Suppression of Inflammatory Responses by Black Rice Extract in RAW 264.7 Macrophage Cells via Downregulation of NF-κB and AP-1 Signaling Pathways

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Abstract

Anthocyanin, a phenolic compound, has been reported to have an anti-inflammatory effect against lipopolysaccharide (LPS) induced changes in immune cells. However, little is known about the molecular mechanisms underlying its anti-inflammatory effects. Few research studies have concerned the anti-inflammation properties of colored rice extract as a functional material. Therefore, the purpose of this study was to examine anti-inflammatory effects of the polar fraction of black rice whole grain extracts (BR-WG-P) that features a high anthocyanin content. Our results showed that BR-WG-P significantly inhibited LPS-induced pro-inflammatory mediators, including production of NO and expression of iNOS and COX-2. In addition, secretion of pro-inflammatory cytokines including TNF-α and IL-6 was also significantly inhibited. Moreover, BR-WG-P and anthocyanin inhibited NF-κB and AP-1 translocation into the nucleus. BR-WG-P also decreased the phosphorylation of ERK, p38 and JNK in a dose dependent manner. These results suggested that BR-WG-P might suppress LPS-induced inflammation via the inhibition of the MAPK signaling pathway leading to decrease of NF-κB and AP-1 translocation. All of these results indicate that BR-WG-P exhibits therapeutic potential associated with the anthocyanin content in the extract for treating inflammatory diseases associated with cancer.

Keywords: Black rice - anti-inflammation - NF-κB - AP-1 - anthocyanin

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Introduction

Inflammation is a protective response of a tissue to injury caused by pathogens, damaged cells, or irritants (Philip et al., 2004). Acute inflammation, the early stage of inflammation, is a rapid response to an injurious agent that activates various type of inflammatory cells, including neutrophils, macrophages, mast cells and lymphocytes to the site of injury (Coussens and Werb, 2002). When acute inflammation is obvious over a short period of time, it has a therapeutic effect. However, if inflammation occurs a long time period, the cellular response becomes a pattern of chronic inflammation. Chronic inflammation is related to various types of chronic diseases, including cancer, diabetes, cardiovascular disease, arthritis and autoimmune disease (Aggarwal, 2004). Moreover, chronic inflammation has been found to be a risk factor of most types of cancer and is involved with tumorgenesis, including tumor initiation, promotion, progression, invasion and metastasis (Coussens and Werb, 2002; Aggarwal et al., 2006; Lu et al., 2006). Therefore, methods to reduce inflammation and prevent these chronic health problems have been widely investigated.

The mechanism by which microbial pathogens including bacterial, viral, and parasitic induce inflammation can be better understood via the innate immune response (Mogensen, 2009). Lipopolysaccharide (LPS), a component within the cell wall of Gram-negative bacteria, could induce activation of monocytes and macrophages resulting in pro-inflammatory cytokine production leading to an acute inflammatory response against the pathogens (Sweet and Hume, 1996). These responses are associated with the activation of Toll like receptors (TLR) especially in subtype 4 (TLR4). The binding between LPS and TLR4 recruits the adapter protein for the activation of the NF-κB and AP-1 signaling pathway. (Akira, 2003; and and J, 2012). The activation of NF-κB and AP-1 leads to the induction of the pro-inflammatory mediator expression including inducible cyclooxygenase-2 (COX-2), inducible nitric oxide synthase (iNOS), nitric oxide (NO), interleukin-1 (IL-1), interleukin-6 (IL-6) and tumor necrotic factor alpha (TNF-α) (Lawrence, 2009). Involvement of the MAPK pathway in the expression of inflammatory mediators through NF-κB and AP-1 activation has been demonstrated (Fang et al., 2004; Kim et al., 2014).

Colored rice extract especially black rice or red rice has been reported to possess an anti-oxidant capacity, anti-invasion and anti-inflammatory effect (Ichikawa et al., 2001; Min et al., 2010; Pintha et al., 2014). However,
the molecular mechanism of the anti-inflammatory effect of black rice extract has not been reported. In the present study, black rice whole grain sample was harvested from an organic farm in Chiang Mai Province in the northern part of Thailand, and was then subjected to the extraction procedure to achieve the polar fraction (BR-WG-P) which was enriched with anthocyanin content. We next investigated the anti-inflammatory effect of the BR-WG-P against RAW 264.7 macrophage cells and found that the anthocyanin-enriched black rice extract could inhibit LPS-induced inflammation through the inhibition of the NF-kB and AP-1 signaling pathway.

Materials and Methods

Chemicals and reagents
RPMI 1640 and penicillin-streptomycin were obtained from Gibco (Grand Island, NY, USA). LPS purified from Escherichia coli and radio-immunoprecipitation assay (RIPA) lysis buffer, protease inhibitors and β-actin antibody were purchased from Sigma-Aldrich (St. Louis, MO, USA). Cytokine ELISA kits were supplied from Biolegend (USA). Antibody for PARP was purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Antibody for iNOS was obtained from Merck-Millipore (Darmstadt, Germany). Antibodies against COX-2, p38, pJNK, JNK, pERK, ERK, NF-kB and c-Jun were purchased from Cell Signaling Technology (Beverly, MA, USA). Antibody for PARP was purchased from Merck-Millipore (Darmstadt, Germany). Antibodies against COX-2, p38, pJNK, JNK, pERK, ERK, NF-kB and c-Jun were purchased from Cell Signaling Technology (Beverly, MA, USA). Antibody for PARP was purchased from Sigma-Aldrich (St. Louis, MO, USA). Cytokine ELISA kits were supplied from Biolegend (USA). Antibody for iNOS was obtained from Merck-Millipore (Darmstadt, Germany).

Plant material
Black rice (Oryza sativa L.) was harvested in wet season from Doi Saket Organic Farm, Chiang Mai Province, Thailand. A voucher specimen number (023149) was certified by the herbarium at the Flora of Thailand, Faculty of Pharmacy, Chiang Mai University, Thailand.

Preparation of BR-WG-P
Briefly, one kilogram of whole grains black rice was finely ground and soaked in 50% ethanol for 24 h. The rice samples were then filtered through filter paper to separate the residue. The filtrated samples were then evaporated using a rotary vacuum evaporator (BUCHI, Switzerland) to obtain the ethanolic fractions. The ethanolic fractions were further partitioned with saturated butanol to obtain the polar fraction of the black rice whole grains extract (BR-WG-P). The polar fraction was then evaporated and freeze-dried to obtain the black rice extract powder.

Cell cultures
The RAW 264.7 macrophage cell line was obtained from American Type Culture Collection (ATCC). The low cell binding dish was used to culture the cells (Thermo Scientific, USA). These cell lines were cultured in RPMI1640 supplemented with 10% FBS, 2 mM L-glutamine, 50 U/mL of penicillin, and 50 μg/mL of streptomycin. The cells were maintained in a 5% CO₂ humidified incubator at 37°C.

Quantification of total phenolic content and phenolic compounds in BR-WG-P
Total phenolic content in BR-WG-P was determined using the modified Folin-Ciocalteau assay as previously described (Jr., 1965). Briefly, BR-WG-P (0.4 mL) was mixed with 0.3 mL of Folin-Ciocalteau reagent and kept in the dark at room temperature for three minutes. After that, 0.3 mL of sodium carbonate was added to the mixture. Then, the mixture was further incubated in a dark at room temperature for 30 min. The absorbance of blue complex was evaluated at 765 nm using UV-visible spectrophotometer compared to a standard curve prepared with various concentrations of gallic acid (GA) solution. The total phenolic content was shown as milligrams of GA equivalents per gram of BR-WG-P (mg GAE/g extract).

The phenolic compounds in BR-WG-P were determined by HPLC analysis compared with standard gallic acid, protocatechuic acid, catechin, chlorogenic acid, vanillic acid, coumaric acid and ferulic acid. BR-WG-P was dissolved in methanol and subjected to HPLC (Agilent Technologies, CA, USA) using reversed-phase C18 column (WATER, MA, USA). The mobile phase was composed of methanol to 0.1 % trifluoroacetic acid (TFA) (1:1). The detection wave length was 280 and 325 nm and the flow rate was set at 1.0 mL/min. The peak area was calculated and compared with the standard to obtain the concentration of all the compounds (mg/g extract).

Quantification of total flavonoid content in BR-WG-P
Total flavonoid content was measured using the aluminium chloride (A1C1₃) colorimetric assay with slightly modified (Zhishen J, 1999). BR-WG-P (0.25 mL) was mixed with 0.125 mL of 5% sodium nitrite (NaNO₂) for 5 min. After that 0.125 mL of 10 % A1C1₃ was added into the mixture. Then one mL of sodium hydroxide (NaOH) was added to the mixture and it was incubated for 15 min at room temperature. The solution was then mixed and the absorbance was measured at 510 nm compared with the standard catechin using a spectrophotometer. The total flavonoid content was expressed as mg catechin equivalents per gram extract (mg CE/g extract).

Quantification of total proanthocyanidin content in BR-WG-P
Total proanthocyanidin content in BR-WG-P was determined using the vanillin assay with slight modification (Herald et al., 2014). Briefly, the BR-WG-P was mixed with 0.1 mL of 1% vanillin in methanol (w/v). Then 0.1 mL of sulfuric acid (H₂SO₄) was added and incubated for 15 min at 30°C. The absorbance of the sample was measured at 490 nm and compared with the standard of catechin. The amount of total proanthocyanidin content was expressed as milligram catechin equivalents per gram of BR-WG-P extract (mg CE/g extract).

Quantification of total anthocyanin content in BR-WG-P
Total anthocyanin analysis was measured using a pH differential method (Wrolstad et al., 2005). The BR-WG-P were dissolved in 0.1% HCl, 80% methanol an incubated for 12 h. The two dilutions of BR-WG-P solution were
Prepared for each developing stage. The first sample (0.25 mL) was diluted with 1 mL of 0.025 M potassium chloride (KCl) buffer pH 1.0 and the second was diluted with 1 mL of 0.45 M sodium acetate (CH₃COONa) buffer pH 4.5. The samples were incubated at room temperature for 15 min. The absorbance of the sample was measured at 520 and 700 nm using UV-visible spectrophotometer.

The total anthocyanin content in the extract was expressed as cyanidin-3-glucoside, which was the most sufficient anthocyanin component. Total anthocyanin content was calculated using the following formula:

\[
\text{Total anthocyanin content} = \frac{A \times Mw \times DF \times 103}{\varepsilon \times 1}
\]

\[
A = (A_{700nm} - A_{520nm}) \text{ pH 1.0} - (A_{700nm} - A_{520nm}) \text{ pH 4.5}
\]

\[
Mw = \text{Molar weight of cyanidin-3-glucoside}
\]

\[
DF = \text{Dilution Factor}
\]

\[
\varepsilon = \text{Molar extinction coefficient} = L \times \text{mol}^{-1} \times \text{cm}^{-1}
\]

\[
L = \text{Cell path length} (1 \text{ cm})
\]

**Determination of Nitric oxide (NO) production**

Nitric oxide production was determined by the detection of the nitrite concentration in cultured medium using the Griess reagent system (Sun et al., 2003). The RAW 264.7 macrophage cells were seeded in a six-well plate at a density of 5.0×10⁵ cells/well for 24 h. Then the cells were pre-treated with or without various concentrations of BR-WG-P (0-150 μg/mL) for 2 h. After that the cells were further incubated with or without LPS (1 μg/mL) for 24 hours and the cultured medium was collected for Griess reaction for the determination of nitrite production. The absorbance was measured at 540 nm using a microplate reader. The nitrite concentration in the cultured medium was calculated and compared with a standard curve of sodium nitrite (NaNO₂).

**Determination of IL-6 and TNF-α secretion**

The secretions of interleukin-6 (IL-6) and tumor necrotic factor-alpha (TNF-α) in cultured medium were determined by an ELISA kit (Biolegend, USA) according to the manufacturer’s instructions. The RAW 264.7 macrophage cells were seeded in a six-well plate at a density of 5.0×10⁵ cells/well for 24 h. Then the cells were co-treated with or without various concentrations of BR-WG-P (0-150 μg/mL) and LPS (1 μg/mL) for 24 h. The cultured medium was collected for ELISA and the absorbance was measured at 450 and 570 nm using a microplate reader. The IL-6 and TNF-α secretion in the cultured medium was calculated and compared with a standard curve of IL-6 and TNF-α.

**Immunoblot analysis**

The cultured RAW 264.7 cells were extracted with a RIPA buffer containing protease inhibitors and were then subjected to 10% SDS-PAGE. The proteins were transferred onto nitrocellulose membranes. The membranes were blocked with 5% non-fat dried-milk proteins in 0.1% TBS-tween, then probed with COX-2, iNOS, p38, p38, pJNK, JNK, pERK, ERK, NF-κB or c-Jun. After washing with 0.3% TBS-tween, membranes were incubated with HRP-conjugated anti-mouse or anti-rabbit IgG antibody. The HRP signal was detected using an enhanced chemiluminescence (ECL) system. Equal protein loading was confirmed by each membrane, which was stripped and reprobed with anti-β-actin or PARP antibody.

**Statistical analysis**

All data are presented as the mean ± standard deviation (S.D.) values. Statistical analysis was analyzed by Prism version 6.0 software using one-way ANOVA, Dunnett’s test or Tukey’s multiple comparison test. Statistical significance was determined at \( p<0.05 \), \( **p<0.01 \) or \( ****p<0.0001 \).

**Results**

**Phytochemical Characterization in BR-WG-P**

After the extraction process, the BR-WG-P was weighed and further used for phytochemical and biological analysis. The percentage of yield of BR-WG-P was 0.307 calculated from one kg of raw material. The phytochemical characterization in BR-WG-P including, total phenolic content, phenolic compound, total flavonoid content, total anthocyanin content, and total proanthocyanidin content were determined (Table 1). The total phenolic content in the extract was 117.6±14.6 mg GAE/g extract. Phenolic acid, one of members in phenolic group, contains the two subgroups including hydroxybenzoic acid and hydroxycinnamic acid. The hydroxybenzoic acid derivatives including, vanillic acid and protocatechuic acid, were found to be 4.2±0.4 and 2.3±0.1 mg/g extract, respectively whereas gallic acid was not found to be present in BR-WG-P. The hydroxycinnamic acid derivatives including, coumaric acid, Ferulic acid, and chlorogenic acid, were detected 0.5±0.2, 1.4±0.0, and 1.7±0.3 mg/g extract, respectively. In terms of the flavonoid content determination, the aluminium chloride (A1C1) colorimetric assay was applied. The flavonoid content was 42.9±2.1 mg CE/g extract. The anthocyanin content was found to be 8.1±1.9 mg C-3-G/g extract. Additionally, the flavonol derivatives, such as catechin and proanthocyanidin, were not found in BR-WG-P.

**Effect of BR-WG-P on NO Production in LPS-induced RAW Macrophage Cells**

<table>
<thead>
<tr>
<th>Compounds (mg/g extract)</th>
<th>BR-WG-P</th>
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<tbody>
<tr>
<td>Total phenolic content</td>
<td>117.6±14.6</td>
</tr>
<tr>
<td>Vanillic acid</td>
<td>4.2±0.4</td>
</tr>
<tr>
<td>Protocatechuic acid</td>
<td>2.3±0.1</td>
</tr>
<tr>
<td>Gallic acid</td>
<td>ND</td>
</tr>
<tr>
<td>Coumaric acid</td>
<td>0.5±0.2</td>
</tr>
<tr>
<td>Ferulic acid</td>
<td>1.4±0.0</td>
</tr>
<tr>
<td>Chlorogenic acid</td>
<td>1.7±0.3</td>
</tr>
<tr>
<td>Total flavonoid content</td>
<td>42.9±2.1</td>
</tr>
<tr>
<td>Anthocyanin</td>
<td>8.1±1.9</td>
</tr>
<tr>
<td>Catechin</td>
<td>ND</td>
</tr>
<tr>
<td>Proanthocyanidin</td>
<td>ND</td>
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</tbody>
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*Values are means±S.D. (n=3), ND=not detectable
To evaluate the anti-inflammatory effect of BR-WG-P, the NO production was then determined by the Griess reagent system (Sun et al., 2003). A non-cytotoxic dose (0-200 μg/mL) of BR-WG-P was measured by MTT assay and showed no growth effects in RAW macrophage cells and was later used in the next experiment (data not shown). NO production could be induced in the LPS-treated RAW macrophage cells. Pretreatment of the cells with BR-WG-P dramatically decreased the LPS-induced NO production in the cells in a dose dependent manner (Figure 1A). A catalytic enzyme iNOS is involved with NO production. The expression of iNOS was upregulated in LPS-induced RAW macrophage cells. However, the expression of iNOS was significantly decreased in a dose dependent manner when the cells were pretreated with BR-WG-P in various concentrations (50-150 μg/mL) (Figure 1B). These results suggested that BR-WG-P extract could reduce iNOS expression leading to a decrease of NO production in LPS-induced RAW macrophage cells.

**Effect of BR-WG-P on COX-2 expression in LPS-induced RAW macrophage cells**

Alteration of COX-2 expression and its enzymatic production, PGE₂, play important roles in the development of inflammation (Ricciotti and FitzGerald, 2011). The expression of COX-2 was dramatically induced in LPS-treated RAW macrophage cells compared with the untreated cells (Figure 2). In contrast, when the cells were pretreated with various concentrations of BR-WG-P, the COX-2 expression was clearly decreased in a dose dependent manner (Figure 2). These results revealed that BR-WG-P could decrease COX-2 expression which was induced by LPS in RAW macrophage cells.

**Effect of BR-WG-P on pro-inflammatory cytokine secretion in LPS-induced RAW macrophage cells**

Pro-inflammatory cytokines including, IL-6 and TNF-α play a key role in the inflammation cascade. These pro-inflammatory cytokine secretions were evaluated using an ELISA kit. The secretions of IL-6 and TNF-α from RAW macrophage cells were induced by LPS. However, when the cells were treated with BR-WG-P the IL-6 and TNF-α secretions from RAW macrophage cells were dramatically reduced in a dose dependent manner (Figure 3).

**Effect of BR-WG-P on NF-κB and AP-1 translocation in LPS-induced RAW macrophage cells**

![Graph showing NO production](image)

**Figure 1. Nitric Oxide Production was Determined Using the Griess Reagent System.** (A). The cells were pre-treated with or without various concentrations of BR-WG-P (0-150 μg/mL) for 2 h and further incubated with or without LPS (1 μg/mL) for 24 h. The nitrite production in the cultured medium was measured at 540 nm using a microplate reader. iNOS expression was determined in RAW macrophage cells pretreated with or without BR-WG-P (0-150 μg/mL) and further incubated with or without LPS (1 μg/mL) using western blot analysis (B). All assays have been demonstrated in triplicate and the mean ± standard deviations are shown as *p<0.05, or **p<0.01 versus LPS treated RAW macrophage cells (1 μg/mL).

![Graph showing COX-2 expression](image)

**Figure 2. COX-2 Expression was Determined in RAW Macrophage Cells that were Pre-treated with or without BR-WG-P (0-150 μg/mL) and Further Incubated with or without LPS (1 μg/mL) using Western Blot Analysis.** This data is representative of three independent experiments.

![Graph showing IL-6 and TNF-α secretion](image)

**Figure 3. The IL-6 and TNF-α Secretion in Cultured Medium was Determined by an ELISA Kit.** The RAW macrophage cells were co-treated with or without various concentration of BR-WG-P (0-150 μg/mL) and LPS (1 μg/mL) for 24 h. The IL-6 or TNF-α in the cultured medium was measured at 540 nm using a microplate reader. All assays have been demonstrated in triplicate and the mean ± standard deviations are shown as *p<0.05, or ****p<0.0001 versus LPS treated RAW macrophage cells (1 μg/mL).
The major transcription factors that control pro-inflammatory cytokine expression were NF-κB and AP-1 (Lawrence, 2009). NF-κB and AP-1 translocation were determined by the detection of NF-κB and AP-1 expression in nuclear extract using western blot analysis. LPS could induce the NF-κB and AP-1 translocation whereas the pretreatment of BR-WG-P (150 μg/mL) obviously decreased NF-κB and AP-1 translocation (Figure 4). Moreover, pretreatment of the cells with anthocyanin (100 μg/mL) that was highly contained in BR-WG-P could significantly decrease the LPS-induced NF-κB and c-Jun (AP-1 heterodimer) translocation (Figure 4). The pretreatment of hydroxybenzoic acid (25 μg/mL) also decreased the LPS-induced NF-κB and AP-1 translocation whereas vanillic acid and protocatechuic acid treatment revealed no differences in the NF-κB expression level while the expression of AP-1 was slightly decreased compared with the LPS-induced RAW macrophage cells (data not shown). These results suggested that anthocyanin and hydroxybenzoic acid which are major compounds in BR-WG-P could inhibit NF-κB and AP-1 translocation in LPS-induced RAW macrophage cells.

Effect of BR-WG-P on MAPK signaling pathway

MAPK signaling pathway is one of the signal transduction that control NF-κB and AP-1 activation (Fang et al., 2004; Kim et al., 2014). Therefore, the MAPK protein phosphorylation and expression including pERK, ERK, pp38, p38 pJNK and JNK were investigated. The level of pERK, pp38, and pJNK were significantly increased in LPS-induced RAW macrophage cells (Figure 5); while the phosphorylation of the MAPK signaling proteins was dramatically decreased in a dose dependent manner after pretreating the cells with BR-WG-P (Figure 5). These data revealed that BR-WG-P could inhibit MAPK (ERK, p38, and JNK) activation resulting in inhibition of NF-κB and AP-1 translocation into the nucleus.

Discussion

The anthocyanin rich fraction that was extracted from several kind of fruits and vegetables including berry, cranberry or grape exhibited antioxidant, anti-proliferative, and anti-inflammatory properties (Olsson et al., 2004; Seeram et al., 2004; Yi et al., 2005). In Thailand, many people interested in the health value of colored rice such as black rice as an anthocyanin rich choice. Whole grains black rice (Oryza sativa L.) were harvested in the wet season from Doi Saket Organic Farm, Chiang Mai Province, Thailand. After that the extraction was performed using 50% ethanol and then partition with saturated butanol to obtain polar fraction of BR-WG-P. Although, BR-WG-P provided a low percentage yield (0.307%), it contains a high phenolic content especially the flavonoids group, anthocyanin (Table 1) which plays major roles in anti-inflammatory effect (Min et al., 2010).

Pathogen induced chronic inflammation was related to an increase in NO production which is a mediator and regulator of inflammatory responses (Korhonen et al., 2005). The LPS induced RAW macrophage cells revealed an increase in NO production whereas after pre-treatment with BR-WG-P, NO production was reduced (Figure 1A). The key enzyme that regulates NO production is inducible nitric oxide synthase (iNOS) which is primarily responsible for NO production in the inflammatory processes (Zamora et al., 2000). In this study iNOS expression was dramatically increased in LPS-induced RAW macrophage cells (Figure 1B) resulting...
in an increase in NO production that was involved with the inflammatory process. Pretreatment of BR-WG-P obviously decreased the iNOS expression which lead to a reduction in NO production. Likewise iNOS expression, COX-2 became upregulated after being stimulated by LPS which was dramatically inhibited when the cells were pre-treated with BR-WG-P (Figure 2). A reduction of COX-2 expression resulted in a decrease of prostaglandin E2 (PGE2) production which is involved in the inflammatory process (Kalinski, 2012) by promoting local vasodilatation and induction of neutrophils, macrophages, and mast cells in the early stages of inflammation (Wang and Lau, 2006).

The pro-inflammatory cytokines, such as IL-1β, IL-6 and TNF-α are significant for controlling the inflammatory process (Moller and Villiger, 2006). IL-6 not only acts as an inducer of acute phase reactions but also plays an important role in chronic inflammation (Gabay, 2006) and TNF-α is also significant in inflammatory disease (Bradley, 2008). Therefore, an investigation into certain natural products, particularly black rice that could reduce cytokines secretion is important. This study found that LPS induced IL-6 and TNF-α secretion from RAW macrophage cells, while the secretion levels were significantly decreased when the cells were treated with BR-WG-P (Figure 3). The mechanisms that are involved in the production of pro-inflammatory mediators and cytokines including iNOS, COX-2, IL-1, IL-6 and TNF-α that play critical roles in inflammatory process may be controlled by the transcription factor NF-kb or AP-1 (Barnes and Karin, 1997). Stimulation occurs through the interaction between LPS and TLR4 that recruit the adaptor protein for the induction of NF-kb or AP-1 translocation into the nucleus via the activation of the mitogen-activated protein kinase (MAPK) signaling pathway (Fang et al., 2004). All of these members in MAPK including extracellular signal-regulated kinase (ERK), p38, and c-Jun NH2-terminal kinase (JNK) play a major role in the control of AP-1 activation (Kiyiakis, 1999). This data is related to the previous results that showed a decrease in NF-kb and AP-1 translocation into the nucleus (Figure 4) and a reduction of the phosphorylation form of ERK, p38 and JNK after pre-treating the cells with BR-WG-P (Figure 5). Moreover, anthocyanin which is enriched in BR-WG-P may be the active compound for inhibition of NF-kb and AP-1, the key transcription factor involved with inflammation (Figure 4).

Another pathway that controls pro-inflammatory mediators and cytokines production may be involved with the Ikk/NF-kb signaling pathway (Kawai and Akira, 2007). The stimulation occurs through the interaction between LPS and TLR4 and leads to the activation of the Ikb kinase (Ikk) complex, which induces the phosphorylation of Iksbα leading to NF-kb translocation and activation (Akira, 2003; Hoesel and Schmid, 2013). Therefore, BR-WG-P not only inhibits the NF-kb transcription factor but also inhibits pro-inflammatory cytokine, IL-6 and TNF-α secretion.

These findings suggested that anthocyanin and hydroxybenzoic acid enriched BR-WG-P could inhibit NF-kb and AP-1 activation that controls the expression of pro-inflammatory mediator including iNOS, COX-2, IL-6 and TNF-α, leading to a decrease in the inflammatory process that is involved with chronic diseases including cancer via the inhibition of the MAPK signaling pathway.

References


