidney.

Immunohistochemistry
For immunohistochemistry, the dewaxed sections were boiled with sodium citrate to retrieve antigen and soaked in 3% hydrogen peroxide to quench endogenous peroxidase, then blocked with normal goat serum for 30 min at 37°C. After that, the sections were incubated with Ki67 (ab15580, Abcam, Cambridge, UK) at 1:500 dilution or iNOS (ab3525, Abcam, Cambridge, UK) at 1:500 dilution overnight at 4°C. After washing with PBS, the sections were incubated with secondary antibody (CoWin Biosciences, Beijing, China) for 30 min at room temperature. Subsequently, the sections were counterstained with diamobenzidine (DAB) and hematoxylin, respectively. All the images were captured with 40× objective under light microscope (TE2000, Nikon, Tokyo, Japan). The number of Ki67 positive cells in the glomerulus of 10 randomly selected nonoverlapping fields was evaluated (n = 4). The integral optical density (IOD) of iNOS of each group (n = 4) was measured by Image-Pro Plus 6.0 software.

CBA Analysis of Inflammatory Factors
Inflammatory factors in serum of every rat were detected as previously described (Wu et al., 2018). Briefly, the levels of IL-1α, IL-2, IL-10 and IFN-γ were measured with a cytometric bead array kit (Becton Dickinson, SanJose, USA) by FACS Canto II (Becton Dickinson, SanJose, USA) according to the manufacturer's instruction.

Cell Culture and Treatment
Human mesangial cells (HMCs) at the fifth passage were purchased from the Advanced Research Center of Central South University (Wuhan, China). HMCs were cultured in RPMI 1640 (Gibco, Grand Island, NY), supplemented with 10% fetal bovine serum (FBS, Biological Industries, Israel), as well as 100 U/ml penicillin and 100 mg/ml streptomycin (Gibco, Grand Island, NY) in a cell incubator with 5% CO₂ at 37°C.

HMCs were treated with 30 μg/ml LPS (Sigma-Aldrich, MO, USA) for 24 h with or without paeoniflorin at 5, 10, 20 and 40 μM for MTT, Edu, cell cycle, RT-PCR and western blot assays. The dose of paeoniflorin for the in vitro experiment was determined referring to Chen's study (Chen et al., 2017). For further western blot detection, HMCs exposed to 30 μg/ml LPS were treated with 20 μM LY294002 (Selleck Chemicals, Shanghai, China), 10 nM insulin-like growth factor 1 (IGF-1, PeproTech, NJ, USA) with or without 40 μM paeoniflorin, for 24 h. Every experiment in vitro was repeated at least three times.

Cell Viability Assay
HMCs (8×10⁴/well) were seeded in 96-well plates until they reached 70-80% confluence, then were treated as described above. After treatments, MTT solution (Biosharp, Hefei, China) at 0.5 mg/ml was added in each well, and the cells were then incubated at 37°C for 4 h. Afterwards, MTT-formazan crystals were solubilized by 150 μl DMSO at room temperature, then the absorbance was measured at 570 nm using a microplate reader (Thermo Fisher Scientific, USA) after thorough mixing by shaking. Cell viability experiment was expressed in “%”, that is the percentage of absorbance in respect to the control (100%).

Edu Assay
KeyFlour 488 Click EdU kit (KeyGEN, Nanjing, China) was used to detect cell proliferation. HMCs were cultured in 96-well plates and treated as described above. After being fixed with 4% paraformaldehyde for 15 min, cells were incubated with 2 mg/ml glycine for 5 min, then washed twice with 3% BSA/PBS (v/v) and permeabilized with 0.5% Triton-X 100 for 20 min at room temperature. Subsequently, HMCs were incubated with Edu solution at 10 μM for 30 min, then stained with DAPI (Sigma-Aldrich, MO, USA) at 5 μg/ml for 5 min. The images were captured with 20× objective under a fluorescence microscope (DMI3000B, Leica, Wetzlar, Hessen, Germany). The rate of Edu-positive cell (%) was assessed by Image-Pro Plus 6.0 software.

Cell Cycle Analysis
HMCs were cultured in 12-well plates (2×10⁵/well) and treated as described above. After 24 h treatment, cells were fixed in 70% ice-cold ethanol/PBS (v/v) overnight at 4°C. Afterwards, HMCs were centrifuged at 800 rpm for 5 min at 4°C and washed with ice-cold PBS twice. After that, HMCs were centrifuged again and resuspended with 500 μl Propidium iodide (PI) staining solution at 50 μg/ml (KeyGEN, Nanjing, China). Samples were then moved to 5-ml BD-Falcon tubes and stored on ice. Flow cytometry analysis was performed by FACS Calibur flow cytometer (Becton Dickinson, SanJose, USA). ModFIT 5.0, a cell cycle analysis software, was used to define the percentage of G0/ G1, S and G2/M phase.

RT-PCR Analysis
RNA of HMCs cultured in 6-well plates (4×10⁵/well) after the same treatments above was prepared using RNAiso Plus (Takara,
Beijing, China). Reverse transcription was performed using Prime Script RT Reagent Kit with gDNA Eraser (Takara, Beijing, China) according to the manufacturer's instruction.

RT-PCR was performed as previously described (Wu et al., 2016), using a detection system (Bio-Rad, Berkeley, CA, USA) with SYBR® Premix Ex TaqTM II qPCR Kit (Takara, Beijing China). The relative levels of IL-1α, IL-2, IL-10 and IFN-γ mRNAs were normalized to GAPDH and calculated according to the 2^(-ΔΔCt) formula. The sequences of the RT-PCR primers were designed by using the Primer Premier 5.0 software (Premier Biosoft, USA) and are shown in Table 1.

Western Blot Analysis
The proteins of renal cortex tissue and HMCs were extracted and quantified as described previously (Liu et al., 2018). Proteins were loaded and separated in 10% SDS-PAGE, then transferred to polyvinylidene difluoride membranes (Millipore, MA, USA) using a Trans-Blot (Bio-rad, Berkeley, CA, USA). Afterwards, the membranes were blocked with 5% non-fat milk powder (w/v) for 1 h at room temperature. The membranes were incubated with primary antibodies of PI3K (#4257), p-AKT (#4060), ACTG (#4691), p-GSK-3β (#5558), GSK-3β (#12456) and Cyclin D1(#2922), which were obtained from Cell Signaling Technology (Danvers, MA, USA) and GAPDH (Lot. IM001-0712, Excell Bio, Shanghai, China) at 4°C overnight. After being washed three times with TBST (Boster, Wuhan, China), the membranes were then incubated with goat anti-rabbit antibody conjugated with horseradish peroxidase at room temperature for 1 h. Finally, these membranes were washed again and detected with an ECL kit (Bio-rad, Berkeley, CA, USA) under a multi-functional imaging system (Tanon 5200, Shanghai, China). All the protein bands were analyzed with Image J 1.48 software.

Statistical Analysis
Data were presented as mean ± SD. Statistical analysis was performed with the SPSS 18.0 software (Chicago, IL, USA). Except for the results of the 24 h urinary protein, which was analyzed by two-way ANOVA, comparisons among groups were made by one-way ANOVA, followed by Duncan's test. P values less than 0.05 were recognized as significant. Protein bands for western blot were normalized using appropriate controls as mentioned in figure legends.

RESULTS

Paeoniflorin Ameliorates Renal Damage in MPGN Model Rats
To confirm the renoprotective effects of paeoniflorin in MPGN model rats, we detected 24 h urinary protein, the levels of Scr, BUN, TC, TG in serum, as well as the renal pathology changes in different groups. Our results showed that paeoniflorin decreased 24 h urinary protein since the seventh week till the end of the animal experiment (Figure 1A). Paeoniflorin also reduced the levels of Scr and BUN, exhibiting an improvement of renal function (Figures 1B, C). Besides, paeoniflorin lessened the contents of TC and TG, which manifested that paeoniflorin played a protective role in dyslipidemia (Figures 1D, E). Furthermore, paeoniflorin decreased the numbers of mesangial cells, improved glomerular structures (black arrows), reduced inflammatory infiltration (green arrows), tubule dilation (blue arrows) and protein cast (yellow arrows) to reverse renal injury (Figure 1F). Prednisone, as the positive drug in the animal study, showed protective effects on MPGN, which was similar to paeoniflorin. These results revealed that paeoniflorin ameliorated renal damage in MPGN model rats through protecting against renal function, dyslipidemia and pathological changes.

Paeoniflorin Inhibits Mesangial Cells Proliferation and Mesangial Matrix Expansion in MPGN Model Rats
As a molecular marker of proliferation, the Ki67-positive cells in the mesangial area were regarded as mesangial cell proliferation (Lu et al., 2016). The immunohistochemical results showed that Ki67 was obviously exhibited in the glomerulus of model rats, while the Ki67 positive cells numbers were reduced markedly in the normal and treatment groups (P < 0.01) (Figures 2A, C). Except for the increased number of mesangial cells, mesangial matrix expansion is one of the characteristics of MPGN. As shown in Figures 2B, D, evident mesangial matrix expansion was visible in the model group, as the mesangial matrix index (MMI) of the model group was significantly elevated in comparison to the normal group (P < 0.01). Fortunately, the increase of MMI was suppressed by paeoniflorin compared with the model group (P < 0.01). Though prednisone distinctly decreased the numbers of Ki67-positive cells in the glomerulus, there was no statistical difference of MMI between the model group and prednisone group. These results demonstrated that paeoniflorin inhibited mesangial cell proliferation and mesangial matrix expansion in MPGN model rats.

Paeoniflorin Prevents Inflammatory Response in MPGN Model Rats
MPGN, as a typical primary glomerulonephritis, is also accompanied by an increase of inflammatory response. Therefore, immunohistochemistry was performed to detect the expression of iNOS in renal tissues of different groups. As shown in Figures 3A, B, both paeoniflorin and prednisone

TABLE 1 | Nucleotide sequences of the primers used for qRT-PCR.

| Genes | Sense primers (5′ to 3′) | Anti-sense primers (5′ to 3′) |
|-------|--------------------------|-----------------------------|
| IL-1α | GAGGTGTATGCGATCAACTCTG  | CTTCCCTCTGAGTCTTTGG    |
| IL-2  | TCCCAACTCCACCGAGTGC    | GTGCTGATTAAGTCCCTGGG    |
| IL-10 | GCAAGCGCTTCTGGAAGATGA  | GCTTCACGGGCCCTGGTCTTG  |
| IFN-γ | CTCGCCTGAAGTGCTGACC    | ACTGGGATGCTCTGACCTC    |
| GAPDH | CAACTGGTCTGAGCTGGACCTG  | GTGCTGCTCTTGAAGTCAAGAG  |
reduced the expression of iNOS in renal tubules \((P < 0.01)\). We also carried out a cytometric bead array to measure the levels of IL-1\(\alpha\), IL-2, IL-10 and IFN-\(\gamma\) in serum. Results showed that the treatment of paeoniflorin and prednisone significantly decreased the contents of the above inflammatory cytokines (Figures 3C–F). Hence, we can summarize that paeoniflorin presented a good effect on anti-inflammation in MPGN model rats.

**Paeoniflorin Mediates PI3K/AKT/GSK-3\(\beta\) Signaling Pathway in MPGN Model Rats**

Recent research reported that enhancing the PI3K/AKT pathway would facilitate mesangial cell proliferation and MPGN (Gao and Han, 2017). Our previous study found that the AKT/GSK-3\(\beta\) pathway is a crucial mechanism of proliferation and inflammatory response in mesangial cells (Wu et al., 2018). In the present study, western blot results showed that the expressions of PI3K, p-AKT and p-GSK-3\(\beta\) are all up-regulated in the model group, which could lead to speculation that the PI3K/AKT/GSK-3\(\beta\) pathway is involved in MPGN pathogenesis (Figure 4). On the contrary, paeoniflorin and prednisone notably down-regulated the levels of PI3K, p-AKT and p-GSK-3\(\beta\) \((P < 0.01)\). These results revealed that the therapeutic action of paeoniflorin in MPGN by ameliorating mesangial cell proliferation and inflammatory response is probably through regulating the PI3K/AKT/GSK-3\(\beta\) pathway.
Both of our previous and recent studies proved that LPS induced HMC proliferation (Sun et al., 2018; Wu et al., 2018). To confirm the role of paeoniflorin in protecting mesangial cells in vitro, HMCs were incubated with 30 μg/ml LPS and different dose of paeoniflorin for 24 h. As shown in Figure 5A, MTT assay results showed that LPS significantly promoted cell vitality of HMCs (P < 0.05), while paeoniflorin (from 5μM to 40μM) inhibited the growth of HMCs (P < 0.01). An EdU incorporation assay was used to detect DNA synthesis, which is regarded as an indicator of cell proliferation. The rate of EdU-positive cells was increased in the LPS group, while paeoniflorin obviously reduced EdU incorporation into HMCs (Figures 5B, C). Since EdU is a cell proliferation marker that incorporates into S-phase proliferating cells, these data suggest that paeoniflorin may suppress HMC cell proliferation by affecting the cell cycle.

To further evaluate the effect of paeoniflorin on HMC proliferation, we next performed flow cytometry for cell cycle analysis. As shown in Figures 6A–D, LPS reduced the percentage of cells in the G0/G1 phase but increased that in the S and G2/M phases, showing that LPS could promote cell cycle progression. Conversely, paeoniflorin at a concentration of 40μM increased the proportion of cells in G1 and decreased that in the S and G2/M phases. These results showed that paeoniflorin could block LPS-induced cell cycle progression by inhibiting G1-S phase transition and arresting cells in G0/G1.

To confirm whether paeoniflorin arrested HMCs on the G1 phase, we also detected the expression of Cyclin D1 by western blot. Cyclin D1 is a key gene in cell cycle regulation which controls the G1-S transition during the cell cycle. A high activity of Cyclin D1 would cause premature cell passage through the G1 checkpoint, which leads to accumulation of DNA damage and finally abnormal cell proliferation (Verim et al., 2013). Our results showed that LPS induced the expression of Cyclin D1, while paeoniflorin decreased it (Figure 6E). The results indicated that LPS promoted HMC proliferation, and paeoniflorin inhibited G1-S phase transition and arrested cells in the G0/G1 phase.

### Paeoniflorin Inhibits HMC Proliferation and Affects the Cell Cycle

Paeoniflorin Reduces the mRNA Expressions of Inflammatory Factors in HMCs

LPS also induced inflammatory response in HMCs (Tsugawa et al., 2017), which is in accordance with the rat model of MPGN. To clarify the anti-inflammatory effects of paeoniflorin on HMCs, the mRNA expressions of IL-1α, IL-2, IL-10 and IFN-γ...
were determined by RT-PCR. As shown in Figure 7, paeoniflorin at concentrations of 20 and 40 μM significantly reduced the up-regulation of the examined cytokines induced by LPS ($P < 0.05$ or $P < 0.01$). These results demonstrated that paeoniflorin could inhibit the LPS-mediated inflammatory response of HMCs.

**Paeoniflorin Mediates PI3K/AKT/GSK-3β Signaling Pathway in HMCs**

To confirm the molecular mechanisms underlying the anti-proliferative and anti-inflammatory effects of paeoniflorin against HMCs induced by LPS, the protein expressions of...
Paeoniflorin Ameliorates MPGN Through PI3K/AKT/GSK-3β Pathway

As shown in Figure 8, the protein expressions of PI3K, p-AKT and p-GSK-3β were increased in the LPS group (P < 0.01). Luckily, paeoniflorin significantly suppressed the expressions of PI3K, p-AKT and p-GSK-3β (P < 0.01), showing that paeoniflorin possibly prevented HMC injury through regulating the PI3K/AKT/GSK-3β pathway.

PI3K is a crucial molecule in regulating the function of mesangial cells, targeting the activation of AKT/GSK-3β signaling in kidneys. To further investigate whether the protective effects of paeoniflorin were mediated by the PI3K/AKT/GSK-3β pathway, we detected the effects of paeoniflorin under the intervention of PI3K inhibitor LY294002, or PI3K agonist IGF-1 in HMCs. As shown in Figure 9, the protein levels of PI3K and its downstream proteins p-AKT and p-GSK-3β were inhibited by LY294002 and increased by IGF-1, compared to the LPS group (P < 0.01 or P < 0.05). Paeoniflorin further suppressed the phosphorylation of AKT and GSK-3β when combined application with LY294002. Remarkably, paeoniflorin reversed the suppression of the PI3K/AKT/GSK-3β signaling pathway induced by IGF-1, which demonstrated the definite inhibition of paeoniflorin on the PI3K.

DISCUSSION

Paeoniflorin is one of the active compounds in *Paenia radix* Alba, which is an important herb used in traditional Chinese medicine to treat chronic kidney disease or other chronic inflammatory diseases (Li et al., 2017). Our previous study found that paeoniflorin ameliorated nephritic syndrome model rats through inhibiting inflammation and podocyte apoptosis (Lu et al., 2017). Some studies reported that paeoniflorin improved oxidative damage and inflammation in mesangial cells (Chen et al., 2017; Zhang et al., 2013). Mesangial cells are the most active intrinsic cells in the glomerulus, while proliferative reaction is their most common response to several stimulations, such as immune complexes, hypoxia and macromolecular substances. MPGN is characterized by mesangial cell proliferation and mesangial matrix expansion. Therefore, suppression of mesangial cell proliferation would be the most important way to treat MPGN. However, whether paeoniflorin could be a potential drug in MPGN still unclear.

In the present study, we observed the protective effects of paeoniflorin on MPGN and explored the underlying mechanism. For this purpose, a MPGN rat model established by BSA and LPS was used, which was also known as chronic serum sickness (CSS), characterized by glomerular immune complex deposition,
hypercellularity and matrix expansion (Bao et al., 2007). Our data demonstrated that paeoniflorin lessened the number of mesangial cells, inhibited mesangial matrix expansion and reduced the expressions of Ki67 in MPGN. We also carried out a HMCs proliferation model induced by LPS to evaluate the effects of paeoniflorin in vitro. Results showed that paeoniflorin suppressed cell viability and reduced proliferative cell staining with Edu. In addition, paeoniflorin blocked cell cycle progression in HMCs induced by LPS and down-regulated the expression of Cyclin D1. These results suggested that paeoniflorin has potent effects against mesangial cells proliferation in vivo and in vitro.

The activated and proliferative mesangial cells in the glomerulus could produce kinds of vasoactive substances and cytokines including interleukin-1 (IL-1), interferon (IFN) and chemotactic factor (CXC), thereby promoting ongoing mesangial cell proliferation (Gao et al., 2017). In fact, inflammation plays a key role in the development of MPGN, including anti-Thy1 nephritis and IgA nephropathy animal models (Hua et al., 2013; Aizawa et al., 2014). Our results showed that the levels of inflammatory factors, IL-1α, IL-2, IL-10 and IFN-γ were augmented in MPGN model rats, as well as in HMCs treated with LPS. Moreover, the expressions of macrophage marker iNOS were increased in tubules in MPGN model rats. Paeoniflorin reversed the inflammatory response by lessening the levels of inflammatory factors and the expressions of iNOS. In consequence, our results demonstrated that paeoniflorin inhibited inflammatory response in MPGN.

Among the three classes of PI3Ks, class IA PI3Ks have been studied most extensively. Class IA PI3Ks are heterodimeric enzymes composed of a regulatory p85 subunit and a p110 catalytic subunit (Chiu et al., 2014). An increase of p85 could stabilize p110 proteins which would augment p85–p110 heterodimers, thereby enhancing PI3K activity (Kok et al., 2009). Moreover, the increased levels of p85 upon cellular stimulation correlates with sustained AKT phosphorylation in cells (Esposito et al., 2008). Recent reports showed that the PI3K/AKT/GSK-3β pathway has been reported involved in cell proliferation (Chan et al., 2018), apoptosis (Xie et al., 2017), and inflammation (Jing et al., 2017).
in various kidney diseases. Currently accumulating evidence supports that the PI3K/AKT signaling pathway contributes to the promotion of proliferation in mesangial cells (Feng et al., 2016; Ying et al., 2017). Additionally, one study reported that phosphorylation of AKT is involved in the initiation and development of mesangial proliferation in anti-Thy1 MPGN model rats (Qiu et al., 2012). GSK-3β, a major downstream signaling molecule of AKT participating in various cellular processes, plays a crucial role in inflammation and proliferation (Hong et al., 2017; Xie et al., 2018). More importantly, GSK-3 inhibitors may emerge as effective drugs for proliferative kidney diseases (Soos et al., 2006). Recent research reported that paeoniflorin inhibits cell proliferation through suppression of PI3K/AKT signaling (Wu et al., 2013; Zheng et al., 2015). In the present study, the PI3K/AKT/
GSK-3β signaling pathway was activated in the MPGN model rats and HMCs induced by LPS, as the expressions of PI3K, p-AKT, p-GSK-3β were markedly increased. After paeoniflorin treatment, the expressions of related proteins were decreased. These results illustrated that paeoniflorin improves MPGN mostly through regulating the PI3K/AKT/GSK-3β pathway.

Similarly, the effects of paeoniflorin on reducing inflammatory cytokines (IL-1α, IL-2, IL-10, IFNγ) may be related to the inhibition of the PI3K/AKT/GSK-3β signaling pathway. It was reported that the PI3K inhibitor LY294002 reduced the level of IL-2 in T lymphocytes through inhibiting the PI3K/AKT/GSK-3β pathway (Braunstein et al., 2008). Methane-rich saline induced the contents of IL-10 and IFN-γ through activating the PI3K/AKT/GSK-3β pathway (Yao et al., 2017). However, the molecular mechanism underlying regulating of anti-inflammatory cytokines of the PI3K pathway in our study is poorly understood. Research illustrated that GSK-3β can activate the transcriptional activity of NF-κB by phosphorylation of its p65 subunit (Chen et al., 2018; Yang et al., 2019). Inhibition of GSK-3β affects macrophage polarization and inhibits NF-κB activation, and that results in potent anti-inflammatory effects in several animal models (Liu et al., 2016; Suber et al., 2017; Tantray et al., 2017; Wang et al., 2018). Meanwhile, GSK-3β gene deletion prevents cell differentiation induced by NF-κB (Hoeflich et al., 2000). GSK-3β inhibitor lithium chloride and SB415286 leads to the reduction of pro-inflammatory molecules by decreasing NF-κB activity (Wang et al., 2013). Furthermore, research showed that LPS could induce cell proliferation and the contents of inflammatory cytokines of HMCs through activation of NF-κB nuclear translocation (Jiang et al., 2011). Therefore, it is reasonable to presume that the anti-inflammatory effects of paeoniflorin on mesangial cells and MPGN are related to the NF-κB pathway by suppressing GSK-3β.

FIGURE 9 | The effects of LY204002 and IGF-1 on HMC proliferation in response to paeoniflorin. HMCs exposed to LPS (30 μg/ml) were treated with LY294002 (20 μM), IGF (10 nM) with or without paeoniflorin (40 μM) for 24 h. (A) The protein expressions of PI3K, p-AKT, AKT, p-GSK-3β and GSK-3β were analyzed using western blot. (B–D) The statistical data of all the proteins were analyzed with Image J 1.48 software. The obtained values of PI3K were normalized to the housekeeping gene GAPDH, AKT for p-AKT, and GSK-3β for p-GSK-3β especially. Data were expressed as mean ± SD, n = 3. *P < 0.05 and **P < 0.01 versus control group, #P < 0.05 and ##P < 0.01 versus LPS group.
LY294002 is a kind of PI3K inhibitor that can inhibit the activity of PI3K and AKT and inactivate GSK-3β (Song et al., 2018). Insulin-like growth factor-1 (IGF-1) is known to bind to the IGF-1 receptor as well as the PI3K agonist (Li et al., 2015). To confirm the regulation of paeoniflorin on the PI3K/AKT/GSK-3β pathway, LY294002 and IGF-1 were administered alone or combined with paeoniflorin in HMCs. Fortunately, we found that paeoniflorin enhanced the inhibitory effects on the activation of the PI3K/AKT/GSK-3β pathway by LY294002. On the other hand, paeoniflorin reversed the effects of IGF-1. These results manifested that paeoniflorin protects HMC injury mostly through down-regulating PI3K.

In conclusion, paeoniflorin displayed a positive therapeutic effect on MPGN in vivo and in vitro by inhibiting mesangial cells proliferation and inflammation response. More importantly, we first demonstrated the potential mechanisms of effective prevention of paeoniflorin on MPGN, namely by down-regulating the PI3K/AKT/GSK-3β signaling pathway. Indeed, paeoniflorin may be a novel therapeutic agent for MPGN.

ETHICS STATEMENT

This study was carried out in accordance with the recommendations of Animal Ethics Committee of Guangzhou University of Chinese Medicine. The protocol was approved by Animal Ethics Committee of Guangzhou University of Chinese Medicine.

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AUTHOR CONTRIBUTIONS

JZ designed the experiments. BL, JL, LB and RL performed in vivo research. PZ, DC and HL performed in vitro research. JS, XL and YW contributed to the preparation of reagents, materials and experimental equipment. CL and JW analyzed the data. BL and YZ wrote the manuscript. All authors read and approved the final manuscript.

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**Conflict of Interest Statement:** 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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