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Edible blue-green algae reduce the production of proinflammatory cytokines by inhibiting NF-κB pathway in macrophages and splenocytes

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Abstract

Background—Chronic inflammation contributes to the development of pathological disorders including insulin resistance and atherosclerosis. Identification of anti-inflammatory natural products can prevent the inflammatory diseases.

Methods—Anti-inflammatory effects of blue-green algae (BGA), i.e., *Nostoc commune var. Sphaeroides Kützing* (NO) and *Spirulina Platensis* (SP), were compared in RAW 264.7 and mouse bone marrow-derived macrophages (BMM) as well as splenocytes from apolipoprotein E knockout ($apoE^{-/-}$) mice fed BGA.

Results—When macrophages pretreated with 100 µg/ml NO lipid extract (NOE) or SP lipid extract (SPE) were activated by lipopolysaccharide (LPS), expression and secretion of proinflammatory cytokines, such as tumor necrosis factor α (TNF α), interleukin 1 β (IL-1 β), and IL-6, were significantly repressed. NOE and SPE also significantly repressed the expression of TNF α and IL-1 β in BMM. LPS-induced secretion of IL-6 was lower in splenocytes from *apoE*^{-/-} fed an atherogenic diet containing 5% NO or SP for 12 weeks. In RAW 264.7 macrophages, NOE and SPE markedly decreased nuclear translocation of NF- κ B. The degree of repression of proinflammatory gene expression by algal extracts was much stronger than that of SN50, an inhibitor of NF- κ B nuclear translocation. Trichostatin A, a pan histone deacetylase inhibitor, increased basal expression of IL-1 β and attenuated the repression of the gene expression by SPE. SPE significantly down-regulated mRNA abundance of 11 HDAC isoforms, consequently increasing acetylated histone 3 levels.

Conclusion—NOE and SPE repress pro-inflammatory cytokine expression and secretion in macrophages and splenocytes via inhibition of NF- κ B pathway. Histone acetylation state is likely involved in the inhibition.

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General significance—This study underscores natural products can exert anti-inflammatory effects by epigenetic modifications such as histone acetylation.

Keywords

blue-green algae; anti-inflammation; RAW 264.7 macrophages; NF-κB; cytokine array; histone deacetylation

1. INTRODUCTION

Chronic inflammation is an underlying cause of several pathological disorders including insulin resistance, dementia, rheumatoid arthritis, atherosclerosis, and cancer [1, 2]. Non-steroidal anti-inflammatory drugs (NSAID) are generally used to treat acute and chronic inflammatory conditions. However, due to their adverse side-effects and increased cardiovascular disease (CVD) risk associated with chronic use of several NSAID [3], there is a critical need to identify natural products with anti-inflammatory properties. Several anti-inflammatory bioactive compounds that have been extensively studied thus far include curcumin, resveratrol, anthocyanin, and green tea polyphenols [4–8].

Nuclear factor κ B (NF- κ B) is a major transcriptional factors responsible for proinflammatory cytokine production, such as tumor necrosis factor α (TNF α), interleukins (ILs), inducible nitric oxide synthase and cyclooxygenase-2, under inflammatory conditions [9, 10]. NF- κ B exists as homo- or heterodimers consisting of five subunits of Rel family, i.e., p50, p52, p65, c-Rel and RelB [11]. In an unstimulated state, NF- κ B is present in the cytoplasm bound with inhibitor of κ B α (I κ B α), which masks the nuclear localization sequence of p65 [12] and NF- κ B activation largely depends on I κ B kinase β (IKK β). Extracellular inflammatory signals including lipopolysaccharide (LPS), pro-inflammatory cytokines, reactive oxygen species, advanced glycation end products, and oxidized lowdensity lipoprotein trigger the phosphorylation of IKK β , which in turn phosphorylates I κ B α for ubiquitination and degradation by proteasome [13–18]. This event liberates NF- κ B to enter the nucleus for the induction of pro-inflammatory gene expression [19–24].

Blue-green algae (BGA), also known as cyanobacteria, are one of the most primitive forms of photosynthetic prokaryotes. They have been consumed as food or medicine for centuries and human consumption of BGA was recorded in the 14th century during the Aztec civilization [25]. They are recognized for their protective effects against viral and bacterial infections, cancer, allergy, diabetes, inflammation and hyperlipidemia [26–29]. At present, *Spirulina platensis* (SP) is the most commonly consumed and commercialized BGA species. We have previously reported that lipid extract of *Nostoc commune var. sphaeroides Kützing* (NO), another BGA species, inhibited NF- κ B DNA binding activity and consequently repressed the pro-inflammatory gene expression in RAW 264.7 macrophages [30]. In this study, we compared anti-inflammatory effects of two BGA species using several model systems: RAW 264.7 macrophages, bone marrow-derived macrophages (BMM), and splenocytes isolated from BGA-fed apolipoprotein E knockout (*apoE*^{-/-}) mice. We found that in all of three systems, NO and SP either as lipid extract or as whole algae repressed pro-inflammatory gene expression and production.

2. METHODS AND MATERIALS

2.1. Preparation of BGA lipid extraction

BGA powder was kindly provided by Algaen Corporation (Winston Salam, NC) for NO and by Earthrise Nutritionals (Irvine, CA) for SP. Lipid extracts of BGA into chloroform/ methanol (1:2) were prepared as previously described [30, 31]. Lipid extracts were stored

under N_2 gas at -20° C for short term or at -80° C for long term. The lipid extracts were dried down under N_2 to remove solvents and then dissolved in cell medium by sonication.

2.2. Bone marrow isolation and macrophage differentiation

BMM were isolated from C57BL/6J mice (Jackson Laboratory, Bar harbor, ME). Briefly, mouse legs were removed from the hip joint and cleaned, after which the femur and tibia were cut at the tip and subsequently bone marrow was collected by centrifugation. The bone marrow cells were differentiated into macrophages in low-glucose DMEM containing 2 mmol/L L-glutamine, 1 mmol/L sodium pyruvate, 30 U/mL penicillin, 300 µg/mL streptomycin, 20% FBS, and 30% L929 cell conditioned media. L929 cells were generously provided by Dr. John Parks (Wake Forest University School of Medicine, Winston-Salem, NC). BMM were incubated in a humidified chamber at 37°C with 5% CO₂ and cell culture medium was changed every 3 days for 7 days until they became confluent.

2.3. Macrophage cell culture and treatment

RAW 264.7 macrophages (ATCC, Manasas, VA) and BMM were incubated in RPMI-1640 containing 10% FBS, 100 U/mL penicillin, 100 μ g/mL streptomycin, 1 x vitamins and 2 mmol/L L-glutamine in a 37°C humidified chamber with a supply of 5% CO₂. Macrophages were incubated with 0–100 μ g/mL of NO lipid extract (NOE) or SP lipid extract (SPE) for 12 h and subsequently activated by LPS (Sigma-Aldrich, St. Louis, MO) at 100 ng/ml concentration for additional 18 h. Cells and medium were collected for pro-inflammatory cytokine expression and secretion.

SN50, a cell permeable inhibitor specific for NF- κ B nuclear translocation, was purchased from Enzo Life Science (Plymouth meeting, PA) and dissolved in sterile water. RAW 264.7 macrophages were incubated with 50 µg/mL of NOE or SPE for 11 h followed by 1 h incubation with 50 µg/mL of SN50. Subsequently, the cells were treated with LPS (100 ng/mL) in the presence of algal lipid extract and SN50 for additional 3 h. For the experiment with trichostatin A (TSA), a pan histone deacetylase (HDAC) inhibitor, RAW 264.7 macrophages were treated with algal lipid extract (100 µg/ml) and 50 nmol/L of TSA for 12 h prior to LPS stimulation for 7 h. All cell culture supplies were purchased from Thermo Scientific Hyclone (Logan, UT), unless stated otherwise.

For all cell culture experiments, cells without exposure to any algal extract were considered as control. Algal extracts were incorporated into cell culture medium by sonication and therefore no solvent was used as a vehicle.

2.4. Splenocyte isolation and culture

Spleens were harvested from male $apoE^{-/-}$ mice (Jackson Laboratory) fed a high fat/high cholesterol (15% fat, 0.2% cholesterol by wt) containing 5% NO or SP by wt for 12 wk from 8 wk of age. After being anesthetized with ketamine HCl (50 mg/kg)/xylazine (10 mg/kg) and subsequently euthanized by cardiac puncture and cervical dislocation, spleen of each mouse was excised aseptically and ground in RPMI-1640 medium containing 10% FBS, 100 U/mL penicillin and 100 µg/ml streptomycin. After removal of red blood cells by a pre-warmed RBC lysis buffer (eBioScience, San Diego, CA), the cells were resuspended in the medium and centrifuged at 2,000 rpm for 5 min. Cell pellet was resuspended in PBS and filtered through 40 µm strainer (BD Biosciences, San Jose, CA). After washing with PBS, cells were resuspended in RPMI-1640 complete media and plated at a density of 1 x 10⁶/0.5 mL for experiments. Cells were incubated with LPS (500 ng/mL) for 20 h and IL-6 concentrations in the media were measure by ELISA (eBioScience) according to the manufacturer's instruction. All procedures were approved by the Institutional Animal Care and Use Committee of the University of Connecticut.

2.5. Quantitative realtime PCR (qRT-PCR)

Total RNA was extracted using TRIzol reagent (Life technology, Carlsbad, CA) and cDNA synthesis was performed as previously described [30, 31]. qRT-PCR was conducted using Bio-Rad CFX96 Real-Time system (Bio-Rad, Hercules CA). Primers were designed according to GenBank database and the sequences were previously published [30].

2.6. Cytokine array

RAW 264.7 macrophages were pretreated with 100 μ g/mL of NOE or SPE for 12 h, after which they were incubated with LPS for 18 h. Conditioned medium was collected and centrifuged at 12,000 x g for 5 min to remove any cell debris or dead cells. Secretion of cytokines was assessed by RayBio Mouse Cytokine Antibody Array (RayBiotech, Norcross, GA) according to the manufacturer's protocol. Signals were captured using ChemiDoc XRS + system (Bio-Rad, Hercules, CA).

2.7. NF-KB translocation by Western blot analysis

RAW 264.7 macrophages incubated with 100 µg/ml of NOE or SPE for 12 h were activated by LPS for 1 or 2 h. Nuclear and cytoplasmic fractions of the cells were separated by using Cayman nuclear extraction kit (Ann Arbor, MI) and Western blot analysis was conduct as we previously described [30, 31]. Polyclonal anti-p65 and anti-GAPDH antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Mouse monoclonal antibody against TATA binding protein was obtained from Abcam (Cambridge, MA) and used as a loading control for nuclear fraction. Protein expression were detected using Westpico horseradish peroxidase chemiluminescence (Pierce, Rockford, IL) and imaged using a Chemidoc XRS+ system (Bio-Rad) and Image Lab software (Bio-Rad).

2.8. Acetylation of histone 3 (H3)

After incubation with 100 μ g/mL of SPE for 12 h and LPS for additional 18 h, RAW 264.7 macrophages were washed twice with ice-cold PBS and cells were collected by centrifugation at 12,000 x g for 5 min. Cells were then resuspended in Triton X extraction buffer containing 0.5% Trition X-100 (v/v), 0.02% NaN₃ (w/v) in PBS and 1 X protease inhibitor cocktail (Calbiochem, San Diego, CA) for 10 min on ice. Subsequently, the cell lysates were centrifuged for 10 min at 4°C to pellet nuclei and the pellets were then acid extracted in 0.2 N HCl overnight at 4 C. After centrifugation at 12,000 x g for 10 min, the supernatants were collected for Western blot analysis as described above. Antibodies against total H3 and acetyl-H3K9 were purchased from Cell Signaling Technology (Danvers, MA).

2.9. Statistical analysis

One-way analysis of variance (ANOVA) and Newman-Keuls pairwise comparisons to detect significant difference between groups were performed using GraphPad InStat 5 (GraphPad Software, La Jolla, CA). P values less than 0.05 were considered significant. Values were expressed as mean \pm SEM.

3. RESULTS

3.1. Pro-inflammatory cytokine expression and secretion were repressed by NOE and SPE in macrophages

We examined the effects of NOE and SPE on pro-inflammatory cytokine expression. In RAW 264.7 macrophages, both algal lipid extracts significantly repressed LPS-induced TNFa, IL-1 β and IL-6 mRNA expression compared with control and their repressive effects showed a similar potency (Figure 1). Significant reductions in the gene expression were observed in a dose-dependent manner as low as 25 µg/mL (data not shown).

Furthermore, NOE and SPE markedly decreased the secretion of TNFa, IL-6, granulocyte colony-stimulating factor, granulocyte macrophage colony-stimulating factor, and monocyte chemoattractant protein 1 (Figure 2).

Consistent with our observations in RAW 264.7 macrophages, BMM treated with NOE and SPE showed significantly lower expression of TNFa and IL-1 β compared with control BMM (Figure 3).

3.2. IL-6 secretion was decreased from splenocytes of mice fed NO and SP supplements

Fresh splenocytes were isolated from $apoE^{-/-}$ mice fed an atherogenic diet containing 5% of NO or SP for 12 wk. When the splenocytes were challenged by LPS for 20 h, cells from NO or SP-fed mice showed significantly lower IL-6 levels in cell medium than control mice (Figure 4).

3.3. Algal lipid extracts inhibited NF-KB nuclear translocation in macrophages

Translocation of NF- κ B p50/p65 dimer to the nucleus is one of the major steps for its activation to increase transcription of pro-inflammatory cytokines. We previously reported that NOE significantly decreased NF- κ B p65 DNA binding activity [30]. To address if the decrease in DNA binding activity is, at least in part, due to the inhibition of p65 nuclear translocation and to compare a potency of the repression between NOE and SPE, RAW 264.7 macrophages pre-treated with algal lipid extracts were stimulated by LPS for 1 or 2 h. At 1 h LPS stimulation, control cells showed markedly increased p65 in the nucleus with p65 being non-detectible in the cytoplasmic fraction (Figure 5). In contrast, macrophages pretreated with NOE and SPE had less p65 translocation to the nucleus and detectible amount of p65 was present in the cytoplasm. The similar trend was shown at 2 h LSP stimulation.

3.4. Inhibition of NF-κB nuclear translocation could not solely account for the repression of pro-inflammatory cytokine expression by NOE and SPE in macrophages

Although translocation of NF- κ B to the nucleus is a major regulatory step for its transcriptional activity, other posttranslational modifications in NF- κ B have been recently recognized as important regulatory mechanisms for NF- κ B activity [32, 33]. To evaluate contribution of nuclear translocation of p65 to the repressive effect of NOE and SPE on TNF α and IL-1 β expression, we used SN50, an inhibitor specific for NF- κ B translocation to the nucleus. SN50 binds to the nuclear localization sequence (NLS) of p65 and therefore p65 cannot be recognized by karyopherin α , resulting in cytoplasmic sequestration. With the treatment of SN50, LPS-induced nuclear translocation of p65 was blocked (data not shown). Although SN50 significantly lowered mRNA levels of TNF α and IL-1 β compared with control, repression of the gene expression was markedly stronger in NOE or SPE-treated RAW 264.7 macrophages than SN50-treated cells (Figure 6A). Interestingly, TSA significantly increased basal expression was attenuated by TSA (Figure 6B). NOE also elicited the similar results (data not shown).

3.5. Down regulation of HDAC expression and subsequent increases in acetylated histone 3 by SPE in macrophages

Inhibition of HDAC by TSA attenuated the repression of pro-inflammatory gene expression by SPE in LPS-stimulated macrophages, suggesting HDAC may play a role in the antiinflammatory function of the algal extract. Therefore, we next evaluated if SPE can alter the expression of 11 HDAC isoforms in RAW 264.7 macrophages. mRNA abundance of all HDAC isoforms was significantly decreased by SPE in both unstimulated and LPS-

stimulated macrophages (Figure 7A). HDAC expression was also significantly lowered by LPS. Consistent with significant decreases in HDAC mRNA levels by SPE in both unstimulated and LPS-stimulated macrophages, SPE increased the level of acetylated H3 by ~2-fold (Figure 7B). Interestingly, although LPS repressed HDAC mRNA levels to a similar extent to SPE, there was only ~40% increase in acetylated H3 in LPS-stimulated cells compared with controls.

4. DISCUSSION

Chronic low-grade inflammation is causally linked to the pathogenesis of several metabolic diseases, such as insulin resistance, metabolic syndrome, type 2 diabetes, CVD and non-alcoholic fatty liver disease. NSAID are commonly used to treat acute and chronic inflammatory conditions, but not without their adverse effects [3]. Therefore, identification of anti-inflammatory natural products with minimal side-effects, yet effective in blocking pathways leading to chronic inflammation is critically needed. We previously reported that NOE exerts ant-inflammatory effects by inhibiting NF- κ B DNA binding activity in macrophages [30]. In the present study, we further demonstrated that SPE inhibits pro-inflammatory mediator expression and secretion to a similar extent to NOE in macrophage cell line as well as primary mouse macrophages. Furthermore, the anti-inflammatory was confirmed in splenocytes isolated from NO or SP-fed *apoE^{-/-}* mice, demonstrating the protection against inflammation by the two BGA species is likely to occur in vivo.

Humans have a long history of BGA consumption as food or medicine for hundreds of years, yet scientific understanding of their health-promoting properties only began ~ 30 years ago. Currently, SP is the most commonly consumed and commercialized BGA species [25]. NO is another edible BGA species with several purported health benefits. However, there has been a concern on safety of BGA for human consumption. Certain BGA species produce toxins, such as microcystins, nodularins, saxitoxins, anatoxins, and β -methylamino-L-alanine and therefore naturally harvested BGA products may contain the toxins due to contamination with toxin-producing algae species during harvest [34-37]. The two BGA species used in the present study are cultivated in the controlled environment to minimize any toxin contamination. We previously demonstrated that SP and NO are free of microcystins, the major algal toxin, and long-term supplementation of the two BGA did not show any adverse side-effects in mice [37]. Furthermore, in 2011, The Dietary Supplements Information Expert Committee assigned a Class A safety rating for SP and permitted the admission of quality monographs for SP in the United States Pharmacopeia and National Formulary [38]. As safety is prerequisite to investigation on health-promoting properties of any natural products, we believe the two BGA species may be safe natural products for human consumption to gain health benefits if their functions are scientifically proven.

Studies have demonstrated that BGA consumption promotes immunity and protects against inflammatory diseases such as colitis, arthritis, and allergic rhinitis in animals and humans [39–42]. In particular, Rasool et al. [42] showed that *Spirulina fusiformis* supplementation minimized inflammatory response to near normal condition in mice injected with inflammatory inducer. We previously showed that NOE repressed the expression and secretion of pro-inflammatory cytokine such as TNFa and IL-1 β in RAW 264.7 macrophages [30]. Consistent with our previous findings, NOE significantly decreased mRNA levels of the pro-inflammatory cytokines not only in RAW 264.7 macrophages but also in BMM. SPE also elicited the same anti-inflammatory effect with similar potency to NOE. Phycocyanin (PC) and γ -linoleic acid are the most well-known anti-inflammatory components in BGA. In particular, C-PC, a natural blue pigment (~14% of SP dry weight) [40, 43, 44], has shown to have a strong inhibitory property against inflammation in macrophages [45, 46] and in animal models [47, 48]. We found that in NOE, unsaturated

fatty acids, which account for ~15% of the extract mass, are partly responsible for the repression of pro-inflammatory cytokine production but other compounds are required to achieve full degree of repression in RAW 264.7 macrophages [30]. Studies on identification of anti-inflammatory compounds in NOE and SPE are currently underway. As both algal extracts elicits marked repression of inflammation, the identified compounds may be used as anti-inflammatory drugs alternative to NSAID.

In addition to the inhibitory role of NOE and SPE in the production of pro-inflammatory mediators shown in vitro using RAW 264.7 macrophages and BMM, we sought to evaluate if the BGA supplementation can accumulate enough amount of putative anti-inflammatory compounds in other immune-modulating cells in mice to inhibit inflammation. It has been shown that spleen is a monocyte reservoir and plays an important role not only in monocyte production but also in monocyte/macrophage clearance during inflammation [49, 50]. Therefore, we isolated splenocytes from $apoE^{-/-}$ mice, a common mouse model for atherosclerosis, that were fed an atherogenic diet supplemented with 5% of NO or SP for 12 wk. When the splenocytes from mice fed a NO or SP-containing diet were stimulated by LPS, IL-6 secretion was significantly lower than cells from control mice. This result suggests that NO and SP supplementation can be beneficial to inhibit inflammation in vivo.

Mounting evidence suggests that activation of NF- κ B pathway is a major underlying event for the development of inflammatory diseases and its inhibition can decrease the disease progression [51–53]. Epidemiological and clinical studies have demonstrated the effect of polyphenol-rich foods, such as green tea and extra virgin olive oils, on the inhibition of NF- κ B activation [54]. We previously demonstrated that NOE repressed pro-inflammatory cytokine production, at least in part, by inhibiting NF- κ B p65 DNA binding activity in RAW 264.7 macrophages [30]. In the present study, both NOE and SPE inhibited nuclear translocation of NF- κ B p65 in LPS-stimulated macrophages. However, of interest is that blockage of p65 translocation to the nucleus by a chemical inhibitor was significantly less effective in repressing TNF α and IL-1 β expression than NOE and SPE. This suggests that mechanisms other than NF- κ B nuclear translocation may exist for the algal extract to elicit the drastic reduction in the expression of pro-inflammatory cytokine expression.

Although NF- κ B nuclear translocation is the best-known mechanism for its activation [12], several lines of evidence indicate that posttranslational modifications in p65 subunit, such as phosphorylation and acetylation, impact its activity by altering DNA binding, interaction with inhibitor of $\kappa B \alpha$ (I $\kappa B \alpha$) in the nucleus, and association with chromatin structure modifiers such as histone acetyltransferases (HAT) and HDAC [32, 33]. It is our particular interest that acetylation of p65 at Lys218, 221 and 310 by p300 and cAMP response element binding (CREB) binding protein (CBP) regulates its transcriptional activity by increasing p65 DNA binding and by inhibiting its association with nuclear IkBa for the prevention of NF- κ B relocation to the cytoplasm [55]. In contrast, deacetylation at Lys310 by corepressor complexes inhibits p65 activity [56, 57]. In the present study, we observed that basal IL-1 β and TNFa mRNA levels were significantly induced by TSA, a pan HDAC inhibitor, in unstimulated macrophages. Bailey and Ghosh [58] suggested that NF- κ B in association with a corepressor complex actively represses pro-inflammatory gene expression in unstimulated macrophages and the corepressor complex is composed of nuclear receptor corepressor, HDAC3 and others. Therefore, the induction of IL-1 β and TNFa expression by TSA in the present study is well in line with their findings that active transcriptional repression of the pro-inflammatory genes occurs when macrophages are not stimulated by inflammatory insults. Furthermore, we observed that repression of the pro-inflammatory gene expression by NOE and SPE was significantly ameliorated by TSA in LPS-activated RAW 264.7 macrophages. The data suggest that HDAC may be involved in basal transrepression of IL-1 β and TNFa when there are no extracellular inflammatory insults, while it mediates the

repressive effect of the algal extracts on the expression of the pro-inflammatory genes in the activated macrophages. We found that the mRNA expression of 11 HDAC isoforms was significantly repressed by SPE both in unstimulated and activated macrophages. Among 11 HDAC isoforms, HDAC 1, 2, 3, 4, 5 and 9 are major isoforms in RAW 264.7 macrophages (data not shown). High levels of acetyl H3 observed in the SPE-treated cells are likely due to the down-regulation of HDAC expression although we cannot rule out possible alterations in HAT by SPE. LPS also repressed HDAC mRNA levels to a similar extent to SPE. Although a degree of repression of HDAC mRNA by SPE and LPS was similar, acetyl H3 levels were markedly higher in SPE-treated cells than LPS-activated cells. It can be speculated that LPS and SPE may differentially regulate HDAC expression at the posttranscriptional levels and/ or HAT activity may also be altered by the treatments to alter H3 acetylation. Conundrum is that both unstimulated cells and SPE treatment in LPS-treated cells showed low proinflammatory gene expression but their acetyl H3 levels were very different. It seems important to understand the effect of SPE and LPS on the expression/activity of each HDAC isoform rather than overall changes in H3 acetylation state for gaining mechanistic insights into critical roles of HDAC in inflammation. Future study is warranted to identify specific HDAC isoforms that are involved in the regulation of the inflammatory gene expression and to determine how their activity is regulated such as cellular localization, posttranslational modification and association with other regulatory proteins. The information can be used to develop effective therapeutic targets to inhibit inflammation.

In conclusion, we found that NO and SP either as a whole or a lipid extract have an antiinflammatory effect using several model systems, i.e., RAW 264.7 macrophages, BMM and mouse splenocytes. Repression of pro-inflammatory mediator expression by the BGA is mediated partly by inhibiting NF- κ B nuclear translocation. In addition, our results also support that HDAC and modulations in chromatin structure are likely involved in the antiinflammatory function of NO and SP. Dysregulation of pro-inflammatory mediator production from macrophages is a major contributor to the development of pathological conditions related to inflammation such as insulin resistance and atherosclerosis. Therefore, the two BGA species tested in the present study may be used as a natural product to prevent inflammatory diseases.

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Abbreviations

apoE ^{-/-}	apolipoprotein E knock out
BGA	blue-green algae
BMM	bone marrow-derived macrophages
CBP	CREB-binding protein
CREB	cAMP-response element-binding protein
C-PC	C-phycocyanin
CVD	cardiovascular disease
HAT	histone acetyltransferase
HDAC	histone deacetylases
IrBa	inhibitor of kappa B a

IL	interleukins
LPS	lipopolysaccharide
NLS	nuclear localization sequence
NO	Nostoc commune var. sphaeroides Kützing
NOE	NO lipid extract
SP	Spirulina platensis
SPE	SP lipid extract
TNFa	tumor necrosis factor a
TSA	trichostatin A

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Highlights

- Anti-inflammatory effects of edible blue-green algae in macrophages
- Inhibition of NF-xB pathway for the anti-inflammatory effects in macrophages
- Repression of IL-6 secretion from splenocytes of mice fed blue-green algae
- Role of histone deacetylases in the anti-inflammatory role of blue-green algae



Figure 1. Repression of pro-inflammatory expression by NOE and SPE in RAW 264.7 macrophages

Cells were pre-treated with 100 μ g/mL of NOE or SPE for 12 h followed by co-incubation of 100 ng/mL LPS and algal extracts for additional 18 h. qRT-PCR analysis was conducted to measure mRNA levels of TNFa, IL-1 β and IL-6. Values are expressed as mean \pm SEM, P < 0.05, n = 15–18.



Figure 2. Inhibition of pro-inflammatory cytokine secretion by NOE and SPE in RAW 264.7 macrophages

Cells were incubated with 50 μ g/mL of NOE or SPE for 12 h and subsequently activated by 100 ng/mL LPS for 18 h. Bottom chart indicates location of cytokines in the array. GCSF, granulocyte colony-stimulating factor; GM-CSF, granulocyte macrophage colony-stimulating factor; IL, interleukins; IFN γ , interferon gamma; MCP-1, monocyte chemoattractant protein 1; RANTES (CCL5), chemokine C-C motive ligand 5; SCF, stem cell factor; sTNFR1, soluble tumor necrosis factor receptor 1; TNF α , tumor necrosis factor alpha; TBP, TNF binding protein; VEGF, vascular endothelial growth factor.



Figure 3. Inhibitory effects of TNFa and IL-1 β expression by NOE and SPE in BMM Bone marrow derived macrophages isolated from C57BL/6J mice were treated with 100 µg/mL of NOE or SPE for 12 h, after which they were co-incubated with 100 ng/mL for 18 h. Values are expressed as mean \pm SEM, P < 0.05, n = 3.







Figure 5. Attenuation of nuclear translocation of NF- κB p65 by NOE and SPE in RAW 264.7 macrophage

Cells were pre-treated with 50 μ g/mL of NOE or SPE for 12 h and subsequently they were activated by 100 ng/mL LPS for 1 or 2 h. Cytoplasmic and nuclear fractions were isolated and Western blot analysis for p65 was conducted. TATA binding protein (TBP) and GAPDH were used as a nuclear and cytoplasmic marker, respectively.



Figure 6. Alternative mechanisms for anti-inflammatory effects of NOE and SPE in RAW 264.7 macrophages

A. Effect of SN50 on pro-inflammatory gene expression. Cells were incubated with NOE or SPE (50 μg/mL) for 12 h. Subsequently, they were incubated with 50 μg/mL of SN50, a NF- κ B translocation inhibitor, for 1 h, after which they were activated with 100 ng/mL LPS for 3 h. qRT-PCR analysis was conducted to measure mRNA levels of TNFa and IL-1 β . Value are expressed as mean ± SEM, P < 0.05, n = 3. **B**. RAW 264.7 macrophages were treated with NOE or SPE (100 µg/ml) and 50 nmol/L of TSA for 12 h and then they were activated by LPS (100 ng/mL) for 7 h. qRT-PCR analysis was conducted to measure mRNA levels of TNFa and IL-1 β . Value are expressed as mean ± SEM, P < 0.05, n = 3. **B**. RAW 264.7 macrophages were activated by LPS (100 ng/mL) for 7 h. qRT-PCR analysis was conducted to measure mRNA levels of TNFa and IL-1 β . Value are expressed as mean ± SEM, P < 0.05, n = 3.



Figure 7. Regulation of HDAC expression and levels of acetyl H3 by SPE in RAW 264.7 macrophages

A. RAW 264.7 macrophages were pre-treated with 100 μ g/mL of SPE for 12 h and subsequently with 100 ng/mL LPS and algal extracts for additional 18 h. qRT-PCR analysis was conducted to detect HDAC isoform mRNA abundance. Values are expressed as mean \pm SEM, P < 0.05, n = 6. **B**. Cells were treated with SPE and LPS as described above and Western blot analysis was conducted to detect total and acetyl H3 levels. One representative blot of 2 separate experiments is shown. Densitometry analysis was conducted for the ratios of acetyl H3K9 to total H3. A ratio of each band and an averaged value of each treatment are listed in the table.