3-Hydroxy-4,7-megastigmadien-9-one, isolated from Ulva pertusa, attenuates TLR9-mediated inflammatory response by down-regulating mitogen-activated protein kinase and NF-κB pathways

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\textbf{ABSTRACT}

\textbf{Context:} Seaweeds are rich in bioactive compounds in the form of vitamins, phycobilins, polyphenols, carotenoids, phycocyanins and polysaccharides; many of these are known to have advantageous applications in human health. 3-Hydroxy-4,7-megastigmadien-9-one (comp) was isolated from Ulva pertusa (U. pertusa) Kjellman (Ulvaeeae), which is a familiar edible green seaweed.

\textbf{Objective:} This study evaluates the anti-inflammatory activity of comp in CpG DNA-stimulated bone marrow-derived dendritic cells (BMDCs).

\textbf{Materials and methods:} For evaluating the effect of comp on cytokines production, BMDCs were treated with doses of comp (0, 0.5, 1, 2, 5, 10, 25 and 50 \textmu M) for 1 h before stimulation with CpG DNA (1 \textmu M). Cytokine production was measured by ELISA. Western blotting was conducted for evaluating effect of comp (50 \textmu M) on MAPKs and NF-κB pathways. Luciferase reporter gene assay was conducted for effect of comp (0, 5, 10 and 25 \textmu M) on transcriptional activity of AP-1 and NF-κB.

\textbf{Results:} Comp exhibited strong inhibition of interleukin (IL)-12 p40, IL-6 and TNF-α cytokine production with IC\textsubscript{50} values of 6.02 \pm 0.35, 27.14 \pm 0.73, and 7.56 \pm 0.21 \textmu M, respectively. It blocked MAPKs and NF-κB pathways by inhibiting the phosphorylation of ERK1/2, JNK1/2, p38 and IκB\textalpha. In addition, it strongly inhibited the transcriptional activity of AP-1 and NF-κB with IC\textsubscript{50} values of 8.74 \pm 0.31 and 12.08 \pm 0.24 \textmu M, respectively.

\textbf{Discussion and conclusion:} Taken together, these data suggest that comp has a significant anti-inflammatory property and warrants further studies concerning the potential of comp for medicinal use.

\textbf{Introduction}

The mammalian immune system consists of two types, innate and adaptive immunity. The innate immunity is the first line of defence in combating invading pathogens and is mediated by phagocytes including macrophages and dendritic cells (DCs). Acquired immunity eliminates pathogens in the late phase of infection and generates immunological memory (Akira et al. 2006). Chronic inflammation results from continuous exaggerated immune reaction to injury or exposure to foreign pathogens (Uronis et al. 2009; Katoh et al. 2013). Microbial pathogen recognition by innate immune system is important for activation of microbialidal effectors and progression of adaptive immunity (Honstetter et al. 2004). Toll-like receptors (TLRs) consist of a family of pattern recognition receptor that sense conserved molecular pattern of microbes. They are crucial for recognition of microbial infection and initiating inflammatory and immune responses (Rakoff-Nahoum et al. 2004; Yuk & Jo 2011). TLR9 recognizes unmethylated DNA having CpG island derived from viruses and bacteria (Takeuchi & Akira 2010). Stimulation of immune cell with microbial product causes the production of pro-inflammatory cytokines, such as TNF-α and various interleukins and is maintained for the duration of inflammatory response (Ma et al. 2013; Shi et al. 2015).

TLR stimulation causes the phosphorylation and activation of extracellular signal regulated kinase (ERK1/2), c-Jun NH2-terminal kinase (JNK) and p38 mitogen-activated protein kinase (MAPKs) signalling pathways (Intayoung et al. 2016). Activator protein (AP-1), a heterodimeric transcription factor composed of c-Fos, c-Jun, ATF and JDP families, is activated by MAPK signalling pathway. AP-1 upregulate the transcription of genes encoding pro-inflammatory mediators (Hossen et al. 2015). NF-κB pathway is one of the well-established inflammatory pathways (Greten et al. 2004; Zaki et al. 2011). Its activation requires phosphorylation and subsequent degradation of IκB, allowing the nuclear translocation of NF-κB, where it causes the transcription and release of pro-inflammatory cytokines (Saba et al. 2015).

In search for new drugs from natural products, marine organisms, including seaweeds and marine microorganisms have been the focus of numerous studies (Khan et al. 2008). Ulva pertusa (U. pertusa) Kjellman (Ulvalveae), an edible green seaweed, is consumed by local inhabitants of Asia (Shi et al. 2013). It has...
been authorized for human utilization by French authorities because of nutritional interests, i.e., rich in dietary fibres, vitamins, oligoelement and minerals (Pengzhan et al. 2003). Previous studies found that ulvan extracted from *U. pertusa* possessed antihyperlipidemic and antioxidant activities (Qi et al. 2012). Sulphated polysaccharides from *U. pertusa* have immune modulatory and anti-avian influenza virus activities (Song et al. 2016). 3-Hydroxy-4,7-megastigmadien-9-one (comp), a norisoprenoid, is the degradation product of carotenoids (Machida & Kikauchit 1996). As part of on-going research on evaluating biological effects, in this study, comp was isolated from *U. pertusa*, and was examined for anti-inflammatory activities for the first time. The detailed mechanisms of anti-inflammatory potential of comp have not been reported yet. Here, we investigated the anti-inflammatory potential of comp, especially on TLR9-mediated inflammatory response in bone marrow derived dendritic cell.

**Materials and methods**

**Preparation of 3-hydroxy-4,7-megastigmadien-9-one**

*U. pertusa* was collected on Jeju Island, South Korea. A voucher specimen (JBR10253) has been deposited in herbarium of the Jeju Biodiversity Research Institute. Dried *U. pertusa* was subjected to solvent extraction in 70% aqueous ethanol. The resultant solvent suspension was extracted with *n*-hexane, ethyl acetate (EtOAc), *n*-butanol and water successively that gave the respective fraction extract. A portion of EtOAc-soluble fraction was subjected to medium pressure liquid chromatography (MPLC) by reserved-phase silica gel through water–methanol gradients to give 45 fractions. Fraction 18 (47.9 mg), by silica gel with *n*-Hex/EtOAc = 1/1 to give 3-hydroxy-4,7-megastigmadien-9-one (comp). Structure of comp was identified by spectroscopic studies and by comparing the obtained data with literature values (Machida & Kikauchit 1996).

**Cell cultures and measurement of cytokine production**

To grow BMDCs, wild-type 6-week-old female C57BL/6 mice were used as previously described (Chae et al. 2013; Manzoor et al. 2014a). All animal procedures were approved by and performed according to the guidelines of the Institutional Animal Care and Use Committee of Jeju National University, Jeju, South Korea (#2010-0028). Briefly, bone marrow cells were differentiated in RPMI 1640 (BD, Grand Island, NY) medium containing granulocyte-macrophage colony-stimulating factor for DCs generation. For BMDCs, on day 6 of incubation the cells were harvested and seeded in 48-well plates at a density of 1 × 10^5 cells/0.5 mL, and then treated with the comp for 1 h before stimulation with CpG DNA. Supernatants were harvested 18 h after stimulation. Concentrations of murine IL-12 p40, IL-6 and TNF-α in the culture supernatants were measured by enzyme linked immunosorbent assay (ELISA) (BD PharMingen, San Jose, CA).

**Cell viability assay**

The cell viability was measured by standard procedure of 3-(4,5-dimethyl-2,5 thiadiazol)-2,5 diphenyltetrazolium bromide (MTT) assay (Chae et al. 2013).

**Western blot analysis**

Bone marrow-derived dendritic cells (BMDCs) were dispensed to 35 mm culture dishes at a concentration of 2 × 10^6 cells/2 mL and incubated for 1 h at 37 °C. The cells were pretreated with or without comp for 1 h before treatment with CpG DNA at the indicated time points. The cells were collected and, then, lysed in lysis buffer (PRO-PREP lysis buffer, iNtRON Biotechnology, South Korea). A protein sample (30 μg) was subjected to electrophoresis in 10% SDS-polyacrylamide gels and transferred to a polyvinylidene fluoride membrane (Bio-Rad, Hercules, CA). The membrane was incubated with 1/1000-diluted rabbit polyclonal antibodies that specifically recognize phospho-ERK1/2, phospho-p38, phospho-JNK1/2, phosphor-IкBα (Cell Signaling Technology, Danvers, MA), and β-actin (Santa Cruz Biotechnology, Santa Cruz, CA). After washing, the membrane was incubated with a horseradish peroxidase-linked goat anti-rabbit IgG (Cell Signaling Technology), and immunoactive bands were detected as previously described (Koo et al. 2012).

**Luciferase assay**

For AP-1 and NF-κB reporter assays, HEK293T cells were plated in 24-well plates and grown overnight as described previously (Manzoor et al. 2013a, 2013b). Cells were transfected using Fugene 6 (Roche, Indianapolis, IN) with a AP-1 or NF-κB reporter gene, pRLnull (Promega, Madison, WI) and pcDNA3 (empty vector) or TLR9-encoding pcDNA3 kindly provided by Prof. R. Medzhitov (Yale University, New Haven, CT). After incubation of 24 h, cells were pretreated with comp for 1 h before stimulation with CpG (1 μM). After further incubation for 18 h, cells were lysed in a passive lysis buffer (Promega), and firefly luciferase vs. Renilla activities were measured using a Dual Luciferase Reporter assay system (Promega, Madison, WI).

**Statistical analysis**

All experiments were performed at least three times, and the data are presented as the mean ± standard deviation (SD) of three independent experiments. One-way ANOVA was used for comparison between the treated and the control groups. *p* < 0.05 was considered statistically significant.

**Results**

**Effect of comp on the cell viability of BMDCs**

The chemical structure of comp isolated from *U. pertusa* is shown in Figure 1. To assess the possible cytotoxicity of comp on BMDCs, cell viability was determined by using MTT assay. The results demonstrate that cell viability was not affected by comp at the indicated doses (Figure 2).

**Inhibitory effect of comp on pro-inflammatory cytokine production in CpG DNA-stimulated dendritic cells**

Mature and activated DCs produce high amounts of pro-inflammatory cytokines, such as IL-12, TNF-α and IL-6 (Wei et al. 2012). The anti-inflammatory potential of comp on the TLR9-activated cytokine production was evaluated by using a Luciferase Model Kit (Promega, Madison, WI). Comp was isolated from *U. pertusa*. The chemical structure of comp isolated from *U. pertusa* is shown in Figure 1. To assess the possible inhibitory effects of comp on TLR9-activated cytokine production, the cells were pretreated with comp for 1 h before stimulation with CpG DNA (1 μM) at the indicated time points. The cytokine production was determined by using the Dual Luciferase Reporter assay system. The results demonstrated that comp inhibited the production of IL-12, TNF-α and IL-6 in a dose-dependent manner (Figure 2).
To explore the anti-inflammatory activity, comp was evaluated for inhibition of IL-12 p40, IL-6 and TNF-α production in CpG DNA-stimulated bone marrow derived DCs. DCs treated with comp alone showed no production of pro-inflammatory cytokines (data not shown). Stimulation of BMDCs with CpG DNA caused significant increase in the production of IL-12 p40, IL-6 and TNF-α (Figure 3). Comp pretreatment exhibited strong inhibition of IL-12 p40, IL-6 and TNF-α production in the CpG-stimulated BMDCs (Figure 3).

**Effects of comp on the phosphorylation of MAPK and IκBα by CpG DNA-stimulated BMDCs**

In response to extracellular signals, the three major MAPKs including p38 kinase, c-jun N-terminal kinase (JNK) and extracellular signal-regulated kinase get activated through phosphorylation leading to the increase gene expression of certain pro-inflammatory cytokines like IL-12, IL-6 and TNF-α (Intayoung et al. 2016). Therefore, through western blot analysis, we evaluated the effects on MAPK phosphorylation and NF-κB activation in CpG DNA-stimulated BMDCs, with and without comp treatment (Figure 4). All three MAPKs get phosphorylated in BMDCs stimulated with CpG DNA. p38, JNK1/2 and ERK1/2 phosphorylation was detected between 15 and 30 min of CpG DNA-stimulation. ERK1/2 and JNK1/2 phosphorylation returned to baseline levels within 60 min of CpG DNA-stimulation. Comp pretreatment in the presence of CpG DNA showed strong inhibition of phosphorylation of ERK1/2, JNK1/2, and p38 (Figure 4(A,B)). Activation of NF-κB requires phosphorylation and subsequent degradation of IκB, which allowed nuclear translocation of NF-κB leading to the expression of certain pro-inflammatory cytokine genes (Tsumagari et al. 2015). Activation of NF-κB pathway was analysed indirectly through phosphorylation of IκBα. CpG DNA stimulation induced IκBα phosphorylation within 15 min get to maximum level at 30 min and return to baseline at 60 min of stimulation (Figure 4(A,B)). Comp pretreatment completely inhibited phosphorylation of IκBα in CpG DNA-stimulated BMDCs (Figure 4(A,B)). Taken together these results demonstrate that comp can inhibit CpG DNA-stimulated phosphorylation of ERK1/2, JNK1/2, p38 and IκBα and consequently inhibited MAPK and NF-κB pathways.

**Comp treatment inhibited AP-1 reporter activity in HEK293T cells**

AP-1 is activated by MAPK signalling pathway, which up regulates the transcription of gene encoding pro-inflammatory cytokines (Lee & Yang 2013; Saba et al. 2015). To investigate whether the comp had an inhibitory effect on CpG DNA-stimulated AP-1 transcriptional activity, an AP-1 reporter gene assay was conducted (Figure 5). The HEK293T cells transfected with empty pcDNA3 showed little AP-1-dependent luciferase activity on CpG DNA-stimulation. On the contrary, the HEK293T cells transfected with pcDNA3-mTLR9 exhibited robust AP-1-dependent luciferase activity on CpG DNA-stimulation. However, comp pretreatment exhibited strong inhibition of AP-1-dependent luciferase activity in HEK293T cells transfected with pcDNA3-mTLR9 (Figure 5). Thus, the data suggest that comp has an
inhibitory effect on TLR9-dependent AP-1 activation on CpG DNA stimulation.

**Comp treatment inhibited NF-κB reporter activity in HEK293T cells**

NF-κB is a family of transcription factors that has a key role in regulation of inflammation and immune response (Pacifico & Leonardi 2010). To determine whether comp inhibits CpG DNA-stimulated NF-κB transcriptional activity, an NF-κB reporter gene assay was performed (Figure 6). The HEK293T cells transfected with pcDNA3-mTLR9 exhibited robust NF-κB-dependent luciferase activity on CpG DNA stimulation. However, comp pretreatment exhibited strong inhibition of NF-κB-dependent luciferase activity in HEK293T cells transfected with pcDNA3-mTLR9 (Figure 6). Thus, these data demonstrate that comp has inhibitory effect on TLR9-dependent NF-κB activation.

**Discussion**

In this study, the anti-inflammatory potential of comp, a constituent of *U. pertusa* was examined. To the best of our knowledge, this is the first study to ascertain that comp, isolated from *U. pertusa* has the potential for medicinal solicitation in inflammation-associated diseases. A large body of evidence indicates that inflammation is major risk factor in the development of various human diseases like chronic asthma, rheumatoid arthritis, inflammatory bowel disease (IBD), multiple sclerosis and psoriasis. Therefore, inhibiting inflammation is critical to prevent or control various diseases (Debnath et al. 2013). The agent that reduced pro-inflammatory mediator and cytokines level is regarded as an effective therapeutic strategy for relieving a chronic inflammatory disease.

The pro-inflammatory cytokine IL-12 is expressed predominantly by activated dendritic and phagocytic cells, and is stimulated by pathogens, TLR ligands and CD40L (Teng et al. 2010).
IL-12 p40, IL-6 and TNF-α production warrant further studies concerning potential in future anti-inflammatory application. ERK, p38 kinase and c-jun-N-terminal kinase (JNK) get activated through phosphorylation in response to extracellular signals and ultimately leads to modulation of gene expression in the nucleus (Manzoor & Koh 2012). Phosphorylation of IkB tag it for proteosomal degradation resulting into release and nuclear translocation of NF-kB and hence activation of NF-kB pathway (Tsumagari et al. 2015). In this study, we found that comp inhibited the phosphorylation and hence activation of all three MAPK kinases including ERK1/2, JNK1/2 and p38 in CpG DNA-stimulated BMDCs. Similarly comp also inhibited NF-kB pathway activation by suppressing the phosphorylation of IkBz in CpG DNA-stimulated BMDCs. Stimulation of TLR9 by bacterial DNA (Koh 2011; Masuda et al. 2013) activates AP-1 and NF-κB transcription factors leading to production of inflammatory cytokines (Debnath et al. 2013; Manzoor et al. 2014b). In this study, we found that comp down-regulated AP-1 and NF-kB in HEK293T cells.

In summary, our study demonstrates that comp significantly suppressed the level of pro-inflammatory cytokines. It has also been demonstrated that comp strongly inhibited MAPK and NF-kB pathways. The anti-inflammatory effect of comp was further confirmed by inhibition of AP-1 and NF-κB reporter gene activity. Taken together, comp-mediated anti-inflammatory activity represents a potential therapeutic use of the comp for inflammatory diseases.

Conclusions

In conclusion, our data suggest that this compound can act as potent anti-inflammatory drug. We found that comp had an inhibitory effect on the production of pro-inflammatory cytokines by down-regulating TLR9-dependent AP-1 and NF-kB activation. These results suggest that comp might be beneficial in the treatment of inflammation and autoimmune-associated diseases. Therefore, further studies are required on the detailed mode of action and in vivo efficacy of comp.

Disclosure statement

The authors report no declaration of interest.

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