Docosahexaenoic acid attenuates cell death and interleukin-1β secretion in THP-1 cells responded to *Aggregatibacter actinomycetemcomitans* invasion.

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**Abstract**

Macrophages play an important role in the innate immune system by producing inflammatory cytokines in response to substances derived from bacteria and viruses. Recently, omega-3 fatty acids have been shown to have beneficial effects in inflammatory diseases. In this study, we investigated the effects of omega-3 fatty acids on inflammatory responses and cell death induced by the Gram-negative bacterium *Aggregatibacter actinomycetemcomitans* in macrophages. *A. actinomycetemcomitans* invasion induced secretion of interleukin (IL)-1β and expression of inflammasome-associated factors, including nucleotide-binding oligomerization domain like receptor protein 3 (NLRP3), adaptor apoptosis-associated speck-like protein containing a caspase recruitment domain (ASC), caspase-1, caspase-4, and cleaved-gasdermin D (GSDMD), in THP-1 cells. Docosahexaenoic acid (DHA), an omega-3 fatty acid suppressed NLRP3, ASC and caspase-1 expression as well as IL-1β secretion by THP-1 cells in response to *A. actinomycetemcomitans* invasion. Moreover, DHA impaired assembly of ASC, suppressed the expression of caspase-4 and cleaved-GSDMD, and suppressed cell death in *A. actinomycetemcomitans*-invaded THP-1 cells. Taken together, our data suggest that DHA attenuates the secretion of IL-1β and cell death induced by *A. actinomycetemcomitans* invasion of macrophages via suppression of inflammasome-associated factors.

**Keywords:** Docosahexaenoic acid, *Aggregatibacter actinomycetemcomitans*, Macrophage, Inflammasome, Interleukin-1β, Cell death.

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**Introduction**

*Aggregatibacter actinomycetemcomitans* is a Gram-negative microorganism that is frequently isolated from patients with periodontal disease [1]. In periodontal tissue, *A. actinomycetemcomitans* induces host immune responses by producing exotoxic and endotoxic virulence factors [2]. Pathogen recognition is essential to activate the innate immune system, and is primarily performed by macrophages and dendritic cells. Periodontopathic bacteria including *A. actinomycetemcomitans* are medically important pathogens, and many have been associated with systemic disorders such as diabetes mellitus [3], arteriosclerotic cardiovascular diseases [4], and infectious diseases [5]. Chronic systemic inflammation is the result of pro-inflammatory cytokine secretion following activation of the innate immune system. Previous experiments showed that *A. actinomycetemcomitans* invasion induced production of interleukin (IL)-1β, a critical inflammatory mediator in macrophages [6].

Inflammation is an essential component of immune responses and can be triggered by tissue damage or bacterial infection. The early events of the innate immune response are mediated by pattern recognition receptors (PRRs) [7]. The inflammasome is a large intracellular signalling complex containing cytosolic PRRs including nucleotide-binding oligomerization domain-like receptors (NLRs) [8]. In NLR inflammasome complexes, the NLR protein 3 (NLRP3) inflammasome plays a notable role in innate immune responses to bacterial infection [9]. The NLRP3 inflammasome, through the adaptor apoptosis-associated speck-like protein containing a caspase recruitment domain (ASC), activates caspase-1 resulting in processing and release of IL-1β. IL-1β is a key cytokine for control of immune tolerance during inflammation.
These pathways are currently designated as “canonical” and “non-canonical” NLRP3 inflammasome activation [11]. Canonical inflammasome activation requires two independent steps: transcription and oligomerization. In the first step, LPS is recognized by TLR4, promoting NLRP3 inflammasome transcription following nuclear factor kappa-light-chain-enhancer of activated B cells (NF-κB) activation. NLRP3 inflammasome oligomerization results in IL-1β release at the second step [12]. By contrast, non-canonical inflammasome activation in macrophages is induced by Gram-negative bacteria such as Escherichia coli, Citrobacter rodentium or Vibrio cholerae[13]. In the non-canonical inflammasome pathway, LPS released by cytosolic Gram-negative bacteria, which is endocytosed by host cells, activates mouse caspase-11 or human caspase-4/5 [14]. Activation of caspase-4 promotes production of pro-inflammatory cytokines and results in cleavage of the newly discovered protein, gasdermin D (GSDMD), which is required for pyroptosis[15]. Pyroptosis is a type of inflammatory programmed cell death and is regulated via caspase-1-dependent or caspase-1-independent mechanisms. Activation of inflammatory caspases causes pyroptosis as well as IL-1β, these functions are also important for eliminating pathogens and maintaining host cell homeostasis [16].

Fatty acids modulate inflammatory processes and contribute to the pathophysiology of diet-related chronic diseases [17]. Fatty acids are composed of carbon, hydrogen, and oxygen, and their physiological activity differs depending on the number of double bonds [18]. Polyunsaturated fatty acids (PUFAs) are classified as essential fatty acids that cannot be synthesized by mammals and therefore must be obtained from dietary sources. There are two main categories of PUFAs: omega-6 and omega-3 PUFAs. Several human nutritional studies have suggested that dietary factors can promote or reduce inflammation [19,20]. Diets high in fat, especially saturated fatty acids appear to increase the risk of developing heart disease [21]. By contrast, foods high in omega-3 PUFAs can have beneficial effects in rheumatoid arthritis and in stabilizing advanced atherosclerotic plaques [22]. Previously, we reported that docosahexaenoic acid (DHA), an omega-3 PUFA, promoted M2 macrophage polarization [23]. M2 macrophages have anti-inflammatory properties mediated through secretion of anti-inflammatory cytokines. However, the relationships between inflammatory responses induced by bacterial infection and omega-3 PUFAs are poorly understood.

In the present study, we examined the ability of A. actinomycetemcomitans to induce expression of inflammatory cytokines and non-canonical inflammasome related factors in macrophages. We show that DHA suppressed macrophage inflammatory responses by inhibiting ASC speck formation, secretion of IL-1β, and prevented macrophage pyroptosis induced by A. actinomycetemcomitans invasion via suppression of the non-canonical inflammasome pathway.

Materials and Methods

Cells

THP-1, human leukemia monocyctic cell lines were purchased from the Japanese Cancer Research Resources Bank (Osaka, Japan). These cells were cultured according to published methods [23]. To differentiate THP-1 monocytes into macrophages, cells were stimulated with phorbol 12-myristate 13-acetate (PMA) (Sigma-Aldrich) overnight in 6-well plates (Iwaki, Chiba, Japan). And then, cells were refreshed to the culture medium [24]. After 48 h, cells were pre-treated with docosahexaenoic (DHA) (50 µM) for 24 h.

Bacterial strains and A. actinomycetemcomitans invasion

A. actinomycetemcomitans strain Y4 was grown according to the method accounted by previous report [6]. THP-1 cells (5×10⁵ cells/mL) were invaded with A. actinomycetemcomitans at a multiplicity of infection (MOI) of 50 in 6-well plates (Iwaki) according to the procedure described by Okinaga et al. By using this method, it was confirmed that bacteria can invade the cells under a microscope [6].

Reagents

DHA was purchased from Cayman Chemical (Ann Arbor, MI, USA). Anti-IL-1β polyclonal antibody was from Novus Biologicals (Centennial, CO, USA). Anti-NLRP3 monoclonal antibody was purchased from AdipoGen Life Sciences (Basel, Switzerland). Anti-caspase-1 monoclonal antibody was from Abcam (Cambridge, UK), and anti-caspase-4 monoclonal antibody was from MBL (Nagoya, Japan). Anti-ASC and anti-GSDMD monoclonal antibodies were from Santa Cruz Biotechnology (Dallas, TX, USA). Anti-β-actin monoclonal antibody was obtained from Sigma-Aldrich (St. Louis, MO, USA). Anti-rabbit anti-IgG horseradish peroxidase (HRP) conjugated antibody and anti-mouse anti-IgG HRP conjugated antibody were obtained from GE Healthcare (Little Chalfont, UK).

Quantitative real time PCR (qRT-PCR)

qRT-PCR was performed in accordance with a published protocol (Kawano et al., 2019). The primer sequences for qRT-PCR were human IL-1β, forward 5’-TCAGCCAATCTTICATTGCTCAA-3’ and reverse 5’-TGGCGAGCTCAAGTACTT CTG-3’; human β-actin, forward 5’-GGCGCGGCTACAGTTCCA-3’ and reverse 5’-CTTAA GTGACCACGCTTTCC-3’.

Enzyme-linked immunosorbent assay (ELISA)

Cell supernatants were assessed for human IL1-β and mouse IL-1β using Quantikine ELISA kits (R&D systems) according to manufacturer’s instructions.
Western blotting
Proteins from treated cells were extracted using sodium dodecyl sulfate (SDS) lysis buffer (50 mM Tris–HCl containing 2% SDS, pH 6.8) and a protease inhibitor mixture (Thermo Fisher Scientific, MA, USA). Western blotting was performed as described in our previous report.

Cell viability and cytotoxicity assays
To evaluate cell viability, a cell counting kit-8 (CCK-8) (Dojindo Molecular Technologies, Kumamoto, Japan) was used. The CCK-8 assay was performed according to manufacturer’s protocol. Cytotoxicity was assessed using a lactate dehydrogenase (LDH) assay kit (Dojindo Molecular Technologies). The assay was performed with recovered supernatants according to the manufacturer’s protocol.

Immunofluorescence
Cells were fixed using the method accounted by Okinaga et al. [6]. The cells were incubated with anti-ASC antibody overnight at 4°C and incubated with secondary Alexa Fluor 488-conjugated goat anti-mouse (Invitrogen, Carlsbad, CA, USA) for 1 h at room temperature. The cells were mounted in VECTASHIELD mounting medium with DAPI (Vector Laboratories, Burlingame, CA, USA). All-in-One Fluorescence microscope system (Keyence, Tokyo, Japan) was used for detection of immunofluorescence images.

Statistical analyses
Data were analysed using Excel (Microsoft, Redmond, WA, USA) and expressed as the means ± standard deviations. One-way ANOVA test were used followed by Tukey’s post hoc test. The difference between two groups was assessed using unpaired Student’s t-tests. A value of p<0.05 was considered statistically significant.

Results
DHA suppressed production of IL-1β induced by A. actinomycetemcomitans invasion
We used PMA-differentiated THP-1 macrophages to investigate the inflammatory effects of A. actinomycetemcomitans invasion. A. actinomycetemcomitans enhanced the expression of IL-1β 6 h post-invasion of THP-1 macrophages; however, DHA pre-treatment suppressed upregulation of IL-1β transcripts (Figure 1A). ELISA showed that DHA significantly downregulated IL-1β secretion 24 h post-invasion with A. actinomycetemcomitans in THP-1 macrophages (Figure 1B). To confirm the effect of other fatty acids on secretion of IL-1β, we used EPA, another omega-3 fatty acid, and PAL, a saturated fatty acid. EPA slightly reduced the secretion of IL-1β induced by A. actinomycetemcomitans invasion, while PAL increased.

DHA suppressed ASC speck formation
To explore the effects of DHA on inflammasome activity, we investigated the expression of NLRP3 and ASC. As shown Figure 2A, DHA suppressed NLRP3 and ASC expression induced 24 h following A. actinomycetemcomitans invasion. In addition, DHA inhibited the assembly of ASC 24 h following A. actinomycetemcomitans invasion (Figure 2B). A. actinomycetemcomitans promoted ASC speck formation in about 50% of the cells, but was rescued by treatment with DHA.

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DHA reduced the expression of caspase-1 and suppressed the non-canonical inflammasome pathway

The influence of DHA on expression of caspases important for inflammasome activity was investigated. DHA treatment suppressed caspase-1 activation induced by A. actinomycetemcomitans invasion for 12-24 h (Figure 3A). We also investigated the effect of DHA on non-canonical inflammasome signalling. As shown in Figure 3B, expression of caspase-4 induced 12-24 h following A. actinomycetemcomitans invasion was inhibited by DHA pre-treatment. In addition, A. actinomycetemcomitans invasion for 12-24 h induced cleavage of GSDMD, cleavage was suppressed by DHA pre-treatment (Figure 3C). The roles of pro-inflammatory mediators during A. actinomycetemcomitans invasion were investigated using siRNA-transfected THP-1 cells. Cells independently silenced for ASC, caspase-1, caspase-4 and GSDMD expressions were prepared. IL-1β secretion by A. actinomycetemcomitans-invaded THP-1 cells was reduced in the absence of ASC, caspase-1, and GSDMD.

DHA attenuated A. actinomycetemcomitans cytotoxicity

Finally, we examined the cell damage induced by A. actinomycetemcomitans and its mitigation by DHA. A. actinomycetemcomitans invasion reduced cell viability and stimulated the release of LDH, but DHA pre-treatment suppressed both effects (Figures 4A and 4B).
Docosahexaenoic acid attenuates cell death and interleukin-1β secretion in THP-1 cells responding to Aggregatibacter actinomycetemcomitans invasion.

**Discussion**

In this study, we showed that DHA suppressed cell death and interleukin-1β secretion in THP-1 cells responded to Aggregatibacter actinomycetemcomitans invasion. Our results may be useful in establishing novel treatment strategies for chronic oral inflammatory diseases using DHA supplementation.

Aggressive periodontitis is primarily caused by *A. actinomycetemcomitans* invasion and excessive inflammatory responses in immune cells, including macrophages [25]. We previously reported that *A. actinomycetemcomitans* invasion of macrophages occurred through a phagocytic process and that cytotoxicity was linked to cytoplasmic factors in macrophages [6]. We showed that DHA suppressed the response to *A. actinomycetemcomitans* invasion in the human monocytic cell line THP-1 (Figure 1A). As shown Figure 1B, secretion of IL-1β into culture media was increased by the noncanonical pathway was activated [41]. In our experiments, DHA significantly reduced the expression of caspase-1, caspase-4, and cleaved-GSDMD induced by *A. actinomycetemcomitans* invasion. Therefore, DHA may inhibit signal transduction by directly affecting ASC-associated upstream events in inflammasome formation. Another study showed that dietary PUFAs suppressed NLRP3 inflammasome formation through autophagy using knockout experiments of autophagy-related factors [32,33]. We previously reported that DHA promoted the differentiation of M2 macrophages with anti-inflammatory activity through autophagy-related mechanisms [23]. Considering this finding, it is possible that autophagy is also partially caused by degradation of NLRP3 and ASC by DHA. By contrast, PAL, a saturated fatty acid, enhanced IL-1β secretion. PAL can promote polarization of specific cells in response to inflammatory stimuli [34]. In addition, PAL attenuates autophagy mechanisms and is involved in development of insulin sensitivity and obesity-related metabolic diseases such as type 2 diabetes [35]. Pro-inflammatory M1 macrophages accumulated in pancreatic islets in a mouse model of type 2 diabetes, and expression of inflammatory cytokines such as IL-1β and tumor necrosis factor-α was also enhanced [36]. Therefore, PAL shows effects opposite to those of the PUFAs DHA and EPA. Further research is needed to investigate whether fatty acids regulate inflammasome activity triggered by autophagy-associated molecules.

Caspase-1 activation in macrophages invaded with Salmonella or *Shigella* species results in cytokine processing and cell death [37]. Caspases-4 and -5, which belong to the family of cysteine proteases, are classified as inflammatory caspases. These enzymes contain a long prodomain containing a caspase activation and recruitment domain (CARD) and are composed of large and small subunits. Human caspase-4 is a regulator of non-canonical inflammasome activation and contributes to defense against Gram-negative bacteria [38]. GSDMD is a critical target of caspase-4 and a key mediator of host responses [39]. The N-terminal fragment of GSDMD mediates pore formation in the plasma membranes of dying cells [40]. Caspase-4 binds directly to cytosolic LPS via its CARD. LPS binding induced oligomerization of caspase-4, indicating that the noncanonical pathway was activated [41]. In our experiments, DHA significantly reduced the expression of caspase-1, caspase-4, and cleaved-GSDMD induced by *A.
Actinobacillus actinomycetemcomitans invasion (Figure 3). These results indicated that DHA prevents cell membrane damage caused by Gram-negative bacterial invasion through suppression of caspase-4-mediated GSDMD cleavage. Thus, we conclude that cleavage of GSDMD was blocked directly or indirectly by DHA treatment. IL-1 family cytokines are released during inflammatory responses in numerous tissues [42]. IL-1β is detected as a cytosolic factor that lacks an N-terminal secretion signal and is therefore not secreted via conventional pathways [43]. These facts suggest that membrane pore formation by the N-terminal fragment of GSDMD is important in extracellular release of IL-1β. Hence, DHA may inhibit the formation of cell membrane pores caused by the cleavage of GSDMD. Furthermore, we speculate that DHA may subsequently block the non-canonical inflammasome signaling pathway by inhibiting the interaction between A. actinomycetemcomitans and caspase CARDs. We confirmed that ASC, caspase-1, and GSDMD silencing inhibited IL-1β secretion, but caspase-4 silencing had no effect (data not shown). In the absence of caspase-4, it is considered that IL-1β secretion is promoted by caspase-4-mediated GSDMD cleavage. Thus, we conclude that DHA can suppress the inflammatory response by altering inflammasome signaling and inhibit pyroptosis by preventing cleavage of GSDMD. These findings demonstrate DHA’s inhibitory effects on secretion of IL-1β and cell death during bacterial invasion. DHA may have beneficial effects on the inflammatory responses caused by periodontal pathogens.

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