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Anti-Inflammatory Effects of Benfotiamine are Mediated Through the Regulation of Arachidonic Acid Pathway in Macrophages

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Abstract

Benfotiamine, a lipid-soluble analogue of vitamin B1, is a potent anti-oxidant that is used as a food supplement for the treatment of diabetic complications. Our recent study indicates a novel role of benfotiamine in the prevention of bacterial endotoxin, lipopolysaccharide (LPS)-induced cytotoxicity and inflammatory response in murine macrophages. Nevertheless, it remains unclear how benfotiamine mediates anti-inflammatory effects. In this study, we investigated the antiinflammatory role of benfotiamine in regulating the arachidonic acid (AA) pathway generated inflammatory lipid mediators in RAW 264.7 macrophages. Benfotiamine prevented the LPSinduced activation of cPLA2 and release of AA metabolites such as leukotrienes (LTB4), prostaglandin E2 (PGE2), thromboxanes 2 (TXB2) and prostacyclin (PGI2) in macrophages. Further, LPS-induced expressions of AA metabolizing enzymes such as COX-2, LOX-5, TXB synthase and PGI2 synthase were significantly blocked by benfotiamine. Furthermore, benfotiamine prevented the LPS-induced phosphorylation of ERK1/2 and expression of transcription factors NF-kB, and Egr-1. Benfotiamine also prevented the LPS-induced oxidative stress and protein-HNE adducts formation. Most importantly, as compared to specific COX-2 and LOX-5 inhibitors, benfotiamine significantly prevented the LPS-induced macrophage death and monocytes adhesion to endothelial cells. Thus, our studies indicate that the dual regulation of COX and LOX pathways in AA metabolism could be a novel mechanism by which benfotiamine exhibits its potential anti-inflammatory response.

Keywords

Vitamin B1; inflammation; arachidonic acid; Cox and Lox

Introduction

Overwhelming inflammatory responses accompanied by deregulated immunological and coagulation functions results in multiple organ failure leading to septic shock a major cause of mortality [1]. Although, treatment with the antibiotics kills bacteria and impedes the infection, the bacterial debris which contains large amounts of endotoxins could trigger the inflammatory responses [2]. Bacterial lipopolysaccharide (LPS), the structural component of

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the Gram-negative bacterial outer cell wall, is a potent initiator of inflammatory response during most commonly seen bacterial infections [3]. Binding of LPS to its cognate CD14 receptor on the monocyte/macrophage cell membranes induces the release of various proinflammatory cytokines and chemokines [4-6], which are implicated in the pathogenesis of the major inflammatory complications such as sepsis [7]. Although a number of nonsteroidal anti-inflammatory drugs (NSAID) and antioxidants recently emerged as potential therapeutic targets for intervention of sepsis complications [8], their success rate at the translational studies is low [9-11]. Therefore, identification of novel therapeutic compounds with both anti-oxidative and anti-inflammatory properties could yield potential therapeutic approaches to sepsis and other related inflammatory complications.

Thiamine (vitamin B1) has been used for decades for the treatment of several disorders such as neurological, diabetic, and cardiovascular complications [12-15]. Since, thiamine is water-soluble, it cannot be stored in the body for so long and is rapidly excreted in the urine. The bioavailability of orally administered thiamine salt is comparatively low when compared with fat-soluble analogs [16-18]. In 1954, variety of lipid-soluble thiamine derivatives were discovered, subsequently they were named as allithiamines since they belong to Allium family of vegetables such as crushed garlic, onions and leeks etc [19]. These lipid-soluble thiamine derivatives have much higher absorption and bioavailability than water-soluble thiamine salts [20, 21]. Benfotiamine, one of the lipid-soluble derivatives of vitamin B1 was found to be the most effective antioxidant and was developed to improve bioavailability for pharmacological administration [22, 23]. Benfotiamine contains an open thiazole ring, which allows it to pass through the cell membrane, and the open ring is closed by undergoing reduction reaction making biologically active thiamine [21]. Oral administration of benfotiamine raises thiamine levels in blood and tissues to a much higher degree than the water-soluble salts [20, 21]. We have recently shown that benfotiamine prevents LPS-induced NF-kB -dependent expression of inflammatory cytokines and chemokines in macrophages. We have also shown that benfotiamine prevents LPS-induced generation of reactive oxygen species (ROS) and macrophage death in vitro [24]. Further, we have also demonstrated that benfotiamine prevents LPS-induced ocular inflammatory complication, uveitis, in rats [25]. Thus, our previous studies identified benfotiamine as a novel anti-inflammatory compound. However, the molecular mechanisms through which benfotiamine exert its anti-inflammatory effect is not clear.

Recent studies indicate that LPS-stimulated macrophages lead to the activation of phospholipase A2 (cPLA2) which causes the release of AA from the cell membrane [26-28]. AA (20:4, n-6) is esterified in glycerol in the lipid bilayer of cell membrane at sn-2 position [26]. Once AA is released it is further metabolized via COX-2 and LOX-5 pathways. COX-2 catalyzed AA pathway leads the generation of prostanoids such as prostaglandins (PGs), prostacyclins (PGI2), and thromboxanes (TXBs) which have been shown to be most potent inflammatory mediators. LOX-5 catalyzed AA pathway result in the generation of leukotrienes (LTBs) involved in several inflammatory diseases such as asthma pathogenesis [29, 30]. Even though, the specific inhibitors of COX-2 and LOX-5 have been shown to be anti-inflammatory, they are not much effective, since they only stunt either COX or LOX pathway moving AA metabolism towards the other pathway [31]. Therefore, in order to identify a possible mechanism by which benfotiamine acts as an anti-inflammatory agent, in this study, we have examined the efficacy of benfotiamine in regulating the AA pathway generated inflammatory lipid mediators in RAW264.7 macrophages. Our results suggest that benfotiamine drastically inhibited the release of AA metabolites by inhibiting the activation of cPLA2. Benfotiamine also prevented the LPS-induced COX-2 and LOX-5 enzymes and generation of PGE2, TXB and LTB4. Further, benfotiamine also prevented the LPS-induced protein-HNE adducts and lipid hydroperoxides formation in macrophages. Thus, our results indicate that benfotiamine could be a novel anti-inflammatory agent which can ameliorate

endotoxin induced inflammation by preventing both COX and LOX pathways in the AA metabolism.

Materials and Methods

Materials

Dulbecco's modified Eagle's medium with 4 mM l-glutamine and 4.5 g/l glucose, phosphate-buffered saline (PBS), penicillin/streptomycin solution, trypsin, and fetal bovine serum (FBS) were purchased from Invitrogen Gibco (Grand Island, NY). [5, 6, 8, 9, 11, 12, 14, 15-³H (N)] arachidonic acid was purchased from NEN Life Science (Boston, MA). RIPA cell lysis buffer, antibodies against LOX-5, PGI₂ synthase and β-actin were purchased from Santa Cruz biotechnology, (Santa Cruz, CA) USA. Antibodies against cPLA₂ were obtained from Cell Signaling Inc. (Beverly, MA). Antibodies against Thromboxane synthase and COX-2 were obtained from Abcam Inc. (Cambridge, MA). Thromboxane B2 and leukotriene B₄ EIA kits and REV-590 were obtained from Cayman chemical Inc. (Ann Arbor, MI). Prostaglandin E₂ (PGE₂) and prostacyclin (PGI2) assay kits were obtained from Assay Designs, Inc (Ann Arbor, MI). Benfotiamine, lipopolysaccharide (*Escherichia coli* 0111:B4), indomethacin and the reagents used in Western blot analysis were obtained from Sigma (St. Louis, MO). All other reagents used were of analytical grade.

Cell Culture

RAW264.7 macrophages obtained from American Type Culture Collection (Manassas, VA) were grown in DMEM containing 10% FBS, 1% penicillin/streptomycin in a 95% air 5% CO₂-humidified atmosphere at 37°C. Macrophages were pretreated with benfotiamine with either 50 or 100 μ M or carrier for overnight in serum-free medium and subsequently stimulated with 1 μ g/ml LPS for indicated time periods.

Release of arachidonic acid and its metabolites in RAW 264.7 cells culture medium

RAW264.7 cells were seeded in 12-well plate at the density of ~ 0.35×10^{6} cells/well, without or with benfotiamine (100 µM) in the complete growth media. Media were removed and replaced with 1 ml of serum-free DMEM containing 0.1 µCi/ml ³H-arachidonic acid and incubated for 16 h. The cells were washed twice with phosphate-buffered saline (PBS) containing 0.1% bovine serum albumin to remove unadsorbed arachidonic acid and stimulated with 1 µg/ml LPS for 1 h. Subsequently, the cells were incubated with fresh medium containing benfotiamine (100 µM) for additional 18 h. The culture medium was collected and centrifuged for 15 min at 10,000 rpm and supernatant was used to measure the radioactivity using a Beckman liquid scintillation counter (Beckman Coulter, Fullerton, CA).

Western Blot Analysis

The confluent macrophages were incubated without or with benfotiamine (50 or 100 μ M), followed by treatment with 1 μ g/ml LPS for 18 h. The cells were washed twice with PBS and lysed in an ice-cold RIPA lysis buffer. The crude lysates were cleared by centrifugation at 12,000g for 10 min at 4°C. Equal amounts of cell lysates (30-50 μ g) were separated on 10 % SDS-PAGE, and transferred to polyvinylidene difluoride membranes (Immobilon; Millipore, Bedford, MA). The membranes were incubated in blocking solution containing 5% w/v dried fat-free milk and 0.1% v/v Tween-20 in Tris-buffered saline. Subsequently, the membranes were incubated with specific antibodies against cPLA2, COX-2, LOX-5, TXB synthase and PGI2 synthase. The blots were then washed, exposed to HRP-conjugated secondary antibodies (1:5,000 dilution) for 1 h, and the antigen-antibody complex was detected by enhanced chemiluminescence (Pierce, Rockford, IL). The membranes were

stripped and probed with antibodies against glyceraldehyde-3-phosphate dehydrogenase (GAPDH) to ensure equal protein loading. Fold changes in the band intensities were quantified by densitometry analysis by using Kodak Image Station software.

Determination of PGE2, 6k-PGF_{1α} (PGI2), TXB2, LTB4 and cPLA2 levels

The RAW cells (~ 0.3×10^6 cells/well in 6-well plates) were growth-arrested in the serumfree medium without or with benfotiamine (100 μ M) followed by incubation with 1 μ g LPS/ ml for another 18 h. The medium was collected from each well and cleared by centrifugation (5000 rpm; 5 min). The levels of PGE2 and PGI2 were determined using respective assay kits according to the manufacturer's instructions (Assay Designs, Inc.). For determining the levels of TXB2, LTB4 and cPLA2 specific ELISA kits were used according to the manufacturer's instructions (Cayman chemical).

Reverse transcription-PCR

The RAW264.7 cells were pre-incubated without and with benfotiamine for overnight followed by incubation with 1 µg/ml LPS for additional 6 h. Total RNA was isolated by using RNeasy® Mini Kit from Qiagen, USA. Equal amount of RNA was used for RT-PCR using Qiagen® one step RT-PCR kit. The gene specific primer (purchased from sigma; Sigma Genosys, USA) sequences were; COX-2, sense: 5'-GGT TAC AAA AGC TGG GAA GC-3', anti-sense: 5'-GGG GGT TCC GAG TTT ATA CT-3'. LOX-5, sence: 5'-GGC ACC GAC GAC TAC ATC TAC-3', anti-sense: 5'-CAA TTT TGC ACG TCC ATC CC-3'; TXB synthase, sense: 5'-ATC AGC CAA GCC TGT GAA CT-3', anti-sense: 5'-CCT CTC TTC TGC TGC TTG CT-3' and GAPDH, sense: 5- TCC ACT CAC GGC AAA TTC AAC G-3', anti-sense 5'- TAG ACT CCA CGA CAT ACT CAG C-3'. The PCR reaction was carried out in a GeneAmp 2700 thermocycler (Applied Biosystems, Foster City, CA) under the following conditions: initial denaturation at 95°C for 15 min; 30 cycles of 94°C 30 s, 62°C 30 s, 72°C 1 min, and then 72°C 5 min for final extension. PCR products were electrophoresed with 1 % agarose-1xTAE gels containing 0.5 µg/ml ethidium bromide. The band densities were analyzed using Kodak image station.

Determination of Lipid hydroperoxides

The lipid hydroperoxides were determined colorimetrically by using a lipid hydroperoxides assay kit obtained from Cayman chemicals as per the supplier's instructions. Briefly, lipid hydroperoxides kit measures the hydroperoxides directly utilizing the redox reactions with ferrous ions. Lipid hydroperoxides were extracted from the samples into chloroform. To 450µl of chloroform-methanol solvent and 50µl of chromogen was added, kept at room temperature for 5 min, and absorbance was read at 500 nm using a 96 well plate reader.

Protein-HNE Adducts formation

RAW264.7 cells were seeded on glass chamber slides without or with benfotiamine (100 μ M) followed by incubation with 1 μ g LPS/ml for 18 h. The slides were fixed with 4% paraformaldehyde for 30 min. After the slides were thoroughly washed with Tris-buffered saline (TBS) containing 0.1% Triton X-100, they were incubated with anti-HNE antibodies for overnight at 4° C followed by washing and incubation with 1:160 diluted secondary antibodies (goat anti-rabbit IgG conjugated with FITC) for 1 h. The slides were mounted and photographs were taken with a Nikon fluorescence microscope.

Macrophage viability and U-937 cell adhesion to HUVECs

The RAW cells (0.3×10^6 cells/well in 6-well plates) were growth arrested in the serum-free medium without or with benfotiamine (100 μ M), indomethacin (COX-2 inhibitor, 10 μ M) and REV 5901 (LOX-5 inhibitor, 10 μ M) for overnight followed by incubation with 1 μ g/ml

LPS for another 24 h. Cell viability was determined using Millipore cell counter. For cell adhesion HUVECs were plated in 96-well plate, grown to 70% confluence, and treated without or with benfotiamine, indomethacin and REV 5901 for overnight in the absence and presence of 1 ug/ml of LPS. Cells were rinsed twice with serum-free medium and U-937 cells were added to each well followed by incubation with 1 µg/ml LPS for additional 12 h. MTT assay was performed to determine cell adhesion at recoding the absorbance at 562 nm.

Statistical analysis

Data presented as mean \pm SE (n=6) and p-values were determined by unpaired Student's t test using Microsoft Office Excel 2007 software. P<0.05 was considered as statistically significant.

Results

Benfotiamine prevents release of arachidonic acid metabolism in LPS-induced macrophages

To investigate the anti-inflammatory effect of benfotiamine against LPS-induced toxicity; at first, we have examined metabolism of AA in macrophages. Macrophages were incubated with ³H-AA followed by stimulation with LPS with or without benfotiamine. Results shown in the Fig. 1 indicate that in the LPS treated macrophages the release of pre-labeled AA and its metabolites were significantly increased to ~5-fold as compared to un-stimulated macrophages. Benfotiamine significantly (~90%) prevented the LPS-induced formation of AA metabolites. Thus, our studies indicate that benfotiamine prevents LPS-induced AA metabolism in macrophages.

Benfotiamine prevents LPS-induced activation of cPLA2 in macrophages

Phospholipase A2, especially cPLA2 is well known to mediate the release of AA from membrane phospholipids; we next examined the potential of benfotiamine in preventing the activation of cPLA2 in LPS-induced macrophages. Stimulation of macrophages with LPS caused an approximately 2-fold increase in the cPLA2 activity and pre-incubation of macrophages with benfotiamine significantly (>70%) prevented the LPS-induced activity of cPLA2 (Fig 2A). Further, the results shown in Fig 2B indicate that LPS-induced approximately 2-fold increase in cPLA2 expression. The LPS-induced increase in the expression of cPLA2 was significantly (>90%) prevented by benfotiamine. Thus, our results suggest that benfotiamine prevents LPS-induced AA metabolism by preventing the expression of cPLA2.

Effect of benfotiamine on LPS-induced production of AA pathway- generated lipid mediators in RAW264.7 macrophages

Since PGs, TXBs, LTBs and 6k-PGF1 α are AA pathway-generated lipid mediators which play a major role in inflammatory complications, we next examined the effect of benfotiamine on LPS-induced generation of PGE2, TXB2, 6k-PGF1 α and LTB4 in cell culture media. LPS caused >3-fold and >20-fold increase in the levels of PGE2 and TXB2, respectively, in the macrophages as compared to untreated macrophages. The increases in PGE2 and TXB2 levels were significantly brought down by benfotiamine (Fig. 3A and B). Since, PGI2 has a very short half-life and is readily converted to 6k-PGF1 α , we next determined 6 k-PGF1 α levels in RAW264.7 macrophages. Macrophages treated with LPS showed a significant (>4 fold) increase in the levels of 6k-PGF1 α which was significantly prevented by benfotiamine (Fig. 3 C). Further, we have also measured the effect of benfotiamine on LPS-induced LTB4 levels in macrophages. As shown in Fig. 3D, RAW264.7 cells stimulated with LPS showed a >3 fold increase in the level of LTB4 which was significantly (>60%) reduced in cells treated with benfotiamine. These results suggest that benfotiamine could efficiently prevent LPS-induced production of AA pathway-generated inflammatory metabolites in RAW264.7 macrophages.

Benfotiamine prevents LPS-induced activation of AA pathway enzymes in macrophages

Since PGs, TXs, LTBs and 6k-PGF1a are catalyzed by COX-2, TXB synthase, PGI2 synthase and LOX-5, respectively in the AA metabolism pathway, we next examined the effect of benfotiamine on LPS-induced expression of these proteins in macrophages. Our results shown in Fig 4 A-D indicate that LPS-induced approximately 6-fold increase in COX-2, >3 fold in LOX-5, 3-fold in TXB synthase and 2.5 fold in PGI2 synthase expressions in macrophages. The increases in COX-2, LOX-5, TXB synthase and PGI2 synthase were significantly abolished in the presence of benfotiamine by 50%, 95%, 95% and 90%, respectively. Further, to examine the effect of benfotiamine on LPS-induced AA metabolism enzymes at the transcriptional levels, we next measured the effect of benfotiamine on the LPS-induced mRNA levels of AA pathway enzymes. The results shown in the figure 5A-C indicate that LPS caused a significant 6-, 3.6- and 3-folds increase in the levels of COX-2, LOX-5 and TXB synthase mRNA levels, respectively. However, in the macrophages pretreated with benfotiamine followed by LPS treatment, the increase in the mRNA levels of COX-2, LOX-5 and TXB synthase were significantly (60 to 90%) decreased. Thus, our studies indicate that benfotiamine prevents the expression of AA pathway enzymes at their transcriptional level.

Effect of benfotiamine on phosphorylation of ERK1/2 and expression of transcription factors NF-kB and Egr-1

Since, transcription factors such as NF-kB and Egr-1 are known to regulate the transcriptional regulation of AA pathway enzymes; we next examined the effect of benfotiamine on LPS-induced activation of these transcription factors. Our results shown in the Fig 6 A-C indicate that benfotiamine significantly prevented the LPS-induced activation ERK1/2 and the transcription factors NF-kB and Egr-1 which transcribe gene for COX-2 and mPGES-1 [32]. Thus, our results suggest that benfotiamine prevents LPS-induced expression of AA pathway enzymes by preventing their transcriptional activation.

Effect of benfotiamine on LPS-induced protein-HNE adduct and lipid hydroperoxides formation

Since transcription factors such as NF-kB, Egr-1 are known to be activated under oxidative stress generated reactive oxygen species, we next examined the effect of benfotiamine on LPS-induced oxidative stress. The levels of protein-HNE and lipid hydroperoxides were used as markers for increased oxidative stress. Our results shown in fig. 7A indicate that the benfotiamine ameliorated protein-HNE adduct formation in LPS-induced macrophages. Further, macrophages challenged with LPS showed significantly (>2.5 fold) increased lipid hydroperoxides levels which were significantly (60%) decreased in the macrophages pretreated with benfotiamine (Fig.7B), suggesting that benfotiamine prevents LPS-induced activation of redox sensitive transcription factors by decreasing cellular oxidative stress in macrophages.

Effect of benfotiamine and specific inhibitors of COX-2 and LOX-5 on LPS-induced macrophage viability and U-937 cell adhesion to HUVECs

Since, our previous results indicate that benfotiamine prevents both COX-2 and LOX-5 enzymes; we next examined the dual inhibitory activity of benfotiamine on LPS-induced macrophage death and compared with the specific inhibitors of COX and LOX enzymes. Benfotiamine, indomethacin (COX-2 inhibitor) and REV 5901 (LOX-5 inhibitor) were pre-

incubated with macrophages followed by stimulation with LPS for another 24 h and cell viability was measured by using an automatic electronic cell counter (Scepter, Millipore). LPS-caused a significant cell death and all three inhibitors prevented the LPS-induced cell death (Fig 8A). However, out of all three agents, benfotiamine significantly prevented (~70%) LPS-induced cell death followed by COX-2 inhibitor (40%) and LOX-5 inhibitor (30%). Further, benfotiamine significantly prevented the LPS-induced monocyte adhesion (~70%) to endothelial cells as compared to COX-2 or LOX-5 inhibitors (Fig 8B). Thus, our results indicate that benfotiamine could be a novel anti-inflammatory agent as compared to specific COX-2 and LOX-5 inhibitors, since it could efficiently prevent LPS-induced generation of inflammatory lipid mediators by inhibiting both the COX and LOX enzymes.

Discussion

During Gram-negative bacterial infections, the pathological condition of inflammation is initiated bacterial outer cell wall components, especially endotoxins such as LPS [33]. LPS can directly activate macrophages by triggering the production of pro-inflammatory cytokines such as TNF [34], IL-1, IL-6, IFN, chemokines such as MCP-1, MIP1 [35], leukotrienes [36] and others such as nitric oxide and prostaglandins [37]. Overproduction of these inflammatory markers amplifies the host immune response and leads to inflammatory complications such as sepsis [38]. Therefore, the pharmacological interventions of LPS-induced production of inflammatory mediators by macrophages are essential events to control a variety of inflammatory related disorders. In this study, we have for the first time shown that benfotiamine, a fat-soluble analogue of vitamin B1, modulates the production of AA pathway generated inflammatory lipid mediators.

Several studies indicate that the use of natural anti-inflammatory products provides an attractive and safe alternative to modulate various inflammatory disorders [39-41]. Due to its therapeutic action in some frequently observed clinical syndromes, thiamine hydrochloride has been advised and used over a long period of time [42-45]. There are no reports of adverse effects of oral thiamine, even at dosages of several hundred milligrams a day [44]. Benfotiamine, a unique derivative of thiamine is the most potent of the allithiamines, a group of lipid-soluble forms of thiamine [46]. Compared to water-soluble thiamine salts, benfotiamine is absorbed better in the intestine reaching maximum plasma levels of thiamine about 5 times higher and increases active metabolite thiamine diphosphate (ThDP) to 120 folds. The bioavailability is at maximum about 4 times as high as that of thiamine hydrochloride and other lipophilic thiamine derivates [47]. In fact, a recent randomized, double blind, placebo-controlled clinical study in Germany indicated that benfotiamine at a dose of 600 mg/day and higher had no side effects [48]. A number of studies have revealed its potential to alleviate diabetic microangiopathy, neuropathy and other oxidative stressinduced pathological conditions in various experimental animal models [49-53]. However, no reports are available that show benfotiamine supplementation prevents sepsis complications in animal or human studies. We have recently reported that the anti-oxidative and anti-inflammatory properties of benfotiamine in preventing the LPS-induced cytotoxicity in macrophages as well as preventing the ocular inflammation in rats [24, 25]. The present study investigates the potential efficacy of benfotiamine in preventing AA pathway dependent inflammatory signals in macrophages. This is an important advance for the use of benfotiamine as an anti-inflammatory food supplement for treating inflammatory disease conditions.

COX-2 and LOX-5 are the critical enzymes of AA metabolism which catalyzes AA to PGs and LKBs, major lipid mediators involved in various inflammatory diseases [32]. In the present study, we have investigated the effect of benfotiamine on LPS-induced AA pathway generated inflammatory lipid mediators. Our results demonstrate that benfotiamine

significantly prevented the AA pathway generated inflammatory lipid mediators as well as the expression of enzymes such as COX-2 and LOX-5, which catalyze the formation of lipid mediators. Thus, inhibition of COX-2 and LOX-5 appears to be responsible for the decreased biosynthesis of pro-inflammatory PGs and LTB4. Septic patients and animals treated with LPS have increased expression of pro-inflammatory cytokines and PGI2 synthase [54]. Since, PGI2 once formed rapidly gets converted into 6-ketoPGF1, and is involved in vasodilation resulting in reduced blood pressure, and can ultimately lead to sepsis [55]. In this study, benfotiamine significantly prevented 6-ketoPGF1 levels and PGI2 expression suggesting that benfotiamine could efficiently prevent PGI2 -induced vasculopathy upon bacterial infections.

Lipoxygenases catalyze the oxygenation of AA [56]. It is known that leukotrienes are important regulators of cancer cell proliferation and cause apoptosis in non-cancer cells [57-61]. Therefore, we examined whether benfotiamine could regulate the expression and synthesis of LOX-5. Our results demonstrate that LPS-induced expression and synthesis of LOX-5 in RAW264.7 cells were significantly prevented by benfotiamine (fig. 4 and 5B). Our results thus indicate that benfotiamine prevents the expression of two key regulatory enzymes of AA metabolism i.e. COX-2 and LOX-5. Indeed, we observed that as compared to specific COX and LOX inhibitors, benfotiamine is more potent in preventing the LPSinduced macrophage death as well as monocytes adhesion to endothelial cells. During bacterial infections and systemic inflammatory pathologies such as severe sepsis, increased production of ROS causes oxidative stress leading to alterations in protein functions and increased susceptibility to protein degradation [58]. In addition to proteins, polyunsaturated fatty acids such as AA, present in membrane lipids are highly susceptible to ROS-mediated damage [59]. Peroxidation of membrane lipids results in the fragmentation of polyunsaturated fatty acids resulting in the production of various cytotoxic and highly reactive aldehydes, such as malonaldehyde and 4-hydroxy-2-nonenal (HNE) [60]. Since benfotiamine prevents the formation of AA acid metabolites by preventing the activation of cPLA2, we next examined if benfotiamine prevents AA induced generation of HNE. Our results indicate that benfotiamine prevents lipid peroxidation as well as subsequent protein-HNE adducts formation. These results suggest that anti-oxidative properties of benfotiamine could prevent LPS-induced lipid peroxidation and HNE formation.

LPS-induced inflammatory signals are known to mediate through ROS, resulting in the activation of protein kinase C (PKC) and mitogen activated protein kinase (MAPK) [60, 62] which can cause inflammation by activating transcription factor such as NF-kB. It has been shown earlier that various antioxidants such as vitamin C and vitamin E inhibit ROS-induced inflammatory responses by preventing the activation of NF-kB [63,64]. Our current studies indicate that benfotiamine prevents activation of NF-kB as well as Egr-1 and phosphorylation of ERK1/2. Thus prevention of LPS-induced AA metabolizing enzymes and formation of inflammatory lipid mediators by benfotiamine could be mediated regulating redox-sensitive transcription factors.

In summary, this study shows that AA pathway generated inflammatory lipid mediators in response to endotoxin exposure could be prevented by a dietary supplement, benfotiamine. Further, benfotiamine inhibited LPS-induced NF-kB and Egr-1 and expression of key regulatory enzymes of AA metabolism such as cPLA2, COX-2 and LOX-5 in murine macrophages. Our studies indicate that benfotiamine is a more potent anti-inflammatory compound as compared to specific LOX and COX inhibitor. Thus, dual regulation of COX and LOX pathways by benfotiamine could be a novel mechanism by which benfotiamine exhibits its anti-inflammatory response and benfotiamine supplementation could be a potential therapeutic approach to prevent inflammatory complications such as sepsis.

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Abbreviations

AA	Arachidonic acid
LPS	Lipopolysaccharide
PGE2	Prostaglandin E2
PGG2	Prostaglandin G2
PGH2	Prostaglandin H2
PGI2	Prostacyclin
ТХВ	Thromboxane
LTB	Leukotriene
cPLA2	Cytosolic phospholipase A2
sPLA2	soluble phospholipase A2
COX-2	Cyclooxygenase-2
LOX 5	Lipoxygenase-5
МАРК	Mitogen-activated protein kinase
ERK	Extracellular signal-regulated kinase
NF-kB	Nuclear Factor-KappaB
Egr-1	Early growth response factor 1
ROS	Reactive oxygen species
NSAID	Nonsteroidal anti-inflammatory drugs

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Highlights

- Vitamin B1 analogue benfotiamine regulates the arachidonic acid metabolism.
- Benfotiamine prevents LPS-induced TXBs, PGs and LTBs.
- Benfotiamine prevents LPS-induced expression of Cox-2 and Lox-5.
- Benfotiamine prevents NF-kB and EGr-1 in macrophages.
- Benfotiamine could be novel anti-inflammatory agent in the prevention of sepsis.



Figure 1.

Benfotiamine prevents the release of arachidonic acid (AA) metabolites from LPSstimulated RAW264.7 cells. The RAW254.7 cells were incubated with ³H-AA (0.1 µCi/ml) as described in methods and stimulated with 1 µg/ml LPS for 1 h. The cells were washed and incubated for 18 h in a fresh medium containing benfotiamine (100 µM). The release of metabolites in the medium was determined by measuring the radioactivity in the culture medium. Data represent mean \pm S.E. (*n*=6). [#]*p* <0.0001 as compared with AA alone, ^{**}*p*<0.0001 as compared with AA+LPS-treated cells; AA; Arachidonic acid, L; LPS, B; Benfotiamine.



Figure 2.

Benfotiamine prevents LPS-induced generation of cPLA2 in RAW264.7 macrophages. The RAW cells were growth-arrested in a medium containing 0.1% serum without or with benfotiamine (100 μ M) and challenged with 1 μ g/ml LPS. The cPLA2 levels and expression were determined in cell homogenate using the monoclonal enzyme immunoassay kit and western blotting, respectively. Data represent mean ± S.E. (*n*=6). [#]*p*<0.0001 Vs control; ^{**}*p*<0.0001 Vs LPS treated cells. Cont; Control, Ben; benfotiamine.



Figure 3.

Effect of benfotiamine on LPS-induced production of PGE2, LTB4, TXB2 and 6k-PGF1 α and in RAW264.7 macrophages. A-D) The RAW cells were growth-arrested in Dulbecco's modified Eagle's medium containing 0.1% serum with or without benfotiamine (100 μ M) for overnight and challenged with 1 μ g/ml LPS for 18 h. The levels of PGE2, LTB4, TXB2 and 6k-PGF1 α were determined in the culture medium by using the monoclonal enzyme immune assay kits as per manufacturer's instructions. Data represent mean ± S.E. (*n*=6). #*p*<0.0001 Vs control cells; ** *p* <0.0001 Vs LPS-treated cells. Cont; control, Ben; Benfotiamine.



Figure 4.

Benfotiamine prevents LPS-induced expression of COX2, LOX-5, TXB synthase and PGI2 synthase in RAW cells. The RAW cells were growth-arrested by incubating in 0.1% FBS medium without or with benfotiamine (100 μ M), and stimulated with 1 μ g/ml LPS for 18 h. Cytosolic extracts were prepared and equal amounts of cytosolic proteins were subjected to Western blot analysis using antibodies against (A) COX-2, (B) LOX-5 (C) TXB synthase (D) PGI2 synthase and (E) GAPDH. A representative blot from each group is shown. Data represent mean \pm S.E. (*n*=3). $^{\#}p$ <0.0001 Vs control; $^{**}p$ <0.0001 Vs LPS treated cells. B-50 and 100; Benfotiamine 50 and 100 μ M.

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Figure 5.

Effect of benfotiamine on LPS-induced COX2, LOX-5 and TXB synthase at transcriptional level in RAW macrophages. Macrophages were growth-arrested in Dulbecco's modified Eagle's medium containing 0.1% serum with or without benfotiamine (100 μ M) overnight and challenged with 1 μ g/ml LPS for 6 h. The total RNA was isolated and reverse transcriptase-PCR analysis was carried out using specific primers for COX-2, LOX-5 and TXB synthase. Equal amount of PCR products were electrophoresed with 1% agarose-TAE gels containing ethidium bromide. GADPH served as control in reverse transcriptase-PCR analysis. A representative blot from each group is shown. Data represent mean \pm S.E. (*n*=3).* *p*<0.0001 Vs control; ** *p*<0.0001 Vs LPS treated cells. Cont; control, Ben; Benfotiamine.



Figure 6.

Effect of benfotiamine on ERK1/2 phosphorylation and avtivation of p65 and Egr-1. Growth-arrested murine macrophages were treated with LPS without and with benfotiamine (100 μ M) for indicated times at 37°C. Cell lysates were prepared and equal amounts of protein were subjected to western blot analysis using antibodies against A) ERK1/2, B) p65 and C) Egr-1. A representative blot from each group is shown. Data represent mean ± S.E. (*n*=3). # *p*<0.0001 Vs control; ***p* <0.0001 Vs LPS treated cells.



Figure 7.

Effect of benfotiamine on LPS-induced protein-HNE adduct and lipid hydroperoxides formation. Macrophages were growth arrested in 0.1 % FBS and with or without benfotiamine (100µM) subsequently, the cells were exposed to LPS (1µg/ml) for 24 h. (A) Protein-HNE adduct formation was evaluated using rabbit anti-protein-HNE antibodies and FITC-labeled anti-rabbit IgG in goat. A representative blot is shown (200 X magnification). (B) Lipid hydroperoxides were measured using a microplate assay kit according to the manufacturer's instructions. Data represent mean \pm S.E. (*n*=6). *#p*<0.0001 Vs control cells; ***p* <0.0001 Vs LPS treated cells. Cont; control, Ben; Benfotiamine.



Figure 8.

Effect of benfotiamine and specific inhibitors of COX-2 and LOX-5 on LPS-induced macrophage viability and monocyte adhesion to endothelial cells. A) Growth-arrested macrophages were pre-incubated in the presence or absence of benfotiamine (100 μ M), COX-2 inhibitor indomethacin (10 μ M) and LOX-5 inhibitor REV 5901 (10 μ M) for overnight at 37°C. (A) Cell viability was determined by trypan blue exclusion method. (B) Serum-starved HUVECs cells were pretreated with benfotiamine (100 μ M), COX-2 inhibitor indomethacin (10 μ M) and LOX-5 inhibitor REV 5901 (10 μ M), COX-2 inhibitor indomethacin (10 μ M) and LOX-5 inhibitor REV 5901 (10 μ M) for 12 h in the absence and presence of LPS (1 ug/ml). The cells were washed and U-937 monocytes were added to each well and the incubation continued for another 12 h. Cell adhesion assays were performed by MTT assay. Data represent mean \pm S.E. (n=4). $^{\#}p$ <0.001 Vs control; $^{*}p$ <0.01 and $^{**}p$ <0.001 Vs LPS-treated cells.