Sulfur-containing amino acids in aged garlic extract inhibit inflammation in human gingival epithelial cells by suppressing intercellular adhesion molecule-1 expression and IL-6 secretion

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Abstract. Aged garlic extract (AGE) contains various biologically active sulfur-containing amino acids, such as S-allylcysteine (SAC), S-1-propenylcysteine (S1PC) and S-allylmercaptocysteine (SAMC). These amino acids have been demonstrated to lower hypertension, improve atherosclerosis and enhance immunity through their anti-inflammatory and antioxidant activities. It was recently reported that the administration of AGE alleviated gingivitis in a clinical trial. In this study, to gain insight into this effect of AGE, the authors examined whether AGE and the three above-mentioned sulfur compounds influence the effects of tumor necrosis factor-α (TNF-α) in inducing intercellular adhesion molecule-1 (ICAM-1) expression and interleukin-6 (IL-6) secretion in Ca9-22 human gingival epithelial cells. It was found that SIPC reduced the level of ICAM-1 protein induced by TNF-α possibly through post-translational levels without affecting the TNF-α-induced mRNA expression. However, SAC and SAMC had no effect. It was also confirmed the inhibitory effect of an antimicrobial peptide [human-β defensin-3 (hβD3)] and found that the inhibitory effects of hβD3 and S1PC were synergistic. On the other hand, the TNF-α-induced IL-6 secretion was attenuated by SAC and SAMC in a dose-dependent manner, whereas S1PC was ineffective. In addition, SAC and SAMC, but not S1PC inhibited the phosphorylation of the transcription factor nuclear factor-κ-light-chain-enhancer of activated B cells (NF-κB), which is involved in the expression of inflammatory molecules, suggesting that the anti-inflammatory effects of SAC and SAMC are mediated, at least partly, by NF-κB. On the whole, the findings of this study suggest that the three sulfur amino acids in AGE function synergistically in alleviating inflammation in human gingival epithelial cells.

Introduction

Gingivitis is a gingival inflammation triggered by periodontal pathogens present in oral biofilms (1). Unless some physical treatments, such as plaque control and scaling are administered, gingivitis becomes more severe and develops into periodontitis with more severe inflammation, leading to the destruction of connective tissues, the absorption of alveolar bones and tooth loss (1,2). It has been demonstrated that excessive inflammation in the gingival tissue, as well as an immune reaction triggered by the increased number of periodontal pathogens, such as Porphyromonas gingivalis (P.g.), plays an important role in the severity and progression of periodontal diseases (3,4). Thus, the control of inflammation may help to delay disease progression or recover severe symptoms in periodontal diseases.

Aged garlic extract (AGE) is prepared by aging garlic (Allium sativum L.) for a period of >10 months in aqueous ethanol and contains pharmacologically active sulfur-containing amino acids, such as S-allylcysteine (SAC), S-1-propenylcysteine (S1PC) and S-allylmercaptocysteine (SAMC). In human and animal studies, AGE and these sulfur compounds have been shown to exert beneficial effects, such as the improvement of atherosclerosis (5,6), decrease of high blood pressure (7,8) and the modulation of immunity (9,10). These effects are possibly

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Abbreviations: AGE, aged garlic extract; ANOVA, analysis of variance; BSA, bovine serum albumin; cDNA, complementary DNA; DAS, diallylsulfide; DMEM, dulbecco's modified eagle's medium; EDTA, ethylenediaminetetraacetic acid; ELISA, enzyme-linked immunosorbent assay; ERK, extracellular signal-regulated kinase; FBS, Fetal bovine serum; hβD3, human β-defensin-3; HCl, hydrochloric acid; HPLC, high performance liquid chromatography; HRP, horseradish peroxidase; HUVECs, human vein endothelial cells; ICAM-1, intercellular adhesion molecule-1; IgG, immunoglobulin G; IL-6, interleukin-6; LC-MS, liquid chromatography-mass spectrometry; LPS, lipopolysaccharide; MMP, matrix metalloprotease; NF-κB, nuclear factor-κ-light-chain-enhancer of activated B cells; NOD, nucleotide binding oligomerization domain-containing protein; PBS, phosphate-buffered saline; PCR, polymerase chain reaction; P.g., Porphyromonas gingivalis; RANKL, receptor activator of NF-κB ligand; ROS, reactive oxygen species; S1PC, S-1-propenylcysteine; SAC, S-allylcysteine; SAMC, S-allylmercaptocysteine; SD, standard deviation; TBS, Tris-buffered saline; TNF-α, tumor necrosis factor-α

Key words: aged garlic extract, gingival epithelial cells, ICAM-1, IL-6, inflammation, sulfur-containing amino acid
ascribed, at least in part, to the anti-inflammatory (9,11) and/or antioxidant (12,13) activities of sulfur compounds in AGE.

It has been reported that several naturally occurring substances, such as curcumin and theaflavin, alleviate the absorption of alveolar bones in animal periodontitis models (4,14). It has also been demonstrated that one of the sulfur-containing compounds present in garlic, diallylsulfide (DAS), reduces inflammatory reactions elicited by P.g.-derived lipopolysaccharide (LPS) in human gingival fibroblast cells (15). Furthermore, DAS and another garlic sulfur compound, alliin, have been shown to inhibit the growth of periodontal microbes (16-18). These findings suggest that sulfur compounds in garlic attenuate moderate or severe inflammation occurring in gingival tissues of gingivitis and periodontitis. Recently, a clinical study demonstrated that the intake of AGE significantly improved gingivitis (1). However, the mechanisms through which AGE attenuates gingivitis remain to be fully elucidated. In this study, the authors examined whether SAC, SIPC and SAMC in AGE exert anti-inflammatory effects against tumor necrosis factor-α (TNF-α)-induced inflammatory reactions in the human gingival epithelial cell line, Ca9-22.

Materials and methods

**Aged garlic extract (AGE) and sulfur compounds.** AGE was prepared as previously described (12). Briefly, AGE was manufactured according to the following steps: Originally grown raw garlic (A sativum L.) was cut into slices, immersed in aqueous ethanol, and extracted for >10 months at room temperature (12). The sulfur-containing amino acids, SIPC, SAC and SAMC, are produced during the aging process of raw garlic in 50% ethanol solution for >10 months (19). The purification and structure determination of sulfur compounds in AGE was performed by high performance liquid chromatography (HPLC) and liquid chromatography-mass spectrometry (LC-MS) systems as previously described (9).

**Reagents.** Dulbecco's modified Eagle's medium (DMEM, D6046) and WST-1 reagent (501594401) were purchased from Sigma-Aldrich. Fetal bovine serum (FBS, 10270), 0.05% trypsin/EDTA solution (25300-054), Halt protease and phosphatase inhibitor single-use cocktail (1861280) and TRIzol reagent (15596018) were from Thermo Fisher Scientific. Human recombinant human TNF-α (100 ng/ml) and 50 mM dithiotheritol in 100 mM Tris-hydrochloric acid (HCl) buffer, pH 6.8), and was heated at 95°C for 5 min. The samples were separated by electrophoresis and were blotted onto a nitrocellulose membrane (1620090, Bio-Rad). The blotted membrane was blocked with 5% non-fat dry milk (999S, Cell Signaling Technology) in Tris-buffered saline (TBS) for 1 h. Following treatment of the cells with TNF-α (100 ng/ml), SIPC, SAC or SAMC (each at 1 to 100 µM) for 6 h, an aliquot of culture medium was collected and frozen at -80°C until use. The IL-6 protein amount in the medium was determined with a human IL-6 Uncoated enzyme-linked immunosorbsent assay (ELISA) kit (88-7066-22, Thermo Fisher Scientific).

**Western blot analysis.** Cells were washed with phosphate-buffered saline (PBS) 5 times and total protein was extracted with RIPA lysis buffer containing protease inhibitor cocktail. The protein amount in the extract was determined with the Pierce BCA Protein Assay kit (23225, Thermo Fisher Scientific) with bovine serum albumin (BSA, A7030, Sigma) as a standard. The protein extract (10 µg) was denatured by the addition of sample buffer [5% sodium dodecyl sulfate, 12.5% glycerol and 50 mM dithiotheritol in 100 mM Tris-hydrochloric acid (HCl) buffer, pH 6.8], and was heated at 95°C for 5 min. The samples were separated by electrophoresis and were blotted onto a nitrocellulose membrane (1620090, Bio-Rad). The blotted membrane was blocked with 5% non-fat dry milk (999S, Cell Signaling Technology) in Tris-buffered saline (TBS)-T buffer (0.1% Tween-20, pH 7.6) for 1 h, and was subsequently incubated in the presence of the rabbit anti-ICAM-1 (4915S, Cell Signaling Technology), anti-NF-κB (sc-109X, Santa Cruz Biotechnology) or anti-phosphorylated NF-κB (3033, Cell Signaling Technology) antibody at a 1:2,000 dilution overnight at 4°C. The membrane was washed 3 times and incubated in the presence of the HRP-conjugated anti-rabbit IgG antibody (7074S, Cell Signaling Technology) for 1 h at room temperature. On the other hand, another blotted membrane was incubated with the HRP-conjugated anti-β-actin antibody (PM053-7, MBL Life Science) at a 1:2,000 dilution for 1 h at room temperature, in order to quantitatively normalize the immunoreactive ICAM-1 protein levels. After washing, immunoreactive proteins on the nitrocellulose membrane were visualized with Armasham ECL Prime peroxidase solution (RPN2232V, GE Healthcare), by using a luminoimage analyzer (Chemidoc™ MP, Bio-Rad). The density of each immunoreactive band was analyzed with Image Lab™ software ver. 4.1 (Bio-Rad).

**Reverse transcription-quantitative polymerase chain reaction (RT-qPCR).** Total RNA was isolated from cells with TRIzol reagent. Contaminated genomic DNA in the total RNA was removed by incubation with gDNA eraser (RR047A, Takara)
Figure 1. Effect of S1PC and hβD3 on the level of ICAM-1 protein induced by TNF-α. (A) The concentration (1 to 100 ng/ml, 12 h) - and (B) time (1 to 24 h, 100 ng/ml) - dependent induction of ICAM-1 protein stimulated by TNF-α. The cells were left untreated (control) or treated with TNF-α in the absence or presence of (C) S1PC (1 to 100 µM), (D) AGE (0.01 to 1 mg/ml), (E) SAC (10, 100 µM), (F) SAMC (10, 100 µM) or (G) hβD3 (1 to 100 nM) for 24 h and total protein was extracted. ICAM-1 proteins in the extract were detected by western blot analysis. Each bar in the graphs represents the mean ± SD of the band intensity relative to β-actin (n=3). *P<0.05, **P<0.01 in comparison to the control (Tukey’s multiple comparison test). &P<0.05 in comparison to TNF-α alone (Student's t-test). S1PC, S-1-propenylcysteine; hβD3, human β-defensin-3; TNF-α, tumor necrosis factor-α; ICAM-1, intercellular adhesion molecule-1; AGE, aged garlic extract; SAC, S-allylcysteine; SAMC, S-allylmercaptocysteine.
for 5 min at room temperature, and total RNA was reverse transcribed into complementary DNA (cDNA) with a PrimeScript RT reagent kit with genomic DNA Eraser (RR047A, Takara). cDNA was amplified with primer pairs using a SYBR Premix Ex Taq II (RR820A, Takara). The level of gene expression relative to the internal control (β-actin) was calculated by performing a quantitative (real-time) PCR with a Real-time PCR system (PikoReal 96, Life Technologies; Thermo Fisher Scientific). The reaction was run using the following program: 30 sec at 95°C, 40 repeats of 10 sec at 95°C, 10 sec of 63°C and 15 sec of 72°C. The sequences of the specific primers were as follows: human ICAM-1 forward, 5'-TGGGACACAAA GCACATATG-3' and reverse, 5'-CAAGGACGAGGAGG AACGC-3'; human IL-6 forward, 5'-TCTCCAACAGCCCT T-3' and reverse, 5'-CTCAGGGCTGAGATG-3'; and β-actin forward, 5'-CCGCGAGATGACCCCAGAT-3' and reverse, 5'-GGTGAAGATCTTTAGTGAAGTGC-3'. The fold change in the relative mRNA level to β-actin was calculated based on the Ct (ΔΔCt) method (20).

Cellular IL-6 protein content. Following treatment of the cells with TNF-α (100 ng/ml) in the absence or presence of AGE (1 mg/ml) or SAC (100 µM) for 3 h, the total protein was extracted with RIPA buffer. The amount of IL-6 protein in the extract was determined by ELISA, as described above. The cellular IL-6 protein content is expressed relative to 1 mg total protein.

Statistical analysis. Data are expressed as the means ± standard deviation (SD). Statistical significance was evaluated using the Student’s t-test for differences between 2 groups or one-way analysis of variance (ANOVA), followed by Tukey’s post hoc test for differences between >3 groups. A P-value <0.05 was statistically significant in comparison to control or the effect of TNF-α alone.

Results

Effect of S1PC and hβD3 on the level of ICAM-1 protein induced by TNF-α. Human gingival fibroblasts and Ca9-22 cells have been shown to express ICAM-1 in response to the periodontal disease bacteria, P. g.-derived LPS or TNF-α (21-25). It was also found that ICAM-1 protein expression in the Ca9-22 cells was profoundly induced by treatment with TNF-α in a concentration-dependent manner (Fig. 1A). In addition, the ICAM-1 protein level began to increase at 3 h, reached peak levels at 12 h and remained elevated until 24 h (Fig. 1B). As shown in Fig. 1C, the presence of S1PC for 3 h inhibited the increased level of TNF-α-induced ICAM-1 protein in a concentration-dependent manner, whereas treatment with AGE exerted minimal inhibitory effects even at the highest concentration (1 mg/ml) (Fig. 1D). Both SAC and SAMC also had no significant effect (Fig. 1E and F).

It has been previously demonstrated that the anti-microbial peptide hβD3 reduces the level of TNF-α-elicited ICAM-1 protein in human vein endothelial cells (HUVECs) (26). Thus, this study examined whether the peptide inhibits the level of ICAM-1 protein in Ca9-22 cells. As shown in Fig. 1G, the increased level of TNF-α-induced ICAM-1 protein was moderately but significantly suppressed by hβD3 at 10 nM, indicating that this peptide is also effective. Furthermore, it was found that the inhibitory effect of hβD3 was enhanced by simultaneously treating the cells in the presence of S1PC (Fig. 2), suggesting the synergistic effect of hβD3 and S1PC.

Effect of hβD3 and S1PC on ICAM-1 gene expression induced by TNF-α. It is conceivable that the inhibitory effect of S1PC on the level of ICAM-1 protein was due to the suppression of ICAM-1 gene transcriptions. Thus, this study then examined the effects of S1PC as well as those of hβD3 over a period of 3 or 24 h treatment on the TNF-α-induced ICAM-1 gene expression by RT-qPCR. As shown in Fig. 3A, TNF-α significantly enhanced the basal ICAM-1 mRNA transcription level with a peak at 3 h. Treatment with S1PC or hβD3 for either 3 or 24 h had no effect on the mRNA expression level of ICAM-1 induced by TNF-α (Fig. 3B and C). These results suggested that the actions of S1PC and hβD3 occurred at the post-transcriptional level, since these compounds did not affect ICAM-1 expression even at the concentrations at which induced a decrease in the protein level.

Effect of S1PC on the level of ICAM-1 protein induced by TNF-α in the presence of cycloheximide (CHX). Since S1PC did not affect ICAM-1 gene expression, we then examined their effect at the translational level using CHX, an inhibitor of protein synthesis. As shown in Fig. 4, the inhibitory effects of CHX on the protein level of ICAM-1 were enhanced in the presence of S1PC in cells treated with TNF-α. This result suggested that the effects of S1PC occurred at the
post-translational level, since this compound induced a further reduction in the ICAM-1 protein level when total protein synthesis was blocked.

Effect of AGE, SAC and SAMC on IL-6 secretion induced by TNF-α. IL-6 is produced in human gingival epithelial cells in response to LPS or TNF-α, leading to the activation of osteoclasts via the upregulation of receptor activator of NF-κB ligand (RANKL) in osteoblasts (21). This study thus examined the effects of TNF-α on IL-6 secretion in Ca9-22 cells. As shown in Fig. 5A, TNF-α stimulated basal IL-6 secretion during 6 h of culture. The enhanced IL-6 secretion induced by TNF-α was reduced by simultaneous treatment with AGE, SAC or SAMC in a concentration-dependent manner (Fig. 5A-C). On the other hand, S1PC had no effect on IL-6 secretion even at the concentration of 100 µM (Fig. 5D).

Figure 4. Effect of S1PC on the level of ICAM-1 protein induced by TNF-α in the presence of cycloheximide (CHX). (A) The cells were left untreated (control) or pretreated with TNF-α (100 ng/ml) for 12 h in the absence or presence of CHX (10 µM) and/or S1PC (10 µM) for 6 h and total protein was extracted. ICAM-1 proteins in the extract were detected by western blot analysis. (B) The quantitative analysis of the band intensity relative to β-actin. Each bar in the graph represents the mean ± SD (n=3). *P<0.01 in comparison to control and **P<0.01 in comparison to TNF-α alone (Tukey's multiple comparisons test). S1PC, S-1-propenylcysteine; hβD3, human β-defensin-3; TNF-α, tumor necrosis factor-α; ICAM-1, intercellular adhesion molecule-1.
shown in Fig. 6B-D, treatment with AGE, SAC or SAMC for 3 h had a minimal effect on the enhanced IL-6 mRNA expression level induced by TNF-α.

**Effect of AGE, SAC and SAMC on the intracellular IL-6 protein content.** In order to examine whether AGE and sulfur compounds affect the IL-6 secretion process, the cellular IL-6
protein content was measured during TNF-α stimulation by ELISA. Treatment with AGE (1 mg/ml) or SAC (100 µM) for 3 h exhibited a tendency to enhance the effects of TNF-α on the cellular IL-6 content, although their effects were not statistically significant (Fig. 7).

**Effect of hβD3, SIPC, AGE, SAC and SAMC on the level of phosphorylated NFκB p65 protein induced by TNF-α.** Stimulation of the cells with TNF-α has been shown to activate a variety of signaling pathways and transcription factors, such as NF-κB, which plays an essential role in the expression of inflammation-related molecules (26-28). In addition, hβD3 has been shown to alleviate periodontitis via the suppression of NFκB (29). Thus, this study examined the effects of hβD3, SIPC, AGE, SAC and SAMC on the phosphorylation (activation) of NFκB induced by TNF-α. As shown in Fig. 8A and C-E, hβD3 (100 nM), AGE (1 mg/ml), SAC and SAMC (each at 100 µM) inhibited NFκB activation, whereas SIPC (100 µM) did not have any significant effect (Fig. 8B).

**Discussion**

In gingivitis and periodontitis, TNF-α secreted from macrophages and neutrophils accumulates in gingival tissues during the progression of inflammation and induces cytokine production, such as IL-6 in gingival cells, leading to an increased RANKL expression in osteoblasts, the activation of osteoclasts and consequently, to the resorption of alveolar bone (30). Therefore, regulating the excess inflammation in gingiva is considered to be a key step to treat the periodontal diseases. In the present study, it was found that sulfur-containing compounds in AGE differentially inhibited the TNF-α-elicited ICAM-1 expression and IL-6 secretion, suggesting that these compounds control inflammation occurring in human gingival tissues under pathological conditions such as gingivitis.

ICAM-1 is expressed by several cell types in response to a variety of stimuli, such as LPS and pro-inflammatory cytokines, and helps to bind lymphocytes to T cells or vascular endothelial cells, leading to the activation of immune responses and the trans-endothelial migration of lymphocytes (31,32). It has been demonstrated that ICAM-1 is highly expressed in inflamed tissues, such as atherosclerotic lesions and is associated with disease progression (32). There are a number of findings indicating that ICAM-1 is highly expressed by treating HUVECs with TNF-α or LPS (33-35). Zhao et al demonstrated that a variety of intracellular signaling molecules, such as p38 MAPK acting on the MAPK pathway, nicotinamide adenine dinucleotide phosphate (NADPH) oxidase and the transduction factor NF-xB, were involved in the ICAM-1 expression in HUVESC (35). In particular, NF-xB plays a crucial role in innate immune responses via enhanced transcription of inflammatory mediators, including interleukins, chemokines and cytokines (27,28). The findings of study suggest that SIPC can modulate the level of ICAM-1 protein in Ca9-22 cells independently of NF-xB. At present, the SIPC-mediated intracellular pathway remains to be elucidated.

In human gingival fibroblast cells, ICAM-1 is expressed in response to pro-inflammatory cytokines, such as TNF-α and IL-1β (21,22) or periodontal pathogens such as P. gingivalis (23) via the activation of NF-xB, p38 MAPK or nucleotide binding oligomerization domain-containing protein (NOD). On the other hand, to the best of our knowledge, there are only a few studies available to date demonstrating the TNF-α-induced expression of ICAM-1 in human gingival epithelial cells, particularly Ca9-22 cells (22,25), and the functional mechanisms have not yet been fully clarified. However, it was previously demonstrated that TNF-α also enhanced the expression or secretion of IL-6 and IL-8 proteins in another human gingival epithelial cell line (OBA9) (36) and the oral epithelial cell line, TR146 (37). These findings indicate that Ca9-22 cells are also available to pharmacologically investigate the anti-inflammatory function of AGE, although Ca9-22 cells have been often used as an oral cancer model. Thus, this study first examined whether TNF-α induces the ICAM-1 expression and found that the levels of both ICAM-1 protein and mRNA were augmented by TNF-α, but only slightly by P.g.-derived LPS, indicating that Ca9-22 cells were less sensitive to LPS. It was also found that SIPC inhibited the level of TNF-α-induced ICAM-1 protein possibly through post-translational modification (Fig. 4), while AGE caused little inhibition, suggesting that AGE may contain some substances to block the SIPC action. Furthermore, it was shown that extracellular signal-regulated kinase-1/2 (ERK1/2), P38 MAPK (data not shown) and NF-xB (Fig. 8B) were not involved in the inhibition by SIPC.
Figure 8. Effect of hβD3, S1PC, AGE, SAC and SAMC on the level of phosphorylated NF-κB p65 protein induced by TNF-α. The cells were left untreated (control) or treated with TNF-α (100 ng/ml) in the absence or presence of (A) hβD3 (10, 100 nM) or (B) S1PC (100 µM) for 24 h, (C) AGE (1 mg/ml), (D) SAC or (E) SAMC (each at 100 µM) for 6 h and total protein was extracted. Total NF-κB (NF-κB) and phosphorylated NF-κB (pNF-κB) p65 proteins in the extract were detected by western blot analysis. Each bar in graphs represents the mean ± SD of the band intensity relative to β-actin (n=3-4). *P<0.05, **P<0.01 in comparison to the control and *P<0.05, **P<0.01 in comparison to TNF-α alone (Tukey's multiple comparisons test). S1PC, S-1-propenylcysteine; hβD3, human β-defensin-3; TNF-α, tumor necrosis factor-α; IL-6, interleukin-6; AGE, aged garlic extract; SAC, S-allylcysteine; SAMC, S-allylmercaptocysteine.
The antimicrobial peptide, hβD3, is mainly produced in gingival epithelial cells and is extracellularly secreted into the oral cavity, and exhibits antimicrobial activity against Gram-positive and negative bacteria, fungi and viruses (29,38). hβD3 also plays an important role in the innate immune response via its immunomodulatory effect, suppressing the periodontal disease progression (29,38). It has been reported that the hβD3 expression level in gingival tissues and crevicular fluid is decreased in patients with periodontitis in comparison to healthy controls (29). It has been found that hβD3 reduced the level of ICAM-1 protein and inhibits the phosphorylation of NF-κB induced by TNF-α, which is consistent with a finding obtained in HUVECs (26). Although it was not statistically significant, SIPC at 10 µM exhibited near the maximal inhibition induced by SIPC at 100 µM on the elevated ICAM-1 protein by TNF-α (Fig. 1C). Similarly, hβD3 exhibited the maximal inhibition at 10 nM (Fig. 1G). Since it seemed to be difficult to detect further inhibition when SIPC and hβD3 at the higher concentrations, such as 100 µM and 10 nM, respectively, are used, the authors used SIPC at 10 µM and hβD3 at 1 nM in the synergistic experiment. As shown in Fig. 2, it was found that the inhibitory effect of hβD3 was enhanced by simultaneous treatment with SIPC, suggesting that SIPC synergistically modulates the action of hβD3 through a signaling pathway different from that of this peptide.

IL-6 is transiently produced and secreted by several cell types in response to infections and tissue injuries under acute inflammation (39). It was well known that IL-6 exerts beneficial pleiotropic effects, i.e., the maturation of B cells into antibody-producing cells and development of effector T-cells (39). On the other hand, the continuous production of IL-6 in inflamed tissues results in the progression of chronic inflammation and autoimmunity (39). Thus, the precise control of IL-6 secretion is crucial to prevent severe inflammation, which is often observed under pathophysiological conditions, including rheumatoid arthritis (39).

In human gingival fibroblasts, the pro-inflammatory cytokine, IL-1β, secreted from macrophages promotes the production and secretion of IL-6 (40). IL-6 facilitates inflammation at higher concentrations via the activation of matrix metalloproteases (MMPs) and osteoclasts, resulting in the gingival tissue destruction and alveolar bone resorption (40). In human gingival epithelial OBA9 cells, the IL-6 mRNA level has been shown to be augmented by stimulation with TNF-α through the phosphorylation of ERK1/2 and p38 MAPK (36). This study found that the enhanced IL-6 secretion by Ca9-22 cells stimulated with TNF-α was significantly inhibited by simultaneously treating the cells with AGE, SAC or SAMC, suggesting their involvement in the regulation of IL-6 secretion. However, these substances did not exhibit any inhibitory effect on the IL-6 mRNA expression levels (Fig. 6), suggesting the post-transcriptional mode of control. In addition, it was found that AGE and SAC tended to augment the content of IL-6 protein in cells induced by TNF-α, supporting their post-translational modifications. Based on these findings, it is possible that the reduced IL-6 release by AGE or SAC may result from the partial blockade of extracellular secretion, but not from the decreased production of the cytokine.

In conclusion, this study found that AGE and its sulfur compounds, SIPC, SAC and SAMC, inhibited inflammatory reactions induced by TNF-α in human gingival epithelial cells. Although their mechanisms of action have not been fully clarified, these data suggest that these sulfur compounds in AGE may directly reduce inflammation occurring in human gingival epithelial cells by suppressing the ICAM-1 expression and IL-6 secretion. Moreover, it is possible that the sulfur compounds in AGE ingested through systemic circulation reduced gingival inflammation to improve gingivitis.

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Availability data and materials

All data generated or analyzed during this study are included in this published article or are available from the corresponding author on reasonable request.

Authors' contributions

MO drafted the manuscript and performed the experiments. TN proposed a part of the experimental ideas and concepts, helped interpret the data and revised the manuscript. Both authors have read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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