Yi-Qi-Jian-Pi Formula Suppresses RIPK1/RIPK3-Complex-Dependent Necroptosis of Hepatocytes Through ROS Signaling and Attenuates Liver Injury in Vivo and in Vitro

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Acute-on-chronic liver failure (ACLF) is described as a characteristic of acute jaundice and coagulation dysfunction. Effective treatments for ACLF are unavailable and hence are urgently required. We aimed to define the effect of Yi-Qi-Jian-Pi Formula (YQJPF) on liver injury and further examine the molecular mechanisms. In this study, we established CCl4-, LPS-, and D-galactosamine (D-Gal)-induced ACLF rat models in vivo and LPS- and D-Gal-induced hepatocyte injury models in vitro. We found that YQJPF significantly ameliorates liver injury in vivo and in vitro that is associated with the regulation of hepatocyte necroptosis. Specifically, YQJPF decreased expression of receptor-interacting protein kinase 1 (RIPK1), receptor-interacting protein kinase 3 (RIPK3) and pseudokinase mixed lineage kinase domain-like (MLKL) to inhibit the migration of RIPK1 and RIPK3 into necrosome. YQJPF also reduces the expression of inflammatory cytokines IL-6, IL-8, IL-1β, and TNF-α, which were regulated by RIPK3 mediates cell death. RIPK1 depletion was found to enhance the protective effect of YQJPF. Furthermore, we showed that YQJPF significantly downregulates the mitochondrial reactive oxygen species (ROS) production and mitochondrial depolarization, with ROS scavenger, 4-hydroxy-TEMPO treatment recovering impaired RIPK1-mediated necroptosis and reducing the expression of IL-6, IL-8, IL-1β, and TNF-α. In summary, our study revealed the molecular mechanism of protective effect of YQJPF on hepatocyte necroptosis,

Abbreviations: ACLF, Acute-on-chronic liver failure; ALP, Alkaline phosphatase; ALT, Alanine aminotransferase; AST, Aspartate aminotransferase; D-Gal, D-galactosamine; ELISA, Enzyme-linked immunosorbent assay, Glb, Globulin, H&E, Hematoxylin and Eosin, HSCs, Hepatic stellate cells, IL-18, Interleukin-18, IL-1β, Interleukin-1β, IL-6, Interleukin-6, LPS, Lipopolysaccharide, MDA, Malondialdehyde, MLKL, Mixed lineage kinase domain-like protein, PBS, Phosphate-buffered saline, RIPK1, Receptor-interacting protein kinases 1, RIPK3, Receptor-interacting protein kinases 3, RT-qPCR, Real-time quantitative polymerase chain reaction, SOD, Superoxide dismutase, TBIL, Total bilirubin, TNF-α, Tumor Necrosis Factor-α, YQJPF, Yiqijianpi Formula, ROS, Reactive oxygen species, Tempol, 4-hydroxy-TEMPO, HBV, Hepatitis B virus, HCV, Hepatitis C virus, Nec-1, Necrotatin, DMEM, Dulbecco’s modified essential medium, FBS, Fetal bovine serum, GAPDH, Glyceraldehyde phosphate dehydrogenase, Co-IP, Co-immunoprecipitation, DCHF-DA, 2′,7′-dichlorodihydrofluorescein diacetate, JC-1, 5.5′,6.6′-tetrachloro-1′,3′,3′-tetraethylbenzimidazolylcarbocyanine iodide, TEM, Transmission electron microscopy, MMP, Mitochondrial membrane potential, Sou-Medrol, Methylprednisolone
INTRODUCTION

Acute-on-chronic liver failure (ACLF) is a syndrome characterized by acute decompensation of chronic liver disease, which has a high short-term mortality rate, and it seriously threatens the lives of patients with chronic liver disease (Xiang et al., 2020). However, the pathogenesis of ACLF remains poorly understood. Among Asian countries, the prevalence of hepatitis in China is still high. There are approximately 97 million hepatitis B virus (HBV) carriers and one million hepatitis C virus (HCV)-infected patients in China. Chronic viral hepatitis is still the basis in ACLF (Xie et al., 2019). Virus replication and bacterial infection are the main causes of liver failure. However, no effective therapy for liver failure beyond supportive treatment is currently available; hence, a new treatment or medicine against liver failure is needed. Previous studies have suggested that ACLF develops in patients with cirrhosis as a consequence of precipitating events leading to acute decompensation and multi-organ failure (Sarin and Choudhury, 2016). Despite variations in the definition of ACLF in different regions, progressive, unrelenting hepatocyte injury and death are common hallmarks of ACLF, which have been well-documented as pathobiology of ACLF. Effective treatments to target hepatocyte injury and death due to ACLF are lacking and thus are urgently required. Unlike apoptosis and pyroptosis, necroptosis is a caspase-independent death program (Asrani et al., 2019). The dying cells exhibit none of the morphological characteristics of apoptosis, instead these cells display the swelling associated with necrosis; the dying cell can trigger an innate immune response, cause the release of inflammatory cytokines, and exacerbate tissue damage (Wang et al., 2014). The inhibitors of necroptosis could have therapeutic potential.

Traditional Chinese medicines have received attention worldwide in last few decades because of their satisfactory therapeutic effects, facile availability, and cost-effectiveness (Bajaj et al., 2018; Engelmann et al., 2020). Our previous study confirmed that the treatment with Yi-Qi-Jian-Pi Formula (YQJPF) composed of Huangqi, Taizishen, Baizhu, Chenpi, Danggui, Fulin, Huangqin, and Licorice can ameliorate HBV-ACLF. Corethrodendron multijugum (Maxim.) B. H. Choi and H. Ohashi (Huangqi in Chinese), Atractylodes japonica (Oliv.) Diels (Danggui in Chinese), Angelica sinensis (Oliv.) Diels (Fulin in Chinese), Angelicae Sinensis Radix, Pseudostellaria heterophylla (Miq.) Pax (Taizishen in Chinese) are Monarch drugs; Angelicae Sinensis Radix, Pseudostellaria heterophylla (Miq.) Pax (Taizishen in Chinese), Scutellariae Radix Scutellariae baicalensis Georgi (Huangqin in Chinese), and licorice, Glycyrrhiza uralensis Fisch. ex DC., G. glabra L., and G. inflata Batalin (Gancao in Chinese) are adjuvants. A retrospective cohort study on 60 patients with HBV-ACLF reported that YQJPF could prevent the development of liver failure and improve the Model for End-Stage Liver disease score (Zhang Z. et al., 2014; Lian et al., 2015). However, the regulation mechanism of its effect has not been elucidated. Atractyloolone is the main sesquiterpene constituent of Atractylodes japonica, which is a necessary component of YQJPF. Atractylodes japonica was used to treat several diseases such as rheumatic diseases, digestive disorders, hepatic protection and influenza for a long history in China. Additionally, atractyloolone was reported to have anti-inflammatory and anti-hepatotoxic effects (Cheng et al., 2019).

In the present study, we developed in vivo and in vitro models with LPS injection that recapitulate some features of clinical ACLF. In these models, we assessed the features of ACLF in laboratory settings. Further, ACLF models were treated with YQJPF to assess its effect on liver injury. Additionally, we explored the role of inflammation infiltration and hepatocyte necroptosis and the potential mechanisms for the treatment of liver injury in ACLF.

MATERIALS AND METHODS

Reagents and Antibodies

Methylprednisolone was obtained from the Nanjing Hospital affiliated to Nanjing University of Chinese Medicine (Nanjing, Jiangsu China) and was dissolved in PBS to a concentration of 30 mg/ml and was stored at 4°C. Necrotatin-1 (Nec-1) and 4-hydroxy-TEMPO (Tempol) were bought from Sigma-Aldrich (St Louis, MO, United States). Dulbecco’s modified essential medium (DMEM), fetal bovine serum (FBS), Opti-MEM medium, phosphate-buffered saline (PBS), and trypsin-EDTA were purchased from GIBCO (Grand Island, NY, United States). The antibodies against β-actin (20536-1-AP), RIPK1 (17519-1-AP), and RIPK3 (17563-1-AP) were purchased from Proteintech Group, Inc (Rosemont, IL, United States). Caspase-3 (#9662), cleaved-caspase-3 (#9664), caspase-8 (#9746), cleaved-caspase-8 (#9748), TNF-α (#11948), IL-18 (#54943), IL-1β (#12242), and IL-6 (#12153) were procured from Cell Signaling Technology (Danvers, MA, United States). Mixed lineage kinase domain-like protein (MLKL; ab243142) were purchased from Abcam Technology (Abcam, Cambridge, United Kingdom).

Preparation and High-Performance Liquid Chromatography of YQJPF for Quality Control

YQJPF was provided by Nanjing University of Chinese Medicine, Nanjing Hospital Affiliated to Nanjing University of Chinese Medicine. YQJPF was prepared from nine commonly used Chinese herbal medicines (Table 1). All the herbal targeting RIPK1/RIPK3-complex-dependent necroptosis via ROS signaling. Overall, our results provided a novel perspective to indicate the positive role of YQJPF in ACLF.

Keywords: acute-on-chronic liver failure, reactive oxygen species, inflammation, hepatocytes necroptosis, Yi-Qi-Jian-Pi formula
constituents were obtained from the Beijing Tong Ren Tang Co. Ltd. (Beijing, China). Huangqi, Taizishen, Baizhu, Chenpi, Danggui, Fulin, Huangqin, and Gancao were mixed in the ratio 30:30:30:10:15:15:3:10:10.

Then, all herbs were decocted twice and the decoction liquids were concentrated to a density of 2.86 g/ml and was stored at 4°C. For quality control, HPLC analysis was used as previously described (Yao et al., 2018). YQJPF were assessed using an Agilent 1,260 liquid chromatography system (United Kingdom). Briefly, 10 μl YQJPF solution was injected into an apparatus with an auto sampler. Chromatographic separation was implemented at a flow rate of 1 ml/min with an Agilent C18 column (4.6 mm × 250 mm, 5 μm). The separation phase was composed of 0.2% phosphoric acid (A) and acetonitrile (B). The linear concentration solution gradually increases from 2 to 72% of solvent B over the course of 45 min. The separation temperature was 40°C, with a detection wavelength of 254 nm. The results are shown in Figure 1.

**Animals and Experimental Design**

A total of 56 male Sprague-Dawley rats, with the body weight of 180–220 g, were purchased from Beijing Vital River Laboratory Animal Technology Co. Ltd (Beijing, China). All animals were housed with laboratory chow and tap water at 22 ± 2°C under 55 ± 5% humidity-controlled environment and a 12-h light–dark cycle. The experimental design was approved by the Institutional and Local Committee on the Care and Use of Animals of Nanjing University of Chinese Medicine (Nanjing, China). All animals were provided care in compliance with the National Institutes of Health (United States) guidelines. A mixture of CCl4 and olive oil [1:1 (w/v)] was used to induce liver cirrhosis in rats (1.5 ml/kg body weight). LPS (100 μg/kg) and D-Gal (400 mg/kg) were
used to induce acute liver injury. The mice were randomly divided into seven groups ($n = 8$ for each group). The experimental groups were as follows (Figure 2):

Group 1: control group; i. p. with olive oil.
Group 2: chronic liver injury model group; i. p. with CCl$_4$ for 12 weeks.
Group 3: acute liver injury model group; i. p. with LPS and D-Gal once.
Group 4: ACLF model group; i. p. with CCl$_4$ for 12 weeks followed by i. p. with LPS and D-Gal once.
Group 5: YQJPF low-dose treatment group; i. p. with CCl$_4$ for 12 weeks and i. g. with YQJPF (14.3 g/kg) for 2 weeks followed by i. p. with LPS and D-Gal once.
Group 6: YQJPF high-dose treatment group; i. p. with CCl$_4$ for 12 weeks and i. g. with YQJPF (28.6 g/kg) for 2 weeks followed by i. p. with LPS and D-Gal once.
Group 7: methylprednisolone treatment group; i. p. with CCl$_4$ for 12 weeks and i. p. with methylprednisolone (15 mg/kg) for 2 weeks followed by i. p. with LPS and D-Gal once.

**Cell Culture**

Human LO2 cell line was obtained from the Cell Bank of Chinese Academy of Sciences (Shanghai, China). Cells were cultured in DMEM with 10% FBS and 1% antibiotics and were incubated in 5% CO$_2$ and 95% air humidified atmosphere at 37°C.

**Liver Histopathology**

Liver hematoxylin and eosin (H&E) and Masson and Sirius Red staining were performed as per the previously described methods (Jin et al., 2016). Representative pictures of liver sections were displayed.

**Serum Biochemistry**

Blood samples were collected and incubated at room temperature for 1 h, and extracted serum was collected after centrifugation. Levels of alkaline phosphatase (ALP), aspartate aminotransferase (AST), and alanine aminotransferase (ALT) in serum were measured using enzyme-linked immunosorbent assay kits (Nanjing Jinting Bioengineering Institute, Nanjing, China) according to the manufacturer’s protocols.

**Determination of Hepatic mtDNA, ATP, Malondialdehyde, and Superoxide Dismutase Levels**

Liver tissues were treated according to the treatment regimen mentioned earlier. After treatment, hepatic mtDNA, ATP, malondialdehyde (MDA), and superoxide dismutase (SOD) were quantified using corresponding kits (Nanjing Jinting Bioengineering Institute, Nanjing, China) according to the manufacturer’s instructions.

**Immunohistochemistry**

Immunohistochemical staining of liver tissue was performed using antibodies against caspase 8, MLKL, RIPK1, and RIPK3, as previously described (Zhang F. et al., 2014). Representative pictures of liver sections were displayed.

**Cell Viability Assay**

To assess the effects of different treatments on cell proliferation, Cell viability analysis was performed using MTT. In brief, The LO2 cells were seeded into 96-well plates at a density of $3 \times 10^4$ cells per well. After incubating for 24 h, the cells were treated with LPS (10 μg/ml) and D-Gal (10 mg/ml) for 4 h for the hepatocyte injury model. Further, the cells were treated with serial concentration gradients of YQJPF and atractylone for another 24 h, with triplicates for each concentration. Then 5 mg/ml MTT was added in cells, and cells were incubated at 37°C for 4 h, and the absorbance was measured at a wavelength of 490 nm. Independent experiments were performed in triplicates.

**Trypan Blue Staining**

Cell death was determined using the trypan blue staining assay (Beyotime Biotechnology, Shanghai, China). In brief, after YQJPF treatment, the cells were digested using trypsin without EDTA...
followed by centrifugation. The cells were further washed twice with PBS and were resuspended in PBS. Cell suspension and trypan blue solution were mixed gently in 1:1 ratio and allowed to stand for 3 min. A few of the stained cells were counted with a hemocytometer, and photographs were taken in a blinded fashion at random fields. Stained blue cells represented dead cells.

RNA Isolation and Real-Time PCR
Total RNA was extracted from liver tissues and LO2 cells by using Trizol reagent according to the manufacturer’s protocol (Sigma-Aldrich, St. Louis, MO, United States) and further reverse-transcribed into cDNA by using PrimeScript RT reagent kit (TaKaRa Biotechnology, Beijing, China). Real-time PCR was performed using the SYBR Green I (TaKaRa Biotechnology, Beijing, China). Real-time PCR was performed using the SYBR Green I fluorescent dye (TaKaRa Biotechnology, Beijing, China), according to the manufacturer’s protocol. Glyceraldehyde phosphate dehydrogenase (GAPDH) served as an invariant control, and mRNA levels were expressed as fold changes after normalizing to GAPDH. The experiment was performed in triplicates. Primers (Genscript, Nanjing, China) are listed in Table 2 below.

### Western Blot

Cells and liver samples were lysed using a mammalian lysis buffer (Sigma St. Louis, MO, United States) and were denatured in SDS loading buffer. Then, western blot analysis was performed according to the manufacturer’s instructions (Bio-Rad, Hercules, CA, United States). Thereafter, Cells and liver samples were incubated overnight with antibodies against caspase 3, cleaved-caspase-3, caspase 8, cleaved-caspase-8, MLKL, RIPK1, RIPK3, IL-6, IL-1β, and TNF-α. Protein detection, band visualization, and quantitation were performed as per the manufacturer’s instructions. Anti-β-actin antibody (Danvers, MA, United States) was used as a loading control.

### Immunoprecipitation Assay

The immunoprecipitation assay was performed using co-immunoprecipitation (Co-IP) kit (Thermo) as per manufacturer’s instructions. Briefly, the cell lysates were incubated with anti-RIPK1 and anti-RIPK3 antibodies. Protein G-agarose beads were added and incubated overnight to precipitate the protein complexes. Western blot analysis was performed to detect the protein complexes expression.

### Immunofluorescence Analysis

LO2 cells were seeded on 24-well plates and cultured in DMEM with 10% FBS. Afterward, they were treated with corresponding reagents for 24 h. Then, the cells were fixed with 4% PFA for 30 min at 37°C, permeabilized with PBS-T (0.1% Triton x-100 dissolved in PBS), and blocked with PBS-B (4% BSA dissolved in PBS). The cells were stained with the corresponding antibody (1:200 dilution) overnight at 4°C, followed by incubation with FITC-labelled goat anti-rabbit IgG (1:100 dilution) for 2 h. Finally, 4′,6-diamidino-2-phenylindole (DAPI) staining was performed by incubation in dark for 5 min, and the fluorescence was observed using a fluorescence microscope (Nikon, Tokyo, Japan) to visualize the nuclei.

### Enzyme-Linked Immunosorbent Assay (ELISA)

Levels of IL-6, IL-8, IL-1β, and TNF-α in serum were measured using ELISA kits (Nanjing Jiancheng Bioengineering Institute, Nanjing, China) according to the manufacturer’s instructions.

### Intracellular ROS Assay

The level of intracellular ROS was determined using an oxidation-sensitive fluorescent probe, 2′,7′-dichlorodihydrofluorescein diacetate (DCFH-DA) (Beyotime Biotechnology, Shanghai, China). Briefly, LO2 cells were seeded in a 24-well plate and pre-treated with ROS scavenger Tempol (10 μM) for 1 h before exposing to YQJPF or selective RIPK1 inhibitor Nec-1. DCFH-DA probe was subsequently incubated with LO2 for 30 min. Subsequently, ROS levels were detected according to the manufacturer’s instructions.

### Determination of Mitochondrial ROS

We determined mitochondrial ROS production by using a fluorescent probe, MitoSOX, which is a red mitochondrial superoxide indicator for live cell imaging (Molecular Probes; Life technologies). Briefly, LO2 cells were treated with YQJPF or selective RIPK1 inhibitor Nec-1. Further, LO2 cells were incubated with MitoSox reagent (5 μM) for 10 min at 37°C and washed with PBS. The red fluorescence density was detected using a fluorescence microscope (Nikon, Tokyo, Japan) with the rhodamine channel.

### Mitochondrial Membrane Potential Assay

The mitochondrial membrane potential (MMP) was determined using 5,5′,6,6′-tetrachloro-1,1′-3,3′-tetraethylbenzimidazolylcarbocyanine iodide (JC-1) (Beyotime Biotechnology, Shanghai, China) staining as per the manufacturer’s protocol. LO2 cells were washed with PBS and incubated with JC-1 working solution at 37°C in dark for 20 min. After removing JC-1 solution, the cells were washed with PBS, and images were taken using a fluorescence microscope (Nikon, Tokyo, Japan) with both red and green channels. The MMP is represented by the average red/green fluorescence intensity ratio.

### Transmission Electron Microscopy

LO2 cells were seeded in six-well plates (14,000 cells/well). After the corresponding treatment, transmission electron microscopy (TEM) images were obtained using a JEM 1010 transmission electron microscope (JEOL, Tokyo, Japan). Briefly, LO2 cells were fixed in Karnovsky fixative solution (2% paraformaldehyde and 2.5% glutaraldehyde in 0.15 mol/L sodium cacodylate buffer, pH 7.1–7.3) during 1 h and then washed three times in a

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**Table 2: Primers used for human LO2 cells.**

| Gene | Forward sequence | Reverse sequence |
|------|------------------|------------------|
| IL-6 | GCTTCCTCAAGATGCTGCTGCTGCTGCT | ATTACTGCGGATGCTGCTGCT |
| TNF-α | CAGACAGATGCTGCTGCTGCTGCTGCT | TCTCAGCGCCCTCCAAGTTCC |
| IL-1β | AAGCAGTGGCAGAAGTACCTG | TGGGCTGATAGCCTGCTGCTG |
| IL-18 | CAGAGGATGCTGCTGCTGCTGCTGCT | TGGGAGTGGCCTGCTGCTG |
| GAPDH | TGTCAGCAGAAGTACCTGCTGCT | GTGCACTGCGGAGCTGCT |

**Gene Forward sequence Reverse sequence**

| Gene | Forward sequence | Reverse sequence |
|------|-----------------|-----------------|
| IL-6 | GCTTCCTCAAGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT | ATTACTGCGGATGCTGCTGCT |
| TNF-α | CAGACAGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT | TCTCAGCGCCCTCCAAGTTCC |
| IL-1β | AAGCAGTGGCAGAAGTACCTG | TGGGCTGATAGCCTGCTGCTG |
| IL-18 | CAGAGGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT | TGGGAGTGGCCTGCTGCTG |
| GAPDH | TGTCAGCAGAAGTACCTGCTGCT | GTGCACTGCGGAGCTGCT |

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Brie Biosciences) was used for detection of cell apoptosis. As indicated. Cells were further incubated for 24 h, in 5% CO2 medium and allowed to attach for 24 h. Then, cells were treated independent experiments. Values and standard deviations were calculated from three independent experiments.

Flow Cytometer Analysis
FITC Annexin V apoptosis detection kit (BD Pharmingen™, BD Biosciences) was used for detection of cell apoptosis. Briefly, LO2 cells were seeded in six-well plates in complete medium and allowed to attach for 24 h. Then, cells were treated as indicated. Cells were further incubated for 24 h, in 5% CO2 at 37°C. After that, cells were trypsinized, washed in PBS and stained according to kit according to the manufacturer’s instructions. Stained cells were analyzed using BD Accuri C6 flow cytometer (BD Pharmingen™, BD Biosciences) and data was processed with FlowJo_v10.6.2 software. Each sample was assessed using a collection of 10,000 events. The mean values and standard deviations were calculated from three independent experiments.

Statistical Analysis
Two-tailed Student’s tests and one-way ANOVA analysis were performed using GraphPad Prism software v. 7.0 (Graph Pad Software Inc., San Diego, CA). The values and data are presented as the mean ± SEM from three independent experiments. A p value of <0.05 was considered statistically significant.

RESULTS
YPJPF Alleviated Liver Injury In Vivo
The ACLF rat model was pre-established through intraperitoneal injection of CCl4 for 12 weeks. Further, acute liver injury was induced by using a combination of LPS and D-Gal. And YQJPF was given for 2 weeks to investigate the protective effect on ACLF rats. In addition, methylprednisolone (Sou-Medrol) was used as a positive control drug, which has been confirmed as a therapeutic agent for ACLF (Zhao et al., 2012; Jia et al., 2020). And the establishment of a new model of ACLF in rats and the experimental procedures as shown in Figure 2. The livers in ACLF model became small and hard accompanied by necrosis, with small nodules on their surface; however, YQJPF and methylprednisolone were found to effectively ameliorate the liver morphology. Pathological examinations were used to manifest the effects of YQJPF on hepatic injury. As shown in H&E staining, YQJPF and methylprednisolone ameliorated the disordered hepatic structure (Figure 3A). The effect of YQJPF on liver injury was evaluated. Detection of serum biochemical indicators indicated improvement in liver injury indices such as TBil, ALT, and AST contents after 2 weeks of YQJPF and methylprednisolone treatment in rats with ACLF (Figures 3B–E). In summary, YQJPF and methylprednisolone significantly ameliorated liver injury in ACLF model rats.

Protective Effect of YQJPF on LPS- and D-Gal-Induced Hepatocyte In Vitro
Generally, massive hepatocyte death is implicated in ACLF. Here, we used LPS (10 μg/ml) and D-Gal (10 mg/ml) to induce liver injury in LO2 cells. MTT assay showed that LPS and D-Gal inhibit cell viability, whereas YQJPF and atractylone could promote cell viability in a dose-dependent manner. YQJPF above 10 μg/ml and atractylone at 5 μM showed a significant effect (Figures 4A,B). Light microscopy indicated that the adherent LO2 cells are swollen, and the cell morphology changed after LPS and D-Gal treatment for 4 h. YQJPF and atractylone could improve microscopic performance (Figure 4C). Levels of AST and ALT were detected to assess the hepatocytic damage. The levels of ALT, AST, and LDH in the supernatant of cultured LO2 cells were detected by ELISA. Results showed significantly increased levels of ALT, AST, and LDH in the model group and decreased levels of those in YQJPF and atractylone treatment groups (Figures 4D–F). Thus, YQJPF and atractylone exerted a protective effect on LPS- and D-Gal-induced hepatocytic injury.

YQJPF Inhibited Necroptosis but Not Apoptosis on LPS- and D-Gal-Induced Hepatocytic Injury In Vitro
Necrosis and apoptosis are the most common pathways of death during liver injury (Schwabe and Luedde, 2018). Hence, we first examined which pathway plays a significant role in LPS- and D-Gal-induced hepatocytic injury. The result of the TUNEL assay indicated that YQJPF (10, 20, and 40 μg/ml) and atractylone (5 μM) have no effect on apoptosis (Figure 5B). The protein expression of key indicators such as caspase 3, caspase 8, and the cleaved caspase-3 and caspase-8 did not change after LPS and D-Gal treatment (Figure 5C). Then, the cell apoptosis was determined by flow cytometry, and results indicated that YQJPF and atractylone have no effect on apoptosis (Figure 5D). Therefore, LPS- and D-Gal-induced cell death was not dependent on apoptosis because no obvious apoptotic features could be observed. YQJPF and atractylone prevented cell death without influencing apoptosis. Next, we assessed whether necroptosis plays a role in this process. Positive result in trypan blue staining indicated that membrane permeability is altered by LPS and D-Gal. YQJPF or atractylone treatment significantly decreased the number of dead cells (Figure 5A). To confirm our findings, we used TEM analysis to observe cell morphology and structure. The cells exhibited a typical necrotic cell death morphology, including swelling of organelles (especially mitochondria), condensation of chromatin into small, irregular patches, and chromatin margination, after LPS and D-Gal treatment (Figure 5E). YQJPF could restore cell morphology...
and structure in a dose-dependent manner. Atractylone exhibited the same effect. Necrotic cells underwent organelle breakdown, leading to the leakage of intracellular contents consequently triggering inflammation (Weinlich et al., 2017). Further, we examined some inflammation-associated biomarkers. mRNA and protein levels of pro-inflammatory cytokines IL-1β, IL-6, TNF-α, and IL-18 were detected using RT-qPCR, ELISA, and western blot analysis (Figures 6A–I). The results showed that pro-inflammatory cytokines are decreased in YQJPF and atractylone treatment groups compared with the LPS- and D-Gal-induced model group. Above all, we demonstrated that LPS and D-Gal induce hepatocyte necroptosis but not apoptosis and that YQJPF and its active ingredient atractylone could protect hepatocytes from necroptosis by reducing the expression of inflammatory cytokines in hepatocytes.

The RIPK1/RIPK3 Complex Is Required for YQJPF to Inhibit Necroptosis In Vitro

The RIPK1/RIPK3 signaling pathway has been reported to be involved in the induction of necroptosis (Lin et al., 2016; Newton et al., 2016). Thus, we investigated using western blot and ELISA whether YQJPF and atractylone alter the expression of RIPK1, RIPK3 and MLKL. As shown in (Figures 7A–E), YQJPF or atractylone treatment downregulated the expression of RIPK1, RIPK3 and MLKL. Immunofluorescence staining of RIPK1 and RIPK3 further validated these results. The association between RIPK1 and RIPK3 was analyzed using immunoprecipitation, which indicated that YQJPF and atractylone could inhibit the interaction between RIPK1 and RIPK3 (Figure 7F). To identify whether YQJPF and atractylone are dependent on RIPK1/RIPK3-mediated necroptosis, we used the RIPK1 kinase inhibitor necrostatin-1 (Nec-1) to verify the effect of YQJPF and atractylone. The results indicated that Nec-1 and YQJPF work together, and they had the most significant inhibitory effect on LO2 necroptosis induced by LPS and D-Gal (Figures 8A–E). We further investigated the interaction between RIPK1 and RIPK3 in YQJPF-treated LO2 cells through immunoprecipitation. RIPK1 and RIPK3 were significantly immunoprecipitated in LPS- and D-Gal-induced cells, which could be impaired by YQJPF and RIPK1 inhibitor Nec-1 (Figure 8F). Overall, YQJPF could downregulate the expression of RIPK1 and RIPK3, inhibit their aggregation to form necrosome, and eventually reduce
the expression of pro-inflammatory cytokines by inhibiting RIPK1/RIPK3-mediated necroptosis.

**YQJPF Inhibited Hepatocyte Necroptosis Through Inhibition of Mitochondrial ROS Generation and Depolarization In Vitro**

Numerous studies have shown that ROS production is necessary for necroptosis in several cell lines such as macrophages (Koike et al., 2019), fibrosarcoma L929, HeLa, and human embryonic kidney (HEK) 293 T (Luedde et al., 2014; Zhang et al., 2017). To investigate whether YQJPF reduces ROS production in hepatocytes, we first analyzed the effect of YQJPF on intracellular ROS levels. Observation using fluorescence microscope revealed that YQJPF treatment decreases the ROS level in LO2 cells (Figures 9B,C). MitoSox Red staining was used to evaluate the mitochondrial ROS production. As shown in Figure 9A, YQJPF caused a significant decrease in mitochondrial superoxide, which had the same effect as selective RIPK1 inhibitor Nec-1. In addition, we found that pre-treatment of cells with YQJPF or Nec-1 significantly reversed LPS- and D-Gal-induced reduction in intracellular GSH and SOD levels. Additionally, YQJPF or Nec-1 inhibited MDA formation (Figures 9D–F). The effect of YQJPF on MMP was assessed using JC-1 staining, where a high ratio of red/green indicates an increase in MMP. The result indicated that YQJPF or Nec-1 upregulates MMP, recovers LPS- and D-Gal-induced MMP dissipation, and ameliorates mitochondrial depolarization (Figure 9G). The overproduction of ROS triggers serious damages in various cells. We observed that when the cells are pre-treated with ROS scavenger Tempol (10 μM), it could restore necroptosis-associated cell morphology and structure and inhibit

**FIGURE 4** | Protective effect of YQJPF on LPS- and D-Gal-treated hepatocytes in vitro. Human LO2 cells were treated with LPS + D-Gal for 4 h and further treated with YQJPF and atractylone at indicated concentrations. (A,B) MTT assay to study LO2 cell viability. (C) Cell morphology assessment. Scale bar, 200 μm. (D–F) ALT, AST, and LDH levels in LO2 cell culture. Data are expressed as mean ± SD (n = 6). *vs. control group, †vs. LPS + D-Gal group; *p < 0.05, †p < 0.01; *p < 0.01, ††p < 0.01.
FIGURE 5 | YQJPF inhibited hepatocytic cell death, which was not by apoptosis, in LPS- and D-Gal-treated LO2 cells in vitro. Human LO2 cells were treated with LPS + D-Gal for 4 h and further with YQJPF and atractylone. (A) Trypan blue staining to detect cell death. Scale bar, 100 μm. (B) TUNEL staining to detect cell apoptosis. (C) Western blot analysis of the protein expression of caspases three and eight and cleaved caspases three and eight in the treated human LO2 cells. (D) cell apoptosis assay by flow cytometer in the treated human LO2 cells. (E) Transmission electron microscopy (TEM) of cells. Black arrows represent swollen mitochondria (M), whereas white arrows represent condensed and marginated chromatin in nuclei (N). Data are expressed as mean ± SD (n = 3). *vs. control group, #vs. LPS + D-Gal group; *p < 0.05, **p < 0.01; *p < 0.05, **p < 0.01.
the expression of pro-inflammatory cytokines IL-1β, IL-6, TNF-α, and IL-18, as revealed by western blot (Figure 9H), which had the same effect as that of YQJPF. This suggested that YQJPF reduces the level of ROS in hepatocytes and thus plays a critical role in inhibiting cell death. Collectively, these results suggested that YQJPF inhibits necroptosis by reducing the level of mitochondrial ROS and stabilizing MMP in hepatocytes.

Disruption of Hepatocyte Necroptosis, Instead of Apoptosis, Was Associated with YQJPF Treatment In Vivo

The results of in vivo studies on the effect of YQJPF on hepatocyte necroptosis were the same as those of in vitro results. Protein levels of key caspases did not change under YQJPF treatment. Therefore, reduction in cell death due to YQJPF was independent of apoptosis. Further, we proceeded to investigate whether YQJPF could disrupt hepatocytic death by regulating necroptosis. Western blot analysis of RIPK1, RIPK3, and MLKL, which governed necroptosis, indicated that necroptosis is increased in ACLF in vivo (Figure 10E). However, it was weakened by YQJPF treatment. Furthermore, immunohistochemical staining of RIPK1, RIPK3, and MLKL showed that YQJOF reduces hepatocytic RIPK1, RIPK3, and MLKL levels, which were upregulated in chronic liver injury, acute liver injury, and ACLF model (Figure 10F). Overall, these results demonstrated that YQJPF decreases the levels of RIPK1, RIPK3, and MLKL and decreases ATP production to ameliorate hepatocyte necroptosis instead of apoptosis in ACLF model. Our previous study revealed that JNK1/2-ROS signaling is involved in HSC necroptosis (Jia et al., 2018). To investigate the role of anti-oxidant response in YQJPF-ameliorated hepatocyte necroptosis, we analyzed the effect of YQJPF on SOD and MDA levels in the liver, which are produced during lipid peroxidation. Observation using SOD and MDA detection kits showed that YQJPF treatment increases the anti-oxidant response in the liver (Figure 10A). Interestingly, the levels of ATP and mtDNA decreased significantly in chronic liver injury, acute liver injury, and ACLF model (Figure 10B).
Collectively, these results suggest that anti-oxidant response contributes to the effect of YQJPF on hepatocyte necroptosis. Similar to the in vitro results, we observed a significant increase in the serum levels of TNF-α, IL-18, IL-1β, and IL-6 in ACLF rats, but the effect was altered in a concentration dependent-manner by YQJPF treatment (Figures 10G–J). These results indicated that YQJPF attenuates hepatic inflammation and liver injury by disrupting hepatocyte necroptosis instead of apoptosis in vivo.

DISCUSSION

The pathobiology of ACLF is characterized by hepatic and systemic inflammation; progressive, unrelenting hepatocyte injury; and hepatocyte death. Hepatocyte death typically follows two patterns: necrosis and apoptosis (Vanlangenakker et al., 2008). In our study, we developed ACLF rat models by administering high dose (1.5 ml/kg body weight) of 1:1 (w/v) mixture of CCl₄ and olive oil for 3 months followed by 100 μg/kg LPS and 400 mg/kg D-Gal injection. Hepatocyte injury and cell death index were obvious in the in vivo model. Through the successful establishment of the disease model, we first confirmed the protective effect of YQJPF on ACLF liver injury. YQJPF could ameliorate CCl₄-, LPS-, and D-Gal-induced liver injury. Liver tissue morphology improved and serum levels of hepatic ALT, AST, and Tbil decreased after YQJPF treatment. The protective effect of YQJPF was consistent in the in vitro experiments. To further clarify the mechanism of action of YQJPF, an in vitro hepatocyte injury model was established using LPS (10 μg/ml) and D-Gal (10 mg/ml). YQJPF and atracytylone (the main component of YQJPF) could restore the damage to cell viability induced by LPS and D-Gal and decrease ALT, AST, and LDH levels in cell supernatant. These data clearly indicated that YQJPF confers protection against ACLF.

Liver failure is characterized by massive loss of parenchymal cells, and cell death can occur in this pathophysiological context (Schwabe and Luedde, 2018). Necroptosis is defined as the third type of programmed cell death in addition to apoptosis and...
necrosis (Galluzzi and Kroemer, 2008). It is also known as programmed necrosis or caspase-independent cell death. Necroptosis is believed to trigger a massive inflammatory response that can cause substantial collateral damage to neighboring cells. Findings of a clinical study indicated that circulating RIPK3 is significantly increased in patients with HBV-ACLF and is associated with a clinical outcome (Chen et al., 2020). However, very few studies on drugs inhibiting necroptosis in the liver are available. This study found that YQJPF could attenuate hepatocyte death but has no effect on the inhibition of caspases three and eight activity. Therefore, YQJPF-induced hepatocytic cell death inhibition did not result from apoptosis. Moreover, a typical cell necrosis morphology could be identified, and RIPK3 mediates cell death and regulates inflammatory responses (Zhao et al., 2017). Our study showed that the levels of TNF-α, IL-1β, and IL-6 were found to be significantly increased in LPS (10 μg/ml)- and D-Gal (10 mg/ml)-induced models. The effect was altered in a concentration-dependent manner after YQJPF treatment. RIPK1, RIPK3, and MLKL were identified as necroptosis regulators (Wang et al., 2012). Further, we assessed expressions of RIPK1 and RIPK3. Western blot analysis, ELISA, and immunochemical staining of RIPK1 and RIPK3 demonstrated that YQJPF decreases hepatocytes necroptosis in the ACLF model. Moreover, YQJPF inhibited the combination expression of RIPK1 and RIPK3 and blocked the formation of necrosome. As expected, YQJPF-induced inhibition of inflammatory cytokines and necrosome formation were enhanced when RIPK1 activity was inhibited by pharmacological inhibitor Nec-1. Obviously, YQJPF inhibited RIPK1/RIPK3-complex-dependent necroptosis in human LO2 cells.

Oxidative stress, a major contributor of liver injury, can be induced by excessive ROS. Studies have shown that ROS may be considerably involved in the pathogenesis of liver diseases (Chen et al., 2016). ROS have a significant effect on necroptosis, particularly the mitochondrial ROS generated in response to TNF/TNFR1 engagement. Conversely, ROS accumulation is also gradually regarded as the executioner and mediator of necroptosis (Zhu et al., 2018). ROS is mainly produced in mitochondria that in turn leads to the damage of organelles or even cell death. In this study, we found significantly decreased levels of intracellular and mitochondrial ROS after YQJPF treatment. Following pre-treatment with the ROS scavenger-Tempol, the YQJPF-induced inhibition of cell inflammatory cytokines was significantly enhanced. Thus, YQJPF could inhibit ROS accumulation. The association between RIPK1 and RIPK3 upregulation and mPTP opening has been illustrated in several studies (He et al., 2011). ROS has been reported as the primary factor that activates mPTP opening and has been implicated in the process of necroptosis. In addition, RIPK1-RIPK3 interaction could upregulate ROS levels and enhance mitochondrial superoxide production. Hence, we identified that the inhibition of RIPK1 with Nec-1 markedly enhances YQJPF-induced alteration of mitochondrial superoxide. Cellular anti-oxidant machinery mainly includes SOD, MDA, and GPx that can effectively scavenge free radicals (Riordan and Williams, 2003). In this study, SOD,
FIGURE 9 | YQJPF inhibited hepatocyte necroptosis through the inhibition of mitochondrial ROS generation and depolarization in vitro. Human LO2 cells were treated with LPS + D-Gal for 4 h and further pre-treated with 50 μM Nec-1 for 1 h, followed by YQJPF or atractylone treatment for 24 h. (A) Mitochondrial superoxide was detected and quantified through immunofluorescence by using MitoSox Red staining. Scale bar, 50 μm. (B) ROS production was assessed and quantified by DCFH-DA staining, Scale bar, 50 μm. (C–E) MDA, SOD, and GSH levels were measured by corresponding test kits. (F) ROS levels were assessed using ROS test kit. (G) Mitochondrial membrane potential was detected by JC-1 staining. Scale bar, 50 μm. (H) Western blot and quantitative analysis of the expression of IL-1β, IL-6, TNF-α, and IL-18. Data are expressed as mean ± SD (n = 3). *vs. control group, *vs. LPS + D-Gal group; †p < 0.05, ‡p < 0.01; *p < 0.05, **p < 0.01.
FIGURE 10 | Disruption of hepatocyte necroptosis, and not apoptosis, was required for the action of YQJPF in vivo. (A–D) MDA, ATP, GSH, and SOD levels in the rat livers were measured. (E) Western blot and quantitative analysis of cleaved caspase 8, MLKL, RIPK1, and RIPK3 expression in the rat livers. (F) Immunohistochemical staining and quantitative analysis of caspase 8, MLKL, RIPK1, and RIPK3 in liver tissues. Scale bar, 100 μm. (G–J) Expression levels of IL-1β, IL-6, TNF-α, and IL-18 in the rat livers were detected by ELISA. Data are expressed as mean ± SD (n = 8) * vs. control group, * vs. ACLF group; *p < 0.05, **p < 0.01; #p < 0.05, ##p < 0.01.
GSH, and MDA levels were assessed, and the results suggested that the anti-oxidant response contributes to the effect of YQJPF on hepatocyte necroptosis. Inhibition of RIPK1 by Nec-1 enhanced the effect. Additionally, these findings were confirmed in the rat models.

Traditional Chinese medicine (TCM) pattern is a physiological regulation network, which contains a regulation center, overall effective target, material basis and functional unit. Its mechanism needs to be indicated by means of TCM pattern prevention and treatment characteristics, multiple organ integration effectiveness, and functional unit network so as to guide clinical practice as well as for inspiring basic research. It is urgently to exploring the efficacy mechanism and regulation with the molecular biology and systems biology. In our research, we conducted a deep study of the pathogenesis, TCM pattern feature and efficacy evaluation of TCM in ACLF. And results showed that YQJPF attenuates liver injury in ACLF. Recently, Chen et al. reported that flavonoids have been to possess a wide variety of biological activities such as antioxidant activity, anti-inflammatory activity, hepatoprotective effect, antibacterial activity, antiviral activity, anticancer activity, and antidiabetic activity (Chen et al., 2018). Chen et al. reported that raspberry treatment could ameliorate H2O2-induced oxidative stress in HepG2 cells via Keap1/Nrf2 pathway (Chen et al., 2019). In our study, we found that atractylone (one component of YQJPF) has the similar effect with YQJPF, and inhibited necroptosis of hepatocytes in vitro. Which revealed the main component of YQJPF could play such effect.

In summary, YQJPF attenuates liver injury by inhibiting hepatic inflammation and necroptosis of hepatocytes in vivo and in vitro. Mechanistically, YQJPF inhibits hepatocyte necroptosis by blocking the RIPK1/RIPK3 complex formation followed by relieving mitochondrial dysfunction, which ultimately prevents hepatocytic cell death. Most importantly, we confirmed that YQJPF can be used as a medicine for ACLF treatment and provided a scientific basis for the research on the application of YQJPF in ACLF.

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DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The animal study was reviewed and approved by the Nanjing University of Chinese Medicine.

AUTHOR CONTRIBUTIONS

All of the authors took part in the experiments. Guarantor of the article: ST, FZ, and SZ. Author contributions: Study concept and design: ST, FZ, and SZ. Acquisition, analysis, and interpretation of data: FW, LT, BL, CJ, LG, YL, ZL, JS, and ZZ. Drafting the manuscript: FW and LT. Obtained funding: ST, FZ, and SZ.

FUNDING

This work was supported by the Leading Talent Project of Jiangsu Province Traditional Chinese Medicine (SLJ0216), the National Natural Science Foundation of China (81870423, 82073914, 82000572), the Major Project of the Natural Science Research of Jiangsu Higher Education Institutions (19KJA310005), the Postgraduate Research & Practice Innovation Program of Jiangsu Province, Grant/Award Number: KYCX20_1493, the Joint Project of Jiangsu Key Laboratory for Pharmacology and Safety Evaluation of Chinese Materia Medica and Yangtze River Pharmaceutical (JKLPSE202005), the Natural Science Foundation of Jiangsu Province (BK20200056).

Frontiers in Pharmacology | www.frontiersin.org April 2021 | Volume 12 | Article 658811
15 April 2021 | Volume 12 | Article 658811
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