Dingchuan tang essential oil inhibits the production of inflammatory mediators via suppressing the IRAK/NF-κB, IRAK/AP-1, and TBK1/IRF3 pathways in lipopolysaccharide-stimulated RAW264.7 cells

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Background: Dingchuan tang (asthma-relieving decoction), a formula of nine herbs, has been used for treating respiratory inflammatory diseases for >400 years in the People’s Republic of China. However, the mechanisms underlying the anti-inflammatory action of dingchuan tang is not fully understood. This study aims to investigate the effects of Dingchuan tang essential oil (DCEO) on inflammatory mediators and the underlying mechanism of action.

Materials and methods: DCEO was extracted by steam distillation. Lipopolysaccharide (LPS)-stimulated RAW264.7 macrophages were used as the cell model. Production of nitric oxide (NO) was determined by the Griess test. Protein secretion and mRNA levels of inflammatory mediators were measured by the enzyme-linked immunosorbent assay (ELISA) and quantitative real-time polymerase chain reaction (qRT-PCR), respectively. Protein levels were examined by Western blot. Nuclear localization of nuclear factor-kappa B (NF-κB) was detected using immunofluorescence analyses.

Results: DCEO significantly reduced LPS-triggered production of NO and prostaglandin E2 (PGE2) and decreased protein and mRNA levels of inducible nitric oxide synthase (iNOS) and cyclooxygenase-2 (COX-2). LPS induced upregulation of protein and mRNA levels of cytokines (interleukin-1β [IL-1β], interleukin-6 [IL-6], tumor necrosis factor-α [TNF-α]), and chemokines (monocyte chemoattractant protein-1 [MCP-1], chemokine [C–C motif] ligand 5 [CCL-5], and macrophage inflammatory protein [MIP]-1α) were suppressed by DCEO treatment. Phosphorylation and nuclear protein levels of transcription factors (activator protein-1 [AP-1], NF-κB, interferon regulatory factor 3 [IRF3]) were decreased by DCEO. Protein levels of phosphorylated IκB-α, IκB kinase α/β (IKKα/β), phosphatidylinositol 3-kinase 3-kinase (PI3K), protein kinase B (Akt), TGF β-activated kinase 1 (TAK1), TANK-binding kinase 1 (TBK1), extracellular signal-regulated kinase (ERK), p38 mitogen-activated protein kinase (p38), and c-Jun N-terminal kinase (JNK) were lowered by DCEO. Moreover, degradation of interleukin-1 receptor-associated kinase 1 (IRAK1) and IRAK4 induced by LPS was inhibited by DCEO treatment.

Conclusion: Suppression of the interleukin-1 receptor-associated kinase 1 (IRAK1)/NF-κB, IRAK/AP-1 and TBK1/IRF3 pathways was associated with the inhibitory effects of DCEO on inflammatory mediators in LPS-stimulated RAW264.7 macrophages. This study provides a pharmacological justification for the use of dingchuan tang in managing inflammatory disorders.

Keywords: traditional Chinese medicine, respiratory diseases, anti-inflammation, molecular pathways


Introduction

Inflammation is a defense mechanism that occurs as a result of exposure of tissues and organs to harmful stimuli such as microbial pathogens, irritants, or toxic cellular components. Chronic inflammation may induce various diseases, including asthma, rheumatoid arthritis, multiple sclerosis, inflammatory bowel disease, and psoriasis. These diseases impose a considerable economic burden on the society and shorten the average life expectancy of patients. Macrophages release inflammatory mediators in response to harmful stimuli, playing a pivotal role in the pathophysiology of inflammatory diseases. Lipopolysaccharide (LPS), the major structural component of the outer membrane of Gram-negative bacteria, is a potent activator of macrophages. Toll-like receptor 4 (TLR4) and its downstream signaling pathways involving interleukin-1 receptor-associated kinase (IRAK)/nuclear factor-kappa B (NF-kB), IRAK/activator protein-1 (AP-1) and TRIF/interferon regulatory factor 3 (IRF3) are activated in LPS-exposed macrophages, leading to the production of inflammatory mediators (eg, nitric oxide [NO], interleukin-6 [IL-6], tumor necrosis factor [TNF-α], and interleukin-1β [IL-1β]) regulated by these pathways. These inflammatory mediators will further activate macrophages, neutrophils, and lymphocytes, initiating cellular injury and tissue damage. Complementary and alternative medicines are commonly used to treat chronic inflammatory diseases, including respiratory inflammatory conditions.

Dingchuan tang (asthma-relieving decoction) is a traditional Chinese medicine formula, consisting of Herba Ephedrae, Semen Ginkgo, Flos Farfarae, Rhizoma Pinellia, Semen Armeniacae amarum, Fructus Perilla, Cortex Mori, Radix Scutellariae, and Rhizoma Glycyrrhizae at the ratio of 3:3:3:3:3:2:2:2:1. This formula has been used for treating respiratory inflammatory diseases for centuries in the People’s Republic of China. It was originally documented in “She Sheng Zhong Miao Fang” (Effective Formulas for Health Conservation, issued in 1550) as a remedy for asthma. In patients with asthma, this formula can improve respiratory function and dissipate phlegm. Pharmacological studies have shown that water extract of dingchuan tang exhibited a strong anti-inflammatory effect in Ovalbumin-induced asthmatic animals. The mechanism underlying this anti-inflammatory effect may be the inhibition of iNOS degradation.

Essential oil isolated from herbal medicines contains various bioactive compounds and possesses anti-inflammatory properties. However, there is no study evaluating the bioactivities of the essential oil isolated from dingchuan tang even though it has been reported that the main volatile compounds in Herba Ephedrae (eg, α-terpineol) and Flos Farfarae (eg, oleic acid) have strong anti-inflammatory activities via suppressing the TLR4 signaling pathways. Macrophages play an important role in inflammatory diseases through secretion of a series of inflammatory mediators. RAW264.7 macrophage line is the most commonly used cell model in inflammation research. Thus, we used RAW264.7 cells to examine the effect of dingchuan tang essential oil (DCEO) on the production of inflammatory mediators and explored the underlying mechanisms.

Materials and methods

Herbal preparation

All the herbs, including Semen Ginkgo (the dried seed of Ginkgo biloba L.), Herba Ephedrae (the dried herbaceous stem of Ephedra sinica Stapf.), Flos Farfarae (the dried flower bud of Tussilago farfara L.), Rhizoma Pinellia (the dried tuber of Pinellia ternata [Thunb.] Breit), Semen Armeniacae amarum (the dried seed of Prunus armeniaca L), Fructus Perilla (the dried seed of Perilla frutescens (L.) Britt), Cortex Mori (the dried root bark of Morus alba L.), Radix Scutellariae (the dried root of Scutellaria baicalensis Georgi), and Radix et Rhizoma Glycyrrhizae (the dried root and rhizome Glycyrrhiza uralensis Fisch.) were purchased from the Mr & Mrs Chan Hon Yin Chinese Medicine Specialty Clinic and Good Clinical Practice Centre in the Hong Kong Baptist University. All of them were authenticated by the corresponding author. Voucher specimens of individual herbs have been deposited at the Centre for Cancer and Inflammation Research, School of Chinese Medicine, Hong Kong Baptist University.

DCEO was prepared using the following procedure: air-dried herbs of Dingchuan tang (1,002 g) were soaked in 10 times of water (weight of herbs/volume of water is 1:10) overnight and then extracted by conventional steam distillation using the Clevenger-type apparatus for 5 hours. After drying with sodium sulfate anhydrous, 400 µL of DCEO was obtained and then kept at 4°C in dark vials until use.

Gas chromatography–mass spectrometry (GC–MS) analysis of DCEO

DCEO was analyzed using GC–MS (model QP 2010; Shimadzu, Kyoto, Japan). DB-5MS column (30 m × 0.25 mm × 0.25 µm, Agilent J&W Scientific) was used; the column flow rate was 1 mL/min, injector temperature was 240°C, and detector temperature was 230°C under operating conditions. Oven temperature program was set as
follows: temperature was maintained at 60°C for 1 min; then consecutively ramped to 140°C at 4°C/min, from 140°C to 200°C at 2.5°C/min and from 200°C to 280°C at 8°C/min; and maintained at a constant temperature for 5 min. In all, 1 μL (2 mg of DCEO in 1 mL of methanol) was then injected. Identification of the compounds was based on the comparison of their mass spectra with the NIST 147 Library GC–MS system. The relative concentration of each compound in DCEO was quantified based on the peak area integrated by the analysis program.

A total of 27 compounds were characterized in DCEO, representing 82.2% of the total essential oil (Figure S1). According to the results obtained from GC–MS analysis (Table S1), the main constituents in DCEO were benzaldehyde (4.37%), cyclodecene (E) (2.36%), α-terpineol (2.96%), β-bisabolene (4.71%), dehydro-β-ionone (28.43%), n-hexadecanoic acid (9.21%), and oleic acid (6.20%).

Reagents
Bovine serum albumin (BSA), 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT), LPS (Escherichia coli O55:B5), Griess reagent (modified) and Dimethyl sulfoxide (DMSO) were purchased from Sigma-Aldrich Co. (St Louis, MO, USA). Fetal bovine serum (FBS), Dulbecco’s Modified Eagle’s Medium (DMEM), penicillin, and streptomycin were obtained from HyClone (Logan, UT, USA). RAW264.7 cell line, BALB/c-derived murine macrophage (ATCC, TIB-71) was purchased from American Type Culture Collection (ATCC; Manassas, VA, USA). IL-1β, IL-6, TNF-α, monocyte chemoattractant protein-1 (MCP-1), macrophage inflammatory protein (MIP)-1α and chemokine (C–C motif) ligand 5 (CCL-5) enzyme-linked immunosorbent assay (ELISA) kits were obtained from eBioscience (San Diego, CA, USA). Prostaglandin E2 (PGE2) ELISA kit was obtained from Enzo Life Sciences (Exeter, UK). NF-κB p65, phospho-NF-κB p65 (Ser536), NF-κB p50, c-Jun, phospho-c-Jun (Ser63), protein kinase B (Akt), phospho-Akt (Ser473), and Pol II (H-224) monoclonal antibodies were not treated by DCEO or LPS) was set as 100%. Six replicate wells were used in this experiment.

Determination of NO
Nitrite (NO₂⁻) levels, an index of cellular NO production, in culture supernatants of RAW264.7 cells were determined using a modified Griess reagent as described previously. Briefly, cells were seeded at 1×10⁶ cells/mL in 24-well culture plates. After a 24 h cultivation, these cells were treated with DCEO at various concentrations for 1 h and then at the presence or absence of LPS (100 ng/mL dissolved in phosphate-buffered saline [PBS]) for 24 h. The absorbance at 570 nm was measured using a microplate spectrophotometer (Bio-Rad Benchmark Plus; Bio-Rad Laboratories Inc., Hercules, CA, USA). Cell viability in the control group (cells were not treated by DCEO or LPS) was set as 100%. Six replicate wells were used in this experiment.
and nitrite production was determined. Four replicate wells were used in this experiment.

ELISA
The production of IL-1β, IL-6, TNF-α, PGE2, MCP-1, MIP-1α, and CCL-5 was determined by ELISA. Briefly, RAW264.7 macrophages were seeded in 24-well culture plates at 1×10⁵ cells/well. After a 24 h cultivation, the cells were treated with DCEO at various concentrations for 1 h and then supplemented with or without LPS (100 ng/mL) for 24 h. Cell-free supernatants were collected for determination of cytokine and chemokine concentrations according to the manufacturer’s instructions. Four replicate wells were used in this experiment.

RNA extraction and quantitative real-time polymerase chain reaction (qRT-PCR)
RAW264.7 cells were seeded in 60-mm-diameter culture dishes (4×10⁵ cells/mL) for 24 h and treated with DCEO at various concentrations. LPS (100 ng/mL) was added 1 h after DCEO treatment, and the cells were incubated at 37°C for 6 h. Total RNA was extracted using TRIzol reagent (Thermo Fisher Scientific, Waltham, MA, USA) and reverse-transcribed with oligo-dT using the Prime-Script™ RT reagent Kit (Takara Bio Inc., Shiga, Japan) according to the manufacturer’s protocol. qRT-PCR was carried out by monitoring the increase in fluorescence of SYBR Green using the ViiA 7 Real-Time PCR System (Thermo Fisher Scientific). Primers were synthesized by Invitrogen (iNOS primers: forward AGCAACTAGTCTGGTGTG and reverse TCTTCAGATTCTGCCATTG, COX-2 primers: forward CTGGGACTGACCTCAGTTT and reverse AGGCCCTTGGCAGCTCTGT, IL-6 primers: forward AGTCGGAGAGGAGACTCTCA and reverse ATTCCACGATTCCCAAGAG, IL-1β primers: forward GAAGAGAGAAGACATCTCTG and reverse TCACTTCGAGGTCTGTAGT, TNF-α primers: forward ATGAGAAGCTTCCCAATTG and reverse CTCCACCTGGTGTGTGTCA, CCL-5 primers: forward CATATGGCTCGGACACCA and reverse ACACACTTTGCGGTTCTC, MCP-1 primers: forward ATCGTGAAGGCCACAGAG and reverse CTGCTCAGATGGTGTGTCA, MAPK primers: forward GGCTTTCGGGTCTCTACC and reverse TGCCCTGCTTACCCACCTTTC, M-CSF primers: forward ACACACTTTGCGGTTCTC and reverse CGACCTTCAGGTCGATG, and GAPDH primers: forward GGCCCTCGCTGTCTCCACC and reverse TGCCCTGCTTACCCACCTTTC, M-CSF). Each sample was amplified in triplicate for quantification. Data were analyzed by relative quantification using the ΔΔCt method and normalized to GAPDH. Three replicate wells were used in this experiment.

Western blot analysis
Cell lysates were prepared as previously described. Briefly, RAW264.7 cells were seeded in 60-mm-diameter culture dishes (4×10⁵ cells/mL) for 24 h and treated with DCEO at various concentrations. LPS (100 ng/mL) was added 1 h after DCEO pretreatment, and the cells were incubated at 37°C for 30 or 60 min. RNA extraction and quantitative real-time polymerase chain reaction (qRT-PCR) was carried out by monitoring the increase in fluorescence of SYBR Green using the ViiA 7 Real-Time PCR System (Thermo Fisher Scientific). Primers were synthesized by Invitrogen (iNOS primers: forward AGCAACTACTGCTGGTGTG and reverse TCTTCAGAGTTCTGCCATTG, COX-2 primers: forward CTGGGACTGACCTCAGTTT and reverse AGGCCCTTGGCAGCTCTGT, IL-6 primers: forward AGTCGGAGAGGAGACTCTCA and reverse ATTCCACGATTCCCAAGAG, IL-1β primers: forward GAAGAGAGAAGACATCTCTG and reverse TCACTTCGAGGTCTGTAGT, TNF-α primers: forward ATGAGAAGCTTCCCAATTG and reverse CTCCACCTGGTGTGTGTCA, CCL-5 primers: forward CATATGGCTCGGACACCA and reverse ACACACTTTGCGGTTCTC, MCP-1 primers: forward ATCGTGAAGGCCACAGAG and reverse CTGCTCAGATGGTGTGTCA, MAPK primers: forward GGCTTTCGGGTCTCTACC and reverse TGCCCTGCTTACCCACCTTTC, M-CSF primers: forward ACACACTTTGCGGTTCTC and reverse CGACCTTCAGGTCGATG, and GAPDH primers: forward GGCCCTCGCTGTCTCCACC and reverse TGCCCTGCTTACCCACCTTTC, M-CSF). Each sample was amplified in triplicate for Quantification. Data were analyzed by relative quantification using the ΔΔCt method and normalized to GAPDH. Three replicate wells were used in this experiment.

RNA extraction and quantitative real-time polymerase chain reaction (qRT-PCR)
RAW264.7 cells were seeded in 60-mm-diameter culture dishes (4×10⁵ cells/mL) for 24 h and treated with DCEO at various concentrations. LPS (100 ng/mL) was added 1 h after DCEO pretreatment, and the cells were incubated at 37°C for 30 or 60 min. The cells were then washed twice in ice-cold PBS and lysed with RIPA buffer consisting of 50 mM Tris–HCl, 1% NP-40, 0.35% sodium deoxycholate, 150 mM NaCl, 1 mM EDTA of pH 7.4, 1 mM phenylmethylsulfonyl fluoride, 1 mM NaF, 1 mM NaVO₄, 10 μg/mL aprotinin, 10 μg/mL leupetin, and 10 μg/mL pepstatin A. The supernatant was collected after the cell lysate was centrifuged at 14,000×g/min for 15 min at 4°C. The protein concentration was measured using the Bio-Rad Protein Assay Kit (Bio-Rad Laboratories Inc.). Equal amount of protein sample (30 μg) was resolved by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) sample loading buffer and denatured for 10 min at 95°C. Subsequently, the samples were electrophoresed on 12% SDS-PAGE mini-gel and transferred onto nitrocellulose membrane at 80 V for 3 h at 4°C. Membranes were probed with indicated primary antibodies over night at 4°C and were then blotted with respective horseradish peroxidase-conjugated secondary anti-mouse or anti-rabbit antibody. The specific proteins were detected using enhanced chemiluminescence (ECL) detection kit (Thermo Fisher Scientific) following the manufacturer’s instruction, and the blots were analyzed using the ImageJ software (National Institutes of Health [NIH], Bethesda, MD, USA).

Extraction of cytoplasmic and nuclear proteins
The cytoplasmic and nuclear proteins were prepared as previously described. Briefly, cells were collected for Western blot as described in Western blot analysis section. Cells were washed three times with ice-cold PBS and then suspended in hypotonic buffer for cytosolic extraction. After incubation on ice for 1 min, 12 μL of 10% (v/v) NP-40 was added to the suspension for another 10 min. The cytoplasmic fraction was prepared by centrifugation for 2 min. The remaining pellet generated after initial centrifugation was washed three times with the hypotonic buffer and suspended in high-salt buffer. The suspended cell pellet was incubated for 30 min on ice and then centrifuged for 10 min to obtain the nuclear
fraction. Soluble cell lysates were immunoblotted with the designated antibodies.

**Immunofluorescence staining**

RAW264.7 cells were directly cultured on chamber slides (Cat. 177399; Thermo Fisher Scientific) for 24 h to detect NF-κB/p65 nuclear localization by immunofluorescence assays using a fluorescence microscope. Briefly, after a 60 min stimulation with LPS in the presence or absence of DCEO, cells were fixed with 4% formaldehyde in PBS for 15 min at room temperature and permeabilized with 0.25% Triton-X for 30 min at 37°C and followed by 10 min at room temperature. After blocking with 3% bovine serum albumin (BSA) for 2 h, the cells were incubated with the primary antibody anti-NF-κB/p65 overnight at 4°C. After washing with PBS for three times, cells were incubated for 1 h at room temperature with Alexa Fluor 488-conjugated secondary antibody, the nuclei were stained with DAPI, and fluorescence was visualized using a fluorescence microscope (Leica Microsystems, Wetzlar, Germany).

**Statistical analysis**

At least three replicate wells were used per experimental setting. All values were expressed as arithmetic mean ± standard error of the mean. Data were analyzed by one-way analysis of variance (ANOVA) using SPSS (version 16.0) statistical analysis program, and then, differences among mean were determined using Dunnett’s multiple comparison test. Differences were considered as significant at P<0.05.

**Results**

**DCEO lowered NO and PGE2 production**

The viability of RAW264.7 cells was not significantly altered by a 24-h incubation of up to 50 μg/mL of DCEO in the presence of 100 ng/mL of LPS (Figure 1A). Concentrations of DCEO of 12.5, 25, and 50 μg/mL were used in subsequent experiments. As shown in Figure 1B and C, significant increases in NO and PGE2 resulting from LPS stimulation in RAW264.7 cells were suppressed by DCEO treatment in a dose-dependent manner.

**DCEO downregulated protein and mRNA levels of iNOS and COX-2**

As NO and PGE2 syntheses are catalyzed by iNOS and COX-2, respectively,27,28 we next examined protein and mRNA levels of iNOS and COX-2 using Western blot and qRT-PCR, respectively. Figure 1D–F showed that treatment with DCEO decreased LPS-triggered upregulation of cellular protein levels of iNOS and COX-2 in a dose-dependent manner. Moreover, DCEO treatment significantly decreased the elevated levels of iNOS and COX-2 mRNA induced by LPS in a concentration-dependent manner (Figure 1G and H).

**DCEO decreased cytokine and chemokine production and gene expression levels**

Activation of macrophages via TLR4 signaling induces production of critical proinflammatory cytokines (eg, IL-1β, IL-6, and TNF-α) and chemokines (eg, MCP-1, CCL-5, and MIP-1α) that are necessary to activate inflammatory responses. Therefore, these cytokines and chemokines are regarded as targets for inhibiting LPS-induced inflammatory responses. Our ELISA data showed that production of TLR4-mediated cytokines (IL-1β, IL-6, and TNF-α) and chemokines (MCP-1, CCL-5, and MIP-1α) was markedly increased upon LPS stimulation, and the increments were concentration-dependently inhibited by DCEO treatment (Figure 2A–F).

We further investigate the effect of DCEO on the gene expression of the corresponding cytokines and chemokines. In line with the ELISA results, stimulation with LPS upregulated mRNA levels of cytokines (IL-1β, IL-6, and TNF-α) and chemokines (MCP-1, CCL-5, and MIP-1α) in RAW264.7 macrophages, and the levels of mRNA of IL-1β, IL-6, TNF-α, MCP-1, CCL-5, and MIP-1α were significantly lowered by DCEO treatment in a concentration-dependent manner (Figure 3A–F). GAPDH was used as an internal control, which has been validated in our facility and in RAW264.7 cells.10

**DCEO affected the components of the IRAK/NF-κB, IRAK/AP-1, and TBK1/IRF3 pathways**

The above data showed that DCEO could inhibit proinflammatory cytokines and chemokines regulated by TLR4 signaling. The involvement of TLR4 signaling in the effects of DCEO was further explored in LPS-stimulated RAW264.7 macrophages. TLR4 signaling cascades involve MyD88-dependent and -independent signaling pathways. The MyD88-dependent pathway involves the early phase of NF-κB and AP-1 activation, which leads to the production of inflammatory cytokines.31 As shown in Figures 4B and 5B, the phosphorylations of NF-κB subunit p65 (ser536) and AP-1 (c-Jun and c-Fos) were obviously elevated after 1 h
Figure 1 DCEO inhibited the production of NO and PGE2 and the expression of iNOS and COX-2 in LPS-stimulated RAW264.7 cells. 

Notes: Cells were incubated with indicated concentrations of DCEO for 24 h with or without LPS. Cell viability was determined by the MTT assay. Results were expressed as percentages of respective controls (A). Cells were incubated with indicated concentrations of DCEO for 1 h and then stimulated with LPS for 24 h. The production of NO was indicated by the level of nitrite formed in the supernatant detected using the Griess reagent. PGE2 production was measured by ELISA (B and C). The protein levels of iNOS and COX-2 were determined by immunoblotting. Representative blots of iNOS and COX-2 are shown (D). Bar graphs show the relative levels of iNOS and COX-2 in RAW264.7 cells (E and F). Cells were incubated with indicated concentrations of DCEO for 1 h and then stimulated with LPS for 6 h. The mRNA levels of iNOS and COX-2 were determined by qRT-PCR (G and H). Data presented in bar charts are mean±SEM values from three independent experiments. **P<0.01 vs unstimulated cells. ***P<0.005 and ****P<0.01 vs LPS-stimulated cells determined by Dunnett’s multiple comparisons test.

Abbreviations: DCEO, Dingchuan tang essential oil; NO, nitric oxide; PGE2, prostaglandin E2; iNOS, inducible nitric oxide synthase; COX-2, cyclooxygenase-2; LPS, lipopolysaccharide; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; ELISA, enzyme-linked immunosorbent assay; qRT-PCR, quantitative real-time polymerase chain reaction; SEM, standard error of the mean.
DCEO inhibits inflammatory mediators via suppressing the TLR4 signaling pathways

LPS stimulation. DCEO treatment suppressed the phosphorylation of p65 and AP-1 (Figures 4B and 5B) in a concentration-dependent manner. Next, regulators upstream AP-1 and NF-κB were also examined. Figures 4A and 5A showed that phosphorylations of three key MAPKs subunits including p38, ERK, and JNK were markedly enhanced by LPS stimulation in RAW264.7 cells, and DCEO treatment inhibited the phosphorylation of p38, ERK, and JNK in a concentration-dependent manner (Figures 4A and 5A). Moreover, DCEO treatment markedly reduced LPS-induced...
phosphorylation of IκBα. Phosphorylations of TAK1 and IKKα/β were significantly enhanced by LPS treatment, which could be suppressed by DCEO treatment in a concentration-dependent manner (Figures 4A and 5A). DCEO treatment also inhibited LPS-induced degradation of IRAK1 and IRAK4 (Figures 4A and 5A). Phosphatidylinositol 3-kinase (PI3K)/Akt signaling also plays a role in LPS-triggered inflammatory response in macrophages. As shown in Figures 4B and 5B, phosphorylated PI3K subunits p85 and Akt (ser473) were remarkably elevated after LPS stimulation in RAW264.7 macrophages, while DCEO suppressed the elevation in a concentration-dependent manner. While the
effects of DCEO on NF-κB and AP-1 were relatively weak, the underlying reasons will be explored in the future.

We also investigated the effects of DCEO on the TBK1/IRF3 signaling pathway in RAW264.7 cells. As shown in Figures 4 and 5, the phosphorylation levels of TBK1 and IRF3 were obviously upregulated after LPS stimulation. DCEO treatment decreased the levels of phosphorylated TBK1 and IRF3 in a concentration-dependent manner, suggesting an inhibitory effect of DCEO on TBK1/IRF3 signaling.

It has been reported that NF-κB, AP-1, and IRF3 are the key transcriptional factors in TLR4 signaling. Nuclear translocation of these transcription factors initiates the transcription of genes encoding proinflammatory mediators and

**Figure 4** DCEO affected the molecular components of the IRAK/NF-κB, IRAK/AP-1, and TBK1/IRF3 pathways in LPS-stimulated RAW264.7 cells.

**Notes:** Cells were treated with DCEO at the indicated concentrations for 1 h and then stimulated with LPS for 30 or 60 min. Total and phosphorylated forms of individual proteins were detected by Western blot (A and B). A representative of three independent experiments is shown.

**Abbreviations:** DCEO, Dingchuan tang essential oil; IRAK, interleukin-1 receptor-associated kinase; NF-κB, nuclear factor-kappa B; AP-1, activator protein-1; TBK1, TANK-binding kinase 1; IRF3, interferon regulatory factor 3; LPS, lipopolysaccharide; p, phosphorylated; JNK, c-Jun N-terminal kinase; ERK, extracellular signal-regulated kinase; p38, p38 mitogen-activated protein kinase; IKKα/β, IκB kinase; p-IRF3, p-IRF3 (47 kDa); TBK1, TBK1 (90 kDa); GAPDH, GAPDH (37 kDa); p-PI3K, p-PI3K (85 kDa); PI3K, PI3K (85 kDa); p-Akt, p-Akt (60 kDa); Akt, Akt (60 kDa); p-IκBα, p-IκBα (41 kDa); IκBα, IκBα (41 kDa); p-p65, p-p65 (65 kDa); p65, p65 (65 kDa); p-c-Jun, p-c-Jun (39 kDa); c-Jun, c-Jun (39 kDa); p-IRF3, p-IRF3 (47 kDa); IRF3, IRF3 (47 kDa); p-c-Fos, p-c-Fos (62 kDa); c-Fos, c-Fos (62 kDa); GAPDH, GAPDH (37 kDa).
As shown in Figure 6A and B, nuclear protein levels of IRF3, NF-κB heterodimer components (p65 and p50), and one of the AP-1 components (c-Jun) were markedly increased 60 min after LPS treatment. DCEO treatment inhibited the increment of nuclear IRF3, p65, p50, and c-Jun. Immunofluorescence staining of p65 confirmed the inhibitory effect of DCEO on NF-κB nuclear localization (Figure 7).
Figure 6 DCEO inhibited nuclear localization of NF-κB, AP-1, and IRF3 in LPS-stimulated RAW264.7 cells.

Notes: Cells were treated with DCEO with indicated concentrations for 1 h and then stimulated with LPS for 60 min. The cytoplasmic and nuclear protein levels of NF-κB (p65 and p50), c-Jun, and IRF3 were determined by Western blot (A). Bar graphs show the relative levels of cytoplasmic and nuclear p65, p50, IRF3, and c-Jun. Data are the mean ± SEM of three independent experiments. **P<0.01 vs unstimulated cells. ***P<0.05 and ****P<0.01 vs LPS-stimulated cells determined by Dunnett’s multiple comparisons test (B).

Abbreviations: DCEO, Dingchuan tang essential oil; NF-κB, nuclear factor-kappa B; AP-1, activator protein-1; IRF3, interferon regulatory factor 3; LPS, lipopolysaccharide; SEM, standard error of the mean.

Discussion

In the present study, 27 compounds were identified in DCEO using the GC–MS analysis. The main constituents of DCEO, such as benzaldehyde, α-terpineol, n-hexadecanoic acid and oleic acid, have been reported to possess strong anti-inflammatory effects.34–37 n-Hexadecanoic acid inhibits the activity of phospholipase A2 to control inflammation.34 α-Terpineol reduces IL-6 formation through binding to the IL-6 receptor and subsequent internalization of the entire ligand–receptor complex.35 Benzaldehyde exhibits anti-asthmatic effect via downregulating the production of proinflammatory cytokines.36 Oleic acid exerts anti-arthritis effect by reducing inflammatory cell infiltration.37 These reports imply that the inflammatory mediator inhibitory effect of DCEO may be, at least in part, due to the presence of these compounds. Further studies are needed to identify active constituents inside DCEO.

It is well known that NO and PGE2 are two key mediators of inflammatory responses.38 Excessive NO production, especially by macrophages, can lead to cytotoxicity, inflammation, and autoimmune disorders.39,40 PGE2 is a metabolite of arachidonic acid that elicits a wide range of physiological and pathological processes, such as gastrointestinal protection, inflammation, and cancer.41 PGE2 mediates the onset
of the symptoms of classic signs of inflammation, including redness, swelling, and pain. Hence, measuring the production of LPS-induced NO and PGE2 can evaluate the progression of inflammation, and inhibition of their production might have potential therapeutic value for controlling inflammatory reactions and diseases. In the present study, we examined the effects of DCEO on the production of NO and PGE2. We observed that DCEO remarkably suppressed LPS-induced release of NO and PGE2 in RAW264.7 cells, suggesting that DCEO has potential anti-inflammatory activity. Since iNOS and COX-2 catalyze the production of NO and PGE2, respectively, in activated macrophages, we determined the expression of iNOS and COX-2 in LPS-stimulated RAW264.7 cells. It was found that upon stimulation, the mRNA and protein levels of iNOS and COX-2 were markedly increased and DCEO lowered mRNA and protein levels of iNOS and COX-2 in a concentration-dependent manner. These results suggest that DCEO suppresses NO and PGE2 production through inhibiting the expression of iNOS and COX-2, respectively. Further studies will be conducted to investigate the effect of DCEO on the activities of these two enzymes. In addition to PGE2 and COX-2, leukotriene and 5-lipoxygenase also play an important role in the arachidonic acid metabolic pathway and are linked to asthma pathogenesis. In the future, we will investigate the impact of DCEO on leukotriene formation and 5-lipoxygenase activity.

LPS-activated macrophages release proinflammatory cytokines and chemokines that exacerbate inflammatory responses and lead to tissue injury. Proinflammatory cytokines, such as IL-1β, IL-6, and TNF-α, play crucial roles in the pathogenesis of various acute and chronic inflammatory conditions including allergic asthma, infection, and rheumatoid arthritis. Chemokines are a family of chemoattractant molecules that have a role in neutrophil, eosinophil, and monocyte migration during acute and chronic inflammation. Our data indicate that DCEO treatment significantly inhibits LPS-induced release of TLR4-regulated cytokines (IL-1β,
IL-6, and TNF-α) and chemokines (MCP-1, MIP-1α, and CCL-5) in RAW264.7 cells. Based on these data, it can be concluded that DCEO has inhibitory effect on inflammatory mediators in LPS-stimulated RAW264.7 cells.

TLR4 is the primary recognition molecule for inflammatory responses initiated by LPS. After ligation with LPS, TLR4 recruits intracellular adapters including MyD88, an adaptor recruits IRAK4, thereby allowing the phosphorylation and degradation of IRAK1. This leads to the ubiquitination of TRAF6 and induces the activation of TAK1, which in turn phosphorylates the IKK complex (including IKKα, IKKβ, and IKKγ). The activated IKK complex then phosphorylates IκB, which leads to its ubiquitination and subsequent degradation. Removal of IκBs liberates...
NF-κB for nuclear translocation and binding to cognate DNA binding sites to regulate the transcription of a number of genes that encode cytokines, chemokines, and stress response proteins.\(^5\) Our results show that DCEO treatment inhibits the phosphorylation and nuclear localization of NF-κB in LPS-stimulated RAW264.7 cells. We also found that the phosphorylation levels of IkBα and IKKα/β were decreased by DCEO treatment, indicating DCEO inhibits IKKα/β/IkBα/NF-κB signaling. IRAK4, IRAK1, and TAK1 take part in the activation of IKKs and MAPKs.\(^6\) We found that the level of phosphorylated TAK1 was downregulated by DCEO treatment. We also found that DCEO treatment obviously reduces LPS-mediated degradation of IRAK1 and IRAK4, indicating an inhibitory effect of DCEO on IRAK/TAK1 signaling.

PI3K and its downstream kinase Akt appear to be important components of LPS-induced activation of the NF-κB signaling pathway.\(^7\) Upon exposure to LPS, PI3K interacts with TRAF6 and activates the downstream target Akt.\(^8\) The activated Akt phosphorylates IKKα, leading to IkB degradation and NF-κB translocation to the nucleus and resulting in specific cytokine gene expression.\(^9\) Consistent with a previous report,\(^10\) our study indicates that the phosphorylated PI3K subunit p85 and Akt (Ser473) are markedly elevated following LPS stimulation. DCEO treatment suppresses the phosphorylation of p85 and Akt in LPS-treated RAW264.7 cells, suggesting that DCEO inhibits LPS-triggered PI3K/Akt signaling, which may, at least partially, contribute to the suppressive effect of DCEO on NF-κB signaling.

Other than NF-κB, the AP-1 complex (composed of c-Fos and c-Jun subunits) also plays a critical role in the transcriptional regulation of genes that encode inflammatory mediators.\(^11\) In addition to the role in posttranslational regulation of NF-κB,\(^12\) MAPKs regulate the transcriptional activity and half-life of AP-1.\(^13\) Therefore, we also investigated the effect of DCEO on MAPKs and AP-1 activities. Results showed that the LPS-stimulated phosphorylation of MAPKs (ERK, p38, and JNK) and AP-1 (c-Jun and c-Fos) was suppressed by DCEO treatment, indicating that DCEO inhibits the MAPKs/NF-κB and MAPKs/AP-1 signaling pathways.

Besides MyD88-dependent pathways, the TRIF-dependent pathway also plays a role in LPS-induced inflammatory response in macrophages.\(^14\) LPS can also trigger TRIF signaling that activates TBK1, which in turn phosphorylates IRF3 and triggers its dimerization and nuclear translocation, leading to the expression of IFN-β and CCL-5.\(^15,16\) In our present study, DCEO treatment inhibits LPS-induced phosphorylation and nuclear localization of IRF3. The results are in agreement with the inhibitory effect of DCEO on CCL-5 production. We then examined the effects of DCEO on the phosphorylation of TBK1 that is upstream of IRF3 and found that DCEO reduced the phosphorylation of TBK1, suggesting that DCEO inhibits TBK1/IRF3 signaling.

**Conclusion**

In this study, we found that suppression of the IRAK/NF-κB, IRAK/AP-1, and TBK1/IRF3 signaling pathways is associated with the inhibitory effect of DCEO on inflammatory mediators in LPS-stimulated RAW264.7 macrophages (Figure 8). These findings provide a pharmacological justification for the use of Dingchuan tang in managing inflammatory disorders, as Dingchuan tang is traditionally used for treating asthma, which is typically characterized by an eosinophilic inflammation.\(^17,18\) In the future, we will use asthma models to determine the impact of DCEO on eosinophilic inflammation.

**Acknowledgments**

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**Disclosure**

The authors report no conflicts of interest in this work.

**References**


Supplementary materials

Figure S1 GC chromatogram of DCEO.
Notes: DCEO was analyzed by GC-MS. Peak information is shown in Table S1.
Abbreviations: GC, gas chromatography; DCEO, Dingchuan tang essential oil; GC–MS, gas chromatography–mass spectrometry; TIC, total ion chromatogram.

Table S1 Percentual chemical composition of DCEO

<table>
<thead>
<tr>
<th>Peak no</th>
<th>Retention time (min)</th>
<th>Compounds</th>
<th>Molecular formula</th>
<th>Relative contents (%)</th>
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<tbody>
<tr>
<td>22</td>
<td>46.735</td>
<td>D-Glucitol, hexaacetate</td>
<td>C_{12}H_{26}O_{12}</td>
<td>2.29</td>
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<tr>
<td>23</td>
<td>47.114</td>
<td>Galactitol, hexaacetate</td>
<td>C_{12}H_{26}O_{12}</td>
<td>0.88</td>
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<tr>
<td>27</td>
<td>48.566</td>
<td>(E)-9-Octadecenoic acid, ethyl ester</td>
<td>C_{18}H_{36}O_{2}</td>
<td>0.91</td>
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<tr>
<td>26</td>
<td>48.403</td>
<td>9,12-Octadecadienoic acid</td>
<td>C_{18}H_{34}O_{2}</td>
<td>0.97</td>
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<tr>
<td>25</td>
<td>47.88</td>
<td>Oleic acid</td>
<td>C_{18}H_{36}O_{2}</td>
<td>6.20</td>
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<tr>
<td>24</td>
<td>47.719</td>
<td>9,12-Octadecadienoic acid (Z,Z)-</td>
<td>C_{18}H_{34}O_{2}</td>
<td>2.07</td>
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<tr>
<td>16</td>
<td>27.979</td>
<td>Acetic acid, 6,6-dimethyl-2-methylene-7-(3-oxobutylidene)oxepan-3-ylmethyl ester</td>
<td>C_{18}H_{36}O_{2}</td>
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<tr>
<td>19</td>
<td>30.415</td>
<td>9,12,15-Octadecatrien-1-ol (Z,Z,Z)-</td>
<td>C_{18}H_{36}O</td>
<td>1.05</td>
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<tr>
<td>14</td>
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<td>Acetate, 2-cyclohexenyl-3-[1-(2-oxopropyl)ethenyl]-2,4,4-trimethyl</td>
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<td>21</td>
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<td>n-Hexadecanoic acid</td>
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<td>18</td>
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<tr>
<td>15</td>
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<tr>
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<td>[β]-Bisabolene</td>
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<tr>
<td>17</td>
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<td>C_{18}H_{24}O</td>
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<tr>
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<td>16.995</td>
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<tr>
<td>4</td>
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<td>2-Hexanoylfuran</td>
<td>C_{18}H_{30}O</td>
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<tr>
<td>13</td>
<td>26.453</td>
<td>7-Oxatetracyclo[4.2.0.0(2,4).0(3,5)]octane, 8-acetyl-8-methyl-</td>
<td>C_{18}H_{24}O_{2}</td>
<td>2.27</td>
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<tr>
<td>2</td>
<td>10.039</td>
<td>1-Undecene</td>
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<td>3</td>
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<td>p-Menth-1-en-8-ol</td>
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<td>5</td>
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<td>Cycloendecene(E)</td>
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<tr>
<td>1</td>
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<td>Benzaldehyde</td>
<td>C_{6}H_{6}O</td>
<td>4.37</td>
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</table>

Notes: A total of 27 components were characterized on the basis of a typical library research and literature data. The components showing matches exceeding 80%, which represented ~82.2% of DCEO, were selected.
Abbreviation: DCEO, Dingchuan tang essential oil.