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Anti-sepsis effect of Xiaochaihu decoction based on the TLR4/ MyD88/NF-κB signalling pathway

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ABSTRACT

The aims of this study were to explore the protective effect of Xiaochaihu decoction in mice with sepsis induced by intraperitoneal injection; to explore its anti-inflammatory effect on the TLR4-MyD88-NF- κ B signalling pathway; and to explore the main material basis of the anti-inflammatory effect of Xiaochaihu decoction, with the aim of supplementing and expanding the associated research and providing a scientific foundation for the clinical use of the decoction. The effects of Xiaochaihu decoction on septic mice were analysed by measurements of white blood cells (WBC) and Platelets (PLT); Nitric Oxide (NO) level in serum; IL-6, IL-1 β and TNF- α levels in serum; RT-PCR; Haematoxylin-Eosin (HE) immunohistochemistry; western blotting (WB). The results showed the excellent in vivo anti-inflammatory effects of Xiaochaihu decoction in LPS-induced septic mice, through down regulation of the gene and protein expression of TLR4, MyD88, TRAF6, IKK, IKB α and p65 and the subsequent reduction in the release of inflammatory mediators IL-6, IL-1 β , TNF- α and NO. Moreover, significant anti-septic effect was observed from high and medium doses of Xiaochaihu decoction, but not from the low dose.

1. Introduction

Sepsis, which is relatively common in the clinic, is mainly a systemic inflammatory reaction caused by pathogenic microbial infection [1]. It is an acute, life-threatening condition that can lead to organ dysfunction or circulatory dysfunction in severe cases [2]. The main symptoms of sepsis are chills and high fever. Patients can also have corresponding clinical symptoms at the site of infection [3]. However, in the face of acute and critical illness, such as sepsis, that endangers human life and health, even with today's highly developed medicine, the effects of treatment are poor. Therefore, there is an urgent need to examine the use of Chinese medicine as a means to assist in the treatment of sepsis.

Xiaochaihu decoction was first proposed by the medical sage Zhang Zhongjing in his *Treatise on Febrile and Other Diseases*. It is highly praised by famous doctors from all dynasties, and has gained the reputation of 'typhoid fever especially wonderful little bupleurum' [4]. The most consistent clinical indications of Xiaochaihu decoction is the concurrent cold and heat of Shaoyang disease, which is especially consistent with the main clinical symptoms of sepsis (chills and high fever), which is also one of the reasons why Xiaochaihu decoction was used to treat sepsis in this study. The indications of Xiaochaihu decoction are very wide. However, in most

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studies, Xiaochaihu decoction has be enlimited to the treatment of liver and bile diseases; there are few reports on the treatment of sepsis [5].

It has been reported that sepsis was related to the activation of TLR4 (6). The activation of TLR4 can initiate the downstream proinflammatory pathway of the MyD88/NF- κ B pathway and then release a large number of inflammatory mediators, leading to sepsis [7]. MyD88 interacts with the IL-1R-associated kinase (IRAK) complex, which recruits tumor necrosis factor receptor-associated Factor 6 (TRAF6). TRAF6 can promote transcriptional activation of nuclear factor-KB (NF-kB) genes and pro-inflammatory factors tumor necrosis factor (TNF- α) and interleukin-6 (IL-6) and other inflammatory factors such asinterleukin-1 β (IL-1 β) [8]. Phosphorylation of IkBa is a critical event after TRAF6 activation [9]. CHUK is an important component of the non-classical NF-kB signaling pathway [10]. Therefore , our aim of detection above indexes was to uncover the mechanism how do Xiaochaihu decoction work.

In this study, we observed the protective effect of Xiaochaihu decoction on mice with lipopolysaccharide (LPS)-induced sepsis, a common model of inflammation, and showed that the mechanism of the anti-sepsis effect. This will allow us to improve knowledge of the anti-sepsis effect of Xiaochaihu decoction and provide ascientific basis for its clinical application (Fig. 1).

2. Materials and methods section

2.1. Preparation of Xiaochaihu decoction

The Xiaochaihu decoction was prepared as per the original prescription in the Treatise on Febrile Diseases: *Bupleurum* chinense DC. (Chaihu) Half jin, *Scutellaria baicalensis* Georgi. (Huangqin) SAN liang, Panax ginseng C.A.Mey. (Renshen) SAN liang, *Pinellia* ternata (Thunb.) Half-litre, Glycyrrhiza uralensis Fisch (Zhigancao) SAN liang, Zingiber officinale Roscoe (Shengjiang) SAN liang and Ziziphus jujuba Mill. (Dazao) 12. For the high-dose group, the decoction administered was, according to the archaeology, 1 liang equal to approximately 15 g; for the medium-dose group, the decoction administered was, according to the clinical experience, 1 liang equal to approximately 5 g; and for the low-dose group, the decoction administered was, according to the teaching material, 1 liang equal to approximately 3 g. Namely, in the high-dose group (XCHD H), treatment comprised: Bupleurum 120 g, Scutellariae 45 g, Ginseng 45 g, Pinellia 45 g, Glycyrrhiza 45 g, Zingiber 45 g and Jujube 45 g (for a total daily dosage of 390 g per 60 kg adult). In the medium-dose group (XCHD M), treatment comprised: Bupleurum 40 g, Scutellariae 15 g, Ginseng 15 g, Pinellia 15 g, Glycyrrhiza 15 g, Zingiber 15 g and Jujube 15 g (for a total daily dosage of 130 g per 60 kg adult). In the low-dose group (XCHD L), treatment comprised: Bupleurum 24 g, Scutellariae 9 g, Ginseng 9 g, Pinellia 9 g, Glycyrrhiza 9 g, Zingiber 9 g and Jujube 9 g (for a total 60 kg adult daily dosage 78 g), the ingredients were presented in Table 1. Eight volumes of the medicinal materials were soaked in water for 30 min, heated and boiled, simmered for 60 min, and the residue was then filtered while hot. This process was repeated 3 times; The filtrate was combined 3 times and concentrated to the required concentration under the condition of 60 °C to obtain the brown viscous extract. High, medium and low doses of Xiaochaihu decoction were obtained and put in the refrigerator until use [11–19].



Fig. 1. Schematic of anti-sepsis effectiveness of Xiaochaihu decoction through the TLR4/MyD88/NF-κB signalling pathway.

Table 1

Dose of medication	XCHD H	XCHD M	XCHD L
Bupleurum	120 g	40 g	24 g
Scutellariae	45 g	15 g	9 g
Ginseng	45 g	15 g	9 g
Pinellia	45 g	15 g	9 g
Glycyrrhiza	45 g	15 g	9 g
Zingiber	45 g	15 g	9 g
Jujube	45 g	15 g	9 g
Total	390 g	130 g	78 g

2.2. Design of animal experiment

Seventy-two healthy male KM mice (as the female mice had the effect of menstruation) were randomly divided according to body weight into each group. 12 mice were randomly allocated to the Normal group, 12 mice to the LPS group, 12 mice to the Dexamethasone (DXM, 5 mg/kg) group [20], 12 mice to the XCHD H (60.7 g/kg) group, 12 mice to the XCHD M (20.2 g/kg) group and 12 mice to the XCHD L (12.1 g/kg) group. The following formula was used: daily dose/mice = dose/kg (human body weight) \times conversion factor 9.1 \times body weight of each mice. The mice in each group were administered with 9.1 times the daily clinical crude drug dosage of Xiaochaihu decoction for a 60-kg adult, and received continuous intragastric administration for 7 days, twice a day, while the Normal group was given equal volume pure water. On day 6, 30min after the last dosing, sepsis modelling was performed. Earlier models of sepsis often involved direct administration of toxins such as lipopolysaccharide (LPS) into the blood, peritoneum, or lung. This induces a strong immediate inflammatory response that mimics activation of the innate immune system in human sepsis. The advantage of using this approach is its technical ease and reproducible response following injection of a quantifiable dose of toxin [21]. Therefore, all mice except those in the Normal group were injected intraperitoneally with LPS 0.1 mL/10 g at an injection dose of 15 mg/kg [22]; the Normal group was administered the same dose of normal saline [23–27].

The glucocorticoids often used for sepsis [28], so DXM was used in our experiment as a comparison group to measure the effects of experimental medicines.

These experiments has been approved by Laboratory Animal Ethics Committee, Shenzhen University Health Science Center. The ethical approval referenced number was 4403050195161. And we stated confirmation that these experiments were conducted according to established animal welfare guidelines.

2.3. Effects of Xiaochaihu decoction on organ index, whole blood, WBC, and PLT in septic mice

At 12 h after intraperitoneal injection of LPS, blood was collected from the orbit. All mice were sacrificed after collecting blood by neck mutilation and their organs were dissected immediately. Then, 20 μ L of whole blood was taken to measure the white blood cell (WBC) count and platelet (PLT) count in the whole blood. The removed organs were weighed on a balance to calculate organ index, which was equal to organ weight divided by the mouse body weight [29–31].

2.4. Effect of Xiaochaihu decoction on serum NO in septic mice

The content of NO in serum was determined by Griess's method, and the content of NO in whole blood (centrifugation at 3500 rpm for 15min) and lung tissue was determined 12 h after modelling [32–34].

2.5. TNF- α , IL-6 and IL-1 β in serum were determined by ELISA

Whole blood of mice was collected from the orbit at 12 h after intraperitoneal injection of LPS and centrifuged at 3000 rpm for 15 min to obtain serum. The serum levels of TNF- α , IL-6 and IL-1 β were determined in accordance with the instructions of the ELISA kit. Finally, OD values were measured at 450 nm, and serum levels of TNF- α , IL-6 and IL-1 β were calculated according to the established standard curve [35–39].

2.6. Preparation of HE pathological sections of lung tissue

Immediately after blood collection, each animal's lung was removed and immersed in a 10% neutral formalin buffer. The paraffinembedded lung tissues were then cut into 5 μ m thick slices, stained with haematoxylin–eosin (HE), and examined usinga digital trimograph microscope (BA400Digital, MacOdis Industrial Group Limited). To quantify the extent of histopathological changes, the number of neutrophils in the alveolar septum was calculated using Image-Pro Plus (version 6.0, Media Cybernetics) software. The infiltration of inflammatory cells was presented as a change in the mean number of neutrophils in each of the six photographs taken at 200 × and 400 × magnifications to provide a more detailed view of the lung lesions [40–44]. Two observers utilize ImageJ software, for staining observations to ensure more objective data analysis. The dried sections were observed under a microscope and photographed [45].

2.7. RT-PCR

Real-time PCR was performed in accordance with the manufacturer's instructions for the total RNA miniprep kit (Axygen, USA), and total RNA was extracted from lung tissue using the Fast Quant RT kit (Tiangen, Beijing, China). The following conditions were used: 95 °C for 15 min, followed by 40 denaturation cycles of 95 °C for 10 s followed by extension at 64.1 °C for 30 s. Then, a 96-well reaction plate (Bio-Rad, Hercules, CA) with a reaction volume of 20 μ L was applied directly for the PCR amplification reaction and the mRNA levels of TLR4, MYD88, TRAF6, IKK, IKKB- α , and p65 were detected.

2.8. The protein expression of TLR4, MYD88, TRAF6, IKK, IKKB-a and p65 in cell lysates were detected by western blotting

The total protein was extracted from the lungs with lysate buffer and the concentration was determined using a BCA protein assay kit. All samples were then adjusted to the same concentration. Proteins in lung tissue were separated on a gel containing 10% polyacrylamide by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to a polyvinylidene fluoride (PVDF) membrane. After transfer, the PVDF membranes were washed in TBST (Tris buffered saline containing 0.1% Tween 20) for 5 min, and then the antibodies were added in skim milk. An appropriate amount of sealing solution was added and left to oscillate slowly at 37 °C for 2 h. The membranes were then incubated overnight at 4 °C with primary antibody to TLR4, MYD88, TRAF6, IKK, IKKB- α , and p65 proteins were diluted in accordance with the manufacturer's instructions. After three washes with TBST, the HRP-labelled secondary antibody (1:1000) was incubated at 37 °C for 1.5 h, and then ECL western blotting reagent was used to detect the secondary antibody. The optical density was measured by the laborator imaging analysis system and the ratio of the optical density of the target protein to the internal reference protein band was used as the result.

The expression of TLR4, MYD88, TRAF6, IKK, IKK α , and p65 in lung tissues was determined by immunohistochemistry [46] Lung tissue from each group was immobilised, dehydrated, and the clear sample was immersed in wax and cut into thin slices



Fig. 2. Effects of Xiaochaihu decoction on the evolution of the spleen and thymus index and WBC, PLT, NO, IL-1 β , IL-6, TNF- α in septic mice as determined by ELISA. Normal, blank group; LPS, model group; XCHD H, Xiaochaihu decoction high-dose group; XCHD M, Xiaochaihu decoction medium-dose group; and XCHD L, Xiaochaihu decoction low-dose group. Compared with the LPS group, p < 00.05, p < 00.01.

(approximately 4–5 μ m in thickness). Next, the sections were routinely dewaxed and soaked in 3% hydrogen peroxide at room temperature for 10 min to inactivate endogenous peroxidase, and then washed three times in distilled water. The water bath was then heated for antigen recovery and sealed with goat serum at room temperature for 20 min. Primary antibodies to TLR4, MYD88, TRAF6, IKK, IKKB- α and p65 were incubated at 4 °C (diluted appropriately according to the manufacturer's instructions; Abcam) overnight. After three washes in PBS, biotinised goat anti-rabbit IgG was added dropwise and then incubated at 37 °C for 30 min. After three washes with PBS,the sections were incubated with horseradish peroxidase (HRP) at 37 °C for 30 min. Then, after four washes with PBS, diaminobenzidine (DAB) (OriGene, Beijing) reaction was performed. After washing with distilled water, haematoxylin was slightly redyed, dehydrated and sealed with neutral glue.

3. Results

3.1. Effects of Xiaochaihu decoction on whole blood WBC and PLT in mice

The results of WBC and PLT in whole blood are shown in Fig. 2. Compared with the LPS group, the spleen index and the thymus index were significantly decreased, and the whole blood white blood cell (WBC) count was significantly decreased in the medium-dose group of Xiaochaihu decoction, suggesting that the decoction could affect the immune defence function in the septicmodel mice and thereby relieve the inflammatory response.



Fig. 3. Effect of Xiaochaihu decoction on mRNA expression of IL-1 β , IL-6, TNF- α , TLR4, MYD88, TRAF6, CHUK, IKB- α and p65 in lung tissue from septicmice. Normal, blank group; LPS, model group; XCHD H, Xiaochaihu decoction high-dose group; XCHD M, Xiaochaihu decoction medium-dose group; and XCHD L, Xiaochaihu decoction low-dose group. Compared with the LPS group, p < 00.05, p < 00.01.

3.2. Effect of Xiaochaihu decoction on serum NO in mice

Compared with the Normal group, the serum NO content of mice in the LPS group was significantly increased (P < 0.01 or P < 0.05). Compared with LPS group, the Xiaochaihu decoction medium-dose group had significantly lower serum NO levels (P < 0.01) (Fig. 2).

3.3. Effects of Xiaochaihu decoction on serum levels of IL-1 β , IL-6 and TNF- α in mice determined by ELISA

See Fig. 2 for the results. Compared with the Normal group, the serum contents of IL-1 β , IL-6 and TNF- α were significantly higher in the LPS group (P < 0.01). Compared with the LPS group, the serum levels of IL-1 β , IL-6 and TNF- α were significantly lower in the Xiaochaihu decoction high-dose group (P < 0.01 or P < 0.05).

3.4. RT-PCR

The RT-PCR results for the analysis of lung tissues for each group of mice are presented in Fig. 3. According to these results, the mRNA expression of IL-1 β , IL-6, TNF- α , TLR4, MYD88, TRAF6, CHUK, IKB- α and NF- κ B (p65) in lung tissue from LPS group was significantly higher than in the Normal group (P < 0.01 or P < 0.05). Compared with the LPS group, the mRNA expression of IL-1 β , IL-6, TNF- α , TLR4, MYD88, TRAF6, CHUK, IKB- α and NF- κ B (p65) in the lung tissue in the Xiaochaihu decoction high-dose and medium-dose groups was significantly reduced. However, the mRNA expression levels of the above genes in lung tissue was significantly decreased in the Xiaochaihu decoction low-dose group (P < 0.05), with a slight or no effect on individual indicators.

3.5. HE staining of lung tissue sections

The HE staining of the sections revealed that, in the Normal group, the lung capsule was complete, the bronchial tubes and blood vessels at all levels were complete, the epithelium of the bronchial mucosa was arranged neatly, no cells were shed in the bronchial lumen, the vascular structure was complete, the epithelium cells were arranged regularly, the blood cell aggregation and plasma deposition were observed in the vascular lumen, the alveolar structure was normal and the alveolar epithelium cells were arranged regularly. Inflammatory cell infiltration into the lung tissue of mice with sepsis was alleviated in the Xiaochaihu decoction high-dose and medium-dose groups as shown in Fig. 4.

3.6. The expression of TLR4, MYD88, TRAF6, IKK, IKK α and p65 in lung tissues was determined by immunohistochemistry

As shown in Fig. 5, Immunohistochemical analysis showed that, compared with the Normal group, the positive staining of TLR4, MYD88, TRAF6, IKK, IKK α and p65 in the lung tissue of mice in the LPS group was significantly increased (P < 0.01). Compared with the LPS group, the Xiaochaihu decoction high-dose and medium-dose groups had lower positive staining of the above indicator proteins (P < 0.05).

3.7. The protein expression of TLR4, MYD88, TRAF6, IKK, IKKB- α and p65 in cell lysate was detected by western blotting

As shown in Fig. 6, compared with the Normal group, the relative protein expression of TLR4, MYD88, TRAF6, IKK, IKK α and p65 in the lung tissue was significantly increased in mice with sepsis (the model group) (P < 0.05). Compared with the LPS group, the relative protein expression of TLR4, MYD88, TRAF6, IKK, IKK α and p65 in the Xiaochaihu decoction high-dose and medium-dose groups of was significantly decreased (P < 0.05). When the MYD88 inhibitor ST2825 was added to the model group (i.e., the LPS + ST2825 group),



Fig. 4. Effect of Xiaochaihu decoction HE staining of the lungs of septic mice in various conditions (400× magnification). Normal, blank group; LPS, model group; XCHD H, Xiaochaihu decoction high-dose group; XCHD M, Xiaochaihu decoction medium-dose group; and XCHD L, Xiaochaihu decoction low-dose group.



Fig. 5. Effects of Xiaochaihu decoction on TLR4, MYD88, TRAF6, IKK, IKKB- α and p65 protein levels in the lung tissues of septic mice determined by immunohistochemical staining (Magnification 400×). Normal, blank group; LPS, model group; XCHD H, Xiaochaihu decoction high-dose group; XCHD M, Xiaochaihu decoction medium-dose group; and XCHD L, Xiaochaihu decoction low-dose group. Data are expressed as the mean \pm SEM of 10 mice per group, compared with the LPS group, p < 00.05, p < 00.01.

TLR4, TRAF6, IKK and IKK α in lung tissue of septicmice was obviously inhibited, whereas MYD88 and p65 were significantly inhibited.

4. Discussion

In this study, the clinical application of Xiaochaihu decoction to the treatment of sepsis was of great promise. In clinical practice, we found that Xiaochaihu decoction could reduce sepsis patients' hyperthermia. Eventually, we reviewed a lot of relevant papers and deduced that Xiaochaihu decoction played its anti-inflammatory and antipyretic effects through TLR4/MyD88/NF-κB signaling pathway. After the occurrence of sepsis, lung was an vulnerable organ that was the "first responders" to sepsis damage. Therefore, we deduced that Xiaochaihu decoction reduce inflammatory cell infiltration in lung tissue of septic mice [47]. The efficacy and mechanism of Xiaochaihu decoction against sepsis in mice were examined. The results showed that [1]: both high-dose and medium-dose Xiaochaihu decoction could significantly reduce hyperthermia in septic mice and ameliorates hivering symptoms [2]. The results of the organ index, whole blood WBC, and PLT count showed that the spleen index and thymus index were significantly decreased in the Xiaochaihu decoction medium-dose group, suggesting that it may have a certain effect on the body's defences and immune function, with a significant reduction in the count of WBC and alleviation of the inflammatory response in septic mice (P < 0.01) [3]. Compared with Normal group, the serum NO content in the LPS group was significantly increased (P < 0.01). Compared with the LPS group, the Xiaochaihu decoction medium-dose group had significantly lower serum NO levels (P < 0.05) [4]. ELISA showed that, compared with LPS group, the serum levels of IL-1 β , TNF- α and IL-6 in the Xiaochaihu decoction high-dose and medium-dose groups were significantly decreased (P < 0.01 or P < 0.05) [5]. The histopathological results showed that Xiaochaihu decoction high-dose and medium-dose groups could both reduce inflammatory cell infiltration in lung tissue of septic mice [6]. Compared with the Normal group, the mRNA and protein expression of TLR4, MYD88, TRAF6, IKK, IKKα and p65 in the lung tissue of mice in the LPS group were significantly increased (P < 0.01 or P < 0.05). Compared with the LPS group, the Xiaochaihu decoction high-dose and medium-dose groups had significantly lower mRNA expression levels and protein expression levels in the lung tissue of septic mice (P < 0.01 or P < 0.05) [7]; Immunohistochemical staining revealed that compared with Normal group, the positive staining rates of TLR4, MYD88, TRAF6,



Fig. 6. Effects of Xiaochaihu decoction on the expression of TLR4, MYD88, TRAF6, IKK, IKKB-α and p65 in lung tissue of mice (refer to Fig. S1-Fig. S7). Normal, blank group; LPS, model group; XCHD H, Xiaochaihu decoction high-dose group; XCHD M, Xiaochaihu decoction medium-dose group; and XCHD L, Xiaochaihu decoction low-dose group. Comparison with LPS group, p < 00.05, p < 00.01.

IKK, IKK α and p65 proteins in the lung tissue of mice in the LPS group were significantly increased (P < 0.01). Compared with LPS group, the Xiaochaihu decoction high-dose and medium-dose groups had lower positive staining rates of the above indicator proteins (P < 0.05). All the evidence suggested that the mechanism and efficacy of the anti-sepsis effects of Xiaochaihu decoction high-dose and medium-dose was through the TLR4/MyD88/NF- κ B signalling pathway. However, due to the complexity of Traditional Chinese Medicine (TCM) ingredients, we don't yet know exactly which ingredients are at work, that's where we're headed in the future.

5. Conclusion

In this study, Xiaochaihu decoction exerted a certain protective effect in septic mice, and this protective effect was notably better with the medium and high doses than with the low dose. Based on the above in vitro experiments, the mechanism and efficacy of the anti-sepsis effects of medicinal serum of medium-dose Xiaochaihu decoction were confirmed. In conclusion, Xiaochaihu decoction works through the TLR4/MyD88/NF-xB signalling pathway.

Data availability statement

The present study was guided by the basic principles of research integrity and research conduct with human participants, as stipulated in many documents. Participants were informed that their anonymity and the confidentiality of data pertaining to them will be retained. Hence, we were abided by this agreement with the participants not to share and deposit data for public use.

CRediT authorship contribution statement

Qingxin Yang: Writing – review & editing, Writing – original draft, Software, Resources, Project administration, Methodology, Investigation, Funding acquisition, Formal analysis, Data curation, Conceptualization. Yulong Wang: Writing – review & editing, Writing – original draft, Validation, Resources, Methodology, Formal analysis, Data curation, Conceptualization. Gefei Cao: Writing – review & editing, Writing – original draft, Validation, Software, Methodology, Formal analysis, Data curation, Conceptualization. Xiaoqing Li: Software, Methodology, Formal analysis, Data curation, Conceptualization. Tinghui Zhao: Writing – review & editing, Writing – original draft, Validation, Supervision, Resources, Project administration, Methodology, Investigation.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

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