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RESEARCH ARTICLE

Silver Nanoparticles Affect the Inflammatory Response in a Lung Epithelial Cell Line

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Abstract:

Background and Objectives:

Silver nanoparticles (AgNPs) have a dual effect showing both inflammatory and anti-inflammatory effects; however, the molecular mechanism of their anti-inflammatory effect is not clearly understood. In this study, we investigated the effect of AgNPs on the inflammatory response.

Methods:

We induced an inflammatory response in a lung epithelial cell line using tumor necrosis factor- α (TNF α) as an *in vitro* inflammatory model. Then the effect of AgNPs on the TNF α -induced inflammatory response was observed.

Results:

The mRNA expression of pro-inflammatory cytokines (IL-1 β and IL-18) showed upregulation of IL-1 β by AgNPs alone. However, AgNPs reduced the TNF α -induced upregulation of IL-1 β and IL-18. AgNPs reduced the TNF α -induced NF-KB response, reactive oxygen species (ROS) generation, Nod Like Receptor Family-Pyrin domain containing 3 (NLRP3) gene expression, and caspase-1 activation, indicating that the anti-inflammatory effect of AgNPs was by inhibition of both NF-KB transcriptional and inflammasome pathways. Conversely, AgNPs alone induced the activation of both NF-KB transcriptional and inflammasome pathways, suggesting their involvement in the molecular mechanism of the inflammatory effect of AgNPs.

Conclusion:

Altogether, these findings show that two different pathways are involved in the molecular mechanism of both the dose-dependent inflammatory effect of AgNPs alone and the anti-inflammatory effect of AgNPs against the TNF α -induced inflammatory response. Understanding this mechanism will help to improve the medical applications of AgNPs and suggest their potential as a TNF α inhibitor to treat TNF α -induced inflammatory diseases.

Keywords: Silver nanoparticles, Tumor necrosis factor, Inflammation, Inflammasome, Lung cell line, Cytokines.

Article History

Received: May 1, 2020

Revised: October 16, 2020

Accepted: October 17, 2020

1. INTRODUCTION

Nanotechnology is an emerging field with significant medical potential. Silver nanoparticles (AgNPs) are widely used in many consumer products ranging from textiles to deodorant sprays and many medical applications [1 - 3].

AgNPs have a dual effect; many *in vitro* studies have reported their cytotoxic effect in different cell lines [4, 5], yet their protective anti-microbial and anti-inflammatory effects are widely known [6, 7]. However, the exact mechanism of the anti-inflammatory effect of AgNPs is not fully understood. Clarification of the AgNPs-induced anti-inflammatory mechanism is important to improve their medical applications.

To investigate the anti-inflammatory effect of AgNPs and its mechanism, an appropriate inflammatory *in vitro* model

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should be used. Tumor necrosis factor- α (TNF α) is a pro-inflammatory cytokine that regulates cellular responses in many pathological conditions such as inflammatory diseases and cancer [8]. Inflammation is a major cellular response of TNF α and is mainly induced by the NF-KB activation pathway [9]. Therefore, the TNF α -induced inflammatory response in lung epithelial cells is a suitable model to study the mechanism of the anti-inflammatory effects of AgNPs.

In our previous research, we investigated the effect of AgNPs on TNF α -induced apoptosis and DNA damage responses and concluded that the expression of tumor necrosis factor receptor 1 (TNFR1) on the cell membrane is reduced by AgNPs, resulting in reduced signal transduction of TNF α , including its apoptotic and DNA damage effect [10, 11]. Due to the potential of AgNPs to interfere with the cellular responses of TNF α , we hypothesized that AgNPs might also affect the inflammatory cellular response by their effect on TNFR1. Therefore, both the NF-KB and inflammasome pathways were investigated to help clarify the molecular mechanism of the AgNP effect.

Inflammasomes are multiprotein complexes that activate the inflammatory response [12]. Inflammasome activation results in the activation of caspase-1 and secretion of the pro-inflammatory cytokines IL-1 β and IL-18 [13]. Inflammasome activation is commonly induced by a wide range of nanofibers and nanotubes such as silica, asbestos, and carbon nanotubes and can drive lung inflammation and fibrosis [14, 15]. Recently, the activation of inflammasomes by particulate nanomaterials has been investigated, and it was found that titanium dioxide and silica nanoparticles induce inflammasome activation in different cell lines [16, 17]. Moreover, AgNPs can induce inflammasome activation in human hepatoma cell lines in a size-dependent manner [18]. Therefore, examining the inflammasome pathway can clarify both the inflammatory and anti-inflammatory effects of AgNPs, along with assessing the NF-KB pathway.

In this study, we investigated the dose-dependent effect of AgNPs on the TNF α -induced inflammatory response in lung epithelial cells, along with the molecular mechanism. We used TNF α to induce the cellular inflammatory response, then the response after exposure to AgNPs to clarify the mechanism of the anti-inflammatory effect was investigated. We also investigated the response of the lung epithelial cell line to the direct effect of AgNPs alone without induction of inflammation by TNF α . Our results revealed that both inflammatory and anti-inflammatory effects of AgNPs could be induced by activation or inhibition of both the NF-KB transcriptional and inflammasome pathways. These findings will help improve the use of AgNPs in medical applications.

2. MATERIALS AND METHODS

2.1. Cell Line and Culture

The human pulmonary epithelial cell line NCI-H292 (ATCC CRL-1848TM) was used in this study. To culture the cells, RPMI-1640 medium (L-glutamine with phenol red, Nacalai Tesque, Japan) supplemented with 10% (v/v) heat-inactivated fetal bovine serum (HFBS, Biowest, USA), 100

μ g/ml penicillin, and 10 μ g/ml streptomycin (Nacalai Tesque) was used. Cells were incubated at 37°C in a humidified atmosphere containing 5% CO₂.

2.2. Silver Nanoparticles (AgNPs)

Polyvinylpyrrolidone (PVP)-coated AgNPs were purchased from Sigma-Aldrich (Cat. No. 576832) with a particle size of less than 100 nm. The mean size was approximately 50-90 nm as measured by transmission electron microscopy (TEM; JEM-2000FX, JEOL, Japan). Electronic light scattering (Zeta-potential & Particle Size Analyzer ELSZ-2000, Otsuka Electronics, Japan) measurements showed that the hydrodynamic diameter of AgNPs was 170 \pm 50 nm, the polydispersity index was 0.20, and the zeta potential was -53.2 mV. AgNPs were spherical in shape with slight aggregations in deionized water.

2.3. Tumor Necrosis Factor- α (TNF α)

Recombinant human TNF α (Peprotech, USA) was reconstituted in water to 100 μ g/ml. The required working concentrations were diluted in sterilized culture medium (RPMI-1640) containing a carrier protein.

2.4. Gene Expression Analysis by Real-time (RT) PCR

For mRNA expression analysis of interleukin 1 beta (IL-1 β), interleukin 18 (IL-18), and Nod Like Receptor Family-Pyrin domain containing 3 (NLRP3), cells were seeded at a concentration of 4 \times 10⁵ cells/60 mm cell culture dish. Cells were then exposed to AgNPs (10 and 100 μ g/ml) and/or TNF α (20 ng/ml) for 8 h. After that, cells were detached, collected, and the total cellular RNA was extracted using an RNeasy kit (Qiagen, USA) according to the manufacturer's protocol. An aliquot (1 μ g) of the extracted total RNA was reverse transcribed into cDNA using a RT2 First Strand kit (SABiosciences/Qiagen, USA). The PCR primers for human IL-1 β , IL-18, and NLRP3 were purchased from SABiosciences/Qiagen. The reaction mixture was prepared following the manufacturer's protocol. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as a house-keeping gene to normalize the data. RT-PCR analysis was performed using an ABI PRISM 7000 sequence detection system (Applied Biosystems, Singapore), and the thermocycling conditions were 95°C for 10 min, followed by 40 cycles of 95°C for 15 s and 60°C for 1 min.

2.5. Dual-luciferase Reporter Assay for NF-KB Response Assessment

2.5.1. Plasmids

pGL3-Control vector (E1741, Promega) was used as an empty control reporter plasmid. NF-KB promoter-reporter plasmid (E8491, Promega) was employed to detect the inflammation. Both reporter plasmids contain SV40 promoters and enhancer sequences that result in strong expression of the luciferase encoding gene (luc+) in different types of mammalian cells. Also, pRL-CMV vector (E2261, Promega), which is a Renilla luciferase-encoding control plasmid, was used as an internal control to determine the transfection

efficiency.

2.5.2. Transfection

pGL3 blank control reporter plasmid or NF-KB promoter-reporter plasmid and pRL-CMV internal control plasmid were co-transfected into NCI-H292 cells. Lipofectamine™ LTX reagent with a Plus™ reagent kit (Invitrogen) was used to perform the transfection according to the manufacturer's protocol. Cells were cultured at a density of 2×10^5 /ml in a 24-well plate and incubated overnight at 37°C. Opti-MEM medium (Life Technologies, USA) was used to dilute the Lipofectamine LTX reagent and plasmids, then the Plus reagent was added to the diluted plasmids. Diluted plasmids with Plus reagent were added to the diluted Lipofectamine LTX reagent at a 1:1 ratio and incubated for 5 min at room temperature. Finally, the plasmid-lipid complexes were added to the cells and incubated at 37°C for at least 24 h before exposure to TNF α and AgNPs.

2.5.3. Assessment of Luciferase Activity

After exposure of transfected cells to TNF α (20 ng/ml) and AgNPs (10 or 100 μ g/ml) separately and together for 12 h, the luciferase activities were assessed using a Dual-Luciferase Reporter Assay System (E1910, Promega) according to the manufacturer's protocol. Cells were lysed using 1 \times passive lysis buffer and gentle shaking for 10 min, then cell lysates were transferred to tubes containing luciferase assay reagent II (LAR II). Firefly luciferase (F) signals were measured, then Stop & Glo reagent was added to the tubes, and the Renilla luciferase (R) signals were measured as well. The firefly and Renilla luciferase signals were recorded using a luminometer (Lumat LB9507, BERTHOLD TECHNOLOGIES, Germany) according to the instrument manual. The changes in luciferase activities were calculated using the following equation:

$$\Delta \text{Fold activity} = \left(\frac{F}{R} \right)_{\text{sample}} \div \left(\frac{F}{R} \right)_{\text{control}}$$

2.6. DCF Assay for Oxidative Stress Determination

To quantify intracellular reactive oxygen species (ROS), a ROS assay kit (OxiSelect™, Cell Biolabs, Inc., USA) was used. This assay is based on the cell permeable fluorogenic probe 2',7'-dichlorodihydrofluorescein diacetate (DCFH-DA), diffuses into cells is deacetylated by intracellular esterases to the nonfluorescent 2',7'-dichlorodihydrofluorescein (DCFH). In the presence of ROS, DCFH is rapidly oxidized to the highly fluorescent 2',7'-dichlorodihydrofluorescein (DCF). The fluorescence intensity is proportional to the intracellular ROS levels.

Cells were cultured at a density of 1×10^4 cells/well in a black 96-well plate and incubated overnight. Subsequently, media were removed, and cells were washed twice gently with DPBS (14190, GIBCO, Invitrogen, USA) and incubated with DCFH-DA/media solution for 30 min in the dark at 37°C. Then, after removing the solution and washing the cells with DPBS, the DCFH-DA-loaded cells were exposed to TNF α (20 ng/ml) and AgNPs (10 or 100 μ g/ml) separately and together for 12 h. Parallel sets of wells containing DCFH-DA-loaded

cells without any further exposure were used as a negative control. The fluorescence of DCF was measured at an excitation/emission wavelength of 480 nm/530 nm using a fluorometric plate reader (Microplate Fluorometer, Twinkle LB 970, BERTHOLD TECHNOLOGIES). The amounts of produced DCF were calculated based on a DCF standard curve.

2.7. Caspase-1 Activity Assay

To monitor the activity of caspase-1, a caspase-1/ICE fluorometric assay kit (Biovision, USA) was used. This assay is based on the detection of the cleavage of the substrate YVAD-AFC, which emits blue light; after its cleavage by caspase-1, free AFC emits a yellow green fluorescence. According to the manufacturer's protocol, cells were exposed to TNF α (20 ng/ml) and AgNPs (10 or 100 μ g/ml) separately and together. After 12 h of exposure, cells were pelleted, resuspended in cell lysis buffer, and incubated on ice for 10 min. Then, 2 \times reaction buffer and YVAD-AFC substrate were added to the cell lysate and incubated at 37°C for 1-2 h. Fluorescence was then measured using a fluorometric plate reader (Microplate Fluorometer, Twinkle LB 970, BERTHOLD TECHNOLOGIES) at an excitation/emission wavelength of 400 nm/505 nm. The fold increase in caspase-1 activity was determined by comparing the activity of the enzyme in the samples with the activity of the untreated cells.

2.8. Immunostaining and Confocal Laser Scanning Microscopy

To analyze the expression and localization of purinergic 2X7 receptor (P2X7R), NCI-H292 cells were seeded in a CELLview cell culture dish (Greiner Bio-one North America Inc., USA) at a density of 1.5×10^4 cells/compartiment and incubated for 24 h. The cells were exposed to TNF α (20 ng/ml) only, or together with AgNPs (10 or 100 μ g/ml). After 12 h of exposure, the cells were washed with 1 \times PBS fixed with 4% formaldehyde solution in PBS (Wako) at room temperature, permeabilized with 0.1% Triton X-100, and then blocked with 10% normal goat serum in PBS for 1 h. The cells were then incubated overnight at 4°C with rabbit polyclonal anti-P2X7 receptor antibody (Alomone, Israel) followed by incubation with labeled goat anti-rabbit IgG H&L (Alexa Fluor 594) (Ab150080, Abcam) for 1 h at room temperature. Nuclear DNA was stained with DAPI (4', 6-diamidino-2-phenylindole) (Dojindo, Japan) for 5 min at room temperature. The stained samples were mounted by antifade solution (Fluoromount, Diagnostic Biosystems, Canada). Microscopic observations and images were acquired using a confocal laser-scanning microscope (Leica TCS SP5, Wetzlar, Germany) with a 63 \times 1.4 Plan-apochromat oil immersion lens.

2.9. Statistical Analysis

Statistical analysis was performed using Student's t test. Differences and significances between means of different groups were determined using one-way ANOVA with Duncan's multiple comparison tests. *P < 0.05, **P < 0.01, ***P < 0.001 were considered statistically different as shown in figures. Data are presented as means \pm standard deviation (SD) with at least three independent replicates (n \geq 3).

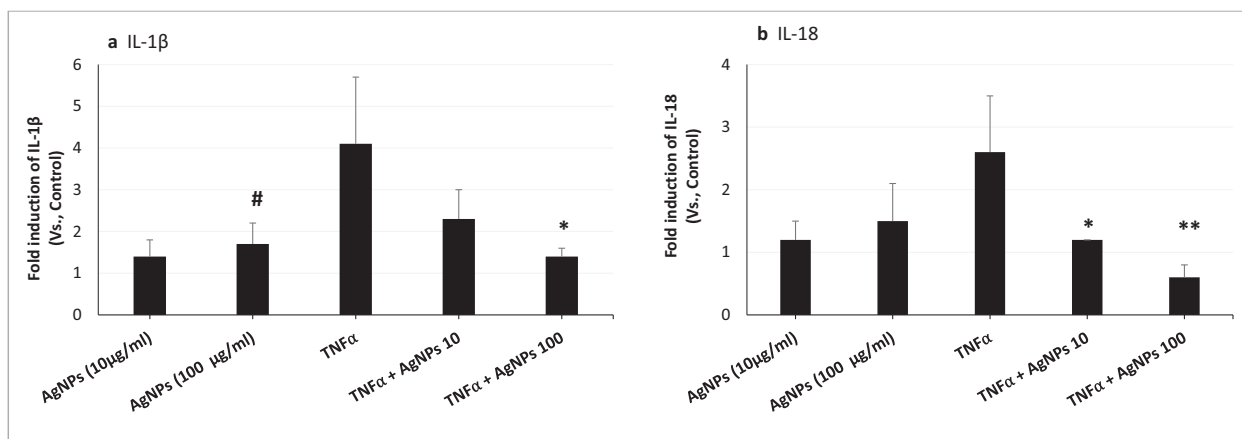


Fig. (1). Suppression of TNF α -induced a) IL-1 β and b) IL-18 mRNA expression levels by AgNPs in NCI-H292 cells. Cells were exposed to 20 ng/ml of TNF α and 10 and 100 μ g/ml of AgNPs separately and together for 8 h. mRNAs expressions were analyzed using real-time (RT) PCR. Control, which absence of both TNF α and AgNPs shows 1.0. The results are figured as means \pm SD, $n \geq 3$, for each group; *, $0.01 < P < 0.05$, representing a significant difference compared to the corresponding TNF α -exposed group. #, $0.01 < P < 0.05$, ##, $P < 0.01$ representing significant differences compared to the control group.

3. RESULTS

3.1. Effect of AgNPs on TNF α -induced pro-inflammatory Cytokines Secretion

IL-1 β and IL-18 are pro-inflammatory cytokines and belong to the IL-1 superfamily. Both are important mediators of the inflammatory cellular response, including fever generation, secretion of other cytokines and chemokines, and attraction of immune cells [19, 20]. TNF α can induce the secretion of IL-1 β [21] and IL-18 [22] in different cell lines. Because of the known anti-inflammatory effect of AgNPs, we investigated the effect of AgNPs on TNF α -induced IL-1 β and IL-18 secretion in NCI-H292 cells.

We conducted a gene expression assay to analyze the IL-1 β and IL-18 mRNA expression levels. As shown in Figs. (1A and B), TNF α -induced IL-1 β and IL-18 gene expressions were significantly reduced by the exposure of cells to both TNF α and AgNPs (100 μ g/ml), from 4.1- to 1.4-, and from 2.6- to 0.6-fold, respectively. Also, the reduction in the expression of both genes was dose-dependent; increasing the concentration of AgNPs from 10 to 100 μ g/ml resulted in greater reduction in TNF α -induced IL-1 β and IL-18 secretions. These results demonstrated the suppression of TNF α -induced IL-1 β and IL-18 mRNA expression levels by AgNPs, suggesting they have an anti-inflammatory effect.

On the other hand, AgNPs alone induced IL-1 β mRNA expression in NCI-H292 cells as shown in Fig. (1A). This result indicated the induction of pro-inflammatory cytokine expression, showing that AgNPs have an inflammatory effect.

3.2. Effect of AgNPs on the TNF α -induced NF-KB Response

Secretion of pro-inflammatory cytokines is regulated at the transcriptional and post-transcriptional levels. The signal transduction of TNF α and its receptor (TNFR1) is the first

signal that might be involved in the secretion of active pro-inflammatory cytokines (IL-1 β and IL-18) *via* the NF-KB transcription pathway [23]. Therefore, we transfected an NF-KB promoter-reporter plasmid to detect the inflammatory response and to check whether the NF-KB pathway is involved in the effect of AgNPs on the TNF α -induced IL-1 β and IL-18 secretion.

As shown in Fig. (2), the NF-KB response in cells exposed to AgNPs alone was dose-dependent. However, when cells were exposed to both TNF α and AgNPs (100 μ g/ml), the fold induction of the NF-KB response was significantly reduced from 2.4 to 1 compared to the TNF α -exposed cells. These results revealed suppression of the TNF α -induced NF-KB response by AgNPs, suggesting that the NF-KB pathway is involved in the anti-inflammatory effect of AgNPs against the TNF α -induced inflammatory response. Also, by activation of the NF-KB pathway, AgNPs can induce their inflammatory effect in a dose-dependent manner.

3.3. Effect of AgNPs on the TNF α -activated Inflammasome Pathway

The second signal involved in the secretion of active pro-inflammatory cytokines (IL-1 β and IL-18) is the activation of caspase-1. Caspase-1 cleaves pro-IL-1 β and pro-IL-18 into their biologically active forms, IL-1 β and IL-18 [24]. Caspase-1 activation can be initiated by NLRP3 inflammasome activation [25], and the generation of ROS can induce NLRP3 inflammasome activation [26]. TNF α has been reported to mediate this inflammasome pathway in a human neuroblastoma cell line [21]. Therefore, we investigated the inflammasome pathway to assess its role in the effect of AgNPs on the TNF α -induced inflammatory response.

ROS generation, NLRP3 mRNA expression, and caspase-1 activity were increased in a dose-dependent manner in cells exposed to AgNPs (10 and 100 μ g/ml) alone as shown in Figs. (3, 4, and 5) respectively. On the other hand, the cells exposed

to both TNF α and AgNPs together showed a dose-dependent significant decrease in both TNF α -induced ROS generation and NLRP3 mRNA expression as shown in Figs. (3 and 4). However, the TNF α -induced caspase-1 activity was significantly reduced only by the higher concentration (100 μ g/ml) of AgNPs as shown in Fig. (5). These results suggested

that the inflammasome pathway is involved in the molecular mechanism of the anti-inflammatory effect of AgNPs against the TNF α -induced inflammatory response. Moreover, AgNPs induced inflammasome activation in a dose-dependent manner, suggesting the dual activities.

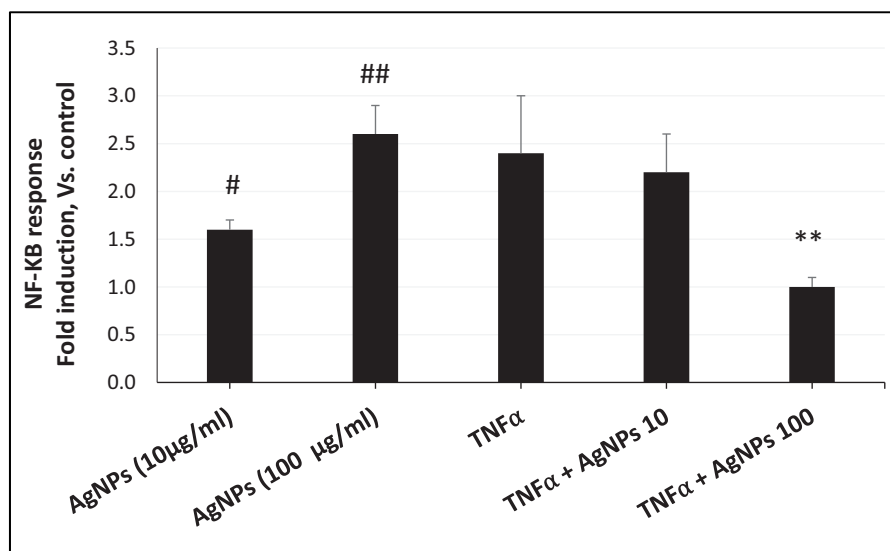


Fig. (2). Suppression of the TNF α -induced NF-KB response by AgNPs in NCI-H292 cells. Cells were transfected with NF-KB promoter-reporter plasmid, and then the transfected cells were exposed to 20 ng/ml of TNF α and AgNPs 10 and 100 μ g/ml of AgNPs separately and together for 8 h. The results are figured as means \pm SD, $n \geq 3$, for each group; ## and ** $P < 0.01$. #, significant difference compared to the control group. **, significant difference compared to the TNF α -exposed group.

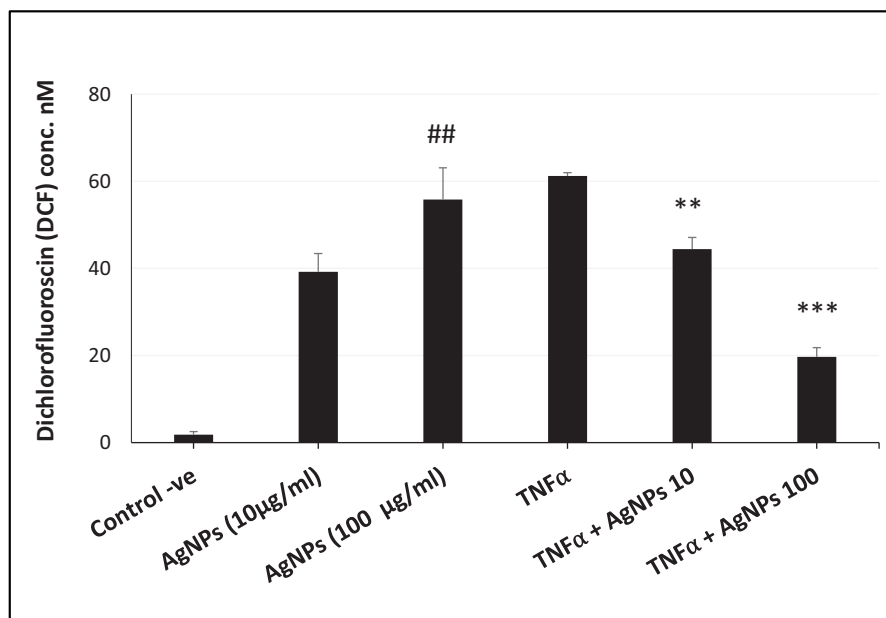


Fig. (3). Suppression of TNF α -induced ROS generation by AgNPs in NCI-H292 cells. Cells were exposed to 20 ng/ml of TNF α and 10 and 100 μ g/ml of AgNPs separately and together for 8 h. The produced DCF amount reflects the generation levels of ROS. The results are figured as means \pm SD, $n \geq 3$, for each group; ## and **, $P < 0.01$; ***, $P < 0.001$. #, significant difference compared to the control group. **, significant difference compared to the TNF α -exposed group.

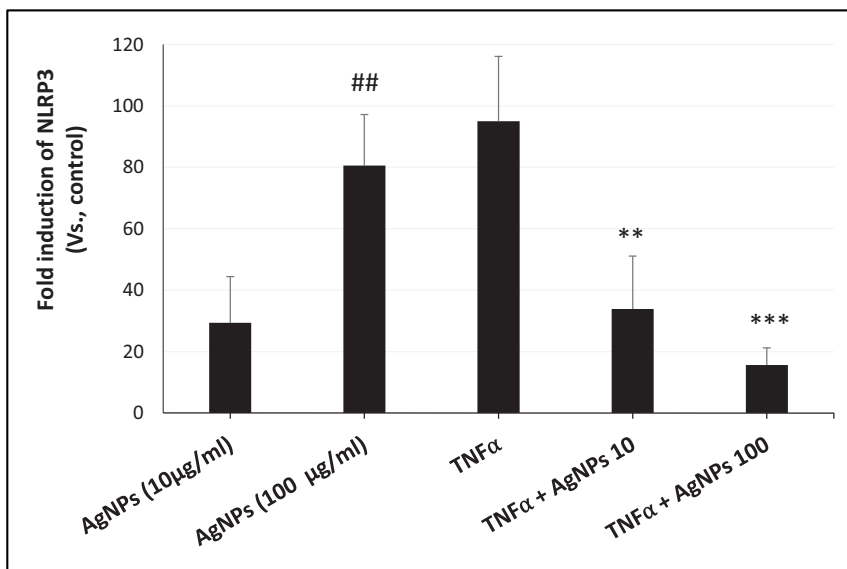


Fig. (4). Suppression of TNF α -induced NLRP3 mRNA by AgNPs in NCI-H292 cells. Cells were exposed to 20 ng/ml of TNF α and 10 and 100 μ g/ml of AgNPs separately and together for 8 h. Expressions of mRNAs were measured using real-time (RT) PCR. The results are figured as means \pm SD, $n \geq 3$, for each group; ## and **, $P < 0.01$. #, significant difference compared to the control group. **, significant difference compared to the TNF α -exposed group.

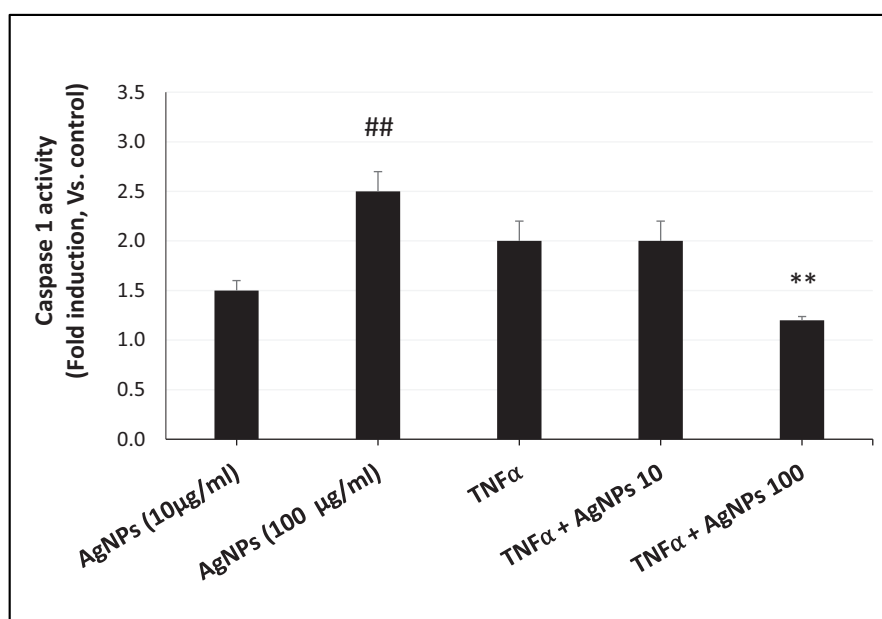


Fig. (5). Suppression of TNF α -induced caspase-1 activity by AgNPs in NCI-H292 cells. Cells were exposed to 20 ng/ml of TNF α and 10 and 100 μ g/ml of AgNPs separately and together for 8 h. Caspase-1 activity in cell lysates was then measured using a caspase-1 fluorometric assay. The results are figured as means \pm SD, $n \geq 3$, for each group; ## and **, $P < 0.01$. #, significant difference compared to the control group. **, significant difference compared to the TNF α -exposed group.

3.4. Expression of Purinergic 2X7 Receptor (P2X7R)

P2X7R is a ligand-gated ion channel that is activated by extracellular ATP [27]. The activation of P2X7R plays an important role in the inflammatory response, including ROS generation [28, 29], NLRP3 inflammasome activation [30, 31], and release of pro-inflammatory cytokines such as IL-1 β and IL-18 [32]. Moreover, P2X7R has been implicated in lung inflammation induced by TiO₂ and SiO₂ nanoparticles *via* inflammasome activation and IL-1 secretion in lung epithelial and macrophage cell lines [33]. Due to the impact of P2X7R,

we hypothesized that the interference of AgNPs with the inflammatory response of TNF α might be affected by P2X7R activation. Therefore, we quantified the expression of P2X7R using immunofluorescence assays and confocal microscopy. As shown in Fig. (6A-D), P2X7 receptors were expressed in NCI-H292 cells after exposure to TNF α alone and to TNF α + AgNPs (10 and 100 μ g/ml). Cells exposed to TNF α + AgNPs (100 μ g/ml) showed a significant decrease in the expression of P2X7 receptors compared to cells exposed to TNF α alone and cells exposed to TNF α + AgNPs (10 μ g/ml).

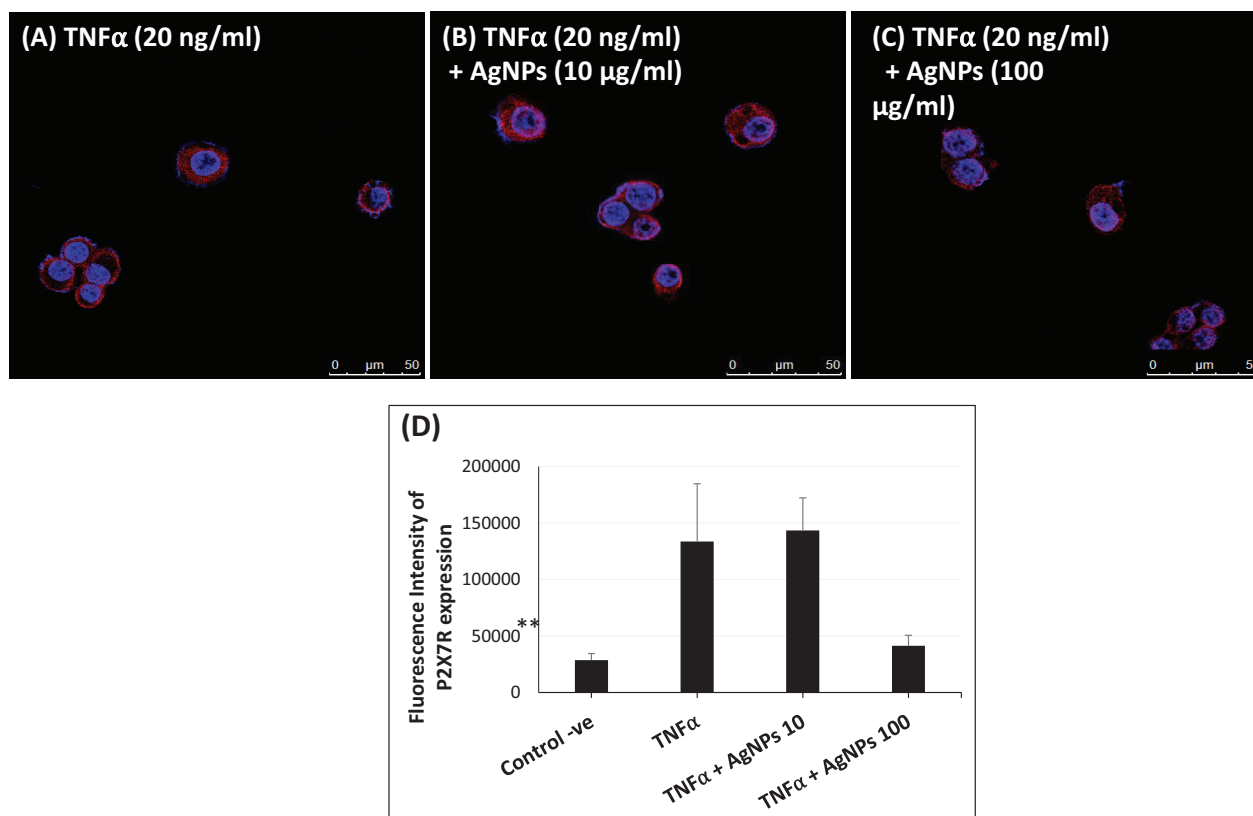


Fig. (6). Expression and localization of Purinergic 2X7 receptor (P2X7R) in NCI-H292 cells using confocal microscopy. Blue indicates the nucleus and red indicates the receptors (P2X7R). (A) NCI-H292 cells were exposed to TNF α (20 ng/ml), showing high expression of P2X7R. (B) NCI-H292 cells were exposed to both TNF α (20 ng/ml) and AgNPs (10 μ g/ml), showing high expression of P2X7R. (C) NCI-H292 cells were exposed to both TNF α (20 ng/ml) and AgNPs (100 μ g/ml), showing lower expression of P2X7R. Exposure was 12 h for all. Scale bar is 50 μ m for all panels. (D) Quantification of P2X7R was determined by the fluorescence intensity. The results are shown as means \pm SD, $n \geq 3$ for each group. **, $P < 0.01$, representing a significant difference compared to the TNF α -exposed group.

These data suggest that AgNPs, together with TNF α , reduced the expression level of P2X7R on the cell membrane, and this resulted in inhibition of P2X7R signal transduction, resulting in reduced TNF α -induced pro-inflammatory cytokine release. This suggests that inhibition of the TNF α -induced P2X7R/NLRP3 inflammasome pathway by AgNPs is a possible molecular mechanism of their anti-inflammatory effect.

4. DISCUSSION

AgNPs are widely known for their antimicrobial effect, which had led to their use in many medical applications such as deodorant sprays, wound dressings, and catheter coatings [34, 35]. However, AgNPs can induce many different cellular responses, such as cell proliferation, inflammation, genotoxicity, and cell death [36 - 38]. Different cellular responses of AgNPs are affected by the different characteristics of AgNPs themselves, such as concentration, size, coating, and surface charge. For example, AgNPs with a size of 3-5 nm can induce inflammation in a neural cell line [39], while synthesized AgNPs with size range of 40-100 nm enhance the anti-inflammatory effect in human peripheral blood mononuclear cells [40], showing the double-edged effect of AgNPs. Therefore, AgNPs should be characterized well to specify the induced cellular response. In this research, we conducted a dose-dependent study of negatively charged, PVP-

coated AgNPs with a mean size of 50-90 nm. PVP-coated AgNPs are recommended for in vitro studies because of their higher stability with less aggregation and dissolution changes as compared to other coatings, such as citrate and polyethylene glycol (PEG) [41].

Because of the widespread medical applications of AgNPs, we hypothesized that the response of healthy cells would be different than that of cells in diseased conditions such as bacterial infection, sepsis, cancer, or inflammatory disease. Therefore, we designed our study to assess the effect of AgNPs on lung epithelial cells in both healthy and diseased cell conditions. For this purpose, we used TNF α as an inflammatory-inducing agent and lung epithelial cells to create an in vitro inflammatory disease model. Then we investigated the effect of AgNPs on lung epithelial cells with and without TNF α . TNF α is known to have autocrine activity, suggesting endogenous TNF α also could induce pro-inflammatory cytokines [42].

The release of pro-inflammatory cytokines (IL-1 β and IL-18) as a mediator of the inflammatory response was analyzed in this study. The results of both IL-1 β and IL-18 mRNA expressions showed a dose-dependent reduction in the TNF α -induced mRNA expressions, revealing an anti-inflammatory effect for AgNPs. However, AgNPs (10 and 100 μ g/ml) induced an inflammatory response in non-treated cells

by inducing pro-inflammatory cytokine expression as shown in Fig. (1A, B). In order to understand the molecular mechanism of the change in TNF α -induced pro-inflammatory cytokines released by AgNPs, we assumed that the NF-KB transcriptional pathway would play a role in this effect. The NF-KB transcriptional pathway is a known signal transducer of TNF α , and is stimulated by the binding of TNF α to its receptor TNFR1. Our results revealed the suppression of the TNF α -induced NF-KB response by AgNPs, especially the higher concentration (100 μ g/ml), as shown in Fig. (2). Moreover, AgNPs alone induced the NF-KB response in a dose-dependent manner, confirming the involvement of the NF-KB transcriptional pathway in both inflammatory and anti-inflammatory effects of AgNPs. We could relate this change in the NF-KB transcriptional pathway to the change in the expression of TNFR1 on the cell membrane. Specifically, we have concluded that the AgNPs can reduce the expression of TNFR1, resulting in a reduction in TNF α -induced signal transduction in a lung epithelial cell line [10]. So, this AgNP effect induced a reduction in the activation of the TNFR1/NF-KB transcriptional pathway, resulting in a reduction in the TNF α -induced inflammatory response as shown in Fig. (7).

The inflammasome activation pathway in this study, in addition to the NF-KB transcriptional pathway, to assess its involvement in both the inflammatory and anti-inflammatory effects of AgNPs was investigated. Inflammasomes are complexes consisting of NLRP3, apoptosis-associated speck-like protein (ASC), and pro-caspase 1 [43]. Many stimuli can

activate inflammasomes such as pathogen-associated molecular patterns (PAMPs) [44], environmental toxins [45], and lipopolysaccharides (LPS) [46]. Nanoparticles can also induce inflammasome activation [47]. The generation of mitochondrial ROS by different stimuli leads to NLRP3 inflammasome activation [48], resulting in caspase-1 activation. Caspase-1 then converts pro-interleukins (pro-IL-1 β and pro-IL-18) to active forms (IL-1 β and IL-18) and initiates the inflammatory response. In this study, we investigated the ROS-mediated inflammasome pathway. Our results showed that ROS generation, NLRP3 mRNA expression, and caspase-1 activity were increased in a dose-dependent manner in non-treated cells exposed only to AgNPs (10 and 100 μ g/ml) as shown in Figs. (3, 4, and 5) respectively. This result shows how AgNPs could activate the inflammasome pathway by ROS generation and then NLRP3 mRNA expression and caspase-1 activation, which lead to pro-inflammatory cytokine release, explaining the inflammatory effect of AgNPs by themselves. On the other hand, the inflammatory cell model (TNF α -induced inflammatory response) showed significant dose-dependent decreases in TNF α -induced ROS generation, NLRP3 mRNA expression, and caspase-1 activity after exposure to AgNPs (10 and 100 μ g/ml), revealing their anti-inflammatory effect against the TNF α -induced inflammatory response. These findings indicate that AgNPs can activate and inhibit the ROS-mediated inflammasome pathway, inducing inflammatory and anti-inflammatory effects respectively depending on the condition of cells (with or without TNF α) as shown in Fig. (7).

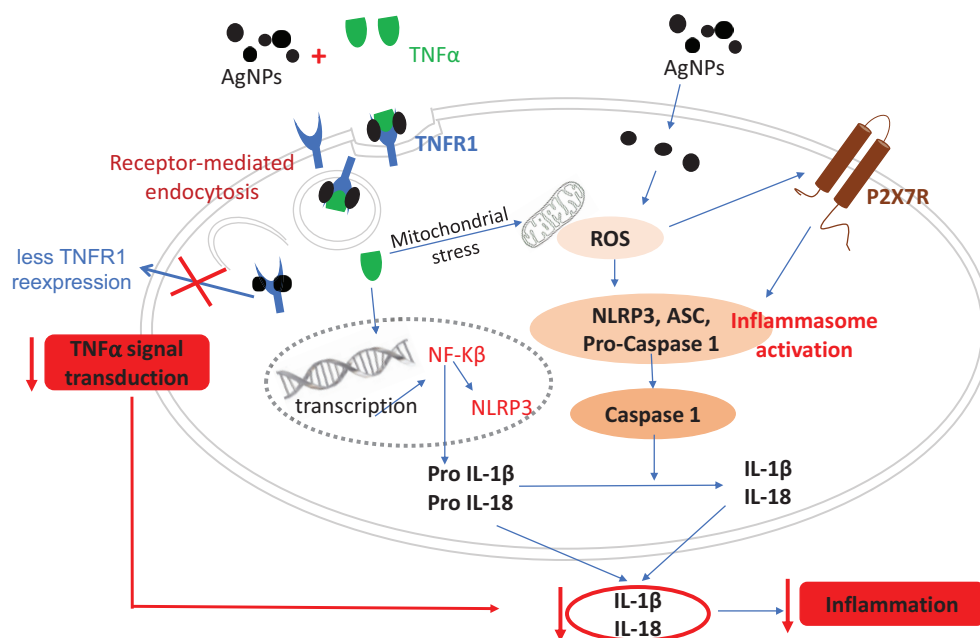


Fig. (7). Molecular mechanism explaining two different possible pathways (TNFR1 and P2X7R) that can reduce the TNF α -induced inflammatory response by AgNPs. Arrows show signal transductions. The red arrows show the reduction of the TNF α -induced inflammatory response. In case of exposure to both TNF α and AgNPs, the nanoparticles non-specifically bind to TNFR1, and TNF α binds specifically with the same receptor, forming a TNFR1-TNF α -AgNPs complex which then enters the cells by receptor-mediated endocytosis. TNF α is then released from the receptor and induces the inflammatory response. The receptors might still bind to AgNPs, thus disturbing the receptor's shape, molecular weight, and characteristics, leading to disturbance of its normal pathway of being recycled to the cell membrane, resulting in less TNFR1 on the cell membrane and more inside the cells.

Since Purinergic 2X7 receptor (P2X7R) plays a key role in the inflammation process, including NLRP3 inflammasome activation [49, 50], we hypothesized that the mechanism of the AgNPs-induced anti-inflammatory effect might be mediated through the expression of P2X7R. P2X7R is activated by extracellular adenosine triphosphate (ATP) and then induces the passage of Ca²⁺, Na⁺, and K⁺ ions across the plasma membrane [51]. Activation of P2X7R induces the generation of ROS [52] and subsequently will induce inflammasome activation and the inflammatory response. The expression of P2X7R was quantified in this study using immunofluorescence assays and confocal microscopy as shown in Fig. (6). The high concentration of AgNPs (100 µg/ml) induced a significant decrease in the TNF α -induced expression of P2X7R compared to the low concentration of AgNPs (10 µg/ml). This result indicates that AgNPs can reduce extracellular ATP production, which then reduces the P2X7R activation. So, we can relate the anti-inflammatory effect of AgNPs to reduced P2X7R expression, which reduces ROS generation, NLRP3 inflammasome activation, caspase-1 activation, and pro-inflammatory cytokines (IL-1 β and IL-18) release; all of these events would finally reduce the inflammatory response as shown in Fig. (7).

Finally, our findings provide details about the possible molecular mechanism of both the inflammatory and anti-inflammatory effects of AgNPs. This molecular mechanism occurs by two different pathways. The first pathway is the NF-KB transcriptional pathway, which is regulated by TNF α /TNFR1 signal transduction, which is inhibited by AgNPs resulting in a reduction in pro-inflammatory cytokine (IL-1 β and IL-18) release, eventually reducing the TNF α -induced inflammation in the *in vitro* disease model. However, this pathway can be activated by AgNPs alone, producing an inflammatory effect in non-treated cells. The second pathway is the Inflammasome pathway, which is regulated by both ROS generation and the expression of P2X7R. In the *in vitro* disease model, the inflammasome pathway would be inhibited by AgNPs *via* a reduction in ROS generation or P2X7R expression, and then a reduction in subsequent events including the NLRP3 inflammasome activation, caspase-1 activation, and pro-inflammatory cytokines (IL-1 β and IL-18) release, eventually reducing TNF α -induced inflammation. However, in non-treated cells, AgNPs could activate the inflammasome pathway, producing an inflammatory effect. In this study, we elucidated some of the details of the anti-inflammatory effect of AgNPs to help to improve their medical applications. Moreover, our findings suggest that AgNPs can be used as a TNF α inhibitor to treat TNF α -induced diseases such as Alzheimer's disease. To further explore the therapeutic potential of AgNPs, *in vivo* studies are needed to complement *in vitro* results of this study.

CONCLUSION

In this study, we found a dose-dependent anti-inflammatory effect of AgNPs against the TNF α -induced inflammatory response and proposed a molecular mechanism for this effect. This mechanism works by inhibition of either the NF-KB transcriptional pathway or the inflammasome pathway, resulting in reduced pro-inflammatory cytokines

(IL-1 β and IL-18) release, eventually reducing TNF α -induced inflammation. Moreover, AgNPs alone induced the inflammatory response in healthy cells by activation of both pathways, revealing that AgNPs are a double-edged sword depending on their concentration and cellular conditions.

ETHICS APPROVAL AND CONSENT TO PARTICIPATE

Not applicable.

HUMAN AND ANIMAL RIGHTS

No human or animals were used in this research.

CONSENT FOR PUBLICATION

Not applicable.

AVAILABILITY OF DATA AND MATERIALS

Not applicable.

FUNDING

None.

CONFLICT OF INTEREST

The authors declare no conflict of interest, financial or otherwise.

ACKNOWLEDGEMENTS

We thank Ms. Hiromi Morita and Mr. Shinya Hattori of the NIMS Molecule & Material Synthesis Platform for assistance in obtaining and analyzing the flow cytometry results.

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