Anti-inflammatory effects of Daehwangmokdantang, a traditional herbal formulation, in lipopolysaccharide-stimulated RAW 264.7 macrophages

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Abstract. Daehwangmokdantang (DHMDT) is a traditional polyherbal formulation that has known antidiarrheal and anti-inflammatory activities. However, the underlying mechanisms of these activities are poorly understood. In the present study, the inhibitory effects of DHMDT on the production of proinflammatory mediators and cytokines in lipopolysaccharide (LPS)-stimulated RAW 264.7 macrophages were investigated. The inhibitory effects of DHMDT on LPS-induced nitric oxide (NO), prostaglandin (PG)\textsubscript{E}\textsubscript{2}, tumor necrosis factor (TNF)-\textalpha and interleukin (IL)-1\textbeta production were examined using Griess reagent and ELISA detection kits. The effects of DHMDT on the expression of inducible NO synthase (iNOS), cyclooxygenase (COX)-2, IL-1\textbeta and TNF-\textalpha, and their upstream signal proteins, including nuclear factor (NF)-\textkappa B, mitogen-activated protein kinases (MAPKs) and RAC-\textkappa B serine/threonine-protein kinase (Akt), a phosphatidylinositol 3-kinase (PI3K) downstream effector, were investigated using western blotting and immunofluorescence staining. The results revealed the pretreatment with DHMDT significantly inhibited the LPS-induced production of NO, PG\textsubscript{E}\textsubscript{2}, TNF-\textalpha, and IL-1\textbeta, and expression of iNOS, COX-2 TNF-\textalpha, and IL-1\textbeta, without any significant cytotoxicity. DHMDT also efficiently prevented the translocation of the NF-\textkappa B subunit p65 into the nucleus by interrupting the activation of the upstream mediator inhibitor of NF-\textkappa B kinase \textalpha/\textbeta. Furthermore, the anti-inflammatory effects of DHMDT were associated with the suppression of LPS-induced phosphorylation of Akt and MAPKs in RAW 264.7 macrophages. Therefore, the results of the present study indicate that DHMDT exhibited anti-inflammatory activity via the inhibition of proinflammatory mediators and cytokines, in which the inactivation of NF-\textkappa B, PI3K/Akt, and MAPKs may be involved. These results suggest that DHMDT may be a potential anti-inflammatory drug candidate.

Introduction

Inflammation is a complex event involving the activation of physiological and pathological immune system processes (1,2). In the normal state, proinflammatory mediators, such as nitric oxide (NO) and prostaglandin (PG)\textsubscript{E}\textsubscript{2}, and proinflammatory cytokines generated from macrophage cells serve an essential role in host survival and tissue repair (1,2). However, when macrophages are activated by specific stimuli, these proinflammatory mediators and cytokines are overproduced, leading to various inflammatory diseases, such as arthritis, inflammatory bowel disease and asthma (3,4). Therefore, inhibition of macrophage activation has been suggested as a potential therapeutic mechanism for mitigating the progression of inflammatory diseases.

Lipopolysaccharide (LPS)-induced activation of RAW 264.7 macrophages is a well-established model of inflammation, with previous studies demonstrating that LPS, a cell wall component of gram-negative bacteria, activates macrophages to produce proinflammatory mediators and cytokines, thereby mimicking an inflammatory reaction (5,6). Studies have also shown that the LPS-induced stimulation of macrophages initiates the toll-like receptor (TLR)-4/myeloid

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differentiation primary response gene 88 signaling pathway, which involves the activation of nuclear factor (NF)-κB, phosphatidylinositol 3-kinase (PI3K)/RAC-α serine/threonine-protein kinase (Akt) and mitogen-activated protein kinase (MAPK) signaling pathways (7,8). This, in turn, has been indicated to upregulate proinflammatory cytokines and inducible enzymes, such as inducible NO synthase (iNOS) and cyclooxygenase (COX)-2 (9,10).

The transcription factor NF-κB plays a major role in the regulation of genes associated with inflammation (11,12). The activation of NF-κB occurs via phosphorylation and degradation of the suppressor protein inhibitor of NF-κB (IκBα) bound to NF-κB, resulting in the translocation of NF-κB into the nucleus and the promotion and induction of the expression of various inflammatory genes (11,12). TLR4-mediated signaling leads to rapid activation of the PI3K/Akt pathway, which has an important role in the regulation of LPS-induced acute inflammatory responses (5,6). Previous studies have indicated that the PI3K/Akt signaling pathway contributes to TLR4-mediated NF-κB activation and cytokine release (7,8). Studies have also demonstrated that MAPKs, such as extracellular signal-regulated kinase (ERK), c-Jun NH2-terminal kinase (JNK) and p38 MAPK, function as a group of signaling molecules that are critical in relaying inflammatory information (13,14). The TLR4-induced activation of MAPKs has been shown to activate the nuclear translocation of NF-κB and to culminate in the initiation of proinflammatory responses (7,8). Thus, inhibition of the NF-κB, PI3K/Akt and MAPK signaling pathways is critical in combating the actions of anti-inflammatory molecules.

Daehwangmokdantang (DHMDT) is a traditional polyherbal formulation that has been used since ancient times in Korea. As described in the Donguibogam, an ancient Korean medical book published in the early 17th century (15), DHMDT is composed of five medicinal herbs: Paeonia suffruticosa Andr., Prunus persica (L.) Batsch, Trichosanthes kirilowii Maxim, Rheum plamatum L. and mirabilite. DHMDT has been used to treat patients with digestive tract cancers and to prevent diarrhea and inflammation (15). Despite the valuable clinical effects of DHMDT on patients, the molecular mechanism of its pharmacological effects has yet to be elucidated. As part of the present research group’s search for novel biologically active substances from traditional medicinal components were then mixed together in the ratios shown in Table 1 to provide a total weight of 42 g. The mixture was boiled with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h. The extract was diluted with distilled water (42 g/500 ml) for 3 h.

Materials and methods

Materials and reagents. Dulbecco’s modified Eagle’s medium (DMEM) and fetal bovine serum (FBS) were purchased from WelGENE Inc. (Daegu, South Korea). LPS (Escherichia coli Serotype 055:B5), MTT and Griess reagent were obtained from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany). PGE2 (cat. no. KGE004B) tumor necrosis factor (TNF)-α (cat. no. MTA00B) and interleukin (IL)-1β (cat. no. MLB000C) ELISA detection kits were purchased from R&D Systems, Inc. (Minneapolis, MN, USA). Antibodies directed against iNOS (1:1,000; cat. no. sc-509), COX-2 (1:500; cat. no. sc-19999), IL-1β (1:100; cat. no. sc-32294), p65 (1:500; cat. no. sc-109), ERK (1:100; cat. no. sc-154), p38 (1:100; cat. no. sc-535), IKK-α (1:100; cat. no. sc-1643), IKKβ (1:1,000; cat. no. sc-8014), Akt (1:1,000; cat. no. sc-8312), lamin B (1:1,000; cat. no. sc-6216), β-actin (1:1,000; cat. no. sc-69879), goat anti-mouse IgG-horseradish peroxidase (HRP) (1:1,500; cat. no. sc-2005), goat anti-rabbit IgG-HRP (1:1,500; cat. no. sc-2004) and bovine anti-goat IgG-HRP (1:1,500; cat. no. sc-2350) were purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). Anti- TNF-α (1:1,000; cat. no. 3707), phospho-JNK (1:500; cat. no. 9255), JNK (1:1,000; cat. no. 9252S), phospho-ERK (1:500; cat. no. 9106S), phospho-p38 (1:500; cat. no. 9211S), phospho-IKKα/β (1:50; cat. no. 2694) and p-Akt (1:500; cat. no. 9271) were purchased from Cell Signaling Technology Inc. (Danvers, MA, USA). All other chemicals were purchased from Sigma-Aldrich (Merck KGaA).

Preparation of the DHMDT extract. All herbs and mirabilite (natrii sulfas) (Table I) were obtained from Dongeui Oriental Hospital, Dongeui University College of Korean Medicine (Busan, South Korea). Each of the four herbal components of DHMDT and mirabilite were cut into small pieces and the components were then mixed together in the ratios shown in Table 1 to provide a total weight of 42 g. The mixture was boiled with distilled water (42 g/500 ml) for 3 h. The extract was filtered with a 0.45 μM filter to remove insoluble materials. The filtrate was lyophilized and then crushed into a thin powder. The extracts were dissolved in dimethyl sulfoxide (DMSO) to a concentration of 100 mg/ml, and the stock solution was then diluted with DMEM to the required concentration prior to use. The final concentration of DMSO was always ≤0.1%.

Cell culture and LPS stimulation. Murine macrophage-like RAW 264.7 cells were obtained from the Korean Cell Line Bank (Seoul, South Korea) and grown in DMEM, supplemented with 10% FBS, glucose (4.5 g/l), sodium pyruvate (1 mM), L-glutamine (2 mM), penicillin (100 U/ml) and streptomycin (100 μg/ml). The cells were cultured in a humidified atmosphere of 5% CO2 at 37°C.

Table I. Components and their quantities in the Daehwangmokdantang decoction.

| Component (pharmacognostic nomenclature) | Quantity of raw material, g (%) |
|-----------------------------------------|-------------------------------|
| Paeonia suffruticosa Andr. (Moutan Cortex) | 10.0 (23.8) |
| Prunus persica L. Batsch (Persicae Semen) | 10.0 (23.8) |
| Trichosanthes kirilowii Maxim (Trichosanthis Fructus) | 10.0 (23.8) |
| Rheum plamatum L. (Rhei Radix) | 6.0 (14.3) |
| Mirabilite (Natrii Sulfas) | 6.0 (14.3) |
| Total | 42 (100) |
streptomycin (100 U/ml) in an incubator at 37°C, with 5% CO₂ and 95% humidity. To stimulate the cells, the medium was exchanged for fresh DMEM, and LPS (500 ng/ml) was added in the presence or absence of DHMDT for various time periods, as described in the following experiments.

Assessment of cell viability. The effects of DHMDT on cell viability were evaluated using an MTT assay. In brief, RAW 264.7 cells were seeded at a density of 1x10⁴ cells/well in a 96-well plate and incubated at 37°C for 24 h, followed by treatment with 800 µg/ml DHMDT alone, 500 ng/ml LPS alone, or 500 ng/ml LPS plus 200, 400 or 800 µg/ml DHMDT for a further 24 h. Untreated RAW 264.7 cells served as the control. Following this, MTT solution was added to each well and the plate was further incubated for 4 h at 37°C. The medium was then discarded, and DMSO was added to dissolve the formazan dye. The optical density was then read at 450 nm using an ELISA reader (Infinite M200; Tecan Group, Ltd., Männedorf, Switzerland).

Measurement of NO production. The accumulation of NO was assayed using Griess reagent (1% sulfanilamide, 0.1% naphthylethylenediamine dihydrochloride and 2.5% phosphoric acid). The cells were pretreated with 200, 400 or 800 µg/ml DHMDT for 1 h and then stimulated with 500 ng/ml LPS for 24 h. Untreated RAW 264.7 cells served as the control group. Subsequently, 100 µl Griess reagent was mixed with an equal volume of cell supernatant and the mixture was incubated at room temperature for 5 min. The optical density at 540 nm was measured, and the concentration of nitrite was calculated according to the standard curve generated from known concentrations of sodium nitrite.

**PGE₂, TNF-α, and IL-1β immunoassay.** The amounts of PGE₂ and proinflammatory cytokines released in the culture medium were measured using mouse PGE₂, TNF-α and IL-1β ELISA kits based on the quantitative sandwich enzyme immunoassay technique. Pretreatment with DHMDT was conducted for 1 h prior to LPS stimulation for 24 h, as described for the NO production assay. The levels of PGE₂, TNF-α and IL-1β in the culture media were quantified using ELISA kits according to the manufacturer's protocol. The absorbance was read at a wavelength of 450 nm using a microplate reader.

**Western blot analysis.** Following treatment with the various concentrations of DHMDT in the presence or absence of LPS, as described for the NO production assay, the cells were washed, scraped into cold PBS and centrifuged at 500 x g for 5 min at 4°C. The cell pellets were suspended in lysis buffer [20 mM sucrose, 1 mM EDTA, 20 µM Tris-HCl (pH 7.2), 1 mM dithiothreitol, 10 mM KCl, 1.5 mM MgCl₂ and 5 µg/ml aprotinin]. Following removal of the cell debris by centrifugation at 13,000 x g for 15 min at 4°C, the protein concentration was determined using Bio-Rad Protein Assay Dye Reagent Concentrate (Bio-Rad Laboratories, Inc., Hercules, CA, USA). In a parallel experiment, nuclear and cytosolic proteins were prepared using nuclear extraction reagents (Pierce; Thermo Fisher Scientific, Inc., Waltham, MA, USA) according to the manufacturer's protocol. Equal amounts of protein (20-30 µg) from each sample were separated by 10% SDS-PAGE and transferred onto nitrocellulose membranes (Schleicher and Schuell; GE Healthcare Life Sciences, Little Chalfont, UK). Nonspecific sites were blocked by incubation of the membranes for 1 h at room temperature with 5% (w/v) non-fat milk powder in Tris-buffered saline containing 0.05% (v/v) Tween-20. Thereafter, the membranes were incubated overnight at 4°C with the corresponding primary antibodies and subsequently incubated with the appropriate HRP-conjugated secondary antibodies for 1 h at room temperature. The specific proteins were detected using an Amersham ECL Western Blotting Detection Reagent (Amersham; GE Healthcare Life Sciences). Densitometric analyses were completed using the ImageJ software (version 1.50i; National Institutes of Health, Bethesda, MD, USA) to quantify protein expression levels. All protein measurements were normalized to β-actin expression and these normalized values were used for all statistical analyses.

**Immunofluorescent staining of NF-κB p65.** RAW 264.7 cells were seeded on glass coverslips in 6-well plates for 24 h. The cells were then treated with DHMDT (200–800 µg/ml) for 1 h, followed by stimulation with 500 ng/ml LPS for 30 min. Untreated RAW 264.7 cells served as the control. The cells were then rinsed twice with PBS and fixed with 3.7% paraformaldehyde in PBS for 10 min at 4°C. Following fixation, the cells were incubated with 0.4% Triton X-100 for 10 min and blocked with 5% bovine serum albumin (Sigma-Aldrich; Merck KGaA) for 1 h at room temperature, followed by probing with rabbit anti-p65 NF-κB antibody (1:50; cat. no. sc-109; Santa Cruz Biotechnology, Inc.) overnight at 4°C. The cells were then incubated with fluorescein isothiocyanate-conjugated donkey anti-rabbit IgG (1:200; cat. no. 711-095-152; Jackson ImmunoResearch Laboratories, Inc., West Grove, PA, USA) for 2 h at room temperature. After washing with PBS, the nuclei were counterstained with DAPI solution (1 mg/ml) for 15 min in the dark, and fluorescence was visualized using a fluorescence microscope (Carl Zeiss AG, Oberkothen, Germany).

**Statistical analysis.** Data from at least three independent experiments are expressed as the mean ± standard deviation (SD). Differences among the groups were evaluated using one-way analysis of variance followed by Duncan's multiple range tests. The statistical analysis was conducted using SPSS 19.0 software (IBM Corp., Armonk, NY, USA). P<0.05 was considered to indicate a statistically significant difference.

**Results**

**Inhibition of the LPS-induced production of NO and PGE₂ by DHMDT in RAW 264.7 macrophages.** The potential inhibitory effects of DHMDT on LPS-induced NO and PGE₂ production in RAW 264.7 macrophages were first examined using Griess reagent and ELISA analyses, respectively. The cells were pretreated with various concentrations of DHMDT for 1 h prior to stimulation with 500 ng/ml LPS for 24 h. As illustrated in Fig. 1, NO and PGE₂ production was significantly increased in the LPS-stimulated RAW 264.7 cells compared with the unstimulated negative control, and pretreatment with DHMDT significantly prevented this increase in a dose-dependent manner. To exclude the possibility of DHMDT having a
cytotoxic effect on the RAW 264.7 cells, the viability of the cells was evaluated using an MTT assay. The concentrations (200, 400 and 800 µg/ml) used for the inhibition of NO and PGE₂ production did not exhibit any cytotoxicity (Fig. 2), confirming that inhibition of NO and PGE₂ production in the LPS-stimulated RAW 264.7 cells was not due to a cytotoxic action of DHMDT.

Inhibition of the LPS-induced expression of iNOS and COX-2 by DHMDT in RAW 264.7 macrophages. To further investigate the mechanism of the inhibitory effect of DHMDT on NO and PGE₂ production, the protein expression of iNOS and COX-2, the enzymes that mediate the synthesis of NO and PGE₂ (9,10), was determined using western blot analysis. As shown in Fig. 3, the protein levels of iNOS and COX-2 were undetectable in the RAW 264.7 cells in the absence of LPS stimulation, and treatment with LPS alone significantly increased iNOS and COX-2 protein levels. However, the pretreatment with DHMDT significantly suppressed the expression of iNOS and COX-2 proteins. These results indicate that DHMDT was able to inhibit the expression of iNOS and COX-2 enzymes, which, in turn, reduced the production of NO and PGE₂, two key mediators of inflammation.

Inhibition of the LPS-induced production and expression of TNF-α and IL-1β by DHMDT in RAW 264.7 macrophages. To examine the effects of DHMDT on the production of proinflammatory cytokines following LPS treatment, the levels of...
TNF-α and IL-1β in the culture media were measured using an ELISA. As shown in Fig. 4, the challenge with LPS alone induced significant increases in the levels of TNF-α and IL-1β, whereas the pretreatment with DHMDT significantly reduced the secretion of TNF-α and IL-1β in the cell media in a concentration-dependent manner. Next, whether the inhibitory effect of DHMDT on TNF-α and IL-1β release into the culture medium was associated with modulation of the protein levels in the RAW 264.7 cells was investigated using western blotting. As shown in Fig. 5, DHMDT significantly suppressed the protein expression of TNF-α and IL-1β compared with that in the LPS only-treated group. The reduced expression of the TNF-α and IL-1β proteins was consistent with the reduction in NO and PGE₂ in the culture media, suggesting that the DHMDT-mediated inhibition of TNF-α and IL-1β production is associated with the downregulation of TNF-α and IL-1β expression.

Attenuation of the LPS-induced nuclear translocation of NF-κB by DHMDT in RAW 264.7 macrophages. As NF-κB plays a pivotal role in the regulation of iNOS, COX-2 and proinflammatory cytokine expression (11,12), the effects of DHMDT on the activation of NF-κB were examined. As displayed in Fig. 6A, LPS stimulation for 30 min caused the translocation of p65, a component of the heterodimer of NF-κB, to the nucleus. However, DHMDT pretreatment effectively blocked the LPS-induced nuclear accumulation of p65 in the cells. These results were confirmed by NF-κB and DAPI co-staining in the LPS-treated RAW 264.7 cells, with DHMDT significantly preventing the nuclear translocation of p65 (Fig. 6B). As the phosphorylation of IκB kinase (IKK) α/β allows IκBα to be phosphorylated and ubiquitinated from the p50/p65 complex of NF-κB, the effects of DHMDT on the inhibition of IKK activity in the RAW 264.7 cells were explored. The western blotting data reveal that DHMDT inhibited the LPS-induced phosphorylation of IKKα/β (Fig. 7A), suggesting that DHMDT effectively inhibits LPS-induced NF-κB pathway activation by blocking the nuclear translocation of NF-κB and the phosphorylation of IKKα/β.

Suppression of the LPS-induced phosphorylation of AKT and MAPKs by DHMDT in RAW 264.7 macrophages. To investigate whether the inhibition of inflammatory reactions by DHMDT was mediated through the PI3K/Akt and MAPK pathways, the effects of DHMDT on LPS-induced phosphorylation of upstream kinases, including Akt, ERK, JNK, and p38 MAPK in the RAW 264.7 cells were examined. As shown in Fig. 7, the phosphorylated levels of Akt and the three MAPKs were significantly increased following treatment for 30 min in the LPS-stimulated cell group, compared with the control group without LPS treatment. DHMDT pretreatment...
Figure 6. Effect of DHMDT on LPS-induced NF-κB nuclear translocation in RAW 264.7 macrophages. The cells were treated with the indicated concentrations of DHMDT for 1 h prior to treatment with 500 ng/ml LPS for 30 min. (A) Nuclear and cytosolic proteins were subjected to 10% SDS-PAGE, followed by western blotting using anti-NF-κB p65 antibody. Lamin B and β-actin were used as internal controls for the nuclear and cytosolic fractions, respectively. (B) ImageJ densitometric analysis of bands expressed in relation to lamin B and β-actin. Data are presented as mean ± standard deviation of the mean. *P<0.05 and **P<0.01 vs. the untreated control; *P<0.05 and ***P<0.01 vs. cells cultured with 500 ng/ml LPS only. (C) The cells were pretreated with 800 µg/ml of DHMDT for 1 h prior to stimulation with LPS for 30 min. Localization of NF-κB p65 was visualized with a fluorescence microscope following immunofluorescence staining with anti-NF-κB p65 antibody (red) and staining with DAPI for the visualization of nuclei (blue). Magnification, x400. DHMDT, Daehwangmokdantang; LPS, lipopolysaccharide; NF-κB, nuclear factor-κB; NE, nuclear extract; CE, cytosolic extract.

Figure 7. Effect of DHMDT on LPS-induced phosphorylation of IKKα/β, Akt and MAPKs in RAW 264.7 macrophages. The RAW 264.7 cells were pretreated with 800 µg/ml DHMDT for 1 h prior to exposure to LPS for 30 min, and total proteins were isolated. (A) The proteins were subjected to SDS-PAGE, followed by western blot analysis. (B) ImageJ densitometric analysis of bands expressed in relation to β-actin. Data are presented as mean ± standard deviation of the mean. *P<0.05, **P<0.01 and ***P<0.005 vs. cells cultured with 500 ng/ml LPS only. DHMDT, Daehwangmokdantang; LPS, lipopolysaccharide; p-, phosphorylated; IKK, inhibitor of nuclear factor-κB kinase; Akt, protein kinase B; ERK, extracellular signal-regulated kinase; JNK, c-Jun NH2-terminal kinase.
clearly attenuated the phosphorylation levels of these kinases compared with those in the LPS only-treated group. These results suggest that the anti-inflammatory effect of DHMDT may be due to modulation of the PI3K/Akt and MAPK signaling pathways in RAW 264.7 macrophages.

Discussion

Macrophages are an important component in the immune defense mechanism. Under normal circumstances, inflammation-associated mediators, including NO and PGE₂, which are generated by iNOS and COX-2 enzymes, respectively, and cytokines generated from macrophages have an essential role in host survival and tissue repair (3,16). However, these inflammatory mediators and cytokines are overproduced in response to stimuli and may cause inflammation-associated disorders (4,17). Moreover, proinflammatory cytokines, including TNF-α and IL-1β, evoke increases in the levels of iNOS and COX-2, followed by significant increases in NO and PGE₂ production (1,2). Therefore, investigation of the suppression of NO and PGE₂ via the inhibition of iNOS and COX-2 is important in the development of anti-inflammatory agents. The results of the present study demonstrated that LPS significantly upregulated the production of NO and PGE₂, and that pretreatment with DHMDT significantly inhibited the LPS-induced release of NO and PGE₂ by the RAW 264.7 macrophages, with no cytotoxicity at the concentrations employed. In addition, the results from the western blot analysis revealed that the application of DHMDT reduced the LPS-induced iNOS and COX-2 levels, and ELISA results demonstrated that DHMDT dose-dependently inhibited LPS-induced TNF-α and IL-1β production by the RAW 264.7 macrophages. Consistent with these findings, it was observed that DHMDT significantly suppressed the protein levels of TNF-α and IL-1β. The results of the present study suggest that DHMDT may elicit anti-inflammatory effects in RAW 264.7 macrophages by reducing the LPS-stimulated production of proinflammatory molecules.

NF-κB is a critical inducible transcription factor, which serves a major role in the regulation of gene expression in response to inflammation (11,12). The activation of NF-κB involves phosphorylation and subsequent proteolytic degradation of the inhibitory protein IkBα, which binds to the NF-κB complex in the cytoplasm. Free NF-κB is then translocated into the nucleus, where it binds to the NF-κB site in the promoter regions of genes for inflammatory proteins, including iNOS, COX-2, TNF-α, and IL-1β. The engagement of LPS activates IKK, which initiates signal-induced degradation of IkB proteins. IKK phosphorylates IkBα and then ubiquitinates p-IkBα from NF-κB, which enables the translocation of NF-κB into the nucleus (8,16). To investigate whether the inhibitory effect of DHMDT on the production and expression of proinflammatory factors was associated with NF-κB pathway activity, the effect of DHMDT on NF-κB nuclear translocation was measured in the present study. The results indicated that LPS alone reduced the levels of NF-κB p65 in the cytoplasm of the RAW 264.7 macrophages and increased those in the nucleus, and that this effect was reversible by pretreatment with DHMDT. Furthermore, additional evidence that the inhibition of NF-κB nuclear translocation by DHMDT resulted from the disturbance of IKKα/β activation was obtained. The present study demonstrated that DHMDT prevented the LPS-induced phosphorylation of IKKα/β in RAW 264.7 macrophages. This suggests that inhibition of the NF-κB signaling pathway by the suppression of IKK activation in RAW 264.7 macrophages may regulate the suppressive effects of DHMDT on inflammatory mediators and cytokines.

Although the role of PI3K/Akt signaling in the regulation of NF-κB transactivation is unclear, studies have shown that phosphorylated Akt promotes the liberation of NF-κB upon the activation of Akt by PI3K and that NF-κB subsequently translocates into the cell nucleus to transcribe inflammatory mediators (13,14). By inducing the nuclear translocation of p65 in macrophages, the MAPK pathway activates pathways leading to the production of proinflammatory mediators and cytokines (18,19). Therefore, PI3K/Akt- and MAPK-targeted therapeutics may be effective for the treatment of inflammatory diseases, given that a wide variety of pharmacological agents reportedly inhibit activation steps in the PI3K/Akt and MAPK signaling pathways (8,20). To investigate whether the anti-inflammatory effects of DHMDT were mediated through the PI3K/Akt and MAPK pathways, the LPS-induced phosphorylation of Akt and various MAPK family proteins, specifically ERK, JNK, and p38 MAPK, were assessed in the present study. DHMDT pretreatment was observed to markedly suppress the LPS-stimulated phosphorylation of Akt, which is a critical step in PI3K activation (7,8). Additionally, DHMDT significantly inhibited the LPS-induced phosphorylation of three MAPKs. These results indicate that DHMDT blocks LPS-induced NF-κB activation via changes in the phosphorylation of Akt and MAPKs. In particular, inhibition of the phosphorylation of Akt or MAPKs may lead to reductions in the production and expression levels of inflammatory enzymes, including iNOS and COX-2, and cytokines in RAW 264.7 macrophages.

In conclusion, the results of the present study indicate that DHMDT has a strong inhibitory effect on the secretion of NO, PGE₂, and inflammatory cytokines, including TNF-α and IL-1β. In the LPS-stimulated RAW 264.7 macrophages, this inhibitory effect was exerted via suppression of iNOS, COX-2, TNF-α and IL-1β protein expression. These inhibitory effects are possibly mediated by the inhibition of NF-κB activation and blockade of the PI3K/Akt and MAPK signaling pathways.

Although further studies are warranted to establish the signaling pathways responsible for the observed effects, the results of the present study suggest that DHMDT may be a useful agent for the prevention or reversal of inflammatory responses.

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