

# Brazilin Suppresses Inflammation via the Down-regulation of IRAK4 in LPS-stimulated Raw264.7 Macrophage

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**Abstract** Brazilin, is a bioactive compound extracted from *Caesalpinia sappan* Linn, has been reported the protective effect of the immune system. Particular attention is now devoted to better understanding of the molecular basis of brazilin anti-inflammatory activity. In the present study, we studied the effect of brazilin on the Raw264.7 macrophage cell lines by a nutrigenomics approaches. Raw264.7 cells were treated with brazilin, then treated with LPS to cause inflammation. The nuclear transcription  $\kappa$ B (NF- $\kappa$ B) promoter activity were analyzed with dual luciferase assay kit. The gene expression and production levels of pro-inflammatory cytokine interleukin (IL)-1 $\beta$ , tumor necrosis factor (TNF) $\alpha$ , and IL-6 were evaluated with semi-quantitative RT-PCR and with ELISA, respectively. We also examined inflammatory signaling, including mitogen-activated protein kinase (MAPK) pathway, iNOS, COX2, and IRAK4. Our findings demonstrated that brazilin down-regulated the expression of IRAK4 protein lead to suppress of c-Jun NH<sub>2</sub> terminal kinase (JNK) signaling, and subsequently inactivation of nuclear transcription  $\kappa$ B (NF- $\kappa$ B), inducible nitric oxide synthase (iNOS) and cyclooxygenase 2 (COX2) thus promoting the expression of the downstream target pro-inflammatory cytokines such as IL-1 $\beta$ , TNF $\alpha$  and IL-6 in LPS stimulated Raw264.7 macrophage cell. Thus, brazilin showed anti-inflammatory activity in Raw264.7 macrophage cell targeting IRAK4 mediated signaling pathway.

**Keywords:** Brazilin, inflammation, NF- $\kappa$ B, IRAK4, MAPK

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## 1. Introduction

Inflammation process is mainly associated with host organisms against exogenous pathogen and incidence of body injuries in healthy condition [1,2]. In contrast, stimulation of inflammation has been recognized as the integral feature of the chronic disease, which includes obesity, diabetes, cancer, and cardiovascular diseases [3,4,5]. Therefore, coordination of inflammatory response results in ameliorating inflammation-associated chronic disease.

Macrophage plays an important role in inflammation and in response to a variety of inflammatory mediators such as lipopolysaccharide and other exogenous products [6]. Toll-like receptors (TLRs) bind to highly conserved leucine-rich repeated sequences expressed by bacterial and viral pathogen-associated molecular patterns (PAMP) [7] and subsequently induce the release the pro-inflammatory cytokines through the myeloid differential factor (MyD88-dependent) and toll-interleukin-1 receptor domain-containing adapter inducing interferon- $\beta$  (TRIF-dependent)

pathways [8, 9]. The pro-inflammatory cytokines such as IL-1 $\beta$ , TNF $\alpha$ , and IL-6 play an important role in inflammation, promoting to induce other pro-inflammatory cytokines, chemokines, and somatic pain [10,11]. Cellular responses to pro-inflammatory cytokines depends on multiple protein kinase cascades including activation of the stress-activated mitogen activated protein kinase (MAPK), c-Jun NH<sub>2</sub>-terminal kinase (JNK), cascade of intracellular kinase (ERK), and p38 MAPK as well as transcription factor nuclear factor  $\kappa$ B (NF- $\kappa$ B) [12,13].

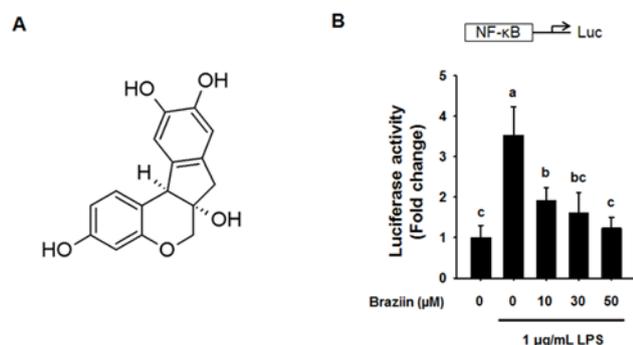
The IL-1 receptor-associated kinase 4 (IRAK4) has been shown to play an important role in MyD88 dependent inflammatory signaling. Moreover, IRAK4 leads to increase of MAPK signaling and I $\kappa$ B kinase, and subsequently activation of NF- $\kappa$ B thus promoting the expression of the downstream target pro-inflammatory cytokines. While, the IRAK4 kinase deficient mice have been shown to be resistant to LPS stimulated inflammation, due to diminish MyD88-IRAK4 association and decrease the pro-inflammatory cytokine productions [14]. Therefore, suppression of IRAK4 is important for prevention of LPS-mediated inflammation.

Brazilin is a major active phenolic component that is isolated from the heartwood of *Caesalpinia sappan* Leguminosae (*C. sappan*). *C. sappan* extracts has been used traditional complementary medicine as anti-inflammation [15,16]. Brazilin has been reported several beneficial biological activities, including modulation of immune function and cancer progression [17,18,19]. However, how brazilin regulates inflammation in macrophage cell still remain unclear. Therefore, this study was performed to determine the molecular mechanism of brazilin on LPS-induced inflammation in macrophage cell. To determine the effect of brazilin on LPS-mediated cellular inflammation, we used Raw264.7 macrophage cell. We analyzed the effect of brazilin on IRAK4 and its downstream targets including iNOS, COX2, NF- $\kappa$ B, MAPKs and pro-inflammatory cytokines in LPS-induced Raw264.7 macrophage cell.

## 2. Materials and Methods

### 2.1. Materials

Lipopolysaccharide (LPS; *Escherichia coli* 0111:B4) was purchased from Sigma (St. Louis, Mo, USA). Brazilin was obtained from dried *C. sappan* L. heartwood (purchased in Seoul, South Korea) methanol extracts, and was purified according to the methods described by Oh *et al.* [20]. The chemical structure of brazilin was identified by the Korea Research Institute of Bioscience & Biotechnology as shown in Figure 1A. The extract was dissolved in dimethyl sulfoxide prior to use. All reagents were purchased from Sigma unless otherwise described.



**Figure 1.** The effect of brazilin on NF- $\kappa$ B luciferase activity in LPS-stimulated Raw264.7 cells. (A) The chemical structure of brazilin. (B) NF- $\kappa$ B luciferase activity of brazilin. Cells were incubated with increasing concentration of the brazilin for 1h and then co-treated with 1  $\mu$ g/mL LPS for 10 h. DMSO was used as a vehicle. Data are mean  $\pm$  SD ( $n=3$ )

### 2.2. Cell Culture and Cell Viability

Raw264.7 macrophage cell lines were purchased from the American Type Culture Collection (ATCC, Rockville, MD, USA). These were cultured in Dulbecco's modified Eagle medium DMEM containing 10% heat-inactivated fetal bovine serum (FBS) (Invitrogen, USA), 100 units/ml penicillin, and 100  $\mu$ g/mL streptomycin (Invitrogen, Carlsbad, CA, USA). Cells were grown at 37  $^{\circ}$ C in a 5 % CO<sub>2</sub>/air environment. To evaluate the cell viability, Raw264.7 macrophage cell lines were plated at a concentration of 5,000 cells/well in 96-well plate and the 2,3-bis-(2-methoxy-4-nitro-5-sulphophenyl)-2H-tetrazolium-5-carboxanilide (XTT) assay was performed as described by Kim *et al* [21]. For

the cell viability analysis, Raw264.7 cells were treated with 0, 10, 30, 50, and 100  $\mu$ M of brazilin.

### 2.3. Western Blotting

Cells were plated overnight in 6-well plates at a density of  $0.7 \times 10^6$  per plate, and further incubated in 1  $\mu$ g/mL LPS for 1 h after treatment. Cells were harvested with ice-cold RIPA buffer (50mM Tris-HCl, 1mM EDTA, 1mM EGTA, 150mM NaCl, 1% NP-40, 0.1% sodium dodecyl sulfate (SDS), 0.25 % sodium deoxycholate, and 5%  $\beta$ -mercaptoethanol) with protease and phosphatase inhibitors. Cell lysates were centrifuged at  $12,000 \times g$  for 5 min at 4 $^{\circ}$ C and the supernatants were then collected. Protein concentrations were determined using the Bradford Protein Assay (Bio-Rad, Richmond, CA, USA). Equal amounts of extracted protein (100  $\mu$ g) were separated by SDS-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to polyvinylidene difluoride membranes. Membranes were blocked with 5% non-fat skim milk phosphate-buffered saline (PBS) containing 0.05 % Tween 20, and were blotted with the indicated primary antibody overnight at 4  $^{\circ}$ C. Bound primary antibodies were detected with a peroxidase-coupled secondary antibody. The reactive bands were visualized by chemiluminescence (Amersham Biosciences, Piscataway, USA).

### 2.4. RNA Extraction and Reverse Transcription Polymerase Chain Reaction (RT-PCR)

Total RNA was extracted from RAW264.7 cells using Trizol® reagent (Invitrogen Corporation, Carlsbad, CA, USA), according to the manufacturer's protocol. Total RNA (1  $\mu$ g) was reverse transcribed to cDNA using the Maxime RT PreMix kit (Intron, Seongnam, South Korea). cDNA was then amplified with an Inno Hot Tap polymerase Kit (Bookyoung SM, South Korea). The primer sequences were as follows: GAPDH, forward (5'-AACTTTGGCATTGTGGAAGG -3') and antisense (5'-ACACATTGGGGGTAGGAACA-3'); TNF $\alpha$ , forward (5'-CTACTCCTCAGAGCCCCCAG -3') and reverse (5'-TGACCACTCTCCCTTTGCAG -3'); IL-1 $\beta$ , forward (5'-CAGGATGAGGACATGAGCACC-3') and reverse (5'-CTCTGCACACTCAAACCTCCAC-3'); IL-6, forward (5'-GTTCTCTGGGAAATCGTGGA-3') and reverse (5'-TGTACTCCAGGTAGCTA -3'). PCR amplification of the resulting cDNA template was conducted under the following conditions. PCR products were analyzed on 1% agarose gels and stained with ethidium bromide. Images were captured with a Gene Fresh ultraviolet (UV) detector (Syngene Bio Imaging, South Korea). Densitometric analysis was performed using the ImageJ program (National Institutes of Health, Bethesda, MD). The results are representative of three independent experiments.

### 2.5. Transient Transfection and Luciferase Assays

For transfection, cells were seeded at a density of  $7 \times 10^4$  cells/well in 48-well plates and incubated for 24 h until approximately 70-80% confluence. NF- $\kappa$ B (2 $\times$ )-luciferase and COX-2 luciferase reporter plasmids or the corresponding empty vector plasmids were co-transfected

to Raw264.7 macrophage cell using SuperFect transfection reagent (Qiagen, Valencia, CA, USA), according to the manufacturer's instructions. Cells were lysed and luciferase activity was determined using the Promega luciferase assay system (Promega, Madison, CA, USA) and a luminometer (Perkin Elmer Cetus, Foster City, CA, USA). Luciferase activity was normalized to that of  $\beta$ -galactosidase.

## 2.6. Nitrite Colorimetric Assay

Raw264.7 cells were seeded at  $7 \times 10^4$  per well in 96-well plates and incubated for 24 h. The cells were pretreated with various concentration of brazilin for 1 h and then treated with LPS (1  $\mu$ g/mL) for an additional 24h. For nitrite determinations, 100  $\mu$ L of culture supernatant was mixed with an equal volume of Griess reagent and the absorbance at 540 nm was measured. The NaNO<sub>2</sub> standard curve was used to determine total nitrite.

## 2.7. Enzyme-linked Immunosorbent Assay (ELISA)

Cells were pretreated with various brazilin concentrations for 1 h and then further stimulated with LPS (1  $\mu$ g/mL) for 24 h. The supernatants were collected and stored at -80°C until cytokine analysis. IL-1 $\beta$ , TNF $\alpha$ , and IL-6 levels in supernatants were determined using ELISA MAX<sup>TM</sup> Kits (BioLegend, San Diego, CA, USA), according to the manufacturer's instructions.

## 2.8. Statistical Analysis

All values are expressed as mean  $\pm$  standard deviation. Data were analyzed by one-way analysis of variance (ANOVA) (9.01, SAS program). *Post-hoc* comparisons were conducted using Tukey's honestly significant difference test. Differences were considered to be statistically significant when *p* values were less than 0.05.

## 3. Results

### 3.1. Brazilin Decreased NF- $\kappa$ B Luciferase Activity in LPS-stimulated Raw264.7 Macrophage Cells

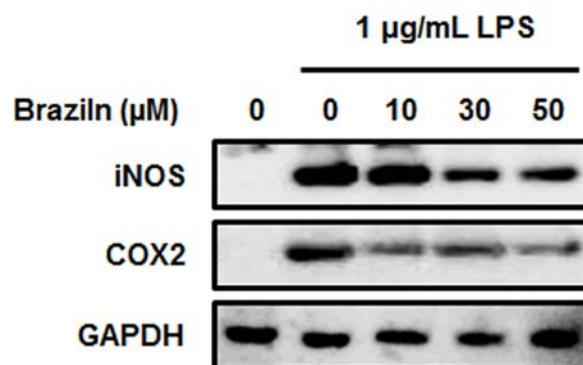
The chemical structure of brazilin is presented in Figure 1A. The effect of brazilin on Raw264.7 cell viability was measured by a XTT assay. As shown in Figure 1B, brazilin decreased cell viability at a concentration of 100  $\mu$ M. Thus, the concentration of 10, 30, and 50  $\mu$ M brazilin was selected for the further investigation.

The induction of inflammatory response is modulated through the activation of NF- $\kappa$ B. Therefore, we evaluated whether brazilin regulates the transcriptional activity of NF- $\kappa$ B. The Raw264.7 macrophage cells were transiently transfected with pNF- $\kappa$ B-Luc plasmid, pre-treated with different concentration of brazilin for 1h and then co-treated with brazilin and LPS. NF- $\kappa$ B transactivation was measured by a dual luciferase assay system. Brazilin significantly suppressed LPS-induced transcriptional activity of NF- $\kappa$ B compared to LPS-treated Raw264.7 with absence of brazilin as shown in Figure 1 (*p*<0.05).

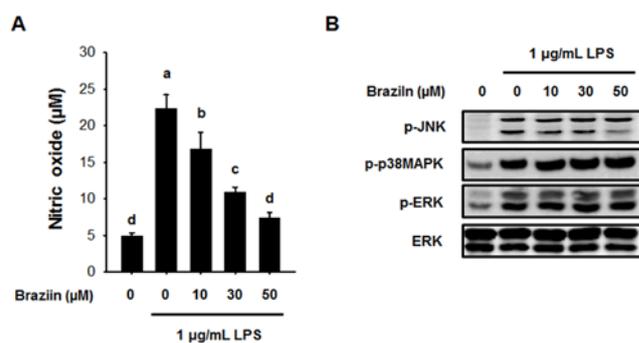
We found that brazilin may show anti-inflammatory activity through suppression of NF- $\kappa$ B pathway.

### 3.2. Brazilin Inhibited the Upstream Target of NF- $\kappa$ B in LPS-stimulated Raw264.7 Macrophage Cell

A number of genes involved in pro-inflammatory response are governed by NF- $\kappa$ B. To better evaluate the anti-inflammatory activity of brazilin on the transcription of NF- $\kappa$ B target gene, the protein level of iNOS and COX2, which depends on NF- $\kappa$ B activity in response to LPS stimulation was investigated. As shown in Figure 2, western blotting analysis showed that LPS caused the elevated expression levels of iNOS and COX2, while both iNOS and COX2 proteins were significantly decreased in Raw264.7 with presence of brazilin.



**Figure 2.** Brazilin down-regulated the expression of iNOS and COX2 protein in LPS stimulated Raw264.7. Raw 264.7 macrophage cells ( $5 \times 10^5$  /well) were pretreated with 10, 30, or 50  $\mu$ M brazilin for 1 h and then induced with 1  $\mu$ g/mL LPS for an 24 h. The expression of iNOS and COX2 protein were detected by western blotting using iNOS, COX2, and GAPDH antibodies. DMSO was used as a vehicle. Data are expressed as mean  $\pm$  SD (*n*=3)

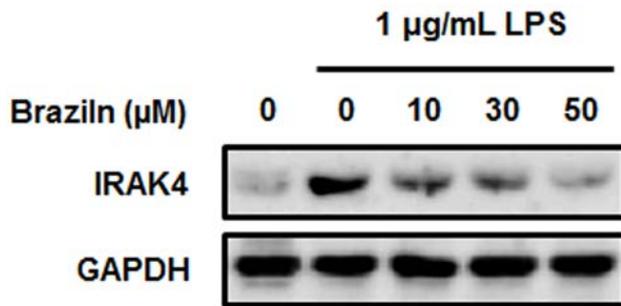


**Figure 3.** Brazilin decreased the nitric oxide production and phosphorylation of JNK in LPS-stimulated Raw264.7 macrophage cell. (A) The production of soluble nitrite oxide. Culture media was subsequently isolated and nitrite concentrations determined. (B) The expression of p-JNK, p-p38MAPK, p-ERK, and ERK in LPS-stimulated Raw264.7 macrophage cell. Cells were pretreated with 10, 30 or 50  $\mu$ M brazilin for 1 h and then stimulated with 1  $\mu$ g/mL LPS for 30 min or 24 h. DMSO was used as a vehicle. Equal amounts of protein in cell lysates were analyzed by Western blot. Data are mean  $\pm$  SD (*n*=3)

### 3.3. The Effect of Brazilin on ROS Production and MAPK Signaling in LPS-Stimulated Raw264.7 Macrophage Cell

ROS and MAPK signaling pathway is crucial role for pro-inflammation through activation of NF- $\kappa$ B pathway,

we examined the intracellular levels of NO using a nitrite colorimetric assay in LPS-induced Raw264.7 with the presence or absence of baseline. As shown in Figure 3A, LPS treatment caused significantly elevated the production of NO levels. However, pre-treatment of brazilin resulted in a decrease of LPS-induced NO levels in Raw264.7 macrophage cell. In addition, we determine the phosphorylation of the MAPK pathway, which are intermediate stage controlling of NF- $\kappa$ B activation. As shown in Figure 3B, the phosphorylation of JNK was decreased by brazilin, but not in the phosphorylation of p38MAPK and ERK.

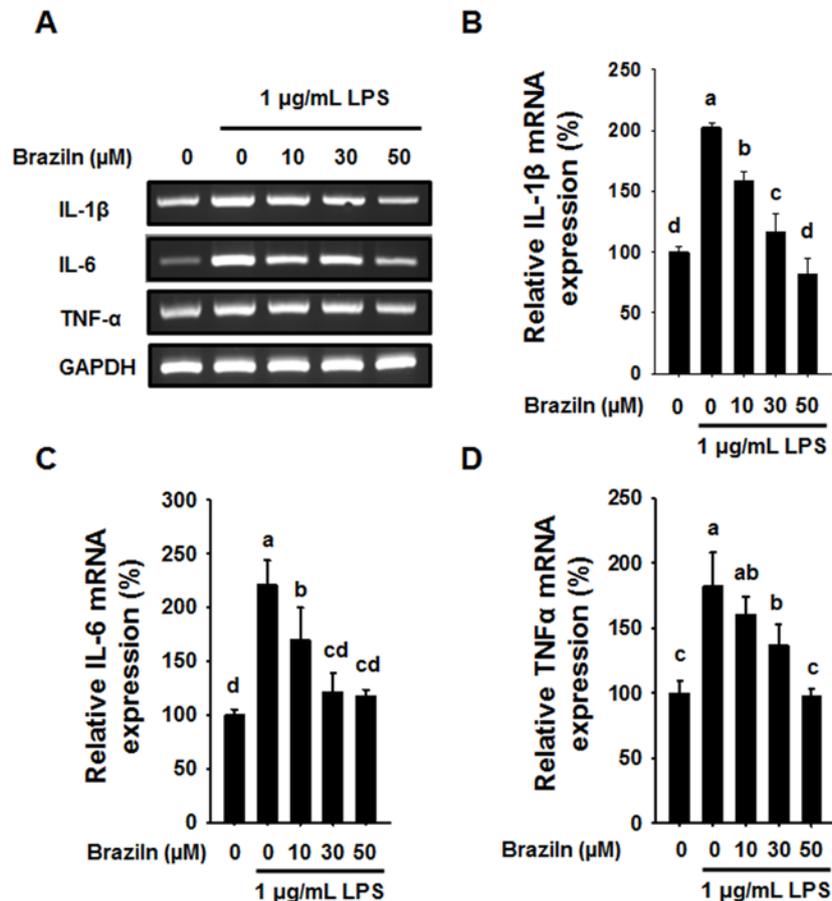


**Figure 4.** LPS increased the expression of IRAK4 protein, whereas brazilin diminished the expression of IRAK4 in Raw264.7 macrophage cell. Cells were pretreated with 10, 30 or 50  $\mu$ M brazilin for 1 h and then stimulated with 1  $\mu$ g/mL LPS for 30 min or 24 h. DMSO was used as a vehicle. Equal amounts of protein in cell lysates were analyzed by Western blot

In fact, IRAK4 is known as the upstream enzymes responsible for phosphorylating MAPK [9]. Next, we determined the expression levels of IRAK4, which are major mediators controlling the NF- $\kappa$ B activation in LPS-induced inflammation. As shown in Figure 4, we sought that brazilin suppressed the expression of IRAK4 in a dosed dependent manner in LPS-induced Raw264.7 compared to the vehicle-treated and LPS-induced Raw264.7 macrophage cell. These results suggest that brazilin could attenuate LPS-induced inflammation in Raw264.7 macrophage, in part, by inhibition of IRAK4 regulated NF- $\kappa$ B activation, and partially affecting the MAPK pathway.

### 3.4. Brazilin Attenuates mRNA Expression of Pro-inflammatory Cytokines in LPS-Stimulated Raw264.7 Macrophage Cell

The pro-inflammatory cytokines is an end product of IRAK4 regulated NF- $\kappa$ B activation and play critical roles in the extent of inflammation. To finalize the analysis of the brazilin on anti-inflammatory activity, we measured the expression of pro-inflammatory cytokine mRNA and the levels of cytokine production in LPS-induced Raw264.7 macrophage cell with presence or absence of brazilin. For this objective, the expression levels of pro-inflammatory transcripts were analyzed by semi-quantitative RT-PCR. The pro-inflammatory cytokine was measured by ELISA.



**Figure 5.** The transcription levels of pro-inflammatory cytokines were down-regulated by brazilin in LPS-induced Raw264.7. Cells were pretreated with 10, 30, or 50  $\mu$ M brazilin for 1 h, stimulated with LPS (1  $\mu$ g/mL), and incubated for a further 24 h. DMSO was used as a vehicle. Data are expressed as mean  $\pm$  SD ( $n=3$ ). (A) Total RNA was amplified by RT-PCR using the indicated primers for IL-1 $\beta$ , TNF $\alpha$ , and IL-6. (B-D) The relative mRNA expression of IL-1 $\beta$ , TNF $\alpha$ , and IL-6 were quantified using the ImageJ program. Data are mean  $\pm$  SD ( $n=3$ )

As shown in Figure 5A-5D, brazilin decreased the expression of genes encoding IL-1 $\beta$ , TNF $\alpha$ , and IL-6 in a dose dependent manner in LPS-treated Raw264.7 macrophage cell. Moreover, pro-inflammatory cytokine production including IL-1 $\beta$  and TNF $\alpha$  were decreased by 30 and 50  $\mu$ M brazilin compared to LPS-induced Raw264.7 macrophage cell as shown in Figure 6A and 6B. In particular, brazilin clearly depressed IL-6 production in LPS-induced Raw264.7 macrophage cell in a dose-dependent manner as shown in Figure 6C. These results indicated that baseline could regulate IRAK4-mediated NF- $\kappa$ B activation pro-inflammatory signaling in LPS-induced Raw264.7 macrophage cell.

## 4. Discussion

Several studies showed that the isomer of brazilin such as brazilein exhibited inflammation suppressive activity and anti-oxidative function [6,22]. However, the molecular targets of brazilin for inflammation response have remained elusive. In this study, we focused on the identification of the molecular target of brazilin and quantification of its effect on anti-inflammation in LPS-induced Raw264.7 macrophage cell.

A previous study demonstrates that TLR/IRAK4 signaling enhances macrophage-associated inflammation [23] and IRAK4 deficient animal model are completely resistant to high dose of LPS [24]. Our results showed that LPS caused not only the pro-inflammatory cytokine production, but also increase of NF- $\kappa$ B promoter activity via activation of IRAK4 protein activity, while brazilin dramatically repressed the expression levels of IRAK4 and subsequently decreased NF- $\kappa$ B activity. NF- $\kappa$ B signaling is required for the transcription of iNOS and COX2, which are thought to be expressed mainly by activated macrophage [25,26]. We also found that brazilin inhibited the expression of NF- $\kappa$ B dependent protein COX2 and iNOS levels as well as their product NO in LPS-induced Raw264.7 macrophage cell.

In addition, stimulation of LPS promotes to release the serum pro-inflammatory cytokines IL-1 $\beta$ , IL-6, and TNF $\alpha$  in animal model [27,28]. Treatment of LPS increased the production of IL-1 $\beta$ , IL-6, and TNF $\alpha$  by Raw264.7 macrophage cell in mouse serum. We observed that LPS-induced production of IL-1 $\beta$ , IL-6, and TNF $\alpha$  were markedly decreased in Raw264.7 macrophage cell with presence of brazilin. In fact, we and others showed that bioactive compounds, such as bebeerine, fucoidan, and hispidin suppressed the production of pro-inflammatory cytokines and subsequent ameliorated inflammation response which is associated with aberrant expression of MAPK signaling [29,30,31].

MAPK signaling promote to enhance the phosphorylation of NF- $\kappa$ B, which induces the function of NF- $\kappa$ B in nucleus [32]. Indeed, bioactive compound shows anti-inflammatory activities through the suppression of p38 MAPK, JNK, and ERK mediated NF- $\kappa$ B pathway in macrophage [29,30,33,34]. Similarly, we found that brazilin significantly reduced the phosphorylation of JNK in a dose dependent manner. However, p38MAPK and ERK were not altered protein expression after brazilin treatment. This result raised the indication that brazilin regulated the phosphorylation of JNK and subsequently

suppressed NF- $\kappa$ B nuclear translocation in LPS-stimulated Raw264.7 macrophage cell.

## 5. Conclusion

We provide that brazilin decreased the expression of IRAK4 levels lead partially to suppression of MAPK signaling involving JNK, which resulted in inhibition of NF- $\kappa$ B activity that is changing the transcription of iNOS and COX2 and subsequently attenuate the inflammatory response end products such as pro-inflammatory cytokines and nitrites. Our data provide proof that brazilin suppress a major source of inflammatory mediators and that macrophage cells are highly responsive to ameliorate a variety of inflammatory factors. Therefore, we suggest that brazilin might be a useful bioactive compound for the prevention of exogenous pathogen-mediated diseases as well as IRAK4-associated inflammatory diseases.

## Conflict of Interest

The authors declare that they have no conflict of interest.

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