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Original Paper

β-Hydroxybutyrate Activates the NFκΒ Signaling Pathway to Promote the Expression of Pro-Inflammatory Factors in Calf Hepatocytes

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Key Words

Oxidative stress • NF-kB pathway • Inflammatory injury • Bovine hepatocytes

Abstract

Background/Aims: β-hydroxybutyrate (BHBA) is the major component of ketone bodies in ketosis. Dairy cows with ketosis often undergo oxidative stress. BHBA is related to the inflammation involved in other diseases of dairy cattle. However, whether BHBA can induce inflammatory injury in dairy cow hepatocytes and the potential mechanism of this induction are not clear. The NF-κB pathway plays a vital role in the inflammatory response. *Methods*: Therefore, this study evaluated the oxidative stress, pro-inflammatory factors and NF-kB pathway in cultured calf hepatocytes treated with different concentrations of BHBA, pyrrolidine dithiocarbamate (PDTC, an NF-kB pathway inhibitor) and N-acetylcysteine (NAC, antioxidant). **Results:** The results showed that BHBA could significantly increase the levels of oxidation indicators (MDA, NO and iNOS), whereas the levels of antioxidation indicators (GSH-Px, CAT and SOD) were markedly decreased in hepatocytes. The ΙΚΚβ activity and phospho-Ικβα (p-ΙκΒα) contents were increased in BHBA-treated hepatocytes. This increase was accompanied by the increased expression level and transcription activity of p65. The expression levels of NF-κB-regulated inflammatory cytokines, namely TNF-α, IL-6 and IL-1β, were markedly increased after BHBA treatment, while significantly decreased after NAC treatment. However, the p-IκBα level and the expression and activity of p65 and its target genes were markedly decreased in the PDTC + BHBA group compared with the BHBA (1.8 mM) group. Moreover, immunocytofluorescence of p65 showed a similar trend. Conclusion: The present data indicate that higher concentrations of BHBA can induce cattle hepatocyte inflammatory injury through the NF-kB signaling pathway, which may be activated by oxidative stress.

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Introduction

The liver is an important metabolic organ and regulates lipid metabolism in animals. Lipid metabolism disorders, such as fatty liver and ketosis, involve liver dysfunction and hepatocyte injury in the later stages of these diseases [1]. In general, dairy cows were subjected to a period of negative energy balance (NEB) during the transition period. Excessive NEB initiates fat mobilization and a subsequent increase in blood non-esterified fatty acid (NEFA) concentration [2, 3]. Large amounts of NEFAs are transported into the liver and are involved in β-oxidation in the hepatocytes to generate more ATP to relieve the NEB. However, large amounts of NEFAs could not be completely oxidized and metabolized into ketones, such as β-hydroxybutyrate (BHBA), the major ketone body. Clinical studies have demonstrated that oxidative stress exists in dairy cows with ketosis [4, 5]. BHBA is touted as an energy substrate that could be generated by the reduction of acetoacetate in the liver of all species and by the oxidation of butyrate exclusively in the ruminal epithelium. Moreover, BHBA is involved in ATP production, energy metabolism and thermogenesis [6]. Some studies have reported that BHBA could activate some signaling pathways as a signaling molecule and is involved in the regulation of food intake, thermogenesis, neuroprotection, and cell viability [6-9]. Even more remarkably, elevated concentrations of BHBA are related to endometritis and mastitis during the transition of dairy cows [10, 11]. Further studies have demonstrated that inflammation could lead to hepatic injury in mice [12, 13]. These studies suggest that BHBA may also be associated with the inflammatory injury of hepatocytes in dairy cattle. However, the underlying mechanisms in dairy cattle are not clear.

The nuclear factor NF-κB pathway plays a vital role in the inflammatory response and immunity [14]. The most classical NF-κB binding form is a p65-p60 that remains associated with the IkB family of inhibitory proteins in the cytoplasm before activation [15]. The cytoplasmic NF-kB/lkB complex is phosphorylated on conserved serine residues of lkB, which leads to the dissociation of IκB from NF-κB and the subsequent translocation of p65 NF-κB to the nucleus. After tyrosine phosphorylation, p65 NF-κB binds to its cognate DNA-binding site in the promoter regions of specific genes, such as the pro-inflammatory factors tumor necrosis factor alpha (TNF-α), interleukin 6 (IL-6) and interleukin 1 beta (IL-1β) [16]. A study reported that oxidative stress-induced hepatic injury could be regulated by the NF-κB signaling pathway in type I diabetic rats [17]. This result indicates that the NF-κB signaling pathway might mediate oxidative stress-induced hepatic injury. However, whether oxidative stress-induced hepatic injury is also regulated by the NF-κB pathway in dairy cattle is not clear. Moreover, oxidative stress is a common mechanism of liver injury [18] and activates the NF-κB-mediated signaling pathway [19]. Dairy cows with severe NEB during early lactation increased oxidative stress [20, 21] and high blood concentrations of BHBA. These results triggered our interest in exploring the relationships among the NF-κBmediated signaling pathway, oxidative stress and hepatocyte inflammatory injury in dairy cattle with NEB.

Therefore, the aim of this study was to investigate whether high levels of BHBA could activate the oxidative stress-mediated NF-κB signaling pathway to promote the synthesis of pro-inflammatory factors in cattle hepatocytes.

Materials and Methods

Ethical approval of the study protocol

All the experiments were performed in accordance with the Guiding Principles in the Use of Animals adopted by the Chinese Association for Laboratory Animal Sciences. All the animal procedures were approved by the Institutional Animal Care and Use Committee of Jilin University.

Isolation of cow primary hepatocytes

Calf (one-day-old, female, 45-50kg, health, fasting) primary hepatocytes were isolated and cultured as previously described [22]. Briefly, a scalpel was used to obtain the caudate lobe of the liver from a Holstein KARGER

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calf that was anesthetized with thiamylal sodium under sterile conditions. The liver was then quickly placed on a sterile bench, and bloodstains on the surface were removed with perfusion solution A (140 mM NaCl, 6.7 mM KCl, 10 mM HEPES, 2.5 mM glucose and 0.5 mM EDTA, pH 7.4, 37°C), revealing the blood vessel on the cross section of the caudate process. A moderate blood vessel was intubated, and the liver was perfused with perfusion solution A (37°C) at a flow rate of 50 mL/min for 15 min. The liver was then perfused with solution B (140 mM NaCl, 6.7 mM KCl, 30 mM HEPES, 2.5 mM glucose and 5 mM CaCl, pH 7.4, 37°C) at the same infusion rate for 3 min. When the liquid became clear, the second step of the perfusion digestion, the addition of perfusate C (37°C, 20 mL/min), was started. Before the final perfusion, all of the other coarse vessels were partially ligated so that the perfusate flowed slowly. Then, 100 mL of RPMI-1640 basic medium was added to the plate in which the digested liver was placed, and the blood vessels, fat and connective tissue were removed. Any parts of the liver caudate lobe that were incompletely digested were cut away, and the remainder of the liver parenchyma was cut into pieces and filtered sequentially through 100 mesh (150 μm) and 200 mesh (75 μm). The hepatocyte suspension was washed twice with basic medium before resuspension in adherent medium. The cell density was adjusted to 2×106 cells/mL with adherent culture medium. The hepatocyte suspension was seeded into a six-well tissue culture plate (2 mL/well) and incubated at 37°C in 5% CO₂. After 4 h, the culture medium was changed to growth medium (containing 10% fetal calf serum, benzylpenicillin and streptomycin). The medium was replaced with fresh medium every 24 h.

BHBA Preparation and Treatment

The concentrations of BHBA used in this study were chosen according to the BHBA concentrations in cow serum [23]. Stock BHBA solution was prepared as follows: 1.206 g of BHBA powder was dissolved in 50 mL of distilled water, sterilized by filtration to obtain a concentration of 0.2 mM, and stored at -20°C. Before the treatment with BHBA, the cells were serum-starved overnight. The hepatocytes were subjected to the following treatments. For the time course experiments, the hepatocytes were treated with BHBA (1.8 mM) for 0, 1, 3, 6, 9, 12 and 24 h. For the dose response experiments, the hepatocytes were treated with BHBA, PDTC (pyrrolidine dithiocarbamate, an IκBα inhibitor) and N-acetylcysteine (NAC, one of the most used antioxidant drugs in liver diseases), respectively. PDTC can inhibit the activation of NF-κB specifically by suppressing the release of the inhibitory subunit IkB from the latent cytoplasmic form of NF-kB. The hepatocytes were treated with 0, 0.6, 1.2, 1.8 and 2.4 mM BHBA, 10 µM PDTC, 10 µM PDTC + 1.8 mM BHBA (PDTC + BHBA) and 10 mM NAC + 1.8 mM BHBA (NAC + BHBA) for 24 h. Each treatment was replicated nine times.

MDA, NO, GSH-Px, SOD and CAT content determination

The hepatocytes were harvested after incubation with different concentrations of BHBA for 24 h and washed twice in ice-cold phosphate-buffered saline (PBS). Then, the cells were lysed using P0013D (Beyotime, Jiangsu, China). The lysate was centrifuged at $12,000 \times g$ for 5 min at 4°C. The supernatant was used to determine the content of malonaldehyde (MDA), nitric oxide (NO), glutathione peroxidase (GSH-Px), superoxide dismutase (SOD) and catalase (CAT) using the appropriate biochemical kits (Beyotime, Jiangsu, China) according to the manufacturer's instructions.

IKKβ activity determination

The hepatocytes were harvested after incubation with different concentrations of BHBA for 24 h and washed twice in ice-cold GENMED clean buffer (Reagent A, GenMed Scientifics Inc., USA) for 5 min at 300 × g at 4°C. Subsequently, the cells were treated with lysis buffer (Reagent B) and incubated for 30 minutes in an ice bath. The lysate was centrifuged for 5 min at $16,000 \times g$ at 4° C, and the IKK β activity in the supernatant was measured using a spectrophotometer and a biochemical kit (Reagent A, GenMed Scientifics Inc., USA) according to the manufacturer's instructions.

RNA extraction and Real-time PCR

Total hepatocyte RNA was extracted with TRIzol reagent (TaKaRa Biotechnology Co., Ltd., Tokyo, Japan) according to the supplier's protocol. The RNA concentration was determined using a K5500 Micro-Spectrophotometer (Beijing Kaiao Technology Development Co., Ltd., Beijing, China) before reverse transcription polymerase chain reaction (RT-PCR). Approximately 5 µg of total RNA was reverse-transcribed to cDNA in 20-µL reactions using PrimeScript Reverse Transcriptase (TaKaRa Biotechnology Co., Ltd., Tokyo, KARGER

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Table 1. The primers sequences used for cDNA generation

Gene	Sequence number	Primer sequences (5'-3')	Length (bp)
NF-кВ p65	NM_001080242.2	For AGGACCAACCAGACCG	240
		Rev TGTCACCAGGCGAGTTAT	
TNFα	NW_003104557.1	For CTGCCGGACTACCTGGACTAT	234
		Rev CCTCACTTCCCTACATCCCTAA	234
IL6	NW_00310889.1	For AACGAGTGGGTAAAGAACGC	144
		Rev CTGACCAGAGGAGGGAATGC	
IL1β	NW 003104294.1	For CTGAACCCATCAACGAAA	190
	NW_003104294.1	Rev ATGACCGACACCACCTGC	
GSH-Px	NM_001101113.2	For GCGGGAGCAGGACTTCTACGA	137
		Rev CCCGATAGTGCTGGTCTGTGAA	
Mn SOD	NM_201527.2	For TTACATAAGGAGCAGGGACG	234
		Rev CAGTGTAAGGCTGACGGTTT	
Cu/Zn SOD	BC 102432.1	For GAAGAGAGGCATGTTGGAGA	220
		Rev CCAATTACACCACGAGCCAA	
CAT	NM_001035386.1	For AGATACTCCAAGGCGAAGGTG	120
		Rev AAAGCCACGAGGGTACCGAAC	
β-actin	BC 142413.1	For GCCCTGAGGCTCTCTTCCA	101
		Rev GCGGATGTCGACGTCACA	

Japan) according to the manufacturer's instructions. The primers were designed using Primer Express software (PE Applied Biosystems, Inc., Foster City, CA, USA; Table 1). The mRNA expression levels were estimated by real-time PCR analysis using FS Universal SYBR Green Real Master (Roche) on the 7500 Real-Time PCR System (Applied Biosystems). The conditions were as follows: 95°C for 3 min, followed by 40 cycles of 95°C for 15 sec and 60°C for 1 min. All the reactions were run in triplicate. The relative expression levels were normalized to β -actin levels.

Western Blotting

The hepatocytes were harvested from the culture plates with PBS, and the total cellular proteins (TP) and nuclear proteins were extracted using a protein extraction kit and a nuclear protein extraction kit (Sangon Biotech Co., Ltd., Shanghai, China), respectively, according to the manufacturer's instructions. Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) separation was performed with 50 μg of protein per lane and with known molecular weight markers (Sangon Biotech Co., Ltd., Shanghai, China). The proteins were then electrophoretically transferred onto polyvinylidene fluoride (PVDF) membranes. Then, the membranes were incubated in blocking solution (5% nonfat milk in Tris-HCl buffer solution [TBS]) for 6 h at 4°C. The blocked membranes were hybridized overnight at 4°C with antibodies against NF-κB p65, phospho-IκBα (p-IκBα), IκBα, histone and β-actin. Next, the membranes were washed three times with TBS containing 0.1% Tween 20 for 5 min at room temperature with shaking, incubated with the appropriate peroxidase-conjugated secondary antibodies for 45 min at room temperature with shaking and then washed four times for 5 min. An immunodetection analysis was performed using an enhanced chemiluminescence solution (ECL, Pierce Biotechnology Inc., Chicago, IL, USA). The relative expression levels of NF- κ B p65, phospho- $I\kappa$ B α (p- $I\kappa$ B α) and $I\kappa$ B α were normalized to the β -actin levels.

Enzyme-linked immunosorbent assay (ELISA)

After the hepatocytes were treated with BHBA, PDTC and NAC, respectively, as described above, the cell-free supernatants were subsequently centrifuged at 3000 rpm for 20 min to extract the upper liquid for use in assays for the pro-inflammatory cytokines TNF- α , IL-6 and IL-1 β with the bovine tumor necrosis factor alpha (TNF-α) ELISA Kit, bovine interleukin-6 (IL-6) ELISA Kit and bovine interleukin-1β (IL-1β) ELISA Kit (IBL, Germany), respectively, according to the manufacturer's instructions.

Electrophoretic mobility shift assay (EMSA)

Nuclear proteins were extracted using a nuclear protein extraction kit (Sangon Biotech Co., Ltd., Shanghai, China) according to the manufacturer's instructions. The special probe recognition sequence for NF-κB is 5'-AGTTGAGGGGACTTTCCCAGGC-3'. The probe was labeled with biotin by incubating at 37°C for 1 h and purified using centrifugal chromatography. The binding reaction was performed using the Lightshift EMSA Optimization and Control Kit (Pierce Biotechnology, Inc., Rockford, IL, USA) according to the manufacturer's instructions. An EMSA/gel-shift binding buffer containing 6 µg of nuclear extract was incubated with the biotin-labeled probe for 20 min at room temperature. The nuclear protein complexes were separated by electrophoresis on non-denaturing 6% polyacrylamide tris/borate/EDTA (TBE) gels and electrotransferred onto a nylon membrane. Then, the membranes were cross-linked using a UV cross-linker

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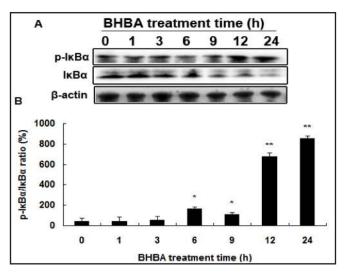
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Fig. 1. Effects of BHBA treatment duration on IkBα phosphorylation in calf hepatocytes. Hepatocytes were treated with 1.8 mM BHBA for 0, 1, 3, 6, 9, 12 and 24 h. (A) Western blots of β-actin, IkBα and p-IkBα. (B) The phosphorylation ratio of IkBα. The data are shown as the mean \pm SEM. *, statistical significance (P<0.05). **, statistical significance (P<0.01). The symbol * and ** indicate statistical differences with the control group (0 h).



(Cany Precision Instruments Co., Ltd., Shanghai, China), and the biotin-labeled probe was detected using an enhanced chemiluminescence solution (ECL; Pierce Biotechnology Inc., Chicago, IL, USA).

Immunocytofluorescence

Calf hepatocytes were grown on glass coverslips to 90% confluency and subjected to BHBA or PDTC treatment as described above. Upon completion of the treatment, the coverslips were washed with PBS, fixed with 4% paraformaldehyde for 20 min at room temperature, subjected to antigen retrieval with EDTA•2Na (95°C, 5 min) and permeabilized with 0.1% Triton X-100 (Sigma-Aldrich, St. Louis, MO). After further washing, the hepatocytes were exposed to the primary antibody NF-κB p65 at 4°C overnight and then incubated with goat anti-rabbit IgG conjugated with cy3 (Beyotime, China) at room temperature for 30 min and counterstained with Hoechst 33258 (Beyotime, China). The coverslips were observed and photographed using laser-scanning confocal microscopy (Fluoview FV1200, Olympus, Japan).

Chemical reagents and antibodies

Collagenase IV, heparin sodium, fetal bovine serum and RPMI-1640 medium were purchased from Gibco (Grand Island, NY, USA). HEPES, insulin, β -hydroxybutyrate powder, NAC and the NF- κ B inhibitor PDTC were provided by Sigma-Aldrich (St. Louis, MO, USA). Vitamin C, dexamethasone acetate, penicillin, streptomycin and other chemicals were purchased from Baoman Biotechnology (Shanghai, China). The β -actin antibody was purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). The p-I κ B α and I κ B α antibodies were provided by Cell Signaling Technology (Danvers, MA, USA). The NF- κ B p65 antibody was supplied by Abcam (Cambridge, MA, USA). Six-well plates and filters were purchased from Corning Incorporated (Corning, NY, USA).

Statistical analysis

The results are presented as the mean \pm SEM and were analyzed using SPSS (Statistical Package for the Social Sciences) 13.0 software (SPSS Incorporated, Chicago, IL, USA). Nine replicate samples were used for each condition. The differences among the groups were compared using independent sample Student's t-tests. A p value lower than 0.05 was considered statistically significant (labeled with one star), and a p value lower than 0.01 was considered highly significant (labeled with two stars).

Results

Effects of the duration of BHBA treatment on $I\kappa B\alpha$ phosphorylation in bovine hepatocytes (time course experiment)

To confirm the optimum time of the BHBA treatment in this study, the protein expression level of IkB α was detected by western blotting. The results showed that the phosphorylation level of IkB α (p-IkB α /IkB α) was higher at 6, 9, 12 and 24 h in the BHBA-treated groups, with the highest level at 24 h compared with the control group (Fig. 1 A, B).

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Table 2. The content of oxidation markers and the activity of antioxidation markers. Oxidation and antioxidation markers in calf hepatocytes. Hepatocytes were treated with 0, 0.6, 1.2, 1.8 and 2.4 mM BHBA. The intracellular levels of the oxidation markers MDA and NO and the antioxidation markers GSH-Px, SOD and CAT were measured using commercial kits. The data are shown as the mean \pm SEM. a, b and c, the same letter indicates no significant difference (P>0.05), different letters mean a significant difference (P<0.05/P<0.01)

markers/ BHBA concentration	MDA (nmol/mg)	NO (μmol/L)	GSH-Px (mU/mg)	SOD (U/mg)	CAT (U/mg)
0 mM	4.1119±0.525a	72.41±3.4881a	1800±13.113ª	18.67±0.5038ª	14.53±0.1503a
0.6 mM	5.0413±0.608 ^a	77.52±3.1272a	1780±12.176 ^a	17.89±0.5769a	14.38±0.1576 ^a
1.2 mM	6.901±0.6899b	88.56±2.265b	1626±11.293b	15.27±0.4093b	13.84±0.1409b
1.8 mM	8.532±0.4876°	93.67±6.573b	1587±19.203b	14.78±0.4208b	13.72±0.1420b
2.4 mM	9.157±0.4291°	89.29±6.328 ^h	1526±12.192°	14.02±0.4073 ^h	13.53±0.1407¢

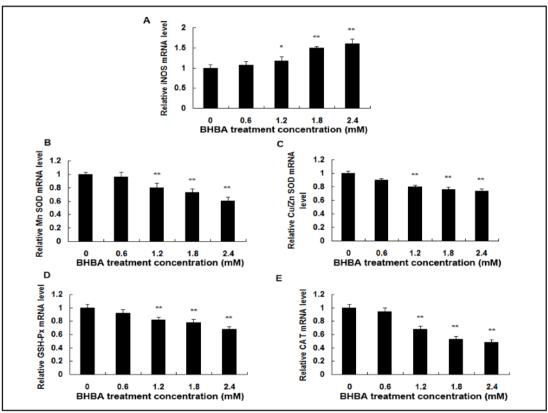


Fig. 2. Effects of different concentrations of BHBA on the oxidation markers MDA and NO and iNOS and the antioxidation markers SOD, CAT and GSH-Px in calf hepatocytes. Hepatocytes were treated with 0, 0.6, 1.2, 1.8 and 2.4 mM BHBA. (A) The mRNA level of iNOS. (B) The mRNA level of Mn SOD. (C) The mRNA level of Cu/Zn SOD. (D) The mRNA level of GSH-Px. (E) The mRNA level of CAT. The data are shown as the mean ± SEM. *, statistical significance (P<0.05). **, statistical significance (P<0.01). The symbol * and ** indicate statistical differences with the control group (no BHBA).

Effects of different concentrations of BHBA on the oxidation markers MDA, NO and iNOS and the antioxidation markers SOD, CAT and GSH-Px (dose response experiment)

The quantities of MDA and NO showed a significant upward trend in all the BHBA-treated groups compared with the control, and the iNOS mRNA expression showed a similar result (Table 2 and Fig. 2 A). Moreover, the mRNA expression and activity of SOD (Mn SOD and Cu/Zn SOD), CAT and GSH-Px showed a significant downward trend in all the BHBA-treated groups (Table 2 and Fig. 2 B, C, D, E).

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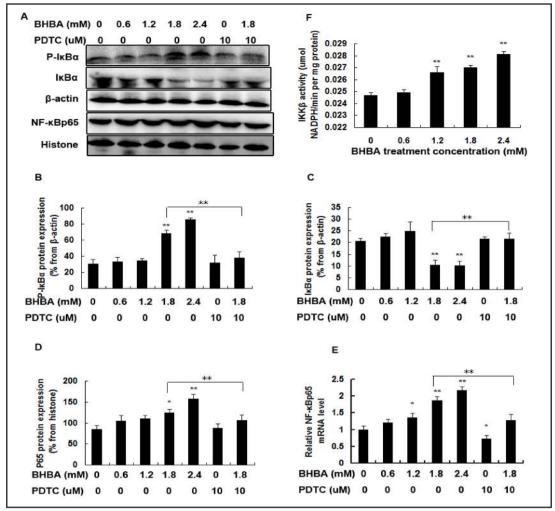


Fig. 3. BHBA activates the NF- κ B signaling pathway in calf hepatocytes. Hepatocytes were treated with 0, 0.6, 1.2, 1.8 and 2.4 mM BHBA, 10 μ M PDTC and 10 μ M PDTC + 1.8 mM BHBA (PDTC + BHBA). (A) Western blots of β-actin, I κ B α , p-I κ B α , NF- κ B p65 and histone. (B-C) Relative protein expression levels of I κ B α and p-I κ B α normalized to β-actin. (D) Relative protein expression levels of p65 normalized to histone. (E) The mRNA levels of p65. (F) The activity of IKK β in response to different concentrations of BHBA. The data are shown as the mean ± SEM. *, statistical significance (P<0.05). **, statistical significance (P<0.01). The symbol * and ** indicate statistical differences with the control group (no BHBA and no PDTC), and the two groups which marked with horiziontal line are statistical differences.

BHBA activates the NF-κB pathway in boyine hepatocytes

The NF- κ B pathway is important for the inflammatory response and immunity. To determine the effect of BHBA on the NF- κ B pathway, the activities of IKK β , I κ B α protein and p65 were measured after the BHBA and PDTC treatments using the activity assay kit, western blotting, real-time PCR, EMSA and immunocytofluorescence. The results showed that the activity of IKK β increased in a BHBA-dependent manner and peaked in the 2.4 mM BHBA treatment group (Fig. 3 F). Similarly, the phosphorylation levels of I κ B α also increased in a BHBA-dependent manner and were significantly higher in the 1.8 mM and 2.4 mM BHBA-treated groups than in the control group but were significantly lower in the PDTC + BHBA group than in the 1.8 mM BHBA-treated group (Fig. 3 A, B, C). Additionally, the mRNA and nucleoprotein levels, as well as the transcription activity of p65, gradually increased after the BHBA treatment, with the highest levels in the 2.4 mM BHBA-treated group, but they significantly decreased in the PDTC + BHBA group compared with the 1.8 mM BHBA-treated

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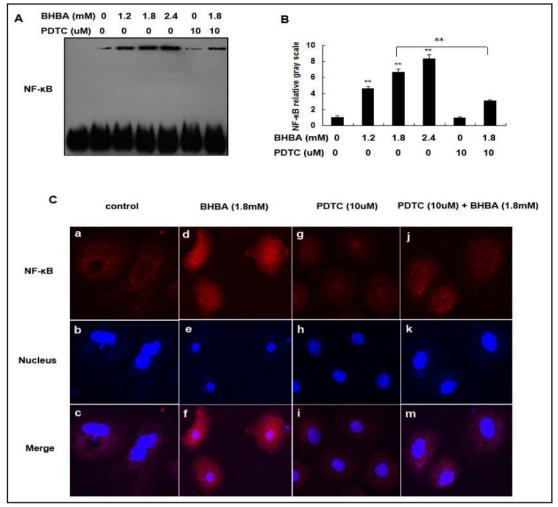


Fig. 4. Transcriptional activity and immunofluorescence staining of NF- κ B p65 in calf hepatocytes. Hepatocytes were treated with 0, 1.2, 1.8 and 2.4 mM BHBA, 10 μ M PDTC and 10 μ M PDTC + 1.8 mM BHBA (PDTC + BHBA). (A) EMSA result of NF- κ B. (B) Relative gray scale of NF- κ B. (C) Immunofluorescence staining of NF- κ B p65 (a, d, g and j) in calf hepatocytes. Images (c, f, i and m) were merged with Hoechst 33258 staining images (b, e, h and k). The data are shown as the mean ± SEM. *, statistical significance (P<0.05). **, statistical significance (P<0.01). The symbol * and ** indicate statistical differences with the control group (no BHBA and no PDTC), and the two groups which marked with horiziontal line are statistical differences.

group (Fig. 3 A, D, E; Fig. 4 A, B). To further confirm that p65 translocates to the nucleus in bovine hepatocytes after BHBA treatment, immunocytofluorescence was performed. Figure 4 C showed a similar trend as the EMSA results. Overall, these results indicate that a certain concentration of BHBA can activate the NF-kB pathway in a dose-dependent manner in cattle hepatocytes.

Effects of BHBA on the pro-inflammatory factors TNF- α , IL-6 and IL-1 β in bovine hepatocytes

The results demonstrated that the mRNA expression of TNF- α , IL-6 and IL-1 β showed an increasing trend after the BHBA treatment and were significantly higher in the 1.8 mM and 2.4 mM BHBA-treated groups than in the control group but showed a decreasing trend in the PDTC + BHBA group compared with the 1.8 mM BHBA-treated group (Fig. 5 A, C, E). The quantities of TNF- α , IL-6 and IL-1 β showed a similar result, which further supports the results of the real-time PCR (Fig. 5 B, D, F). These results suggest that BHBA can significantly

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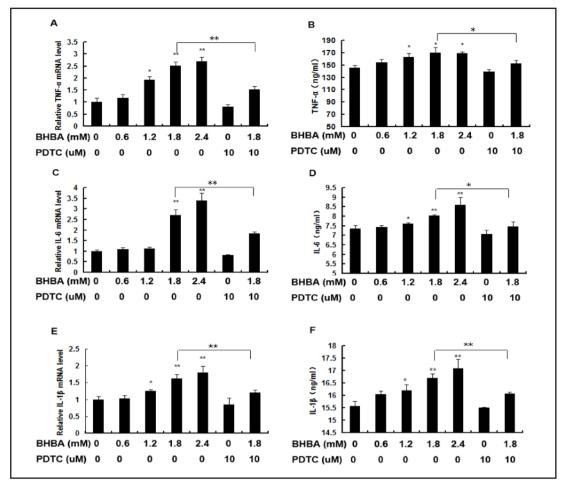


Fig. 5. BHBA-activated NF- κ B signaling pathway activates the expression of pro-inflammatory factors in calf hepatocytes. Hepatocytes were treated as described in Figure 3. (A) The mRNA level of TNF- α . (B) ELISA of TNF- α . (C) The mRNA level of IL-6. (D) ELISA of IL-6. (E) The mRNA level of IL-1 β . (F) ELISA of IL-1 β . The data are shown as the mean \pm SEM. *, statistical significance (P<0.05). **, statistical significance (P<0.01). The symbol * and ** indicate statistical differences with the control group (no BHBA and no PDTC), and the two groups which marked with horiziontal line are statistical differences.

upregulate the mRNA expression of pro-inflammatory factors and increase the release of pro-inflammatory factors in bovine hepatocytes. Taken together, these results demonstrate that BHBA can promote the expression of pro-inflammatory factors by activating the oxidative stress-medicated NF-κB pathway in cattle hepatocytes.

Effects of NAC on the pro-inflammatory factors TNF- α , IL-6 and IL-1 β in bovine hepatocytes In order to identify that pro-inflammatory responses are induced by oxidative stress, the quantities of TNF- α , IL-6 and IL-1 β were measured by ELISA after the BHBA and NAC treatments. The results demonstrated that the quantities of TNF- α , IL-6 and IL-1 β significantly decreased in the NAC + BHBA group compared with the 1.8 mM BHBA-treated group (Fig. 6 A, B, C).

Discussion

High concentrations of BHBA can induce post-partum diseases, poorer reproduction and lower milk production in transition dairy cattle [2]. In the present study, we have shown

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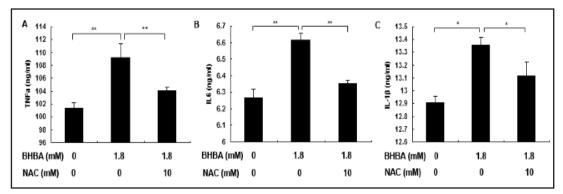


Fig. 6. NAC suppressed the expression of pro-inflammatory factor in calf hepatocytes. Hepatocytes were treated with 0, 1.8 mM BHBA, 10 mM NAC + 1.8 mM BHBA (NAC + BHBA). (A) ELISA of TNF- α . (B) ELISA of IL-6. (C) ELISA of IL-1 β . The data are shown as the mean \pm SEM. *, statistical significance (P<0.05). **, statistical significance (P<0.01). The symbol * and ** indicate that the two groups which marked with horizontal line are statistical differences.

that high concentrations of BHBA could increase the expression and synthesis of the proinflammatory factors TNF- α , IL-6 and IL-1 β in primary calf hepatocytes. These results indicate that the inflammatory response is stimulated by high concentrations of BHBA in hepatocytes, which is in agreement with some studies that reported such a positive influence of elevated concentrations of BHBA on inflammation in endometritis and mastitis in dairy cattle [10, 11, 24], supporting our results that BHBA can raise pro-inflammatory factors in hepatocytes.

The NF-κB transcription factor family is composed of the most critical regulators of the immediate transcriptional responses in inflammatory situations [25]. IκBα phosphorylation is dependent on IKKβ activity, which is important for the release of active NF-κB [16]. To further demonstrate that the release of the pro-inflammatory cytokines TNF-α, IL-6 and IL-1β after BHBA treatment may occur through the NF-κB pathway, the activity of IKKβ, phosphorylated IκBα and NF-κB p65 were detected in BHBA-treated cattle hepatocytes. In this study, we demonstrated that BHBA increased IKKβ activity, induced IκBα phosphorylation and promoted the transfer of NF-kB p65 to the nucleus. However, in the PDTC + BHBA group, p-IκBα and NF-κB p65 were inhibited. Taken together, these results demonstrate that BHBA can activate the NF-kB pathway in cattle hepatocytes. Here, we also demonstrated that the mRNA expression and release of TNF-α, IL-6 and IL-1β were markedly inhibited in the PDTC + BHBA group, which further demonstrated that BHBA could activate the NF-κB pathway to regulate the release of the pro-inflammatory factors TNF- α , IL-6 and IL-1 β in cattle hepatocytes. IL-6 was markedly increased in cows with ketosis and played a central role in the impairment of normal liver functions of transition cows [26]. IL-1β was markedly increased in xanthine oxidase-induced liver injury in type I diabetic rats [17]. TNF- α could promote liver triglyceride accumulation and increase the risk of fatty liver in late-lactating Holstein cows [27]. These findings indicate that greater concentrations of BHBA activate the NF- κ B pathway to increase the expression of TNF- α , IL-6 and IL-1 β , thereby inducing hepatocyte injury. Thus, we speculate that high concentrations of BHBA can induce liver injury by activating the NF-κB signaling pathway-mediated expression of pro-inflammatory factors in dairy cattle with ketosis.

NF- κ B activation is highly dependent on oxidative stress, which is associated with the immune response [17, 28]. An imbalance in oxidants/antioxidants, an excess of oxidants and/or a depletion of antioxidants, can lead to oxidative stress [21, 29]. Oxidative stress is not a classical disease and does not exhibit a specific clinical symptoms, it is one of the basics of some metabolic diseases such as ketosis. Dairy cows undergoing ketosis experience oxidative stress which can be measured in biological fluids and tissues [5, 30, 31]. MDA is one of the final products of polyunsaturated fatty acid peroxidation, which is caused by

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increasing level of reactive oxygen species (ROS) and is commonly known as a marker of oxidative damage [32]. NO acts intracellularly, and excess NO can cause oxidative stress [33]. iNOS mainly catalyzes NO formation, which can induce oxidative stress [34]. Superoxide dismutases (SOD: Mn SOD and Cu/Zn SOD) and catalase (CAT) are two important antioxidant enzymes involved in the removal of ROS [35, 36]. GSH-Px is involved in catalyzing the GSHdependent reduction of hydrogen peroxide [36]. In this study, we have demonstrated that BHBA increased the levels of MDA, iNOS and NO. Moreover, the levels of SOD, CAT and GSH-Px were significantly decreased in the BHBA-treated hepatocytes.

These results demonstrate that BHBA can increase the oxidants status and decrease the antioxidants status. Indeed, as it was shown, greater concentration of BHBA could lead to the higher levels of indices of oxidative status: reactive oxygen metabolites (ROM) and thiobarbituric acid reactive substances (TBARS), and lower levels of antioxidants in the plasma of dairy cows thereby leading to oxidative stress [5]. Pedernera et al. [21] measured indicators of energy balance (non-esterified fatty acids, β-hydroxybutyrate, glucose and insulin-like growth factor-1) and indicators of oxidative stress (reactive oxygen metabolites and biological antioxidants) in the first 5 weeks of lactation. They found that oxidative stress was related to NEB in dairy cow. Additional reports found that ketone bodies served as oxidative fuel and lipogenic precursors in cells, oxidative fuel could improve mitochondrial respiration and ATP production, and then increase mitochondria piece production [8, 37]. Excessive fat mobilization might impair liver function due to intracellular lipid accumulation [38], and cows with hepatomegaly have reduced capacity to produce antioxidants [39]. These reports supported our findings that BHBA could induce oxidative stress in the hepatocytes of dairy cattle. Additionally, Czaja [18] reported that the common mechanism of liver injury was oxidative stress. This study has shown that BHBA activated NF-κB-induced hepatocyte injury by regulating the expression of TNF- α , IL-6 and IL-1 β in hepatocytes. Additionally, we found that the quantities of pro-inflammatory factors TNF- α , IL-6 and IL-1 β were significantly reduced by treating for antioxidant NAC in cattle hepatocytes. It demonstrated that the pro-inflammatory responses were induced by oxidative stress. Therefore, we infer that oxidative stress most likely plays a role in modulating the inflammatory response through the activation of the NF-κB pathway. Indeed, the important oxidative marker MDA had been reported to correlate significantly with IκBα phosphorylation and p65 NF-κB migration to the nucleus [40]. However, the particular mechanism of how oxidative stress activates the NF-κB pathway in dairy cattle is worthy of further study.

The finding that BHBA-induced oxidative stress activates the NF-κB pathway in cattle hepatocytes is novel. However, how BHBA induces the oxidative stress in cattle hepatocytes is unknown. Additionally, whether other signal pathways are involved in inducing inflammation or whether NF-κB has another effect on cattle hepatocytes are also unknown. Therefore, future studies should aim to confirm the specific mechanism of BHBA-induced oxidative stress and confirm the relationship between NF-κB and other signaling pathways in cattle hepatocytes.

Conclusions

Taken together, these results indicate that BHBA can induce the activation of the NFκΒ signaling pathway in cattle hepatocytes by the upregulation of oxidative markers (iNOS, MDA and NO) and downregulation of antioxidative markers (SOD, CAT and GSH-Px). NF-κB induces hepatocyte injury by translocating to the nucleus and increasing the DNA binding activity to increase the transcription of the pro-inflammatory factors TNF-α, IL-6 and IL-1β. The current study identifies a mechanism by which BHBA can regulate inflammatory injury in the hepatocytes of dairy cattle.

Conflict of Interest

None to declare.

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