Abstract

Purpose: To explore the protective effect of glycyrrhizin in rats with nephrotic syndrome (NS) induced by adriamycin (ADR).

Methods: 36 Sprague Dawley (SD) male rats were divided into control, untreated and glycyrrhizin treatment groups. The NS rat model was established by injecting ADR twice in the untreated and in the glycyrrhizin treatment groups. Rats in the glycyrrhizin treatment group were fed glycyrrhizin by intragastric administration for 7 days. Changes in the following indices were observed in the three groups before and 4 weeks after the treatment: 24h urine protein quantitation (UPr), serum cholesterol (Ch), serum albumin (Alb), blood urea nitrogen (BUN), serum creatinine (sCr), laminin (LN), fibronectin (FN), collagen (Col), transforming growth factor β1 (TGFβ1) and connective tissue growth factor (CTGF); histopathology by light and electron microscope. Expression of LN, FN, Col, TGFβ1 and CTGF in the cortex of the kidney were detected by semi-quantitive immunohistochemical analysis. Expression of TGFβ1 and CTGF in the cortex of the kidney was detected by Fluorescein Based Quantitive RT-PCR. Macrophage infiltration was evaluated by the immunoperoxidase staining.

Results: Compared with the control group, 24h UPr, Ch, BUN and sCr of rats in the untreated group were increased. Glycyrrhizin reduced 24h Upr, Ch, BUN, sCr, LN, FN, Col, TGFβ1, CTGF, and mean arterial blood pressure. Pathological changes in the kidney, the expression of LN, FN, Col, TGFβ1 and CTGF in the cortex of the kidney in the glycyrrhizin treatment group were decreased compared with the untreated group. Glycyrrhizin also suppressed macrophage infiltration in the kidneys of NS rat models.

Conclusion: Glycyrrhizin exerts protective effects in rats with NS, reducing the excretion of Upr, Ch, BUN, sCr, and mean arterial blood pressure, and also decreasing expression of LN, FN, Col, TGFβ1 and CTGF in the kidney. Renal function is improved and the severity of NS is lessened.

Nephrotic syndromes (NS) may develop in many mammalian species as a result of primary diseases such as minimal change disease in humans, inflammatory diseases such as membranous nephropathy or immune glomerulonephritis in humans or laboratory animals, or exposure to toxic substances such as adriamycin (ADR) in rats. Besides proteinuria and
podocyte foot process effacement, the development of edema and ascites are permanent clinical symptoms in the majority of sufferers.\(^2\) The pathogenesis of NS is unclear. Some reports have indicated that NS can lead to glomerular sclerosis and renal failure. Mesangial cell proliferation and extracellular matrix (ECM) accumulation are the main pathological features of NS. Transforming growth factor \(\beta 1\) (TGF\(\beta 1\)) and connective tissue growth factor (CTGF) participate in the process of glomerular disease and ECM accumulation, which may result in glomerular sclerosis.\(^3\) Inhibition of cytokines related to glomerular sclerosis and ECM accumulation is the key factor in preventing glomerular sclerosis.

For thousands of years, many herbs have been used as natural remedies for the prevention and/or treatment of renal diseases. Various herbs and herbal products are believed to have kidney protective functions and are widely used in clinical practice in the West as well as in the East. Glycyrrhizin, the major active component extracted from licorice (Glycyrrhiza glabra) roots, is one of the oldest and most commonly prescribed herbs in Eastern traditional medicine, and has been used to treat tuberculosis, peptic ulcers, and liver injury in a number of clinical disorders in human patients.\(^4,5\) Glycyrrhizin may scavenge reactive oxygen species (ROS) and has an anti-inflammatory action.\(^6,7\) Recent publications have reported that glycyrrhizin could down-regulate expression of transcription factor \(\kappa B\), CTGF and TGF\(\beta 1\) in rats with obstructive nephropathy, and inhibit the occurrence of renal interstitial fibrosis.\(^8\)

ADR causes NS in rats that corresponds to minimal change nephropathy or focal segmental glomerulosclerosis.\(^9-10\)

In the present study, we produced rat models of NS by intravenous injection of ADR. Rats in the glycyrrhizin treatment group were fed glycyrrhizin for 7 days. The following indices were observed before and 4 weeks after the treatment: urine protein quantitation (UPr) of 24h, serum cholesterol (Ch), serum albumin (Alb), blood urea nitrogen (BUN), serum creatinine (sCr), laminin (LN), fibronectin (FN), collagen IV(CollIV), Transforming Growth Factor \(\beta 1\) (TGF\(\beta 1\)), Connective Tissue Growth Factor (CTGF) and the mean arterial blood pressure. Histopathological changes were detected by light and electron microscopy on the cortex of the kidney tissues. Expression of LN, FN and Coll in the renal cortex was detected by immunohistochemistry. Expression of TGF\(\beta 1\) and CTGF in the kidney tissues was detected by fluorimetry based on quantitative RT-PCR. Finally, the changes of these markers were compared among the three groups to determine the protective effect of glycyrrhizin in rats with NS.

Materials and Methods

Experimental animals

The study group consisted of 36 adult male Sprague-Dawley (SD) rats weighing 240-260g, which were obtained from the Department of Laboratory Animal Science, Guangzhou Medical School, Guangdong province of P.R.China. All animals were treated humanely and the Medical Laboratory Animal Management Committee of Guangzhou Medical School approved all animal procedures. The rats were housed in a temperature controlled room (24± 1°C) on a 12 hour light and dark cycle, with free access to food and water.

Materials

Glycyrrhizin (Stronger Neo Minophagen C, #H20030184) was purchased from Jian’an Medicine Ltd of Shenzhen, Guangdong province, P.R.China. Adriamycin (doxorubicin HCL, #0408E1, ADR) was supplied by Wanle Medicine Ltd of Shenzhen, Guangdong province, P.R.China. Rabbit anti-LN, -FN, -Col polyclonal antibodies and Streptavidin-biotin peroxidase complex immunohistochemical assay kit (SABC) were obtained from Wuhan BOSTER Ltd., Hubei province, P.R.China. Metabolic cages
(#3700M071) were bought from Tecniplast Gazzade S.ar.l Ltd.

**Preparation of animal models and drug treatment**

36 SD rats were randomly divided into three groups: control, untreated and glycyrrhizin treatment groups, with 12 rats in each group. NS was induced by a single dose of ADR (5mg/kg) injected into the tail vein of conscious rats in the untreated model and the glycyrrhizin treatment groups twice, once every 7 days. After modeling, rats in the glycyrrhizin treatment group were fed with glycyrrhizin for 7 days (200mg/kg glycyrrhizin by intragastric administration), whereas rats in the control and the untreated groups were fed normal saline solution for 7 days. Twelve rats of each group were chosen for biochemical measurement, while 10 rats were selected for the morphological evaluation. The survival rate of the glycyrrhizin treatment group was 93.86%.

**Quantitative detection of 24h UPr and blood biochemical detection**

24h UPr (urinary protein in mg/ml glomerular filtrate in 24h) was determined by spectrophotometry, after 3% sulfosalicylic acid precipitation of urine collected from rats individually housed in metabolic cages for 24h before ADR administration and 4 weeks after drug treatment. Blood samples (3ml from inguinal vein) were collected and centrifuged to separate the plasma. The biochemical parameters (sCh, Alb, BUN, sCr, LN, FN, Col, TGFβ1 and CTGF) levels in the plasma were measured according to the users’ instruction of diagnostic kits (Roche AG, Switzerland) and by the biochemical analyzer Cobas Mira plus (Roche AG, Switzerland) in Guangzhou Medical School, Guangdong province of P.R.China. The parameters were detected by OLYMPUS AU2700 Automatic Biochemical Analyzer. Blood pressure was monitored continuously until 60 min after the administration of glycyrrhizin and stored.

**Histopathological Detection**

Four weeks after drug treatment, rats were anaesthetized with 10% chloral hydrate (0.5ml/100g) by ip injection. One kidney was removed and fixed in 10% formalin, embedded in paraffin and examined in multiple consequent sections. Histopathologic study was carried out using PAS staining and observed by light microscopy. Additionally, other partial kidney tissues were fixed by 3% glutaraldehyde and 1% osmic acid in order, and stained by uranyl acetate and citric acid. Then, the histopathological changes in the renal cortex were observed by electron microscopy. To evaluate glomerular size, 100 glomeruli/group were examined. Glomerular surface area (µm²) was determined in digital images by a single blinded observer using the Soft Imaging System (Olympus, London, UK).

**Immunohistochemistry Assay**

Expression of LN, FN, Col, TGFβ1 and CTGF in kidney tissue was analyzed by immunohistochemical staining. Tissues were fixed in 10% buffered formalin and embedded in paraffin. Commercially available monoclonal antibodies to LN, FN, Col, TGFβ1 and CTGF were used. Immunohistochemical staining was carried out using the avidin biotin method and a commercially available kit. One paraffin-embedded block of kidney tissue was selected from each case and cut into 4µm sections. Deparaffinized sections were treated with methanol containing 3% hydrogen peroxide for 10 min before conducting antigen retrieval using a microwave oven at 95°C for 5 minutes and cooling at 25°C for 2 hr. After washing with phosphate buffered saline (PBS), blocking serum was applied for 10 min. The sections were incubated with an anti LN monoclonal antibody (1:300), anti FN monoclonal antibody (1:300), anti Col monoclonal antibody (1:50), anti TGFβ1 monoclonal antibody (1:500) and anti CTGF monoclonal antibody (1:25) overnight at 4°C. Negative control sections were incubated with PBS instead of the primary antibody. After washing in PBS, a biotin marked secondary anti-
body was applied for 10 min followed by a peroxidase marked streptavidin for an additional 10 min. The reaction was visualized using 3, 3’-diaminobenzidine-tetrahydrochloride. The nuclei were counterstained with hematoxylin. Positive and negative immunohistochemistry controls were routinely used. Reproducibility of staining was confirmed by reimmunostaining via the same method in multiple, randomly selected specimens. Two experienced pathologists, blinded to the description of the slices, examined the stained specimens independently.

Images were collected by Olympus DD70 BX51 (Olympus, Japan) and analyzed by IMAGE-PRO plus 4.1 software (Media Cybernetics, USA). Ten visual fields in each section were randomly selected and the value of the relative optical density (OD) was measured. Five to ten representative replicate sections were assessed per animal and a minimum of 6 animals were analyzed per group. The extent of immunohistochemical staining of LN, FN, Col, TGFβ1 and CTGF was semi-quantified by the mean value of OD in every section.

Fluorimeter Based Quantitative RT-PCR Assay

The mRNA expression of TGFβ1 and CTGF in rat kidney tissue in each group was detected by fluorescein based quantitative RT-PCR. Total RNA was extracted from the cortex of the kidney with a phenol/guanidine isothiocyanate based reagent (TrizolTM, Gibco BRL, Scotland.). Briefly, 1 ml Trizol and 200µl chloroform were added to each tube. The RNA was then precipitated with 500µl isopropanol (v/v) and washed with 75% ethanol and air dried. The purified RNA was then dissolved in 10 µl diethyl pyrocarbonate (DEPC) treated water. The optical density was measured at 260/280 wavelength and stored at -80°C prior to being used in the synthesis of cDNA.

RNA was reverse transcribed using the Super scriptε™ Preamplification System (Life Technology). 500ng total RNA were mixed with 500ng oligo (dT) primers and incubated at 70°C for 10 min. The mixture was then chilled on ice and incubated with a 1× reverse transcriptase buffer (50 mM Tris-HCl, pH 8.4, 75mM KCl), 3 mM MgCl , 500µM of each deoxynucleotide, 10mM dithiothreitol and 200U of Superscript II RT reverse transcriptase at 42°C for 50 min. The 20µl reactions were further incubated at 70°C for 15 min and 2 U RNase H was added to each tube. Following final incubation at 37°C for 20 min, the cDNAs were stored at -80°C until being used.

PCR was performed in MJ Opticon Monitor 2.0 (MJ Ltd, USA) using SYBR Green I (Biogene) as fluorescein. The following pairs of primers were used: β-actin (258bp): 5’GAC CTT CAA CAC CCC AGC CA 3’ (sense), 5’ GTC ACG CAC GAT TTC CCT CTC 3’ (antisense); TGFβ1 (209bp): 5’ GTG GAC CGC AAC AAC GCA 3’ (sense), 5’ ACC AAG GTA ACG CCA GGA AT 3’ (antisense); CTGF (133bp) 5’ GCG TAA AGC CAG GGA GTA 3’ (sense), 5’ AGC AGT TAG GAA CCC AGA TT 3’ (antisense). All primers were synthesized by Sangon Co. (Shanghai, China). The PCR profile consisted of an initial melting step of 2 min at 94°C, followed by 39 cycles of 45 s at 94°C, 20 s at 62°C and 20 s at 72°C, and a final elongation step of 10 min at 62°C.

Sample quantification was carried out with Se quence Detection System (SDS2.0) software constructing a standard calibration curve using serial dilutions of β-actin DNA of known concentration, from which the concentration of an unknown sample could be determined. In order to analyze interassay variation, a least square curve fitting test was used to construct a fit, and the standard error of the data points to the curve was estimated by plotting the log of the sample concentration against the number of cycles the PCR machine undertook to yield a set of fluorescent data. The standard error was estimated to be 15% on the log and was within the values described for β-actin quantification using competitive PCR.
TABLE 1. Quantitative results of 24hUPr, sCh, Alb, BUN, sCr in each group (mean±SD) (n=12)

<table>
<thead>
<tr>
<th>Group</th>
<th>24 UPr (mg/24h)</th>
<th>Alb (g/L)</th>
<th>sCh (mmol/L)</th>
<th>sCr (mmol/L)</th>
<th>BUN (mmol/L)</th>
<th>Blood pressure (mmHg)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>16.7±3.8</td>
<td>34.2±3.5</td>
<td>1.7±0.2</td>
<td>33.6±5.0</td>
<td>4.7±1.0</td>
<td>126.2±9.3</td>
</tr>
<tr>
<td>Untreated</td>
<td>60.6±4.7</td>
<td>20.5±1.6</td>
<td>4.0±0.8</td>
<td>75.1±6.6</td>
<td>11.8±1.6</td>
<td>132.8±10.1</td>
</tr>
<tr>
<td>Glycyrrhizin</td>
<td>46.4±8.3</td>
<td>28.1±2.9</td>
<td>2.3±0.9</td>
<td>52.6±9.0</td>
<td>8.8±1.1</td>
<td>127±5±9.6</td>
</tr>
</tbody>
</table>

All results were expressed as mean±SD. (n=12)

1 P<0.05, 2 P<0.01 vs. the control group, 3 P<0.05, 4 P<0.01 vs. the untreated model group.

TABLE 2. Quantitative changes in levels of LN, FN, ColIIV, TGFβ1 and CTGF in blood plasma in different groups (mean±SD) (n=12)

<table>
<thead>
<tr>
<th>Group</th>
<th>LN (µg/L)</th>
<th>FN (µg/L)</th>
<th>ColIIV (µg/L)</th>
<th>TGFβ1 (ng/L)</th>
<th>CTGF (ng/L)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>42.5±7.5</td>
<td>81.5±3.2</td>
<td>23.7±12.7</td>
<td>22.7±10.5</td>
<td>18.6±14.7</td>
</tr>
<tr>
<td>Untreated</td>
<td>83.6±9.0</td>
<td>196.9±2.1</td>
<td>58.2±14.8</td>
<td>46.6±11.3</td>
<td>40.6±15.0</td>
</tr>
<tr>
<td>Glycyrrhizin</td>
<td>64.4±8.4</td>
<td>134.3±4.1</td>
<td>42.5±13.5</td>
<td>33.3±10.2</td>
<td>30.5±15.5</td>
</tr>
</tbody>
</table>

All results were expressed as mean±SD. (n=12)

1 P<0.05, 2 P<0.01 vs. the control group, 3 P<0.05, 4 P<0.01 vs. the untreated model group.

**Immunoperoxidase Staining of Macrophages**

To evaluate the infiltration of macrophages, immunoperoxidase staining using an ABC kit (Vector Laboratories, Burlingame, CA) was performed. Briefly, fresh-frozen kidney sections were used. Non-specific binding was blocked by incubation with 10% normal sheep serum in Tris-buffered saline for 20 min to reduce background interference. Non-specific staining was blocked by 15-min incubation with avidin and then biotin using an avidin-biotin blocking kit (Vector Laboratories). Endogenous peroxidase activity was inhibited by 20 min incubation with methanol containing 0.3% H2O2. Sections were first incubated with a monoclonal antibody against rat monocytes/macrophages (ED1) for 12 h at 4°C. The sections were then incubated with biotin-labeled goat anti-mouse IgG for 30 min. Biotinylated horseradish peroxidase was applied for 30 min. Peroxidase activity was developed in 3, 3-diaminobenzidine. Mayer’s hematoxylin was added as a counterstain.

Intraglomerular ED1-positive cells were counted in 200 glomeruli/group by two independent observers with no prior knowledge of the experimental design. The average number per glomerulus was used for the estimation.

**Calculations and statistics**

Data obtained from the above experiments were expressed as mean±S.D. (mean±SD) and handled by SPSS12.0 software. Data was analyzed by one way ANOVA with the post hoc Tukey’s test applied for paired comparisons and Kruskal Wallis for non parametric test. A difference was considered significant if the P < 0.05.

**Results**

**Quantitative Detection of 24h UPr, Biochemistry and Arterial Blood Pressure**

Four weeks after the ADR administration, compared with control, levels of 24UPr, sCh, sCr, BUN, LN, FN, ColIIV, TGFβ1, and CTGF in the untreated group were increased while the level of Alb in plasma had decreased (Tables 1 and 2).

Seven 7 days of glycyrrhizin treatment protected the rats from NS by reducing levels of 24UPr, sCh, sCr and BUN while increasing Alb. The levels of LN, FN, Col, TGFβ1 and CTGF in the rats’ plasma also decreased after glycyrrhizin administration. NS rats developed an increase in mean arterial blood pressure which was reduced by Glycyrrhizin treatment (Table 1).
Histopathologic Changes in Renal Cortex

Light microscope photographs show, focal areas of mesangial cell proliferation and vacuolar degeneration, and tubulointerstitial inflammation in kidney tissues in the untreated model group of rats at the time of sacrifice (Figure 1A-C). The number of tubulointerstitial cells in the untreated group was greater than in the other two groups. Histopathological changes of mesangial proliferation and tubulointerstitial inflammation were reduced in rats who received glycyrrhizin treatment as were the number of tubulointerstitial cells. In the untreated group, 2 weeks after modeling, inflammatory cell infiltration appeared in the renal interstitium, renal tubules had atrophied a little and the vessel walls were thickened. Four weeks after modeling, these changes were more severe and fibrous degeneration had occurred in the renal interstitium. Furthermore, electron microscopy in untreated rats showed proliferation of intercapillary cells, increase of ECM and collagen fiber-like filaments, and fusion of foot processes (Figure 1D-E). After glycyrrhizin, mesangial cell proliferation, foot process confluence, and basement membrane thickening decreased compared with that in the untreated group.

Glomeruli from untreated NS animals tended to be larger than those from normal controls, whereas glomerular surface area was decreased by glycyrrhizin compared with the untreated rats (Figure 1F). Collectively, these results demonstrate that glycyrrhizin decreased glomerular hypertrophy in NS animals.
Expression of LN, FN, Col, TGFβ1 and CTGF Protein in Renal Cortex

There was no expression in the kidneys of animals in the control group (Figure 2A, D and G). In rats with ADR-induced NS, LN, FN and Col were expressed strongly along with the aggravation of NS. Glomcruli showed sclerosis and were occupied completely with LN, FN and Col (Figure 2B, E and H). After glycyrrhizin treatment, expression of LN, FN and Col was weakly positive (Figure 2C, F and I).

LN, FN, Col, TGFβ1 and CTGF expression in rats with ADR induced NS was increased compared with normal rats. After glycyrrhizin treatment, expression
Table 3: Semi-quantitative analysis of the expression levels of LN, FN, ColIV, TGFβ1 and CTGF in kidney sections (n=12, x̄±s)

<table>
<thead>
<tr>
<th>Markers</th>
<th>Control group</th>
<th>Untreated group</th>
<th>Glycyrrhizin group</th>
</tr>
</thead>
<tbody>
<tr>
<td>LN</td>
<td>1.88±0.16</td>
<td>2.90±0.18</td>
<td>2.21±0.18</td>
</tr>
<tr>
<td>FN</td>
<td>1.98±0.14</td>
<td>3.27±0.23</td>
<td>2.38±0.30</td>
</tr>
<tr>
<td>Col IV</td>
<td>0.24±0.04</td>
<td>0.42±0.06</td>
<td>0.35±0.04</td>
</tr>
<tr>
<td>TGFβ1</td>
<td>1.20±0.08</td>
<td>3.10±0.17</td>
<td>2.32±0.13</td>
</tr>
<tr>
<td>CTGF</td>
<td>0.23±0.03</td>
<td>0.34±0.05</td>
<td>0.29±0.05</td>
</tr>
</tbody>
</table>

1 P<0.05 vs. the control group; 2 P<0.01 vs. the control group; 3 P<0.05 vs. the untreated model group; 4 P<0.01 vs. the untreated model group.

Table 4: Expression levels of TGFβ1 and CTGF mRNA in kidney tissue of rats in different groups (x̄±s, ng/ml, n=10)

<table>
<thead>
<tr>
<th></th>
<th>Control group</th>
<th>Untreated model group</th>
<th>Glycyrrhizin treatment group</th>
</tr>
</thead>
<tbody>
<tr>
<td>TGFβ1</td>
<td>3.7±0.8</td>
<td>17.2±1.4</td>
<td>5.1±0.3</td>
</tr>
<tr>
<td>CTGF</td>
<td>0.4±0.1</td>
<td>4.1±0.3</td>
<td>1.9±0.2</td>
</tr>
</tbody>
</table>

All results were expressed as mean±SD. (ng/ml, n=10) 1 P<0.05, 2 P<0.01 referred to comparison with the control group, 3 P<0.01 referred to comparison with the untreated model group.

The number of macrophages (ED1-positive cells) in glomeruli was higher in the untreated group than in the control group. Glycyrrhizin treatment reduced the number of glomerular infiltrating macrophages (Figure 3).

Discussion

In this study, 24UPr, sCh, Alb, BUN, sCr and the mean arterial blood pressure were increased in the rats with NS. In the group receiving glycyrrhizin treatment, these makers were decreased and renal function was ameliorated. Proteinuria effects cellular infiltration. Protein overload may induce functional alterations of tubular cells, overexpressing proinflammatory mediators. Thus, tubulointerstitial inflammation in NS models corresponds to the time of overt proteinuria, as in this study.

Glycyrrhizin is a glycoside of glycyrrhetinic acid with two molecules of glucuronic acid. It is one of the active ingredients of licorice, roots of Glycyrrhiza glabra and G. ularensis. It has been prescribed fre-
quent in a herbal Chinese medicine formula to treat various diseases, such as chronic hepatitis C by decreasing in serum alkaline aminotransferase levels in patients \(^9\), kidney ischemia-reperfusion injury in animal models \(^10\) and allergic diseases \(^11\) because of its anti-inflammatory and antioxidant activities. However, excessive glycyrrhizin may induce peripheral edema, hypokalemia, myopathy, and hypertension, which are symptoms of pseudoaldosteronism. It is potentially fatal.\(^{15,16}\) The frequency of pseudoaldosteronism caused by glycyrrhizin depends on the dose and the duration of treatment and is thought to be due to inhibition of type 2 \(11\beta\)-hydroxysteroid dehydrogenase (\(11\beta\)-HSD2) in renal tubular epithelial cells by glycyrrhetic acid.\(^7\)

ADR, an anticancer drug, forms a complex with DNA, inhibiting synthesis of both DNA and RNA.\(^{18}\) Kidney injury induced by ADR in a range of laboratory animals is an analogue of NS in humans, the appearances of which are minimal change nephropathy, focal and segmental glomerulosclerosis.\(^{19}\) Our histopathological findings showed tubulointerstitial changes with cellular infiltration and vacuolar degeneration in the untreated group, by light and electron microscopy. Interstitial inflammation is considered to be an important determinant of the outcome of glomerular inflammation. Several studies suggested that myofibroblasts of the interstitium may play a crucial role in the pathogenesis of fibrosis in glomerular diseases.\(^{20,21}\) Thus, the tubulointerstitial cellular response and vacuolar degeneration seen in our rat model with NS emphasizes the severity of the NS. On the other hand, attenuation of interstitial inflammation and other renal injuries are secondary to inhibition of proteinuria by glycyrrhizin. Glycyrrhizin may delay progression of NS.

Immunohistochemical staining and Fluorimeter based quantitative RT-PCR showed changes in expression of LN, FN, Col, TGF\(\beta\)1 and CTGF in different groups. We interpreted this as due to ADR administration and glycyrrhizin treatment. ECM accumulation, caused by disequilibrium of composition and degradation, is considered a principal pathological feature of NS and may lead to glomerular sclerosis, the key factor in the development of NS.\(^{22}\) LN, FN and ColIV are the ingredients of ECM, while TGF\(\beta\)1 and CTGF are important participants in the process of ECM accumulation.\(^{23,24}\) We also found that glycyrrhizin suppressed the infiltration of macrophages in the kidneys of NS rats. Therefore, we believe that glycyrrhizin may inhibit ECM accumulation, prevent glomerular sclerosis and relieve the renal injury of NS.

In conclusion, glycyrrhizin exerts protective effects on ADR-induced NS in rats. Glycyrrhizin reduces proteinuria, 24UPr, sCh, Alb, BUN, sCr, and the mean arterial blood pressure. Also, it decreases expression of LN, FN, Col, TGF\(\beta\)1 and CTGF in kidney tissues, improves renal function and reduces the severity of glomerulosclerosis and retards the development of NS.

Acknowledgments

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