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ORIGINAL RESEARCH

Protective effect of Huang Gan formula in 5/6 nephrectomized rats by depressing the Wnt/ β -catenin signaling pathway

Liqian Mo¹ Xiaoyan Xiao¹ Shaolian Song¹ Hui Miao² Shiting Liu¹ Dan Guo¹ Xiao Li³ Can Bu¹ Lianbing Hou¹ Xixiao Yang¹

¹Department of Pharmacy, Nanfang Hospital, Southern Medical University, ²School of Pharmaceutical Sciences, Sun Yat-sen University, Guangzhou, ³Department of Pharmacy, The Affiliated Hospital of Qingdao University Medical School, Qingdao, People's Republic of China

Correspondences: Lianbing Hou; Xixiao Yang

Department of Pharmacy, Nanfang Hospital, Southern Medical University, No 1838, Guangzhou Boulevard (North), Guangzhou 510515, People's Republic of China

Tel +86 20 6164 2175; +86 20 6278 7215 Fax +86 20 8770 1797 Email hlianbing@163.com; yxixiao@ hotmail.com

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Abstract: Huang Gan formula (HGF) is a new traditional Chinese herbal medicine created according to the basic theory of traditional Chinese medicine. The aim of this study is to evaluate the effects of HGF on chronic kidney disease and determine the mechanisms of action. The extract of HGF was prepared, and qualitative and quantitative determination of phytochemical was performed with quadrupole time-of-flight mass spectrometer and high-performance liquid chromatography. Sprague-Dawley rats (n=72) were submitted to 5/6 nephrectomy (Nx), and then respectively treated with uremic clearance granule, losartan, HGF low dose, HGF middle dose, and HGF high dose once per day for 12 weeks. The sham group of operated rats (n=22) was treated with normal saline or HGF middle dose as a background control group. Blood and urine biochemical parameters, renal tissue morphology, and mRNA and proteins of Wnt/β-catenin signaling pathways were investigated. The results showed that the quality of the extraction process could be controlled, and a total of eight major compounds were identified and quantified. HGF could decrease the level of serum creatinine, blood urea nitrogen, and urine protein and increase the renal index and creatinine clearance rate in a dose-dependent manner. HGF also remarkably reduced the glomerulosclerosis and tubulointerstitial fibrosis by blocking the Wnt/ β -catenin signaling pathway through inhibiting the Wnt1, β -catenin, transcription factor 4, and fibronectin 1 expressions, simultaneously measured through mRNA and protein levels in the remnant kidney. These results suggest that extraction of HGF could improve remnant renal function and possibly ameliorate glomerulosclerosis and tubulointerstitial fibrosis by depressing the Wnt/ β -catenin signaling pathway.

Keywords: Huang Gan formula, traditional Chinese medicine, chronic kidney disease, Wnt/β-catenin signaling pathway

Introduction

Chronic kidney disease (CKD) was defined in The National Kidney Foundation Kidney Disease Outcomes Quality Initiative (KDOQI) clinical practice guidelines as kidney damage or glomerular filtration rate (GFR) lower than 60 mL/min per 1.73 m² of tissue for 3 months or longer.¹ Prevalence of CKD is estimated to be 8%–16% worldwide, and its incidence is increasing in developed and developing nations.² CKD has become a major public health problem on a global scale because of its serious complications and poor prognosis, and it presents huge socioeconomic burdens to patients, families, and societies.³

There are many risk factors leading to CKD, including hypertension, diabetes mellitus, obesity, infections, xenobiotics, herbal or environmental toxins, mechanical obstruction, and deposition of immune complexes.^{2,4} Despite the different factors leading to CKD, the kidney will eventually exhibit the same pathologic characteristics, including glomerulosclerosis and tubulointerstitial fibrosis, which show main

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histopathological features such as myofibroblast proliferation and excessive accumulation of extracellular matrix (ECM).⁵ Molecular mediators involved with fibrogenic pathway include transforming growth factor (TGF- β), bone morphogenic protein, connective tissue growth factor, plateletderived growth factor, etc. These pathways represent new potential targets for novel strategies to inhibit or retard the progress of kidney fibrosis.^{5,6}

The Wnt/ β -catenin signaling pathway is the canonical pathway, was recently discovered and plays an important role in mediating the renal fibrosis.7,8 In the absence of the Wnt ligand, β -catenin is bound to the scaffold proteins axis inhibition protein (Axin) and adenomatous polyposis coli, and then glycogen synthase kinase (GSK)-3 β is phosphorylated to form a degradation complex, which mainly degrades β-catenin to prevent entry into the nucleus. Upon Wnt activation, Wnt ligands bind to and activate two distinct cell surface receptors named frizzled and low density lipoprotein receptor-related protein (LRP) 5 or 6 and induce the phosphorylation of LRP6, which recruits Axin to the plasma membrane, resulting in the inhibition of β-catenin phosphorylation/degradation. Accumulated β -catenin translocates to the nucleus and complexes with transcription factor/lymphoid enhancer factor (TCF/LEF), moderating the canonical Wnt-target gene transcription.9,10 A simplified schematic of the pathway is shown in Figure S1. General effects of canonical Wnt-pathway activation include the moderation of cell proliferation, differentiation, and cell fate determination. The different Wnt regulators and regulatory mechanism of Wnt signaling pathway have previously been described (www.stanford.edu/~rnusse/ wntwindow.html). Hyperactive Wnt signaling pathway may lead to the development and progression of renal tubular and glomerular diseases and eventually result in CKD.⁶⁻⁸ Thus, the Wnt/ β -catenin signaling pathway may be a potential and novel target for therapeutic intervention of CKD.

Uremic clearance granule (UCG) was developed by our department, which was the first Chinese herbal medicine in the field on prevention of CKD progress in People's Republic of China. Because of the problems and deficiencies in the production and promotion of UCG, our research group intended to work for the secondary development and simplify the prescription according to the theoretical basis of traditional Chinese medicine (TCM), and eventually got the new production called Huang Gan formula (HGF), which had been demonstrated could significantly decrease the serum creatinine (Scr) and blood urea nitrogen (BUN), inhibit oxidative stress in CKD rat models induced by adenine, and delay the progress of renal fibrosis. In this study, we intend to use high-performance liquid chromatography (HPLC)– quadrupole time-of-flight mass spectrometer (Q-TOF-MS) techniques to identify and detect the main ingredient and content of HGF. Based on identified effective substance, the therapeutic effect of HGF then would be evaluated by prepared 5/6 Nx rat model of CKD. Finally, we investigate the changes in Wnt/ β -catenin signaling pathway in renal, to explore the possible anti-renal fibrosis mechanisms of HGF.

Materials and reagents

The herbs of rheum, Radix Bupleuri, Rhizoma Zingiberis, Radix Glycyrrhizae, and Radix Aconiti Lateralis Preparata, were purchased from Kangmei Pharmaceutical Co, Ltd (Guangdong, People's Republic of China) and authenticated by Professor Hongwei Zhang of School of Traditional Chinese Medicine in Southern Medical University. Losartan was purchased from Merck & Co., Inc. (Whitehouse Station, NJ, USA), and the UCG was purchased from Consun Pharmaceutical Group Limited (Guangzhou, People's Republic of China). Both the one-step reverse transcriptase-polymerase chain reaction (RT-PCR) kit (DRR036A) and the SYBR Premix Taq kit (DRR820A) for real-time PCR were obtained from TaKaRa Biotech Co. (Dalian, People's Republic of China). Rabbit polyclonal antibody Wnt1 (ab85060; Abcam, Inc., Cambridge, MA, USA), β-catenin (BS1982; Bioworld Technology, Inc., St Louis Park, MN, USA), GSK-3β (sc9166; Santa Cruz Biotechnology, Santa Cruz, CA, USA), p-GSK-3ß (sc-135653; Santa Cruz Biotechnology Inc., Dallas, TX, USA), transcription factor 4 (TCF4) (BS6172; Bioworld Technology, Inc.), dickkopf 1 (Dkk1) (BS7731; Bioworld Technology, Inc.), fibronectin 1 (Fn1) (ab2413; Abcam, Inc.), and glyceraldehyde 3-phosphate dehydrogenase (GAPDH) (ab9485; Abcam, Inc.) were purchased. Reference substances such as liquiritin, tectoridin, luteolin, aloe-emodin, rhein, emodin, chrysophanol, and physcion were provided by National Institutes for Food and Drug Control (Beijing, People's Republic of China).

Extract preparation

Radix et Rhizoma Rhei (307.1 g) and Radix Bupleuri (614.3 g) were macerated with 70% ethanol (1:7, m/v) for 2 hours and then extracted by refluxing for a period of 1.5 hours for three times. Rhizoma Zingiberis (229.9 g), Radix Glycyrrhizae (307.1 g), and Aconiti Lateralis Radix Preparata (245.7 g) were macerated with water (1:7, m/v) for 2 hours and decocted twice every 2 hours each time. Two-time water decoction was blended together, filtered with three-layer gauzes, and concentrated to 200 mL. The concentrated

solution was precipitated with a final concentration of 80% ethanol for 24 hours. The supernatant merged with the alcohol extraction of Radix et Rhizoma Rhei and Radix Bupleuri for rotary evaporation concentration. The extraction process was performed ten times repeatedly according to the above methods. The yield of extract reached 21.7% (m/m). Similarity indices of the HPLC chromatograms of ten batches of extracts were evaluated by the professional software, Similarity Evaluation System for Chromatographic Fingerprint of Traditional Chinese Medicine (Version 2004A), which was recommended by Chinese Pharmacopoeia Commission and State Food and Drug Administration. Concentrated extract was placed in 4°C for use.

Phytochemical analyses

In all, 50 mg vacuum drying HGF was dissolved in 50 mL acetonitrile/water (1:1, v/v) and sonicated for 15 minutes. The solution was filtered through a 0.45 μ m nylon filter membrane, and then 20 μ L samples were submitted for HPLC analysis (LC-20A; Shimadzu, Tokyo, Japan). The reference substance stock solutions (0.1 mg/mL) were prepared in acetonitrile and used for identification and creation of a standard curve. HPLC with PDA detector was performed to scan the maximum absorption wavelength of HGF samples.

HPLC conditions are as follows: column, ZORBAX Eclipse Plus-C18 (4.6 mm, 150 mm, 5 μ m, Agilent Technologies, Santa Clara, CA, USA); mobile phase, consisted of water phase A (0.1% formic acid) and organic phase B (acetonitrile); column, eluted with a gradient of 85%–77% A at the beginning of 5 minutes, 77%–70% A at 5–30 minutes, 70%–55% A at 30–40 minutes, 55%–50% A at 40–60 minutes, 50%–30% A at 60–70 minutes, 30%–27% A at 70–85 minutes, 27%–20% A at 85–90 minutes; flow rate, 0.8 mL/min; column temperature, 30°C; and detection, 254 nm.

The HPLC–Q-TOF-MS spectrometry was performed for qualitative research. The operating parameters for the electrospray ion (ESI) were as follows: the capillary voltage for positive mode (ESI+) and negative mode (ESI–) ionization was set to ± 1.59 kV. Drying gas (N₂) flow rate was set to 1.5 L/min at a temperature of 200°C. The collision energy was set at 15–40 V. The LC–MS accurate mass spectra were recorded in the 100–1,000 m/z range.

Animals and treatment

All animal procedures were approved by the Animal Experiment Committee of Southern Medical University. Male 12-week-old Sprague-Dawley rats (n=94) weighing 180–200 g were purchased from the Southern Medical

University Animal Experiment Center. All animals were housed in an environmentally controlled room at 24°C±2°C with a relative humidity of 55%±5% and a 12-hour light/dark cycle. Food and water were available ad libitum.

After 1 week of adjustment, 5/6 Nx were performed according to the technique described previously11 under anesthesia with 10% chloral hydrate (1 mL/300 g weight). In all, 72 rats were taken for surgical resection of two-third of the left kidney and the remaining 22 rats had a sham operation performed. One week later, the survival ones among the 72 rats were subjected to surgical removal of the right kidney and the sham-operated rats to an identical procedure but without removal of any part of the right kidney. One week after the operation, the surviving 5/6 Nx rats (n=68) were randomized by the percent of the remnant kidney weight removed (right kidney weight - weight of two poles of left kidney)/(right kidney weight $\times 100$), divided into six subgroups (n=11-12) in each subgroup). The sham-operated rats (n=22) were randomly divided into the sham-operated group (Sham) and the background control group (BC group).

The dose administration was calculated according to the rat body surface area compared to human. After 1 week of acclimatization, Scr and BUN levels were determined, which indicated that the 5/6 Nx model was successful if the levels were double the sham group. Then, intragastric administration was performed in each group as follows: losartan group (losartan; 30 mg/kg/day), UCG (3.6 g/kg/day), HGF low dose (HGF-L; 3.6 g/kg/day, clinical equivalency dose) group, HGF middle dose (HGF-M; 7.2 g/kg/day) group, and HGF high dose (HGF-H; 14.4 g/kg/day) group. BC group was given HGF-M (7.2 g/kg/day), while the sham and 5/6 Nx groups were given the same volume of saline solution. All animals were treated by oral gavage every 12 hours continuously for 12 weeks.

Sample collection and biochemical marker detection

After 12 weeks of treatment, each rat was fasted and placed in an individual metabolism cage for 24 hours urine collection. The serum was collected from abdominal aortic arterial puncture while under anesthesia. The levels of the Scr, BUN, urine creatinine (Ucr), and urine protein (UPr) were detected using an automatic biochemical analyzer (Olympus AU5421; Olympus Corporation, Shinjuku, Tokyo, Japan). The remnant renal tissue was divided into three parts after weighing. One-third was fixed in 10% neutral-buffered formalin for histopathological and immunohistochemistry examination, while the other two-third of the cortex (100 mg each) was stored at -80°C for RT-PCR and western blot (WB).

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Histology and immunohistochemistry staining

The fixed renal tissues underwent conventional alcohol dehydration, were embedded in paraffin and took serial sections (4 μ m for each section), stained with hematoxylin–eosin (HE) and Masson's trichrome, and examined under a light microscope (Olympus). A semi-quantitative analysis of optical density was performed using Image-Pro Plus 6.0 image system (Media Cybernetics, Inc., MD, USA) for Masson staining pictures with the following conditions: ten non-overlapping renal interstitial vision under 400× magnification. The semiquantitative analysis was determined by the percentage of the positive staining area accounted for over the entire visual field. All measurements were performed in a blinded fashion.

Immunohistochemical staining was performed using routine protocol.¹² The sections were stained with the following rabbit polyclonal antibodies: Wnt1, β -catenin, GSK-3 β , TCF4, Dkk1, and Fn1. The IgG isotype negative control was stained by replacing the primary antibody with rabbit serum. The average optical density values were semi-quantified by Image-Pro Plus 6.0 under 400× magnification.

Real-time quantitative PCR

Total RNA was extracted from the renal cortex tissue (n=6) using TRIzol RNA isolation system (Thermo Fisher Scientific, Waltham, MA, USA). The cDNA was synthesized, and PCR was carried out using a standard PCR kit. PCR primers are shown in Table S1. RT-PCR was performed using Light-Cycler[®] 480 (Hoffmann-La Roche Ltd., Basel, Switzerland), which expresses the number of threshold cycle (C_t). Relative expression levels of target genes were determined by calculating 2^{-DDC_t} .¹³

WB analysis

Western blotting was conducted as previously described.¹⁴ Proteins were extracted from approximately 100 mg renal cortex tissues and quantified with bicinchoninic acid assay kit (KeyGen Biotech, Nanjing, People's Republic of China). The proteins of six randomly selected samples were mixed in equal amounts, and total protein was analyzed in triplicates. Antibodies used were the same as the immunohistochemical staining kit, except for internal control GAPDH.

Statistical analyses

Data are presented as the mean ± standard deviation. Statistical analyses were performed using one-way analysis of variance (SPSS13.0, Chicago, USA) test followed by Tamhane's test to compare three or more groups or by Student's *t*-test to compare two groups. P < 0.05 was considered to be statistically significant.

Results Quality control and phytochemical analyses

Figure 1 shows the HPLC fingerprint chromatogram of ten different batches of HGF extract. The standardized characteristic chromatogram was generated by the similarity evaluation system according to the ten chromatograms (shown in Supplementary materials). The similarity indices were calculated and compared with the standardized characteristic chromatogram, which was 0.929, 0.987, 0.982, 0.965, 0.993, 0.974, 0.994, 0.997, 0.942, and 0.981 from S1 to S10 in sequence. All of the similarity indices were greater than 0.92, indicating the components of different batches of extracts were similar and stable. The reference fingerprint of HGF (shown in Figure S2) was produced according to HPLC chromatograms of the ten batches of HGF extracts by Similarity Evaluation System for Chromatographic Fingerprint of Traditional Chinese Medicine (Version 2004A). A total of 21 common peaks were marked, and the maximum peak was set to 18, which was the reference chromatographic peak, while the retention time and peak area were set to 1. The relative retention time and relative peak area of common peaks compared to peak 18 are listed in Tables S2 and S3. Fingerprint results showed that the ten batches of HGF possessed well consistency and the extraction method was stable. Using HPLC fingerprint can better control overall quality of HGF extract, and the quality of the extraction process could also be controlled.

The HPLC–Q-TOF-MS spectrometry was used to characterize the chemical compositions within the HGF extract. The total ion current chromatograms of the HGF extract is shown in Figure 2. A total of eight major compounds were identified and quantified by comparing reference standards. Their chemical names and contents (m/m) were liquiritin (0.167%), tectoridin (1.98%), luteolin (1.03%), aloe-emodin (3.78%), rhein (3.13%), emodin (1.54%), chrysophanol (2.72%), and physcion (1.55%). The results of MS qualitative analysis of HGF is shown in Table S4.

Assessment of renal function

As shown in Table 1, the 5/6 Nx group renal index, Scr, BUN, and UPr levels were significantly higher than the sham and BC groups, and the creatinine clearance rate (Ccr) was significantly lower. The levels of renal index in medicated

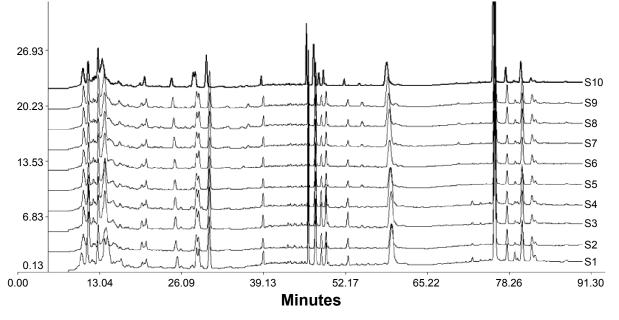


Figure I Fingerprint chromatogram of ten batches of HGF extracts.

Notes: S1–S10 were the UV–HPLC chromatograms of ten batches of HGF extracts, which were detected under the same HPLC conditions. **Abbreviations:** HGF, Huang Gan formula; UV, ultraviolet; HPLC, high-performance liquid chromatography.

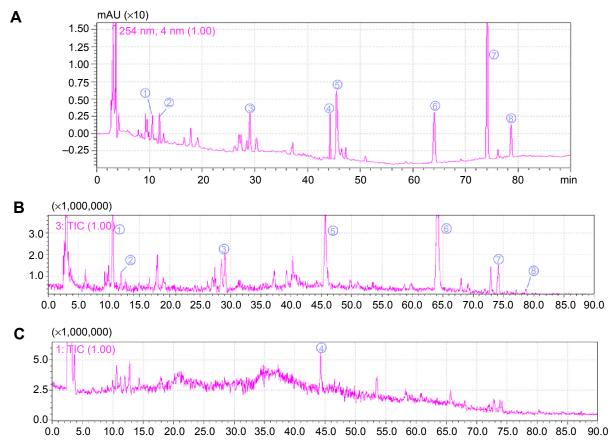


Figure 2 HPLC and Q-TOF-MS chromatograms of HGF extract.

Notes: UV-HPLC/chromatogram (254 nm) (A); the total ion current in the negative ion mode in MS (B); and the total ion current in the positive ion mode (C). The same numbers between HPLC and MS chromatograms indicate the same substances. As follows: ① liquiritin, ② tectoridin, ③ luteolin, ④ aloe-emodin, ⑤ rhein, ⑥ emodin, ⑦ chrysophanol, and ⑧ physcion.

Abbreviations: HPLC, high-performance liquid chromatography; Q-TOF-MS, quadrupole time-of-flight mass spectrometer; HGF, Huang Gan formula; UV, ultraviolet; min, minute.

groups, Scr and BUN in HGF-M and HGF-H groups, and UPr among HGF three dosage groups were all decreased significantly compared with the 5/6 Nx group, while the level of Ccr increased in HGF-L and HGF-H groups. All medicated groups had no significant difference with each other, except within the levels of BUN, Ccr, and UPr in HGF groups compared with losartan groups. There was no significant difference noted between sham and BC groups for all biochemical markers above.

Renal histopathology analysis

HE and Masson staining showed significant glomerulosclerosis and interstitial fibrosis in the 5/6 Nx group 12 weeks following 5/6 Nx and presented proliferation of mesangial cells, tubular atrophy, and infiltration of large numbers of inflammatory cells in renal interstitium. However, these pathological changes were not observed in the sham and BC groups, and the three HGF groups attenuated these pathological changes to varying degrees (Figure 3, HE/Masson). The semi-quantitative analysis of Masson trichrome staining showed strikingly increased interstitial fibrotic areas in the 5/6 Nx group compared with the sham and BC groups, and HGF groups had remarkably reduced interstitial fibrosis (P<0.05) (Table 2, Masson).

Expression and distribution of Wnt/ β-catenin signaling-associated proteins by immunohistochemistry

Figure 4 IHC shows the representative micrographs of Wnt1, β -catenin, TCF4, GSK-3 β , Dkk1, and Fn1 staining within the renal cortex sections from the 5/6 Nx group (micrographs of other groups not shown). All proteins were rarely expressed in normal kidneys, but were distributed in different regions. Wnt1, TCF4, and Dkk1 were predominantly localized within the glomeruli and renal tubules in the diseased kidneys, while β -catenin and GSK-3 β were mainly in the glomeruli and distal convoluted tubules. Fn1, the fibrosis related factor, was mostly observed in the glomerular mesangial area and tubulointerstitium. To quantify protein expression, semi-quantitative analysis of average optical density values was performed and showed that all of the proteins expressed significantly higher in the 5/6 Nx group than the sham and BC groups (P<0.05). HGF inhibited the expression of Wnt1, β -catenin, TCF4, and Fn1 markedly, but had no effect on expression of GSK-3 β and Dkk1, which showed no significant difference compared to the 5/6 Nx group (P>0.05) (Table 2). Fn1 was also expressed much less in HGF groups, consistent with the Masson staining (Figure 3).

The mRNA and protein expression of Wnt/β -catenin signaling by RT-RCR and WB

To further investigate the quantified expressions of Wnt/ β -catenin signaling-associated proteins, RT-PCR and WB were performed in the diseased kidney. As shown in Figure 5, the mRNA expressions of Wnt1, β -catenin, TCF4, GSK-3 β , Dkk1, and Fn1 in the 5/6 Nx group were increased dramatically when compared with the sham group (P<0.01). UCG, losartan, and HGF inhibited the mRNA expression of Wnt1, β -catenin, TCF4, and Fn1 significantly. Simultaneously, the HGF-L, HGF-M, and HGF-H groups also showed inhibition in a dose-dependent manner, though had no significant difference compared with UCG and losartan groups. The GSK-3 β and Dkk1 levels were reduced in HGF groups, but had no significant difference compared to the 5/6 Nx group (P>0.05).

The results of WB corroborated the RT-PCR findings (Figure 6). The expression levels of Wnt1, β -catenin,

Table I Biochemical markers of each group, $\overline{x} \pm s$ (n=9)

Group	RI	Scr (µmol/L)	BUN (mmol/L)	Ccr (mL/min)	UPr (mg/24 h)
Sham	3.23±0.29	41.4±6.0	6.9±1.3	0.57±0.16	3.3±1.4
BC	3.33±0.34	43.0±5.6	7.0±0.8	0.54±0.17	3.7±1.3
5/6 Nx	3.77±0.44#	83.0±5.8#	15.6±0.9#	0.34±0.12 [#]	5.1±0.8#
UCG	3.15±0.29*	76.6±5.4 [#]	13.1±0.9 ^{#,*}	0.53±0.18*	4.1±1.6
Losartan	3.25±0.19*	65.2±5.5 ^{#,*}	20.2±1.9 ^{#,*}	0.31±0.11#	2.2±1.1*
HGF-L	3.20±0.37*	75.5±10.3#	13.5±1.7 ^{#,∆}	0.55±0.17*,∆	4.2±1.0*,∆
HGF-M	3.37±0.25*	71.6±6.1 ^{#,*}	I 3. I±I.0 ^{#,∗,∆}	0.41±0.21	3.2±1.3*
HGF-H	3.33±0.18*	60.0±13.4*	I 3.2±I.0 ^{#,∗,∆}	0.57±0.16*,∆	3.8±0.7*,∆

Notes: Compared with Sham and BC groups, AP <0.05; compared with the 5/6 Nx group, P <0.05; and compared with the losartan group, AP <0.05. RI = renal weight (mg)/ body weight (g). Ccr = urine creatinine (μ mol/L) × urine volume per minute (mL)/Scr (μ mol/L).

Abbreviations: RI, renal index; Scr, serum creatinine; BUN, blood urea nitrogen; Ccr, creatinine clearance rate; UPr, urine protein; Sham, sham-operated group; BC, background control group; Nx, 5/6 nephrectomy; UCG, uremic clearance granule; HGF-L, Huang Gan formula low dose group; HGF-M, Huang Gan formula middle dose group; HGF-H, Huang Gan formula high dose group.

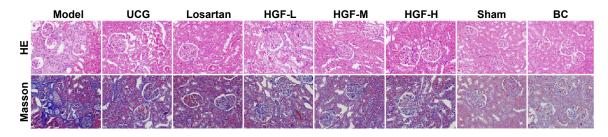


Figure 3 Representative micrographs of HE and Masson staining in the rat renal cortex sections (400×).

Abbreviations: HE, hematoxylin–eosin; UCG, uremic clearance granule; HGF-L, Huang Gan formula low dose group; HGF-M, Huang Gan formula middle dose group; HGF-H, Huang Gan formula high dose group; Sham, sham-operated group; BC, background control group.

TCF4, GSK-3 β , Dkk1, and Fn1 were upregulated in the 5/6 Nx group when compared with those in the sham group (*P*<0.01), while UCG, losartan, and HGF downregulated the expressions of these proteins except for GSK-3 β and Dkk1, which as the inhibitors of Wnt/ β -catenin signaling were both kept at high levels in drug treatment groups, but had no significant difference compared with the 5/6 Nx group (*P*>0.05). The RT-PCR and WB results were approximately consistent with the immunohistochemistry results. All of these indicators showed no significant difference between sham and BC groups.

Discussion

The pathogenic mechanisms of CKD are varied, including systemic and glomerular hypertension, excessive expression of TGF- β , the renin–angiotensin–aldosterone system, podocyte loss, dyslipidemia, etc.^{5,15} However, renal fibrosis is the common end point of virtually all progressive CKDs. Currently, drug treatments of CKD are mainly confined to control renal hypertension and improve the glomerular "high perfusion, high internal pressure and high filtration" state, thereby reducing further lesions of the kidney tissue.

Such drugs used primarily for the angiotensin converting enzyme inhibitors and angiotensin receptor blockers play an important role in control of blood pressure and reduction of proteinuria and TGF- β .¹⁶ Additionally, other treatment strategies, such as lipid-lowering therapy, correction of acidosis, prevention of atherosclerotic events, and modification with low salt and protein diets, have been implemented. But unfortunately,² these therapies only have limited efficacy confronting the complex pathogenic factors and multiple complications of CKD.

UCG as the first and dominant TCM used for treatment of CKD has been employed in People's Republic of China more than a decade, exhibiting positive effects on improving renal function and delaying the progression of chronic renal failure.^{17,18} Losartan, an angiotensin receptor blocker, is recommended by Kidney Disease: Improving Global Outcomes as a pharmacologic therapy for CKD with proteinuria and hypertension.¹⁹ Both are representative effective drugs commonly used in clinical CKD treatment. We choose these two drugs as the positive control groups, UCG as the Chinese medicine control and losartan as the western medicine control. The present study demonstrated that HGF improved

Table 2 Average optical density values of protein expressions by immunohistochemistry staining in renal tissues of rats in different groups, $\bar{x} \pm s$ (n=3)

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Group	Masson (%) [⊽]	Wntl	β-catenin	GSK-3β	Dkkl	TCF4	Fnl
Sham	5.1±0.4	3.31±0.69	1.05±0.08	4.0±0.7	0.42±0.06	2.5±0.1	2.60±0.44
BC	5.3±0.3	3.37±0.04	1.45±0.83	2.7±0.4	0.35±0.06	2.2±0.1	2.90±0.55
5/6 Nx	15.8±1.4 [#]	8.92±0.88 [#]	4.99±1.00#	8.9±1.1#	4.70±0.12#	11.2±1.2#	6.71±0.38#
Losartan	10.9±1.7 ^{#,*}	8.76±1.54#	3.50±0.35#,*	8.3±1.3#	4.29±0.70	6.7±0.3 ^{#,*}	5.26±0.11#,*
UCG	9.9±0.8 ^{#,*}	6.70±0.41 ^{#,*}	2.54±0.62 ^{#,*}	7.0±1.4 ^{#,} *	3.69±0.68	4.3±0.3 ^{#,*}	5.36±0.17 ^{#,*}
HGF-L	8.1±0.5 ^{#,∗,∆}	6.34±0.24 ^{#,*,∆}	2.97±0.82#,*	6.3±0.5 ^{#,*}	3.78±1.15	6.7±0.5 ^{#,*}	5.56±0.37#,*
HGF-M	7.3±0.9 ^{#,∗,∆}	5.95±1.11 ^{#,∗,∆}	2.49±0.57#,*	7.0±1.4 ^{#,*}	4.96±1.46	6.4±0.5 ^{#,*}	4.72±0.70 ^{#,*}
HGF-H	5.6±1.2*,∆	4.32±0.22*,∆	1.83±0.16*,∆	8.8±0.4#	6.62±1.03	6.0±0.5 ^{#,*}	4.16±0.49 ^{#,∗,∆}

Notes: All data are showed after multiplying by 10^2 . Compared with Sham and BC groups, #P<0.05; compared with the 5/6 Nx group, *P<0.05; and compared with the losartan group, $^{\Delta}P$ <0.05. $^{\nabla}$, the proportion of positive area.

Abbreviations: GSK, glycogen synthase kinase; Dkk1, dickkopf 1; TCF4, transcription factor 4; Fn1, fibronectin 1; Sham, sham-operated group; BC, background control group; Nx, 5/6 nephrectomy; UCG, uremic clearance granule; HGF-L, Huang Gan formula low dose group; HGF-M, Huang Gan formula middle dose group; HGF-H, Huang Gan formula high dose group.

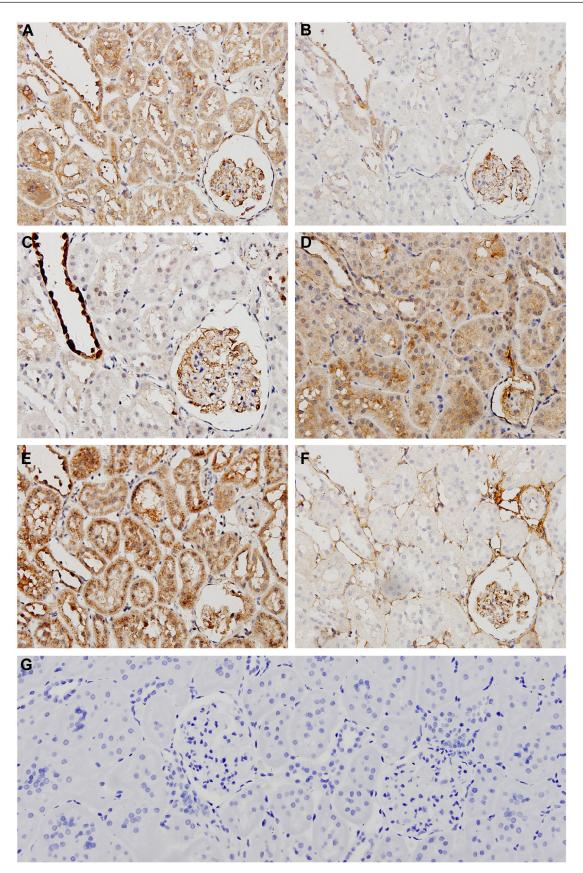


Figure 4 Representative micrographs of immunohistochemistry staining in the 5/6 Nx group rat renal cortex sections (400×). **Notes:** Immunohistochemistry staining of 5/6 Nx group for Wnt1 (**A**), β-catenin (**B**), GSK-3β (**C**), Dkk1 (**D**), TCF4 (**E**), Fn1 (**F**), and IgG negative control (**G**). **Abbreviations:** Nx, 5/6 nephrectomy; GSK-3β, glycogen synthase kinase-3β; Dkk1, dickkopf 1; TCF4, transcription factor 4; Fn1, fibronectin 1.

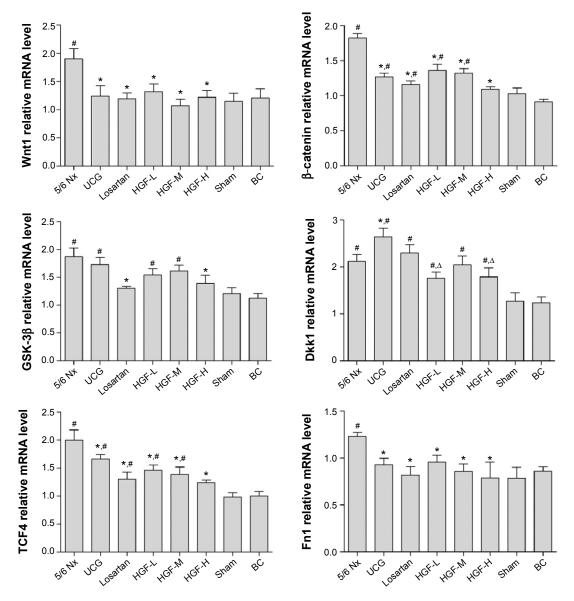


Figure 5 Wnt1, β-catenin, TCF4, GSK-3β, Dkk1, and Fn1 mRNA expression in renal tissues (n=6). **Notes:** Compared with the 5/6 Nx group, *P<0.05; compared with Sham and BC groups, *P<0.05; and compared with the losartan group, *P<0.05. **Abbreviations:** TCF4, transcription factor 4; GSK-3β, glycogen synthase kinase-3β; Dkk1, dickkopf 1; Fn1, fibronectin 1; Nx, 5/6 nephrectomy; Sham, sham-operated group; BC, background control group; UCG, uremic clearance granule; HGF-L, Huang Gan formula low dose group; HGF-M, Huang Gan formula middle dose group; HGF-H, Huang Gan formula high dose group.

remnant renal function and alleviated glomerulosclerosis and tubulointerstitial fibrosis showing similar efficiency with UCG. The level of BUN in HGF groups was significantly lower than losartan and the Ccr was much higher than that, which demonstrated that the HGF showed better renal protective effect than losartan. But losartan was the best in decreasing the UPr, which had been confirmed in the published literature. BC group, which has no excision in any part of renal, was given HGF-M. The goals while setting this negative control group were first to evaluate its toxic effects and second to exclude the inflammation, stress reaction, or renal damage induced by HGF self, which may lead to the false negative results. The research results showed that the biochemical and pathological indicators of the BC group had no significant difference with the sham group, which fitted the expected goal and demonstrated that HGF had no obvious toxic effect on the experimental rats.

TCM formulas often contain multiple medicinal herbs that lead to the action on multiple targets and thereby produce an overall systemic effect.²⁰ TCM based on unique theories and long-term practical experiences is very popular in People's Republic of China, especially in delaying renal disease progression and protecting renal function. It has become an indispensable means of treatment of CKD.²¹

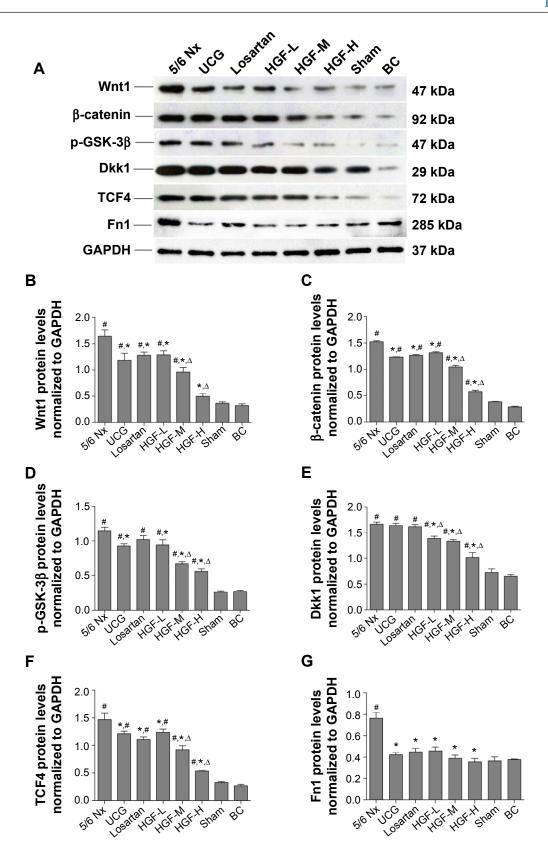


Figure 6 Western blot analysis for detection of Wn1, β-catenin, TCF4, GSK-3β, Dkk1, and Fn1 expression in the kidney.

Notes: Western blot bands of different proteins (**A**); semi-quantitative analysis of the bands (from (**B**) to (**G**) are Wnt1, β -catenin, GSK-3 β , Dkk1, TCF4, and Fn1 in sequence). Compared with 5/6 Nx group, *P<0.05; compared with Sham and BC groups, #P<0.05; and compared with the losartan group AP <0.05.

Abbreviations: TCF4, transcription factor 4; GSK-3β, glycogen synthase kinase-3β; Dkk1, dickkopf 1; Fn1, fibronectin 1; Nx, 5/6 nephrectomy; Sham, sham-operated group; BC, background control group; UCG, uremic clearance granule; HGF-L, Huang Gan formula low dose group; HGF-M, Huang Gan formula middle dose group; HGF-H, Huang Gan formula high dose group; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

Many studies have reported the prescriptions of a single herb, decoctions of multiple herbs, or patent medicines that have been used to treat patients with CKD, which exhibit effects including promotion of diuresis, reduction of proteinuria, and improvement of renal function.²² The results of this study indicating the components and content of HGF are stable and can be controlled, and the main components are aloe-emodin, rhein, and chrysophanol extracted from rheum. The efficacy of most of these constituents has been verified in vitro and vivo. Similar to emodin, it can reduce glucoseinduced ECM synthesis in human peritoneal mesothelial cells.23 Rhein therapy markedly ameliorated renal interstitial fibrotic lesions, and the primary target was TGF-B1.²⁴ In this study, we also proved that HGF could decrease the level of Scr, BUN and UPr and ameliorate tubulointerstitial fibrosis in CKD rats, which may act through the Wnt/ β -catenin signaling pathway.

The Wnt/ β -catenin signaling pathway is the canonical Wnt signal pathway and was recently discovered and plays an important role in mediating the renal fibrosis.^{7,8} In unilateral ureteral obstruction rat models, all members of the Wnt family except Wnt5b, Wnt8b, and Wnt9b were upregulated in the fibrotic kidney with distinct dynamics. Numerous Wnt/ β-catenin target genes (c-Myc, Twist, lymphoid enhancerbinding factor 1, and fibronectin) were induced.⁷ Oppositely, when administered Dkk1, a Wnt-pathway antagonist, could significantly reduce free β -catenin in cytoplasm and inhibit the downstream target gene transcription. ICG-001, a specific blocking β -catenin-mediated gene transduction, successfully blocked the expression of PAI-1, Col I, Fn, and α -SMA induced by TGF-β1 in HKC-8 cells. In vivo study, ICG-001 could improve the renal fibrosis, inhibit overexpression of fiber-related protein, and reduce kidney damage.25

In this study, the Wnt/ β -catenin pathway was activated and Wnt1 was highly upregulated in 5/6 Nx models. Dkk1 as a upstream antagonist could bind to LRP5/6 receptor and induce LRP rapid endocytosis in participation of transmembrane protein Kremen1/2, which eventually inhibited the Wnt signal transduction into cytoplasm.²⁶ The activity of Dkk1 was significantly high in the 5/6 Nx group and in the HGF group was lower than it, which indicated that Dkk1 may possess a feedback regulation. The same characteristic was proved by using unilateral ureteral obstruction rat model, in which the expression of all the DKK family members increased remarkably.^{7,27} As the Wnt signal transduction into cell, β -catenin phosphorylation was then inhibited. So the level of β -catenin in the 5/6 Nx group was high, and HGF could regulate it down. So the downstream target TCF4 was also inhibited in the HGF group.^{27,28} The Fn1, which was the main component of ECM and the target protein of Wnt pathway, was also decreased in the remnant kidney in HGF groups, which demonstrated that HGF could improve renal fibrosis by inhibiting the Wnt/ β -catenin pathway.

Wnt signaling pathway is silent within normal mature kidneys.²⁹ However, a small amount of free β-catenin exists in the cytoplasm, and most of them combine with E-cadherin to form complexes in cell membrane, playing a crucial role in preventing cell migration and maintaining the same type of cell adhesion.³⁰ While the Wnt pathway is activated, the degradation of free β -catenin is frustrated, leading to aggregation within the nucleus causing complexes to form with TCF families of LEF1/TCF, eventually accelerating the progression of epithelial-mesenchymal transition.³¹ Epithelialmesenchymal transition was the direct contributor to the kidney myofibroblast population and renal fibrosis.^{32,33} The GSK-3 β mainly degrades β -catenin by forming degradation complex with adenomatous polyposis coli and Axin. It will be upregulated while the Wnt pathway is activated,^{34,35} which was also demonstrated in this study. However, HGF could not stimulate the expression of these two antagonists. Recent studies have shown that blockade of Wnt/ β -catenin signaling can simultaneously repress multiple renin-angiotensin system genes, which play an essential role in the pathogenesis of CKD and cardiovascular disease,36 and can also ameliorate proteinuria and kidney injury.³⁷ So Wnt/β-catenin signaling pathway becomes more and more important in the progress of CKD. Therefore, it can be used as the new therapeutic target for CKD.

Conclusion

In summary, the Wnt/ β -catenin signaling pathway has become more significant and may be a potential novel target for therapeutic intervention of CKD. The present study demonstrated that HGF improved remnant renal function and alleviated glomerulosclerosis and tubulointerstitial fibrosis potentially by depressing the Wnt/ β -catenin signaling pathway. In future studies, the spectrum–effect relationship of HGF will be identified, and the mechanism will be defined by gene silencing of Wnt/ β -catenin signaling pathway in vitro.

Acknowledgments

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Disclosure

The authors declare no conflicts of interest in this work.

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Supplementary materials

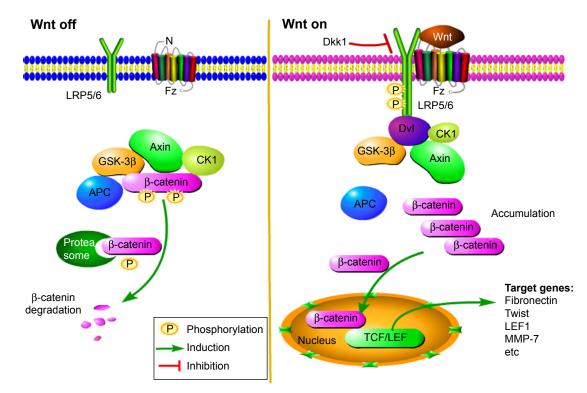


Figure SI Simplified schematic of the canonical Wnt/β -catenin signaling pathway.

Notes: When Wnt is off, β -catenin is phosphorylated by a complex composed of APC, Axin, GSK-3 β , and CK1, which lead to the degradation of β -catenin via the proteasome pathway; When Wnt is on, Wnt ligands bind to and activate Fz and LRP5/6 and induce the phosphorylation of LRP6, which recruits Axin through Dvl to the plasma membrane resulting in the inhibition of β -catenin phosphorylation and degradation. Accumulated β -catenin translocates to the nucleus and complexes with the TCF/LEF, moderating canonical Wnt-target gene transcription.

Abbreviations: APC, adenomatous polyposis coli; GSK-3β, glycogen synthase kinase-3β; CKI, casein kinase I; Fz, frizzled; LRP5/6, low density lipoprotein receptorrelated protein 5 or 6; Dvl, disheveled; TCF/LEF, transcription factor/lymphoid enhancer factor; DkkI, dickkopf I; LEFI, lymphoid enhancer factor I; MMP-7, matrix metalloproteinase 7.

Gene ID	Gene symbol	Sequence (5'-3')	Size of PCR products (bp)	Amplificatior efficiency
24881	Wntl	Forward primer: CTTCGGCAAGATCGTCAACC	51	101.05
		Reverse primer: GCGAAGATGAACGCTGTTTCT		
84353	β -catenin (Ctnnb1)	Forward primer: TGAAGGTGCTGTCTGTCTGCTC	143	100.8
	,	Reverse primer: TGCATCGGACAAGTTTCTCAGA		
84382	TCF4	Forward primer: CCAGTGGAATGAATCAGCCC	190	98.9
		Reverse primer: GAGGTGCTGTAATGGTTTGTGC		
84027	GSK-3β	Forward primer: ATCTTTGGAGCCACCGATTACA	160	98.6
		Reverse primer: TTTGCTCCCTTGTTGGTGTTC		
293897	Dkkl	Forward primer: CGAGGAAGGCATCATTGAAAAC	143	100.3
		Reverse primer: CTGTGGCGCAGTCTGATGATC		
25661	Fnl	Forward primer: TGACAACTGCCGTAGACCTGG	72	99.7
		Reverse primer: TACTGGTTGTAGGTGTGGCCG		
81822	β-actin (Actb)	Forward primer: GGGAAATCGTGCGTGACATT	76	105.7
	/	Reverse primer: GCGGCAGTGGCCATCTC		

Table SI Primers used for RT-PCR a	assay
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Abbreviations: RT-PCR, reverse transcriptase-polymerase chain reaction; TCF4, transcription factor 4; GSK-3β, glycogen synthase kinase-3β; Dkk1, dickkopf 1; Fn1, fibronectin 1.

Table S2 Relative retention time of the common peaks of ten batches of HGF extracts

	Peak I	Peak 2	Peak 3	Peak 4	Peak 5	Peak 6	Peak 7	Peak 8	Peak 9	Peak 10
SI	0.1370	0.1483	0.1691	0.1820	0.1915	0.2592	0.2683	0.3269	0.3731	0.3996
S2	0.1376	0.1485	0.1689	0.1852	0.2004	0.2604	0.2690	0.3314	0.3740	0.4011
S3	0.1379	0.1486	0.1692	0.1825	0.1997	0.2604	0.2692	0.3307	0.3745	0.4015
S4	0.1377	0.1482	0.1691	0.1823	0.1997	0.2605	0.2694	0.3303	0.3744	0.4015
S5	0.1378	0.1484	0.1692	0.1828	0.1998	0.2602	0.2693	0.3296	0.3745	0.4017
S6	0.1380	0.1484	0.1693	0.1824	0.1997	0.2604	0.2695	0.3291	0.3749	0.4020
S7	0.1380	0.1483	0.1693	0.1823	0.1917	0.2602	0.2695	0.3279	0.3749	0.4020
S8	0.1380	0.1483	0.1694	0.1820	0.1915	0.2603	0.2698	0.3268	0.3753	0.4024
S9	0.1378	0.1481	0.1694	0.1821	0.1908	0.2599	0.2695	0.3250	0.3749	0.4021
S10	0.1376	0.1484	0.1692	0.1778	0.1906	0.2575	0.2669	0.3235	0.3737	0.3970

Abbreviation: HGF, Huang Gan formula.

Table S3 Relative areas of the common peaks of ten batches of extracts of HGF

	Peak I	Peak 2	Peak 3	Peak 4	Peak 5	Peak 6	Peak 7	Peak 8	Peak 9	Peak 10
SI	0.14	0.17	0.18	0.45	0.06	0.03	0.05	0.07	0.10	0.20
S2	0.19	0.28	0.17	0.43	0.17	0.05	0.05	0.07	0.11	0.22
S3	0.18	0.23	0.20	0.44	0.10	0.05	0.06	0.08	0.12	0.23
S4	0.17	0.17	0.18	0.51	0.12	0.03	0.04	0.07	0.10	0.20
S5	0.20	0.20	0.24	0.46	0.12	0.04	0.06	0.09	0.13	0.26
S6	0.21	0.20	0.26	0.59	0.07	0.04	0.06	0.07	0.11	0.23
S7	0.13	0.12	0.15	0.38	0.12	0.03	0.04	0.07	0.08	0.19
S8	0.22	0.17	0.21	0.48	0.14	0.03	0.05	0.07	0.09	0.21
S9	0.19	0.18	0.22	0.54	0.07	0.04	0.06	0.08	0.12	0.26
S10	0.17	0.18	0.27	0.40	0.07	0.03	0.07	0.06	0.09	0.21

Abbreviation: HGF, Huang Gan formula.

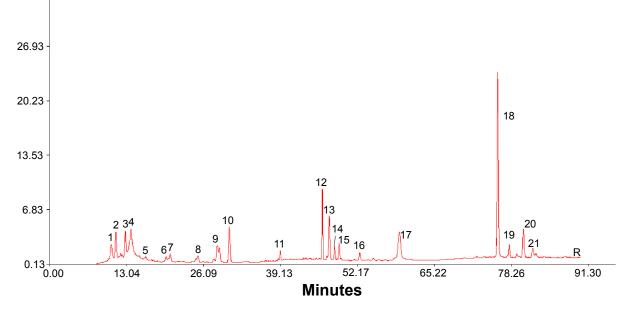


Figure S2 The reference fingerprint of HGF.

Notes: The reference fingerprint of HGF, which is produced according to HPLC chromatograms of the ten batches of HGF extracts by Similarity Evaluation System for Chromatographic Fingerprint of Traditional Chinese Medicine (Version 2004A). A total of 21 common peaks are marked, and the maximum peak 18 is set as the reference chromatographic peak, while the retention time and peak area are set to 1. The relative retention time and relative peak area of common peaks compared to peak 18 are calculated. Fingerprint results showed that the ten batches of HGF possessed well consistency and the extraction method was stable. HPLC fingerprint can better control the overall quality of HGF extract.

Abbreviations: HGF, Huang Gan formula; HPLC, high-performance liquid chromatography.

Peak I I	Peak 12	Peak 13	Peak 14	Peak 15	Peak 16	Peak 17	Peak 18	Peak 19	Peak 20	Peak 21
0.5119	0.6085	0.6232	0.6353	0.6451	0.6898	0.7771	1.0000	1.0259	1.0577	1.0781
0.5143	0.6090	0.6238	0.6364	0.6462	0.6919	0.7831	1.0000	1.0257	1.0575	1.0781
0.5144	0.6089	0.6239	0.6361	0.6460	0.6917	0.7826	1.0000	1.0256	1.0574	1.0780
0.5146	0.6089	0.6239	0.6361	0.6461	0.6917	0.7818	1.0000	1.0256	1.0574	1.0780
0.5145	0.6091	0.6240	0.6363	0.6463	0.6919	0.7806	1.0000	1.0256	1.0575	1.0782
0.5146	0.6091	0.6240	0.6362	0.6462	0.6918	0.7797	1.0000	1.0256	1.0575	1.0782
0.5147	0.6090	0.6241	0.6362	0.6462	0.6918	0.7788	1.0000	1.0257	1.0576	1.0782
0.5150	0.6090	0.6240	0.6363	0.6463	0.6920	0.7775	1.0000	1.0256	1.0576	1.0782
0.5148	0.6091	0.6241	0.6362	0.6462	0.6918	0.7757	1.0000	1.0256	1.0575	1.0783
0.5123	0.6071	0.6224	0.6332	0.6428	0.6865	0.7759	1.0000	1.0262	1.0579	1.0788

Peak I I	Peak 12	Peak 13	Peak 14	Peak 15	Peak 16	Peak 17	Peak 18	Peak 19	Peak 20	Peak 21
0.05	0.19	0.24	0.05	0.07	0.04	0.35	1.00	0.06	0.19	0.07
0.04	0.24	0.18	0.08	0.07	0.05	0.29	1.00	0.07	0.13	0.06
0.05	0.34	0.25	0.10	0.12	0.07	0.40	1.00	0.07	0.19	0.05
0.05	0.19	0.24	0.05	0.07	0.04	0.40	1.00	0.05	0.19	0.05
0.05	0.20	0.19	0.06	0.07	0.04	0.27	1.00	0.08	0.12	0.08
0.05	0.21	0.25	0.06	0.08	0.04	0.36	1.00	0.05	0.19	0.07
0.03	0.13	0.20	0.03	0.05	0.03	0.38	1.00	0.03	0.18	0.05
0.04	0.17	0.23	0.05	0.06	0.03	0.36	1.00	0.07	0.18	0.06
0.05	0.19	0.25	0.06	0.07	0.04	0.39	1.00	0.09	0.18	0.07
0.04	0.18	0.20	0.05	0.06	0.03	0.29	1.00	0.07	0.13	0.03

Table S4 The MS qualitative analysis results of HGF (the mass spectrometry data of main eight ingredients)

Peak number	Retention time (min)	Compound	Molecular formula	Molecular weight (MW)	Measured value	Error value (ppm)	Matching score	Adduction
I	10.633	Liquiritin	C ₂₁ H ₂₂ O ₉	418.3940	417.1185	-1.44	91.84	[M–H]–
2	12.758	Tectoridin	C ₂₂ H ₂₂ O ₁₁	462.4035	461.1089	0.00	96.79	[M–H]–
3	39.25	Luteolin	C ₁₅ H ₁₀ O ₆	286.23	285.0402	-1.05	89.06	[M–H]–
4	44.243	Aloe-emodin	C ₁₅ H ₁₀ O ₅	270.23	271.3651	2.41	87.80	[M+H]+
5	45.598	Rhein	C ₁₅ H ₈ O ₆	284.22	283.0246	-0.7 I	89.20	[M–H]–
6	64.0553	Emodin	C ₁₅ H ₁₀ O ₅	270.23	269.0447	-2.97	92.08	[M–H]–
7	74.158	Chrysophanol	C ₁₅ H ₁₀ O ₄	254.23	253.0498	-3.16	90.23	[M–H]–
8	78.692	Physcion	C ₁₆ H ₁₂ O ₅	284.27	283.0606	-2.12	82.12	[M-H]-

Abbreviations: MS, mass spectrometry; HGF, Huang Gan formula.

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