



Inhibitory effect of Gardenblue blueberry (*Vaccinium ashei* Reade) anthocyanin extracts on lipopolysaccharide-stimulated inflammatory response in RAW 264.7 cells*

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Abstract: Blueberries are a rich source of anthocyanins, which are associated with health benefits contributing to a reduced risk for many diseases. The present study identified the functional Gardenblue blueberry (*Vaccinium ashei* Reade) anthocyanin extracts (GBBAEs) and evaluated their capacity and underlying mechanisms in protecting murine RAW 264.7 cells from lipopolysaccharide (LPS)-stimulated inflammation in vitro. Enzyme-linked immunosorbent assay (ELISA) kit results showed that GBBAEs significantly inhibited the production of nitric oxide (NO), prostaglandin E₂ (PGE₂), interleukin-6 (IL-6), IL-1 β , and interferon- γ (INF- γ). Real-time polymerase chain reaction (PCR) analysis indicated that the mRNA expression levels of IL-6, IL-1 β , tumor necrosis factor- α (TNF- α), monocyte chemoattractant protein-1 (MCP-1), and cyclooxygenase 2 (COX-2) were suppressed in LPS-stimulated RAW 264.7 cells. Additionally, Western blot analysis was used to evaluate the relative protein expression levels of COX-2 and nuclear factor- κ B p65 (NF- κ Bp65). All these results suggested the potential use of GBBAEs as a functional food for the treatment of inflammatory diseases.

Key words: Gardenblue blueberry (*Vaccinium ashei* Reade) anthocyanin extracts (GBBAEs), Anti-inflammatory, RAW 264.7, Cyclooxygenase 2 (COX-2), Nuclear factor- κ B p65 (NF- κ Bp65)

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1 Introduction

Inflammation response is a complex and crucial process mediated by both several inflammatory cytokines (interferon- γ (INF- γ), interleukin-1 β (IL-1 β), and IL-6) and related inflammatory mediators such as prostaglandin E₂ (PGE₂) and nitric oxide (NO), which are produced by cyclooxygenase 2 (COX-2) and inducible nitric oxide synthase (iNOS), respectively (Kim *et al.*, 2013; Cheng *et al.*, 2014). COX-2 and

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nuclear factor- κ B (NF- κ B) activations are well-known biological markers for inflammatory responses. Inflammatory cells such as monocytes and macrophages can produce a large repertoire of cytokines and participate in the pathogenesis of diseases (Yan *et al.*, 2013). So it is valuable to recognize anti-inflammatory mechanisms in order to find treatments and adjuvant treatments of many diseases.

Lipopolysaccharide (LPS) is an endotoxin, an integral outer membrane component used to assess Gram-negative bacteria, and triggers the most potent microbial initiators of inflammatory response, including septic shock, fever, and microbial invasion (Dobrovolskaia and Vogel, 2002). LPS induces NF- κ B activation via several signal transduction pathways, all of which inhibit the phosphorylation of NF- κ B (inhibitor of NF- κ B (I κ B)) (Kim *et al.*, 2010). The LPS-stimulated RAW 264.7 cell model is one of the typical virus-induced inflammation models *in vitro*.

Blueberries are rich in a variety of functional components, such as flavone, organic acid, anthocyanin, superoxide dismutase (SOD), polysaccharides, and mineral elements, contributing to the treatments for many diseases (Almeida *et al.*, 2007). Although Gardenblue blueberry (*Vaccinium ashei* Reade) anthocyanin extracts (GBBAEs) are best recognized for their antioxidant properties with their mechanisms being studied thoroughly (Prior *et al.*, 1998; Sun *et al.*, 2012; Li *et al.*, 2013), recent studies indicated that it is the anti-inflammatory aspect that is responsible for treating cancer and cardiovascular diseases (Johnson *et al.*, 2013; Esposito *et al.*, 2014). Varieties of polysaccharides, glycosides, and alkaloids have proven to have the best anti-inflammatory activity (Hsu *et al.*, 2013; Wang L. *et al.*, 2014). Also, there are studies on the anti-inflammatory effect of blueberries (Cheng *et al.*, 2014; Esposito *et al.*, 2014), but the detailed capacity and underlying mechanisms in GBBAEs are still to be elucidated. There were only a few reports on the verification of the anti-inflammatory capacity and mechanisms of the RAW 264.7 cell model. Also, the mechanism of the NF- κ B pathway in GBBAEs was evaluated for the first time. Therefore, the objectives of the current study were: (1) to evaluate the inhibitory effect of inflammatory cytokines (NO, PGE₂, IL-6, IL-1 β , and INF- γ) release; (2) to evaluate the mRNA gene expression levels of the IL-6, IL-1 β , tumor ne-

crolysis factor- α (TNF- α), monocyte chemoattractant protein-1 (MCP-1), and COX-2 in LPS-stimulated RAW 264.7 cells; (3) to evaluate the relative protein expressions of COX-2 and NF- κ Bp65; (4) to determine the anti-inflammatory mechanism of GBBAEs.

2 Materials and methods

2.1 Materials and chemicals

The blueberry we used is Gardenblue blueberry. They are obtained from Hubei Cowherd Blueberry Technology Co., Ltd., whose planting base was located in Huangpi District, Hubei Province, China. The cell culture medium (GNM-12800), trypsinase (GMN-15400), and penicillin-streptomycin were purchased from Jinuo Biotechnology Co., Ltd. (Hangzhou, China). Fetal bovine serum (FBS) was purchased from Hangzhou Tianhang Biotechnology Co., Ltd. (Hangzhou, China) and the Dulbecco's modified Eagle's medium (DMEM) was purchased from Hyclone (USA). 3-(4,5-Dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT) and LPS were obtained from Sigma (USA). Enzyme-linked immunosorbent assay (ELISA) kits for mouse PGE₂, IL-6, IL-1 β , and INF- γ were purchased from Nanjing Jiancheng Bioengineering Institute (Nanjing, China). TRIzol reagent was obtained from Invitrogen Life Technologies (Germany) and the RevertAid First Strand cDNA Synthesis Kit was obtained from Thermo (Germany). FastStart Universal SYBR Green Master (Rox) was purchased from Roche (Switzerland). All other reagents were of analytical grade.

2.2 GBBAEs' preparation and high-performance liquid chromatography-mass spectrometry/mass spectrometry analysis

The GBBAEs were obtained from fresh-frozen Gardenblue blueberry. Briefly, the 50 g of fresh-frozen Gardenblue blueberry was weighed out and placed in a capacity tube, followed by the addition of 60% alcohol (15 ml/g) as extracting solvent into the sample tube. Then 0.1 mol/L HCl was added to adjust the pH to 3. After being mixed briefly, the sample tubes were placed on magnetic stirrers at 40 °C for 2 h. The extraction process on magnetic stirrers was repeated twice. Then 5 mg/ml aqueous solution of the

crude extracts was loaded onto an AB-8 resin (weak polarity macroporous resin, 0.3–1.25 mm particle size, Nankai Hecheng Science & Technology Co., Tianjin, China) column (25 mm×100 mm). After that, the column was water-washed. Finally the fraction eluted with 60% ethanol was collected. After the eluant was evaporated and freeze-dried, GBBAEs were obtained and then stored at $-20\text{ }^{\circ}\text{C}$ in sealed aluminum bags for future experiment. pH-differential method and high-performance liquid chromatography (HPLC) with a C18 column were applied to the total anthocyanin content using cyanidin 3-glucoside as adequate standards to calculate the equivalent per 100 g in the GBBAEs (Zhang *et al.*, 2015).

The individual anthocyanins were separated by an Agilent 1100 HPLC equipped with a diode array detector connected to an Agilent 6300 mass spectrometer (Agilent Technologies Inc., Santa Clara, CA, USA), and a $5\text{ }\mu\text{m}\times 250\text{ mm}\times 4.6\text{ mm}$ TSKgel C18 column (Tosoh Corp., Tokyo, Japan). The temperature of the column was set at $30\text{ }^{\circ}\text{C}$ during the HPLC analysis. The injection volume of the solution (1 mg/ml) of GBBAEs was 10 μl . Solvent A was 5% (v/v) formic acid in water, and solvent B was 100% methanol. The solvent gradient was: 0 to 5 min, 100% to 75% A; 5 to 25 min, 75% to 60% A; 25 to 40 min, 60% to 50% A; 40 to 42 min, 50% to 55% A; 42 to 44 min, 55% to 40% A; 44 to 46 min, 40% to 35% A; 46 to 52 min, 35% to 30% A. The individual anthocyanins were obtained by HPLC-diode array detection (DAD) at 525 nm. For mass spectrometry (MS) acquisition, mass spectra in the m/z range of 100–1000 were obtained by electrospray ionization in positive-ion mode. For MS acquisition, a capillary voltage of 4.5 kV, a pressure of nebulizing nitrogen of 35 psi (1 psi=6.895 kPa), a capillary temperature of $325\text{ }^{\circ}\text{C}$, and dry gas flow at 10 L/min were applied. The compounds were presented by the relative contents by comparing the peak area of HPLC-MS/MS.

2.3 Cell culture

Culture RAW 264.7 cell lines were gifts from the College of Animal Science and Technology, Huazhong Agricultural University (Wuhan, China). RAW 264.7 cells were cultured in DMEM supplemented with 10% FBS and 1% penicillin-streptomycin at $37\text{ }^{\circ}\text{C}$ in a 5% CO_2 humid atmosphere (Thermo, Germany), until they reached 90% confluence.

2.4 Cell viability assay

The cytotoxicity of GBBAEs against the RAW 264.7 cells was assessed by an MTT assay (Lopes *et al.*, 2012). RAW 264.7 cells were plated into 96-well plates in triplicate at a density of 2×10^4 cells/well and incubated for 24 h. After 24 h, the culture medium was replaced by 200 μl serial dilutions of GBBAEs (50, 100, 200, 400, 800, 1200, 1600 $\mu\text{g/ml}$, each dilution owns six duplicate wells) and further incubated for 24, 48, and 72 h at $37\text{ }^{\circ}\text{C}$ in a 5% CO_2 humid atmosphere. Sterile filtered MTT stock solution (5 mg/ml) was diluted to 0.5 mg/ml by phosphate-buffered saline (PBS, pH=7.4) and then was added to each well to reach the reaction volume of 200 μl . After 4-h incubation, the supernatant medium was removed, and dimethyl sulfoxide (DMSO) (150 μl) was added to each well to dissolve the formazan crystals. Then the plates were read immediately at 490 nm on a microplate reader (BioTek, Vermont, USA). The cell viability was calculated by the following formula: cell viability= $(A_e - A_{bc}) / (A_c - A_{bc}) \times 100\%$, where A_e , A_c , and A_{bc} are the mean absorbances of experiment, control, and blank control, respectively.

2.5 NO production assay

The NO levels in the cultured media, which reflect NO activity, were determined by the Griess reaction (Hsu *et al.*, 2013). RAW 264.7 cells were plated into 24-well plates at a density of 1×10^6 cells/well and incubated for 24 h. After the culture period, cells were exposed to different concentrations of GBBAEs (0, 400, 800, 1200, 1600 $\mu\text{g/ml}$, diluted by GNM-12800 culture medium, each dilution owns three duplicate wells) for 1 h. After 1 h, 20 μl LPS (50 $\mu\text{g/ml}$) was added to each well to reach the final concentration of 1 $\mu\text{g/ml}$ and cells were maintained in culture for 24 h under the same conditions. An assay without LPS was prepared. After incubation, 100 μl of the cell culture supernatant was mixed with an equal volume of Griess reagent (0.1% (1 g/L) *N*-(1-naphthyl) ethylenediamine dihydrochloride and 1% (0.01g/ml) sulfanilamide in 5% (v/v) phosphoric acid). The samples were incubated at room temperature for 15 min, and then the absorbance was measured at 540 nm using a microplate reader. The quantity of NO was determined from a sodium nitrite standard curve. Also, the supernatant was removed and stored at $-20\text{ }^{\circ}\text{C}$ for later analysis.

2.6 Determination of PGE₂ production

The GBBAEs were diluted with DMEM prior to treatment. Cells were treated for 24 h with LPS (1 µg/ml) to permit cytokine production. The PGE₂ concentration in the culture medium was quantified using a competitive enzyme immunoassay kit (R&D Systems, USA) in accordance with the manufacturer's instructions. The production of PGE₂ was measured relative to that observed after control treatment.

2.7 Measurement of proinflammatory cytokine (INF-γ, IL-1β, and IL-6) production

The GBBAEs were diluted with DMEM prior to treatment. The inhibitory effects of GBBAEs on the production of proinflammatory cytokines (INF-γ, IL-1β, and IL-6) from LPS-treated RAW 264.7 cells were determined as described previously. The supernatants were subsequently employed for the proinflammatory cytokine assays using a mouse ELISA kit (R&D Systems, USA).

2.8 Real-time PCR analysis

The total RNA was extracted from RAW 264.7 cells and collected after GBBAEs and LPS treatments using TRIzol reagent (Invitrogen, USA) according to the manufacturer's instructions. Two microgram of total RNA was added to the tubes with 1 µl of oligo (dT) 15 primers and added deionized water without RNase to 20 µl. These tubes were kept at 70 °C for 5 min and cooled rapidly. Then 4 µl of 5× buffer, 2 µl of 10 mmol/L dNTPs, 1 µl of RNA inhibitor, and 1 µl of inverse transcriptase in each were added. Then the tubes were kept at 42 °C for 30 min, and the reactions were stopped by heating the RNA solution at 80 °C for 5 min. Complementary DNA (cDNA) was generated using the RevertAid First Strand cDNA Synthesis Kit (Thermo, Germany). cDNA was used as a template to amplify for real-time polymerase chain reaction (RT-PCR) in triplicates with FastStart Universal SYBR Green Master (Rox) and gene-specific primers. PCR analysis was performed on an ABI (Step-One-Plus) Real-Time PCR system. PCR primers used in this study were as follows: M-GAPDH-S, 5'-AGG AGC GAG ACC CCA CTA ACA-3'; M-GAPDH-A, 5'-AGG GGG GCT AAG CAG TTG GT-3'; M-MCP-1-S, 5'-AGG TGT CCC AAA GAA GCT GTA GT-3'; M-MCP-1-A, 5'-TTT GGT TCC GAT CCA GGT TTT-3'; M-IL-6-S, 5'-GAA ATG

ATG GAT GCT ACC AAA CTG-3'; M-IL-6-A, 5'-GAC TCT GGC TTT GTC TTT CTT GTT-3'; M-TNF-α-S, 5'-TAC TGA ACT TCG GGG TGA TCG-3'; M-TNF-α-A, 5'-GGG TCT GGG CCA TAG AAC TGA-3'; M-IL-1β-S, 5'-TCA AAT CTC GCA GCA GCA CAT C-3'; M-IL-1β-A, 5'-CGT CAC ACA CCA GCA GGT TAT C-3'; M-COX-2-S, 5'-GAA GTT GAT AAC CGA GTC GTT C-3'; M-COX-2-A, 5'-CCA TAG AAT AAC CCT GGT CG-3'.

The PCR amplifications were performed with initial heat denaturation at 95 °C for 10 min and the PCR cycles were repeated 40 times under the following conditions: denaturation at 95 °C for 15 s, annealing at 60 °C for 60 s, and the last extension at 60 °C for 5 min. The dissociation curve was completed with one cycle from 75 to 95 °C (20 s/°C). The expression levels relative to the control were estimated by calculating $\Delta\Delta C_T$ and subsequently analyzed using $2^{-\Delta\Delta C_T}$ method (Kim *et al.*, 2013).

2.9 Western blot analysis

Anti-inflammatory protein expression was evaluated by Western blot analysis according to standard procedures. The cells were pretreated with GBBAEs and stimulated with LPS for indicated periods at 37 °C. After incubation, the cells were washed with Tris-buffered saline (TBS) twice, then collected and lysed in ice-cold radio-immunoprecipitation assay (RIPA) lysis buffer (50 mmol/L Tris-HCl (pH 7.4), 150 mmol/L NaCl, 1% Nonidet P-40 (NP-40), 0.1% (1 g/L) sodium dodecyl sulfate (SDS), and 100 mmol/L phenylmethanesulfonyl fluoride (PMSF)), containing a protease inhibitor 50× cocktail (Roche, Basel, Switzerland) for 30 min on ice, then centrifuged at 12 000g for 5 min to obtain the total protein solution. The protein concentration of extracts was estimated by the Bradford protein assay (Bio-Rad, CA, USA). Also, after quantification of protein concentration, equal amounts of proteins were subjected to SDS-polyacrylamide gel electrophoresis (PAGE) and transferred onto polyvinylidene fluoride (PVDF) membranes. Skim milk (5%) in TBS with 0.1% Tween 20 (TBST) was used to block the membranes. Then the membranes were incubated with each specific primary antibody against COX-2 and NF-κBp65 overnight at 4 °C. The membranes were washed three times for 10 min each between each step and incubated with horseradish peroxidase (HRP)-conjugated

secondary anti-bodies for 30 min at room temperature, and then washed again as before. The protein bands were detected by the Amersham ECL system (Amersham-Pharmacia Biotech, Arlington Heights, IL, USA). The relative expression of proteins was quantified using the software Alpha (Alpha Innotech, ProteinSimple, USA) and calculated by the reference bands of β -actin.

2.10 Statistical analysis

Data are expressed as the mean \pm standard deviation (SD) values with at least three replications. The one-way analysis of variance (ANOVA) with treatment as a factor was used to evaluate the significant differences between groups using SPSS Version 18.0 (Esposito et al., 2014). $P < 0.05$ was considered to be significant.

3 Results

3.1 Major compounds of GBBAEs

The major compounds in GBBAEs were explored by HPLC-MS/MS analysis. As presented in Fig. 1 and Table 1, the major compounds were identified by comparison of their mass spectra while the relative contents were calculated by the peak areas. The GBBAEs consist of 11 major compounds. The relative contents of all compounds range from 3% to 22%. Malvidin-3-galactoside ((19.5656 \pm 0.1746)%) and malvidin-3-glucoside ((21.5313 \pm 0.0298)%) are the two higher level compounds. The ranges of petunidin-3-glucoside, cyanidin-3-glucoside, delphinidin-3-glucoside, delphinidin-3-galactoside, cyanidin-3-galactoside, and petunidin-3-galactoside are almost stable, from (6.9430 \pm 0.0220)% to (9.9280 \pm 0.0618)%. Malvidin-

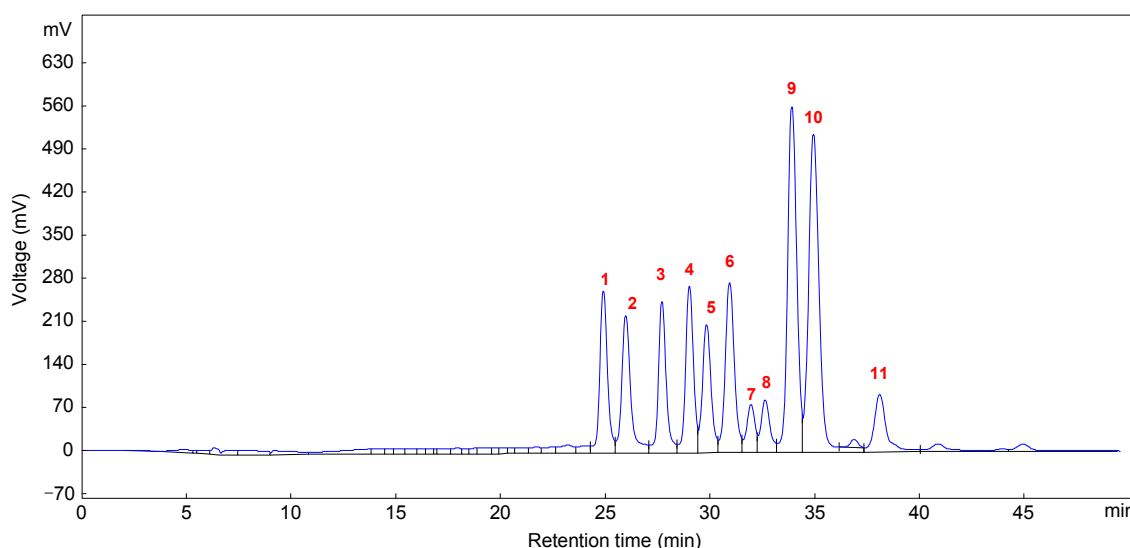


Fig. 1 HPLC analysis of the GBBAEs at 525 nm
Compounds 1–11 are shown in Table 1

Table 1 Identification and analysis of major compounds in GBBAEs by HPLC-ESI-MS/MS detected at 525 nm

Peak	Rt (min)	[M] ⁺ (m/z)	MS/MS (m/s)	Identification	Relative content (%)
1	24.870	465	303	Delphinidin-3-galactoside	7.9711 \pm 0.1313
2	25.939	465	303	Delphinidin-3-glucoside	8.0819 \pm 0.1535
3	27.669	449	287	Cyanidin-3-galactoside	7.4760 \pm 0.0717
4	28.977	449	287	Cyanidin-3-glucoside	8.1975 \pm 0.0607
5	29.798	479	317	Petunidin-3-galactoside	6.9430 \pm 0.0220
6	30.897	479	317	Petunidin-3-glucoside	9.9280 \pm 0.0618
7	31.918	419	287	Cyanidin-3-arabioside	2.6159 \pm 0.0283
8	32.600	463	301	Peonidin-3-galactoside	3.0492 \pm 0.0437
9	33.671	493	331	Malvidin-3-galactoside	19.5656 \pm 0.1746
10	34.903	493	331	Malvidin-3-glucoside	21.5313 \pm 0.0298
11	38.065	463	331	Malvidin-3-arabioside	4.6404 \pm 0.4218

Rt: retention time. Relative content values are expressed as mean \pm SD (n=3)

3-arabinoside, peonidin-3-galactoside, and cyanidin-3-arabinoside are three smaller parts. The cyanidin-3-arabinoside content is only $(2.6159 \pm 0.0283)\%$.

3.2 Effect of GBBAEs on RAW 264.7 cell viability

We first measured the cytotoxicity of GBBAEs in RAW 264.7 cells using the MTT assay. Fig. 2a shows that GBBAEs-treated groups showed markedly increased absorbance values in the employed concentrations (400, 800, and 1200 $\mu\text{g/ml}$), when the processing time was 24 h, reaching the maximum levels (141.78%, 147.87%, and 151.79%; $P > 0.05$). However, the number of cells in each group was significantly decreased after 48 h. When the processing time was 72 h, the lowest concentration was significantly reduced. This condition may be related to the differentiation and proliferation of GBBAEs in vitro and the short survival time. Therefore, 24 h was considered to be the optimal inoculation time for the time-dependent analysis.

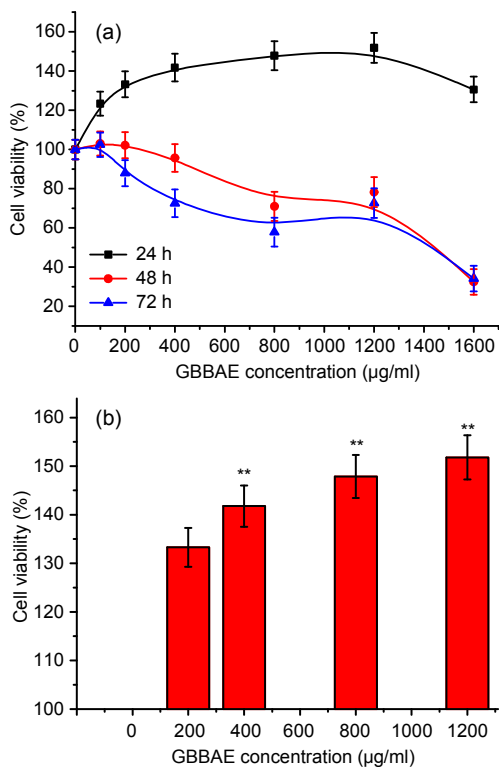


Fig. 2 Effects of GBBAEs on cell viability of murine RAW 264.7 cells

(a) Cells were treated 0, 200, 400, 800, 1200, and 1600 $\mu\text{g/ml}$ GBBAEs for 24, 48, and 72 h, respectively. Cell viability was quantified spectrophotometrically by the MTT assay. (b) Cells were treated 200, 400, 800, and 1200 $\mu\text{g/ml}$ GBBAEs for 24 h, respectively. The reported values are the mean \pm SD ($n=6$). ** $P < 0.01$ versus 200 $\mu\text{g/ml}$ GBBAEs

As shown in Fig. 2b after GBBAE administration for 24 h, the cell viability of all groups increased. Comparing the concentrations used, the growth of RAW 264.7 was promoted by GBBAEs in a dose-dependent manner. GBBAEs did not influence the cytotoxicity of RAW 264.7 cells at the concentrations used (400, 800, and 1200 $\mu\text{g/ml}$). Therefore, in all subsequent experiments, we used concentrations of GBBAEs ranging from 400 to 1200 $\mu\text{g/ml}$.

3.3 Effect of GBBAEs on the NO production

NO is a signaling molecule involved in a broad spectrum of pathophysiological processes, such as inflammation, apoptosis, regulation of enzyme activity, and gene expression. High NO levels are generated in response to inflammatory stimuli and mediate proinflammatory effects (Korhonen *et al.*, 2005). As shown in Fig. 3, the NO production of LPS-induced RAW 264.7 was markedly increased to 100% when 1 $\mu\text{g/ml}$ LPS was added, compared with 3.15%–5.53% for untreated control cells. The NO production decreased significantly in a dose- and time-dependent manner. When GBBAE administration was 1200 $\mu\text{g/ml}$ for 24 h, the NO production decreased by 33.11% ($P < 0.05$), compared with 18.40% in 12 h and 19.54% in 48 h. According to previous studies (Yan *et al.*, 2013), the NO inhibitory effect is similar to the fingered citron. Although the anti-inflammatory effect of blueberry extracts has been explored in animal experiments (Shi *et al.*, 2014), that in NO production of LPS-induced RAW 264.7 has not been reported.

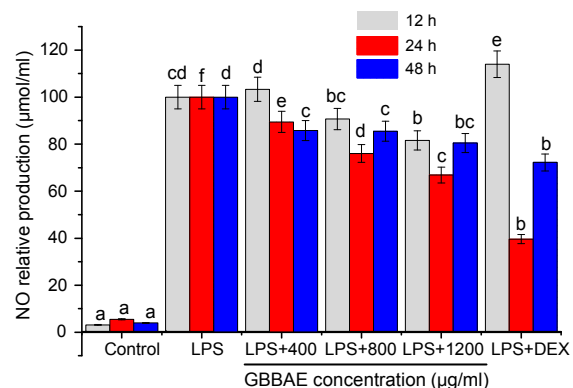


Fig. 3 Effects of GBBAEs on NO production in LPS-stimulated murine RAW 264.7 cells

Cells were pretreated with LPS (1 $\mu\text{g/ml}$) before stimulated with 400, 800, and 1200 $\mu\text{g/ml}$ of GBBAEs in the presence for 12, 24, and 48 h, respectively. As a control, cells were incubated with vehicle alone. The reported values are the mean \pm SD ($n=3$). Different letters indicate significant differences ($P < 0.05$). DEX: dexamethasone

3.4 Effects of GBBAEs on the secretions of PGE₂, INF-γ, IL-1β, and IL-6

To determine the effects of GBBAEs on the secretions of the proinflammatory cytokines PGE₂, INF-γ, IL-1β, and IL-6, RAW 264.7 cells were pretreated with various concentrations of GBBAEs (400, 800, and 1200 μg/ml) and stimulated with 1 μg/ml of LPS for 24 h. As shown in Figs. 4a–4d, the levels of PGE₂, INF-γ, IL-1β, and IL-6 were significantly increased in the LPS-treated cells when compared with the untreated cells. However, the levels of PGE₂, INF-γ, IL-1β, and IL-6 were significantly reduced in the cells pretreated with GBBAEs in a dose-dependent manner. These results demonstrate that GBBAEs inhibited the production of PGE₂, INF-γ, IL-1β, and IL-6 effectively.

3.5 Effects of GBBAEs on the mRNA expressions of COX-2, MCP-1, IL-6, IL-1β, and TNF-α

As shown in Fig. 5, the effects of GBBAEs on LPS-stimulated MCP-1, COX-2, IL-6, IL-1β, and TNF-α protein expressions in RAW 264.7 cells were

examined by RT-PCR. Figs. 5a and 5b show the gene expression levels of COX-2 and MCP-1, the only cells chosen as the control (mRNA fold production is 1). The two gene expression levels of RAW 264.7 were significantly inhibited by GBBAEs in a dose-dependent manner ($P < 0.05$) especially 1200 μg/ml. As for IL-1β and IL-6 shown in Figs. 5c and 5d, the lower GBBAE concentrations showed significant differences ($P < 0.05$) in IL-6 gene expression levels but no effect in IL-1β ($P > 0.05$). With the increase of GBBAE concentration, the two gene expression levels were completely inhibited in 1200 μg/ml, which indicated that a suitable concentration of GBBAEs can completely inhibit IL-1b and IL-6 secreted from RAW 264.7 cells. Although there were no effects ($P > 0.05$) on TNF-α gene expression in Fig. 5e, there was a tendency toward the TNF-α level decline by the increase of GBBAE concentration.

3.6 Effects of GBBAEs on the relative protein expressions of COX-2 and NF-κBp65

The relative protein expressions are shown in Fig. 5f. LPS at 1 μg/ml induced a significant increase

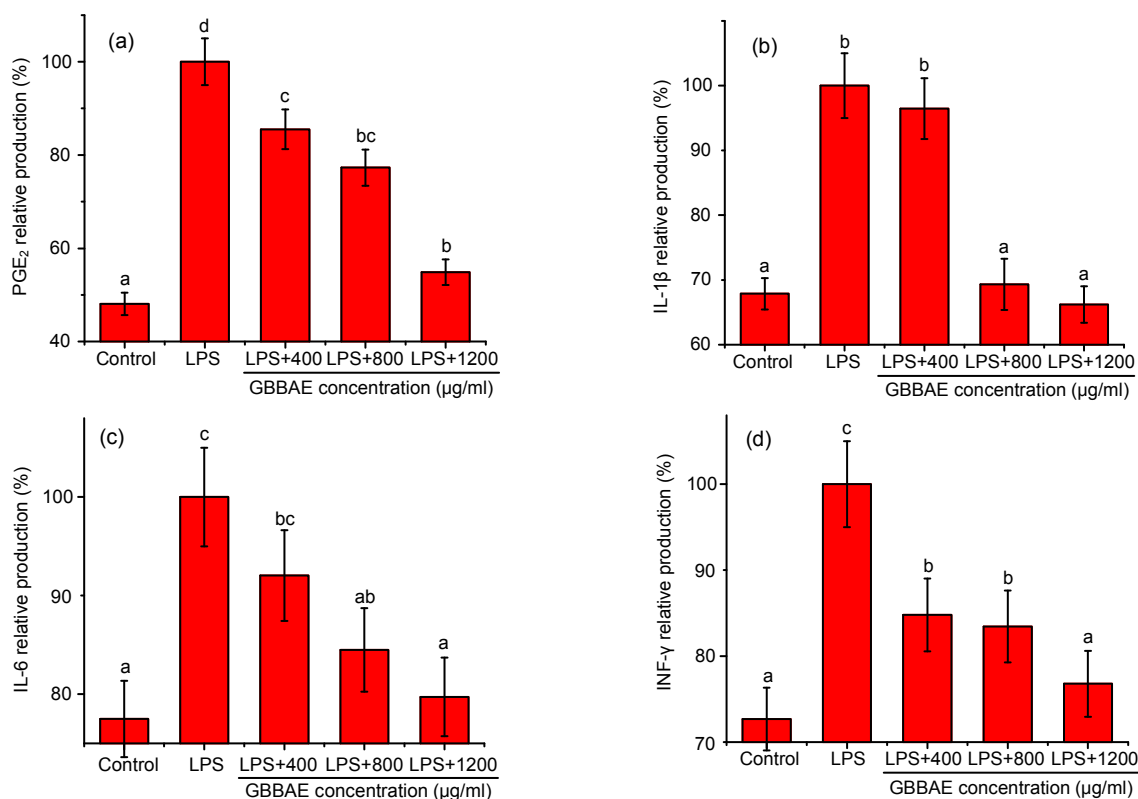


Fig. 4 Effects of GBBAEs on secretions of PGE₂ (a), IL-1β (b), IL-6 (c), and INF-γ (d) in LPS-stimulated murine RAW 264.7 cells. Cells were pretreated with LPS (1 μg/ml) before stimulated with 400, 800, and 1200 μg/ml of GBBAEs in the presence for 24 h, respectively. As a control, cells were incubated with vehicle alone. The reported values are the mean±SD (n=3). Different letters indicate significant differences ($P < 0.05$)

in COX-2 protein expression, compared with the control. In the range of 400, 800, and 1200 $\mu\text{g/ml}$, pretreatment with GBBAEs resulted in the inhibition of this LPS-induced COX-2 protein expression in a dose-dependent manner. Although the inhibitory effect was not significant at a low concentration, a significant

effect appeared when the concentration reached 1200 $\mu\text{g/ml}$. The inhibitory ratio ranged from 15.04% to 46.90%, the latter of which is close to that of the positive group (45.13%). This finding is in accordance with the mRNA expression of COX-2 in Fig. 5a.

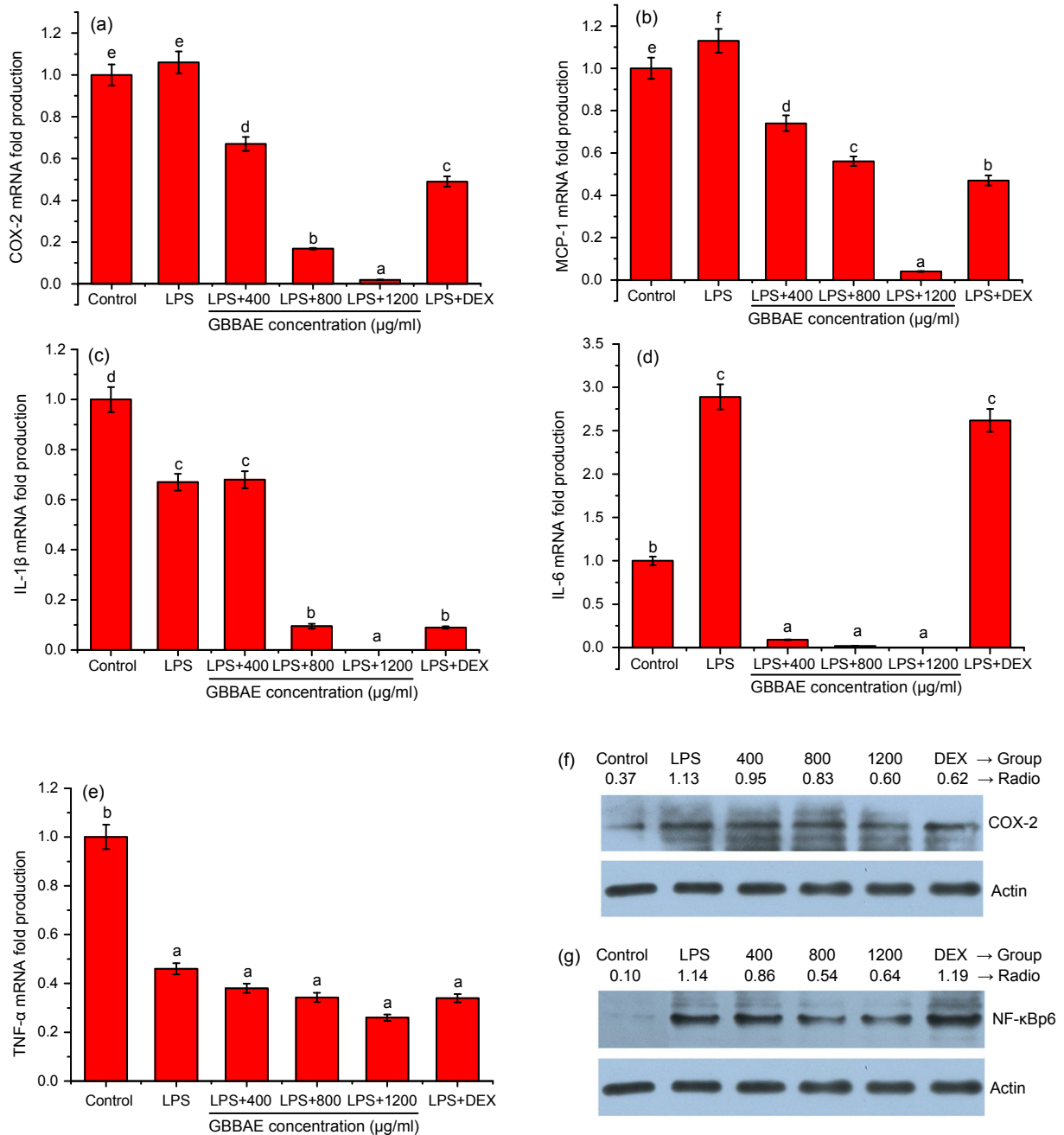


Fig. 5 Effects of GBBAEs on mRNA expression levels of COX-2 (a), MCP-1 (b), IL-1 β (c), IL-6 (d), and TNF- α (f) and on the protein expressions of COX-2 (f) and NF- κBp6 (g) in LPS-stimulated murine RAW 264.7 cells

Cells were pretreated with LPS (1 $\mu\text{g/ml}$) before stimulated with 400, 800, and 1200 $\mu\text{g/ml}$ of GBBAEs in the presence for 24 h, respectively. As a control, cells were incubated with vehicle alone. The reported values are the mean \pm SD ($n=3$). Different letters indicate significant differences ($P<0.05$). DEX: dexamethasone

As is known, NF-κB subunits are retained in the cytoplasm by binding to the inhibitory IκB protein in unstimulated cells. Upon stimulation, dephosphorylation and subsequent degradation of IκB liberates NF-κB subunit p65 to enter the nucleus and bind to DNA target sites (Wang T.Y. et al., 2014). We examined the phosphorylation of p65 by Western blot analysis because of the primary subunit of NF-κB activated by LPS in macrophages. Western blot was also used to examine whether GBBAEs affect the binding of NF-κB. As presented in Fig. 5g, in the range of 400, 800, and 1200 μg/ml, pretreatment with GBBAEs resulted in the inhibition of this LPS-induced NF-κBp65 protein expression in a concentration-dependent manner. The inhibitory ratio ranged from 24.56% to 43.86%, but the positive group showed no significant inhibition. In summary, the results indicate that the GBBAEs suppress inflammatory response via the NF-κB pathway. The flowsheet of the anti-inflammatory effect of GBBAEs through NF-κB is presented in Fig. 6.

4 Discussion

Anthocyanins with different aglycones and sugar moieties have different bioavailability and potential

health effects (Wu et al., 2006). This study not only showed some similar effects to the former studies but also differences. Although the LPS-induced model to study the anti-inflammatory activity in BV2 microglial cells was explored by Lau et al. (2007), the model in RAW 264.7 cells has not yet been explored. The present study indicates that the RAW 264.7 cells can be successfully set up to be a platform of virus-induced inflammation as an in vitro model.

In terms of the compounds of the GBBAEs, Yousef et al. (2013) showed the richest species of 18 anthocyanins in blueberry. In the reported 18 species, the content of delphinidin 3-O-(6"-acetyl) glucoside, cyanidin 3-O-(6"-acetyl) glucoside, malvidin 3-O-(6"-acetyl), petunidin 3-O-(6"-acetyl) glucoside, and malvidin 3-O-(6"-acetyl) glucoside, which do not exist in our study, is only 4.6%. Compared with previous study (Yousef et al., 2013), our study found only 11 species which constitute the majority of the anthocyanins. Thus the abundance of other species is in accordance with most studies (Cheng et al., 2014). There are not many differences in the varieties of anthocyanins but there were in the relative content of anthocyanin compounds of different resources. Based on the differences of instruments and methods, the results obtained were in the range of allowable error.

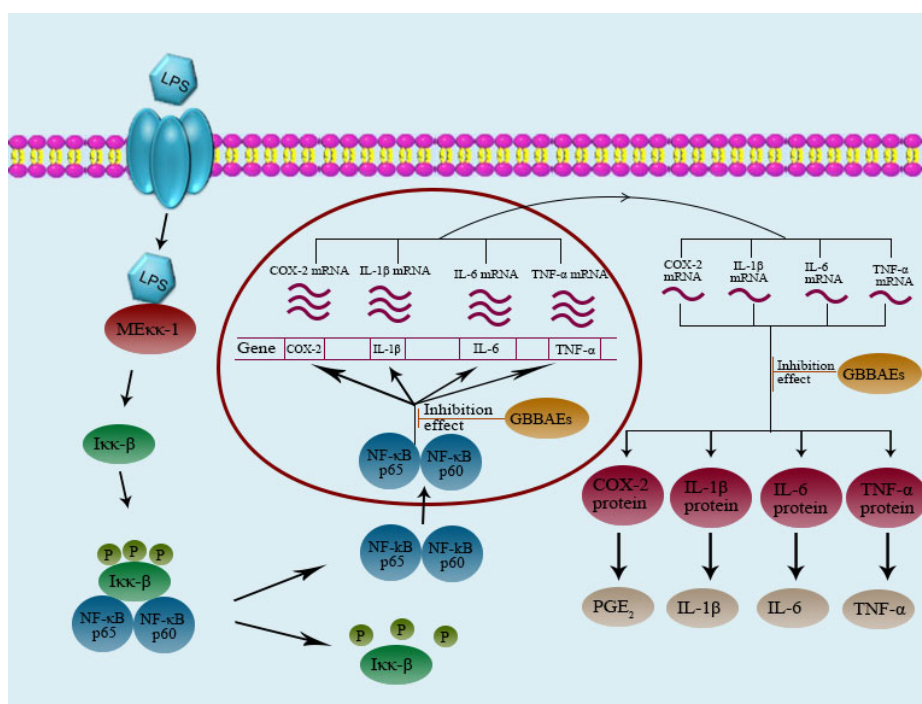


Fig. 6 Schematic diagram about the pathway of GBBAEs in LPS-stimulated murine RAW 264.7 cells

As to the NO inhibitory effect, multiple studies suggest that elevated NO production and oxidative stress in activated macrophages are closely related to inflammation, the development of atherosclerosis and cancer (Berliner *et al.*, 1995; Iqbal *et al.*, 2002; Lee *et al.*, 2003). Excessive proinflammatory cytokine and NO production through NF- κ B activation play an important role in inflammatory diseases (Southan and Szabo, 1996). The underlying mechanisms of blueberry anthocyanins have not been completely studied. In this study, the GBBAEs show a similar effect to many polysaccharides, which are widely used to prevent the inflammatory diseases. It appears that GBBAEs have the potential to be an immunopotentiator.

As to the secretions of relative cytokines, various studies concentrated on different cytokines. We chose the typical cytokines such as PGE₂, INF- γ , IL-1 β , and IL-6 for analysis. IL-1 is well-known to be a proinflammatory cytokine which is constitutively secreted and activated. It mediates macrophage chemotaxis, angiogenesis, and sustained melanoma growth. IL-1 β is the neutral isoform of IL-1 and plays a significant role in tumor development and progression (Okamoto *et al.*, 2010). IL-6 is a multifunctional cytokine that controls many biological processes such as inflammation, differentiation of immune cells, and hepatic regeneration in response to diverse stimuli (Heinrich *et al.*, 2003). Mann *et al.* (2005) reported that PGE₂ secretion associated with COX-2 can promote tumor growth and metastasis. Comparing with previous studies in various plant extracts, GBBAEs showed similarly significant inhibition effects on IL-6 and PGE₂ especially in polyphenol composition. Although the inhibition tendencies of INF- γ and IL-1 β are the same, the effect presented showed little difference.

The relative mRNA and protein expressions were evaluated further. When the same dose of GBBAEs was used on LPS-induced RAW 264.7 cells, the results demonstrated greater suppressive effects on the relative mRNA expressions of COX-2, MCP-1, and IL-6 release than on that of TNF- α and IL-1 β , which implies that GBBAEs showed selective inhibitory effects on COX-2, MCP-1, and IL-6 production. Comparing the differences with the study of Lau *et al.* (2007), all those results further confirmed the activity in various species of anthocyanins. In particular, the protein expression showed that the inhibitory effects of GBBAEs on COX-2 and NF-

κ Bp65 were owing to the attenuation of NF- κ B nuclear translocation. Upon activation, NF- κ B translocates into the nucleus to initiate the expression of over 200 immune, growth, and inflammation genes (Aggarwal, 2004). Thus the cell-signaling properties in this study may be independent of their traditional antioxidant activities. The anti-inflammatory activity contributes to the results.

Given the complexity of the inflammatory reaction, the precise mechanism underlying the neuroprotective properties of GBBAEs is unclear. Based on our results, it is important first to prove the anti-inflammatory effects by building the LPS-induced RAW 264.7 model *in vitro*. In the future, we plan to investigate further the possible role and biomedical significance of GBBAEs in NF- κ B activation as well as other stress signaling cascades.

5 Conclusions

In conclusion, we isolated 11 kinds of anthocyanins in the GBBAEs, of which the major compounds are malvidin-3-glucoside and malvidin-3-galactoside. By successfully establishing an LPS-induced inflammatory RAW 264.7 model *in vitro*, the study shows that GBBAEs significantly inhibited the release of typical cytokines, the levels of relative inflammatory mRNA and protein expressions. Furthermore, the preliminary results further elucidated the molecular mechanisms governing the anti-inflammatory properties through the NF- κ B pathways, indicating the protection of inflammatory cells. This is in accordance with the function groups of anthocyanins in former studies. We suggest that GBBAEs may present a powerful means for the development of anti-inflammatory nutraceuticals to human benefit.

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Compliance with ethics guidelines

Wei XU, Qing ZHOU, Yong YAO, Xing LI, Jiu-liang ZHANG, Guan-hua SU, and Ai-ping DENG declare that they have no conflict of interest.

This article does not contain any studies with human or animal subjects performed by any of the authors.

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中文概要

题目: 蓝莓花青素对 RAW 264.7 细胞的抗炎活性研究及其机理初探

目的: 以园蓝为研究材料, 鉴定园蓝花青素提取物 (GBBAEs) 中的功能性成分的结构, 建立脂多糖 (LPS) 诱导的体外炎症模型, 并评价其抗炎作用和初步机制。

创新点: 首次探究了蓝莓花青素对建立的 LPS 诱导体外炎症模型的营养干预作用, 并初步探究了发挥抗炎机制的作用通路。

方法: 将 RAW 264.7 细胞分为对照组 (不作处理) 和实验组 (1 $\mu\text{g/ml}$ LPS 刺激建模)。实验组进一步分为 3 个不同浓度组: 400 $\mu\text{g/ml}$ GBBAEs 组、800 $\mu\text{g/ml}$ GBBAEs 组和 1200 $\mu\text{g/ml}$ GBBAEs 组。用酶联免疫吸附测定 (ELISA) 试剂盒检测一氧化氮 (NO)、前列腺素 E₂ (PGE₂)、白细胞介素 1 β (IL-1 β)、白细胞介素 6 (IL-6)、干扰素 γ (INF- γ) 等炎症因子的释放量; 用实时定量聚合酶链反应 (RT-PCR) 分析 IL-1 β 、IL-6、TNF- α 、环氧合酶-2 (COX-2) 及单核细胞趋化蛋白-1 (MCP-1) 的炎症相关基因 mRNA 的表达水平; 用蛋白质印迹法 (Western blot 法) 测定相关炎症蛋白 COX-2 和 NF- κ Bp65 表达水平。

结论: 试验结果表明, 通过 ELISA 法测定 GBBAEs 可以显著性抑制 NO、PGE₂、IL-1 β 、IL-6、INF- γ 等炎症因子的释放; RT-PCR 分析阐明在 LPS 诱导的单核-巨噬细胞 RAW 264.7 中, GBBAEs 可以显著性抑制 IL-1 β 、IL-6、TNF- α 、COX-2 及 MCP-1 的炎症相关基因 mRNA 的表达水平。此外, Western blot 法进一步显示 GBBAEs 对相关炎症蛋白 COX-2 和 NF- κ Bp65 表达具有明显抑制作用, 进一步证实 GBBAEs 通过 NF- κ B 机制通路来发挥抗炎作用。

关键词: 园蓝花青素; 抗炎; 单核-巨噬细胞 RAW 264.7 细胞; 环氧合酶-2 (COX-2); NF- κ Bp65