

Research Paper

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Caffeine Inhibits NLRP3 Inflammasome Activation by Suppressing MAPK/NF-KB and A2aR Signaling in LPS-Induced THP-1 Macrophages

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Abstract

Excessive inflammation induced by various risk factors is associated with the development of bronchopulmonary dysplasia (BPD). Caffeine exerts potent anti-inflammatory effects as a clinical preventive medicine for BPD. Recently, NLRP3 inflammasome activation has been demonstrated to be essential for the pathogenesis of BPD. In the present study, we aimed to investigate the effects of caffeine on NLRP3 inflammasome activation in LPS-induced THP-1 macrophages and to explore the underlying the detailed mechanism. We found that caffeine significantly reduced NLRP3 expression, ASC speck formation, and caspase 1 cleavage and therefore decreased IL-1β and IL-18 secretion in THP-1 macrophages. Caffeine also markedly decreased the phosphorylation levels of MAPK and NF-KB pathway members, further suppressing the translocation of NF-KB in THP-1 macrophages. Moreover, silencing of the caffeine-antagonized adenosine A2a receptor (A2aR) significantly decreased cleaved caspase 1 expression in THP-1 macrophages by reducing ROS production. Given these findings, we conclude that caffeine inhibits NLRP3 inflammasome activation by suppressing MAPK/NF-KB signaling and A2aR-associated ROS production in LPS-induced THP-1 macrophages.

Key words: Caffeine, NLRP3 inflammasome, MAPK, NF-KB, A2aR, THP-1 macrophages

Introduction

Bronchopulmonary dysplasia (BPD), the initial definition of chronic pulmonary insufficiency of prematurity [1], is a severe pulmonary complication with multifactorial pathogenesis [2]. Emerging evidence has indicated that multiple factors, including intrauterine chorioamnionitis, oxygen therapy, mechanical ventilation and nosocomial infection, lead to continuous injury in premature lungs that is mediated by inflammatory mechanisms [3-5]. Excessive secretion of proinflammatory cytokines IL-1β predominantly including by alveolar macrophages is thought to be responsible for the decreased alveolarization observed in BPD patients [6, 7]. Therefore, to be most effective, intervention to prevent or ameliorate BPD should focus on suppressing inflammation to protect the injured lungs

[8].

In recent years, caffeine, which is extensively used for the treatment of apnea in preterm infants, has gained attention for its preventive effect on BPD [9]. Initially, researchers attributed the efficacy of caffeine in shortening ventilation time to its stimulation of the respiratory center, which further diminishes inflammation [10]. In addition, it has been reported that caffeine also exerts diuretic effects [11], provides neuronal injury protection [12] and induces surfactant protein B (SP-B) transcription [13]. Recently, growing evidence has suggested that caffeine can inhibit the secretion of proinflammatory cytokines, especially IL-1 β , to reduce inflammation in hyperoxia and lipopolysaccharide (LPS)-induced rodent models [14-16]. However, the molecular mechanisms

underlying the anti-inflammatory function of caffeine remain poorly understood.

A growing number of studies have revealed that activation of the nucleotide-binding domain and leucine-rich repeat protein 3 (NLRP3) inflammasome is related to the pathogenesis of acute lung injury [17, 18]. When macrophages sense external danger signals such as LPS, NLRP3 interacts with apoptosis associated speck like protein containing a CARD (ASC) and the precursor of caspase 1 to form the NLRP3 inflammasome [19, 20]. The activated NLRP3 inflammasome functions as a supramolecular platform for the caspase-1-dependent maturation and secretion of the proinflammatory cytokines IL-1ß and IL-18 in macrophages [21, 22]. A recent study has shown that NLRP3-deficient mice exhibit reduced inflammation and improved lung alveolarization function under hyperoxia [23].

In addition, the hyperactivation of the NLRP3 inflammasome and elevated IL-1 β levels in preterm infants with BPD also demonstrate the major role of NLRP3 in this condition [24]. However, it remains unclear whether caffeine inhibits NLRP3 inflammasome activation in macrophages under LPS stimulation. In this study, we aimed to assess the effects of caffeine on NLRP3 inflammasome activation and to explore the potential underlying mechanisms.

Materials and Methods

Cell culture and treatment

A human monocyte leukemia cell line (THP-1) was obtained from the Cell Bank of the Type Culture Collection of the Chinese Academy of Sciences. The cells were cultured at a density of 5×10⁵ cells/mL in RPMI 1640 medium supplemented with 10% FBS and 1% penicillin/streptomycin solution (Biological Industries, Israel) at 37 °C in a 5% CO2 incubator. In all experiments, THP-1 monocytes were cultured in 6-well plates treated with 100 nM phorbol 12-myristate 13-acetate (PMA) for 24 h to transform into adherent macrophages.

Caffeine, ATP, LPS, PMA and Poly-L-lysine solution were purchased from Sigma-Aldrich (St. Louis, USA). Nigericin was purchased from InvivoGen (San Diego, USA).

Cytotoxicity assay

The cytotoxicity of caffeine toward THP-1 macrophages was evaluated by Annexin V-FITC and PI staining and flow cytometry (BioLegend, USA) according to the manufacturer's instructions. A total of 1×10^5 THP-1 macrophages were seeded in a 12-well plate and incubated with caffeine (100-800 μ M) for 6 h.

ELISA for cytokine measurements

THP-1 macrophages were seeded in a 6-well plate at a density of 5×10^5 cells/mL. The cells were pretreated with caffeine (100-800 µM) or vehicle for 1 h and then treated with LPS (1 µg/ml) for 3 h and ATP (5 mM) for 30 min. The cell culture supernatants were centrifuged at 1500 rpm for 15 min at 4 °C, and the pellets were discarded. Secreted IL-1 β and IL-18 were measured by using commercially available ELISA kits (Abclonal, China) following the instructions provided by the manufacturers.

Quantitative real-time PCR (qRT-PCR)

Total RNA was extracted from cultured cells with TRIzol reagent (Life Technologies, USA) and reverse transcribed into cDNA using a Takara Prime Script RT Reagent Kit (Takara, Japan). qRT-PCR was performed using a TB Green PCR Kit (Takara, Japan) with a Roche Light Cycler 96 Real-Time PCR System (Roche, Switzerland). The PCR thermal cycling parameters were as follows: 95 °C for 600 s; 45 cycles of 95 °C for 10 s, 60 °C for 15 s and 72 °C for 15 s; and melting curve formation from 65 to 97 °C to ensure amplification of a single product. The GAPDH gene was used as an endogenous control. The relative gene expression was calculated using the 2– $\Delta\Delta$ CT method. The sequences of the primers used for qRT-PCR analysis are given in the table.

Table	1.	Primers	used	in	this	study
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Gene	GenBank accession no.	Primer sequence (5'-3')	Product size (bp)
GAPDH	NM_001115114.1	F-GGACTCATGACCACAGTCCA R-TCAGCTCAGGGATGACCTTG	157
NLRP3	NM_001079821.2	F-AACATGCCCAAGGAGGAAGA R-GGCTGTTCACCAATCCATGA	115
Caspase 1	NM_001257118.3	F-GCACACGTCTTGCTCTCATT R-GCCTCCAGCTCTGTAGTCAT	169
ASC	NM_013258.5	F-GTCACAAACGTTGAGTGGCT R-AAGTCCTTGCAGGTCCAGTT	152
IL-1β	NM_000576.2	F-TGCTCAAGTGTCTGAAGCAG R-TGGTGGTCGGAGATTCGTAG	189
IL-18	NM_001562.4	F-TCGGGAAGAGGAAAGGAACC R-TTCTACTGGTTCAGCAGCCA	96
A2aR	NM_000675.6	F-AGGCAGCAAGAACCTTTCAA R-CTAAGGAGCTCCACGTCTGG	236

siRNA transfection

Gene knockdown experiments were performed serum-free Opti-MEM using HiPerFect in Transfection Reagent (Qiagen, Germany). Cells were transfected for 12 h with 5 nM human A2aR small interfering RNA (siRNA) (Qiagen, Germany) or irrelevant nontargeting negative control (NC) siRNA with Alexa Fluor 488 (Qiagen, Germany) or were treated with transfection reagent only (for mock transfection) according to the manufacturer's protocol. The cells were then further incubated for 18 h under standard growth conditions. At 36 h post transfection, depletion of A2aR was confirmed by qRT-PCR and western blotting.

Western blotting

Western blotting was performed using a standard protocol, as previously described [25, 26]. Briefly, total protein was extracted and separated by 10% SDS-PAGE using Mini-PROTEAN TGX gels (Bio-Rad, USA), and the separated proteins were transferred to PVDF membranes. The membranes were blocked with 5% nonfat milk at room temperature for 2 h and then incubated with specific primary antibodies overnight at 4°C. The primary antibodies, including antibodies against NLRP3, Caspase 1 (p20), P-p65 (Ser536), ERK, P-ERK (Thr202/Tyr204), JNK, P-JNK (Thr183/Tyr185), p38, P-p38 (Thr180/Tyr182), ІкВ-а, Р-ІкВ-а (Ser32), cleaved Caspase 3 and ASC, were purchased from Cell Signaling Technology (Boston, USA). Antibodies against Caspase 1, Caspase 3, p65, A2aR, and GAPDH were purchased from Abclonal (Wuhan, China). Next, the membranes were incubated with appropriate secondary antibodies for 1 h at room temperature, and with the bands were visualized enhanced chemiluminescence (ECL) reagent (Thermo Scientific, USA). All the western blotting images for this study are representative of at least 3 independent biological experiments.

Immunofluorescence staining

For NF- κ B staining, THP-1 macrophages were incubated with 800 μ M caffeine or vehicle prior to treatment with LPS (1 μ g/ml) for 3 h. Then, the cells were analyzed by immunocytochemistry as previously described [27]. The primary antibody against NF- κ B p65 was from Abclonal (Wuhan, China).

The staining method used to explore ASC speck formation has been described in the literature [28]. Cells were incubated with 800 μ M caffeine or vehicle prior to treatment with LPS (1 μ g/ml) for 3 h and nigericin (10 μ M) for 90 min. ASC specks were detected with a primary anti-ASC antibody (Abcam, UK). Images were captured using an Olympus FV1000 confocal microscope (Olympus, Japan) at 100× magnification. The nuclei were stained with DAPI (Beyotime, China). All the immunostaining images for this study are representative of at least 3 independent biological experiments.

Measurement of cellular reactive oxygen species

ROS production in THP-1 macrophages was detected using a Reactive Oxygen Species (ROS) Detection Assay Kit (BioVision, USA) and flow cytometry according to the manufacturer's instructions. The data were analyzed with CytExpert software.

Statistical analysis

The results are presented as the mean \pm SD for at least three independent experiments. Student's t-tests were used for comparisons between two groups. Differences among more than two groups were analyzed using one-way ANOVA. Statistical analysis was performed using Graph Pad Prism 7.0 software, p < 0.05 was considered to indicate statistical significance.

Results

Effects of caffeine on cell survival and apoptosis

To test the cytotoxicity of caffeine toward THP-1 macrophages, we detected cell survival by flow cytometry. The results showed that none of the concentrations of caffeine had significant impacts on cell survival compared with the control (Figure 1A and B). To assess the effect of caffeine on apoptosis, we detected the cleaved Caspase 3/Caspase 3 ratio by Western blot analysis. The data indicated that treatment of THP-1 macrophages with caffeine did not alter the cleaved Caspase 3/Caspase 3 ratio at the protein level compared with the control treatment (Figure 1C and D). Afterward, we used 100, 200, 400, and 800 µM caffeine in the following experiments.

Caffeine suppresses IL-1β and IL-18 secretion in LPS/ATP-induced THP-1 macrophages

To test whether caffeine affects inflammasomespecific cytokine expression, we measured the secretion levels of IL-1 β and IL-18 in the cell culture LPS/ATP-induced supernatants of THP-1 macrophages. The data showed that induction of THP-1 macrophages with LPS and ATP significantly increased the secretion of IL-1 β and IL-18 (Figure 2A) and B). Pretreatment with caffeine suppressed the release of both proinflammatory cytokines in a dose-dependent manner (Figure 2A and B). To further determine whether caffeine inhibits inflammatory cytokine expression, we detected the mRNA levels of IL-1 β and IL-18. The results showed that LPS and ATP treatment increased the mRNA levels of IL-1 β and that this effect could be attenuated by caffeine pretreatment (Figure 2C). On the other hand, the mRNA levels of IL-18 were relatively stable; they were affected by neither LPS/ATP stimulation nor caffeine pretreatment, indicating the continuous expression of this cytokine (Figure 2D). Based on the above findings, we assumed that caffeine inhibited NLRP3 inflammasome-specific cytokine secretion and that this effect was achieved by suppression of IL-1 β gene expression.

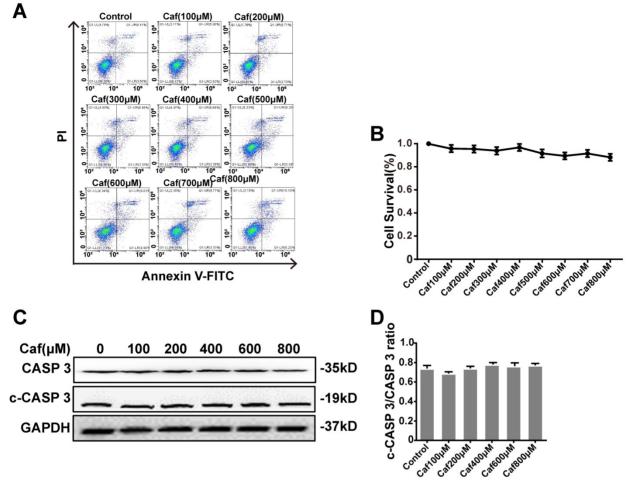


Figure 1. Effects of caffeine on THP-1 macrophage survival and apoptosis. (A) Cell survival was measured in THP-1 macrophages treated with caffeine (100-800 µM) for 6 h using Annexin V-PI staining and flow cytometry. (B) Bar graph showing the survival rates of all cell lines. (C) THP-1 macrophages were treated with caffeine (100-800 µM) for 6 h. The protein levels of cleaved Caspase 3 and Caspase 3 were analyzed by western blotting. (D) Grayscale analysis was used to quantify the relative expression of cleaved Caspase 3 and Caspase 3. The results represent the mean ± SD (n=3).

Caffeine attenuates the activation of the NLRP3 inflammasome

To verify whether the increased secretion of IL-1 β and IL-18 under LPS stimulation contributed to activation of the NLRP3 inflammasome in THP-1 macrophages, we further examined the effect of caffeine on NLRP3 inflammasome activation. The results showed that upon LPS and ATP stimulation, the expression of NLRP3 mRNA and protein, which is the rate-limiting step for inflammasome activation [29], was significantly increased (Figure 3A-C). Moreover, the increased mRNA and protein expression of NLRP3 could be reversed by caffeine pretreatment (Figure 3A-C). The upregulation and activation of NLRP3 promoted Caspase 1 activation (Caspase 1-p20 levels), which in turn promoted the processing and secretion of IL-1 β and IL-18. The PCR results also indicated that caffeine significantly reduced caspase 1 activation rather than Caspase 1 mRNA and protein expression (Figure 3A, B and E). When the NLRP3 inflammasome is activated, the

adaptor protein ASC assembles into a large protein complex named an ASC speck or pyroptosome, which is considered a typical sign of inflammasome activation [30]. Our results showed that caffeine did not change ASC mRNA and protein expression (Figure 3A, B and D) but reduced the number of ASC specks stimulated by LPS and nigericin (Figure 3F). These observations suggest that caffeine can inhibit the activation of the NLRP3 inflammasome and the recruitment of ASC.

Caffeine inhibits NLRP3 inflammasome activity via the MAPK/NF-KB pathway

Emerging experimental evidence has indicated that both MAPK and NF- κ B play important roles in the responses to inflammation stress and NLRP3 activation [31-33]. To investigate whether caffeine suppressed NLRP3 inflammasome activation by inhibiting MAPK/NF- κ B signaling, we analyzed the protein levels and nuclear localization of NF- κ B pathway molecules. The western blot results showed that caffeine pretreatment decreased the levels of phosphorylated IkB-a and p65 upon LPS stimulation (Figure 4A-C). We also found that pretreatment with caffeine significantly inhibited the nuclear localization of p65, implying its blocked activity (Figure 4D). Likewise, we detected decreased phosphorylation levels of JNK, ERK and p38 in cells stimulated by LPS following pretreatment with caffeine (Figure 5). The above results suggest that caffeine inhibits LPS-induced NLRP3 inflammasome activation by inhibiting MAPK/NF-κB signaling in THP-1 macrophages.

Caffeine inhibits NLRP3 inflammasome activation by suppressing A2aR-induced ROS production

Recent studies have shown that xanthines, including caffeine, exert their anti-inflammatory effects mainly by antagonizing A2aR [34] to reduce the production of ROS [35], which are required for activation of the NLRP3 inflammasome [36]. We first silenced A2aR expression by siRNA transfection of LPS and ATP stimulated THP-1 macrophages. After silencing, the protein levels of A2aR were significantly decreased, and notable downregulation of Caspase 1 (p20) expression was observed (Figure 6A and B). The expression of NLRP3, ASC and Caspase 1 was only slightly different in each treatment group compared with the negative control (NC) group (Figure 6A and B). In addition, ROS production in the A2aR siRNA group was lower than that in the NC group (Figure 6C and D). Then, we investigated the interaction between caffeine and A2aR in THP-1 macrophages. The results indicated that caffeine inhibited the expression of A2aR in a dose-dependent manner (Figure 6 E-G) and also inhibited ROS generation (Figure 6H and I). In summary, these findings indicate that caffeine inhibits ROS-mediated NLRP3 inflammasome activation at least partially by decreasing A2aR expression.

Discussion

Notably, extensive studies on the lung-protective effects of caffeine in BPD have been conducted, but whether caffeine can alleviate NLRP3 inflammasome changes in BPD has remained unknown. In the present study, NLRP3 inflammasome levels in LPS-induced THP-1 macrophages were significantly reduced after pretreatment with caffeine. Moreover, MAPK/NF-KB signaling was inhibited by caffeine. RNA silencing of A2aR resulted in effects on the NLRP3 pathway similar to those of caffeine. Our study indicated that caffeine could inhibit the NLRP3 inflammasome in THP-1 macrophages by affecting MAPK/NF-KB signaling, in part by inhibiting A2aR-activated ROS generation (Figure 7). We thus propose a new mechanism for the lung-protective effects of caffeine.

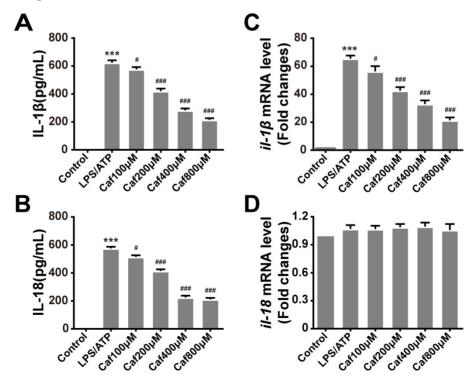


Figure 2. Caffeine attenuates production of the cytokines IL-1β and IL-18 in THP-1 macrophages stimulated by LPS and ATP. (A) IL-1β levels were evaluated in THP-1 macrophages pretreated with caffeine for 1 h and then incubated with LPS (1 µg/ml) for 3 h and ATP (5 mM) for 30 min. (B) IL-18 levels were evaluated in THP-1 macrophages pretreated with caffeine for 1 h and then incubated with LPS (1 µg/ml) for 3 h and ATP (5 mM) for 30 min. (C) The mRNA levels of *il-1β* were analyzed by qRT-PCR. (D) The mRNA levels of *il-18* were analyzed by qRT-PCR. The results represent the mean \pm SD for three experiments. *** p < 0.001 vs. the control group. # p < 0.05, ## p < 0.01 and ### p < 0.001 vs. the LPS and ATP group.

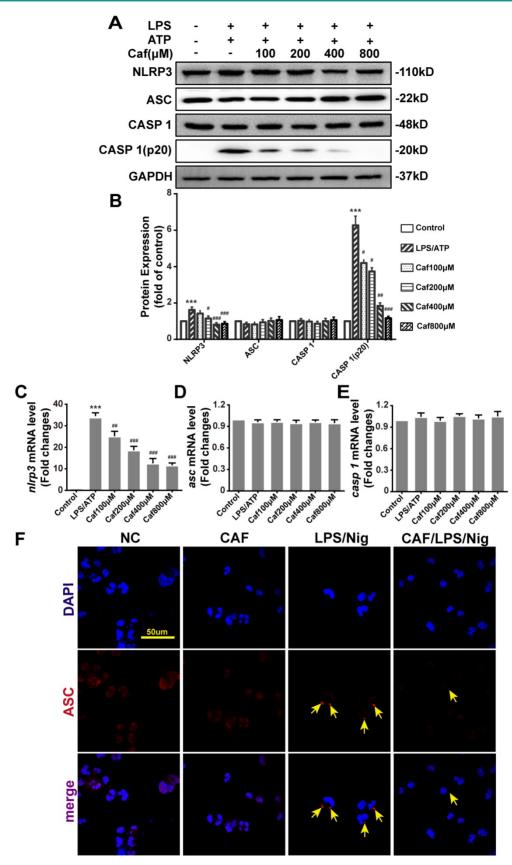
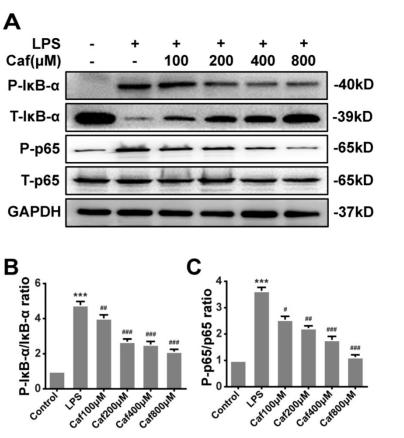


Figure 3. Caffeine inhibits NLRP3 inflammasome activation in THP-1 macrophages stimulated by LPS and ATP/nigericin. (A) The expression of NLRP3 inflammasome component proteins was evaluated by western blotting in THP-1 macrophages pretreated with caffeine for 1 h and then incubated with LPS (1 µg/ml) for 3 h and ATP (5 mM) for 30 min. (B) Densitometric analysis was used to quantify the levels of NLRP3 inflammasome protein expression. The mRNA levels of *nlrp3* (C), *asc* (D) and *caspase l* (E) were analyzed by qRT-PCR. The results represent the mean \pm SD for three experiments. *** p < 0.001 vs. the control group. # p < 0.05, ## p < 0.01 and ### p < 0.001 vs. the LPS and ATP group. (F) ASC speck formation was visualized by immunofluorescence analysis with an anti-ASC (red) antibody in THP-1 macrophages pretreated with caffeine (800 µM) for 1 h and then treated with LPS (1 µg/ml) and nigericin (10 µM) for 90 min. DAPI (blue) was used as a nuclear marker. Scale bar: 50 µm.





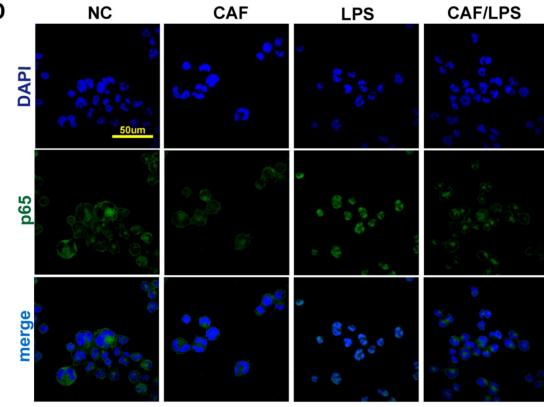


Figure 4. Caffeine inhibits NF-KB activation in THP-1 macrophages stimulated by LPS. (A) The protein levels of IKB- α , p65 and their phosphorylated forms (P-IKB- α and P-p65) were analyzed by western blotting in THP-1 macrophages pretreated with caffeine for 1 h and then incubated with LPS (1 µg/ml) for 3 h. (B) Densitometric analysis was used to quantify the phosphorylation of p65 at Ser536. The results represent the mean \pm SD for three experiments. *** p< 0.001 vs. the control group. # p<0.05, ## p<0.01, and ### p< 0.001 vs. the LPS group. (D) p65 nuclear localization was visualized by immunofluorescence analysis with an anti-p65 (green) antibody in THP-1 macrophages pretreated with caffeine (800 µM) for 1 h followed by LPS (1 µg/ml) for 3 h. DAPI (blue) was used as a nuclear marker. Scale bar: 50 µm.

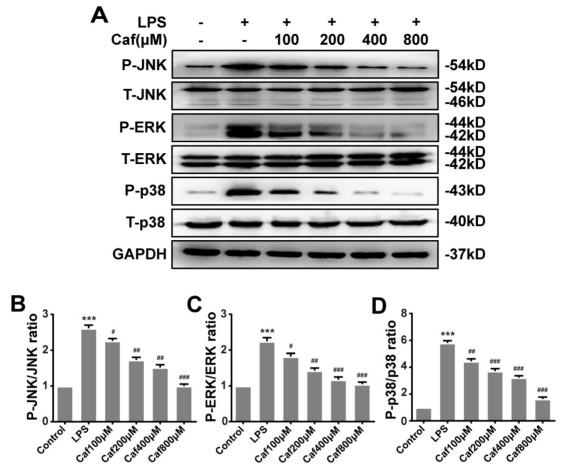


Figure 5. Caffeine suppresses MAPK pathway activation in THP-1 macrophages stimulated by LPS. (A) The protein levels of JNK, ERK, p38, and their phosphorylated forms (P-JNK, P-ERK, and P-p38) were evaluated by western blotting in THP-1 macrophages pretreated with caffeine for 1 h and then incubated with LPS (1 μ g/ml) for 3 h. (B) Densitometric analysis was used to quantify the phosphorylation of JNK at Thr183/Tyr185. (C) Densitometric analysis was used to quantify the phosphorylation of p38 at Thr180/Tyr182. The results represent the mean \pm SD for three experiments. **** p < 0.001 vs. the control group. # p < 0.05, ## p < 0.01, and ### p < 0.001 vs. the LPS group.

To date, a growing number of in vivo and in vitro studies have indicated that alveolar macrophages can be activated by prenatal chorioamnionitis combined with postpartum mechanical ventilation, oxygen therapy and nosocomial infection to secrete proinflammatory cytokines, including IL-1 β , and thus further accelerate the progression of BPD [37-39]. In addition, investigators have verified that the release of IL-1 β by alveolar macrophages is tightly controlled by activation of the NLRP3 inflammasome [40, 41]. Recently, Liao et al. [23] found that NLRP3-/- mice do not exhibit an inflammatory response under hyperoxia treatment and maintain normal alveolarization. Moreover, ventilated preterm baboons also show activation of the NLRP3 inflammasome and increased IL-1 β levels. However, the effects of regular caffeine treatment on the NLRP3 inflammasome in activated macrophages during BPD remain poorly understood. In our study, caffeine treatment of THP-1 macrophages followed by LPS/ATP stimulation downregulated NLRP3, Caspase 1 and IL-1 β . Our findings deepen

understanding of the anti-inflammatory mechanism of caffeine and emphasize the need for further *in vivo* study.

It is well known that activation of the NLRP3 inflammasome is tightly regulated at the transcriptional and posttranslational levels. Previous studies have shown that in macrophages, LPS stimulates MAPK and NF-KB pathway activation as the first signal to transcriptional upregulating of IL-1 β and IL-18 [42-44]. For NF- κB pathway, I κ B- α is phosphorylated and thus marked for degradation by ubiquitin-dependent proteasomes upon LPS stimulation, which allows p65 phosphorylation and disassembled from the heterodimer. NF-KB p65, a representative subunit of the NF-kB heterodimer, is commonly phosphorylated at the Ser536 position and then translocate to the nucleus to upregulate NLRP3 and IL-1 β mRNA expression [45]. In our study, caffeine markedly reduced the phosphorylation of IkB-a to prevent its degradation in LPS-stimulated THP-1 macrophages. At the same time, phosphorylation of p65 was decreased, as was

activation and translocation of NF- κ B. The MAPK pathway also participates in the activation of the NF- κ B pathway as well as the upregulation of IL-1 β and IL-18 [46, 47]. Our results demonstrated that caffeine inhibited the phosphorylation of JNK, ERK, and p38 induced by LPS. Taken together, the above results indicate that pretreatment with caffeine inhibits the transcriptional upregulation of IL-1 β by interfering with MAPK-mediated NF- κ B activity.

Caffeine has been reported to function as a nonselective adenosine receptor antagonist with a relatively high affinity for A2aR [48]. A2aRs are G protein-coupled receptors whose activation stimulates a series of downstream pathways, including the PKA, CREB, and MAPK pathways [49]. Recent experiments have shown that treatment with A2aR antagonists either prevents or reverses the ability of inflammatory stimuli to activate human alveolar macrophages [31]. In addition, A2aR acts as a key regulator of ROS production and is required for ATP-induced sustained NLRP3 inflammasome activation [50, 51]. Our results showed that knockdown of A2aR significantly impaired NLRP3 inflammasome activation and ROS production in LPS/ATP-stimulated THP-1 macrophages. Moreover, caffeine significantly inhibited LPS/ATP-induced A2aR gene transcription and protein expression in THP-1 macrophages, indicating that A2aR inhibition could act as a mediator by which caffeine suppresses NLRP3 activation.

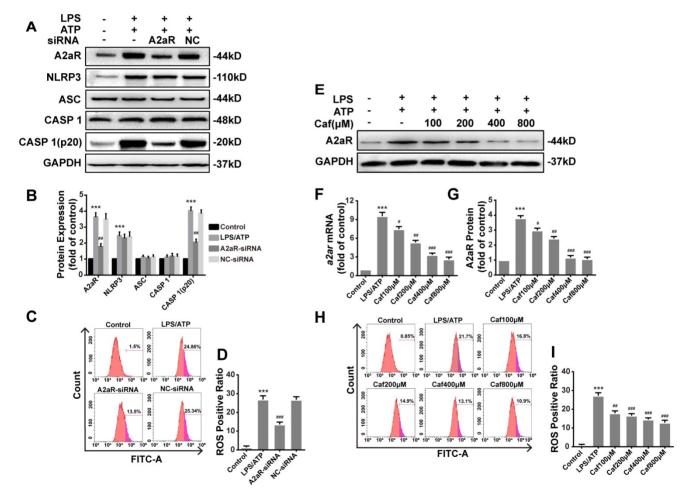


Figure 6. Caffeine inhibits A2aR activity to reduce the generation of ROS required for NLRP3 activation in THP-1 macrophages. (A) The protein levels of A2aR, NLRP3, ASC, Caspase 1 and Caspase 1 (p20) were evaluated by western blotting in THP-1 macrophages preincubated with A2aR-siRNA or negative control (NC)-siRNA for 12 h and then treated with LPS (1 μ g/ml) for 3 h and ATP (5 mM) for 30 min. (B) Densitometric analysis was used to quantify the levels of A2aR, NLRP3, ASC, Caspase 1 and Caspase 1 (p20). (C) THP-1 macrophages were preincubated with A2aR-siRNA or NC-siRNA for 12 h and then treated with LPS (1 μ g/ml) for 3 h and ATP (5 mM) for 30 min. (B) Densitometric analysis was used to quantify the levels of A2aR, NLRP3, ASC, Caspase 1 and Caspase 1 (p20). (C) THP-1 macrophages were preincubated with A2aR-siRNA or NC-siRNA for 12 h and then treated with LPS (1 μ g/ml) for 3 h and ATP (5 mM) for 30 min. ROS production was detected by flow cytometry. (D) Densitometric analysis was used to quantify the levels of ROS. (E) The protein levels of A2aR were evaluated by western blotting in THP-1 macrophages pretreated with caffeine for 1 h and then treated with LPS (1 μ g/ml) for 30 min. (F) The mRNA levels of *a2ar* were detected by qRT-PCR. (G) Densitometric analysis was used to quantify the levels of A2aR protein expression. (H) THP-1 macrophages were preincubated with caffeine for 1 h and then treated with LPS (1 μ g/ml) for 3 h and ATP (5 mM) for 30 min. ROS production was detected by flow cytometry. (I) Densitometric analysis was used to quantify the levels of ROS. The results represent the mean \pm SD for three experiments. *** p < 0.001 vs. the control group. # p < 0.05, # p < 0.01, and # # p < 0.001 vs. the LPS and ATP group.

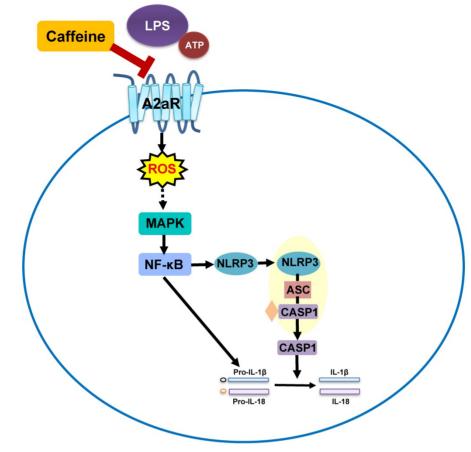


Figure 7. Schematic diagram showing the effect of caffeine on LPS-stimulated NLRP3 inflammasome activation in THP-1 macrophages.

Conclusion

In summary, our findings reveal that caffeine exerts anti-inflammatory effects by inhibiting NLRP3 inflammasome activation in THP-1 macrophages. These findings provide new insights and support a comprehensive understanding of the molecular mechanisms underlying the effects of caffeine treatment in BPD.

Abbreviations

A2AR: Adenosine A2A receptor; ATP: adenosine triphosphate; ASC: Apoptosis associated speck like protein containing a CARD; BPD: bronchopulmonary dysplasia; CAF: caffeine; CREB: cAMP response element-binding protein; DAPI: 4',6-diamidino-2phenylindole; ERK: extracellular-regulated protein kinase; FBS: fetal bovine serum; IkB: inhibitor of kB; IL: interleukin; JNK: c-Jun N-terminal kinase; LPS: lipopolysaccharide; MAPK: mitogen-activated protein kinase; NC: negative control; NF-kB: nuclear factor kappa-light-chain-enhancer of activated B cells; NLRP3: nucleotide-binding domain, leucine-richcontaining family, pyrin domain-containing-3; PMA: phorbol 12-myristate 13-acetate; PKA: protein kinase A; ROS: reactive oxygen species; RPMI: Roswell Park Memorial Institute; siRNA: small interfering RNA; SP-B: surfactant protein B.

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Authors' contributions

Z.W.M, M.L and G.X.H conceived and designed the study. Z.W.M and M.L performed the experiments and analyzed the data. Z.W.M, M.L, C.C and G.X.H wrote and revised the manuscript. All authors read and approved the final manuscript.

Competing Interests

The authors have declared that no competing interest exists.

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