Maxing Shigan Decoction Mitigates *Mycoplasma pneumoniae*-Induced Pyroptosis in A549 Cells via the NLRP3 Inflammasome

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**Background:** *Mycoplasma pneumoniae* is a predominant cause of community-acquired respiratory infections. We recently discovered the clinical efficacy of Maxing shigan decoction (MXSG) in *M. pneumoniae* infection and designed a study to explore the mechanism of action.

**Methods:** Serum IL-1β, IL-18, and TNF-α, and transcript expression of the NLR Family, Pyrin Domain Containing Protein 3 (NLRP3) were measured in the peripheral blood mononuclear cells (PBMCs) of 30 children with *M. pneumoniae* infection and 30 healthy donors. An in vitro model of *M. pneumoniae* infection in A549 cell culture was used to explore the curative effects and mechanisms of MXSG. Pyroptosis was measured by flow cytometry with activated caspase-1 and propidium iodide staining. IL-1β, IL-18, and TNF-α, and NLRP3 transcript expression were measured by qRT-PCR. Protein expression of NLRP3, Caspase-1, pro-caspase-1, IL-1β, pro-IL-1β, and GSDMD-N was determined by Western blotting. Experimental confirmation was performed in NLRP3-overexpressing A549 cells and in the presence of an NLRP3 inhibitor, INF39.

**Results:** *Mycoplasma pneumoniae* infection-induced IL-1β, IL-18, TNF-α, and mRNA expression of NLRP3 in PBMCs and promoted pyroptosis in A549 cells. It also induced IL-1β, IL-18, TNF-α, and up-regulated NLRP3, ro-IL-1β, Caspase-1, Pro-Caspase-1, and GSDMD-N in culture. Similar to the NLRP3 inhibitor INF39, MXSG (0.1, 0.2, and 0.4 mg/mL) suppressed pyroptosis induced by *M. pneumoniae* infection and decreased IL-1β (*P < 0.001), IL-18, TNF-α in culture. MXSG down-regulated NLRP3, pro-IL-1β, Caspase-1, pro-Caspase-1, and GSDMD-N in infected cultures and mitigated NLRP3 overexpression-induced pyroptosis.

**Conclusion:** MXSG mitigates *M. pneumoniae*-induced pyroptosis in A549 cells via the NLRP3 inflammasome.

**Keywords:** *Mycoplasma pneumoniae*, pyroptosis, Maxing shigan decoction, NLRP3 inflammasome

**Introduction**

*Mycoplasma pneumoniae* is one of the smallest self-replicating microorganisms. A harmful human pathogen, *M. pneumoniae* is one of the primary causes of community-acquired respiratory infections, leading to life-threatening tracheobronchitis and pneumonia.1,2 Although the diagnosis of *M. pneumoniae* has dramatically improved, the available treatments (macrolides antibiotics, quinolones, and glucocorticoids) are not fully effective against this pathogen in the respiratory tract.2-5 Alternative remedies for *M. pneumoniae* respiratory infections are urgently needed. In recent years, numerous pharmacological studies have shown that traditional
Chinese medicine (TCM) can effectively treat several intractable diseases, particularly inflammatory diseases and infections caused by drug-resistant bacteria. 6,7 Viral and bacterial infections result in an inflammasome-mediated immune response. The inflammasome induces the release of pro-inflammatory cytokines such as IL-1β, TNF-α, and IL-18, leading to Caspase-1 activation and pyroptosis. 8,9

Maxing shigan decoction (MXSG) is a classic TCM formula composed of whole Ephedra sinica herb, Prunus armeniaca seeds, Glycyrrhiza uralensis roots, and gypsum. MXSG is commonly used to treat respiratory tract diseases, including upper respiratory tract infection, acute bronchitis, pneumonia, chronic obstructive pulmonary disease, and bronchial asthma. 10-12 The active components of MXSG are ephedrine, pseudoephedrine, methylephedrine, amylglycoside, amylglycoside prunasin, glycyrhrizic acid, liquiritin, glycyrrhizin, and CaSO4. 10,13,14 MXSG is also an effective therapy for M. pneumoniae infection, 15,16 although its mechanism of action is unknown. This study aimed to explore the mechanism of MXSG action in M. pneumoniae infection to support further pharmaceutical development of Maxing shigan decoction. A549 cell cultures have been used to explore the effects of M. pneumonia infection on IL-8 release and pyroptosis in human lung epithelial cells, so the same cell line was used in the current study.

Materials and Methods

Ethics Statement

Ethical approval for the study was provided by the independent Ethics Committee of Pudong New Area People’s Hospital. Written informed consent was obtained from the guardians of all participants prior to enrollment. This study was conducted in accordance with the Declaration of Helsinki.

Serum Samples and Isolation of Peripheral Blood Mononuclear Cells

We enrolled 30 children (mean age 31 months, 17 male, 13 female) with M. pneumoniae infection admitted to the Department of Pediatrics, Pudong New Area People’s Hospital (Shanghai, China). Nasopharyngeal aspirate samples were collected for M. pneumoniae detection. Patients were excluded if they had chronic lung disease, immunodeficiency, congenital heart disease, or bronchopulmonary dysplasia. Fasting blood samples were obtained before treatment. Fasting blood samples were collected from 30 age-matched healthy donors as controls. Serum samples were obtained by centrifugation and stored at 80°C. Peripheral blood mononuclear cells (PBMCs) were isolated by centrifugation at 300 × g for 30 min with lymphocyte isolation reagent.

Real-Time Quantitative PCR

Total RNA was extracted using Trizol reagent according to the manufacturer’s protocol (Invitrogen Life Technologies), and 1 µg of each sample was reverse transcribed using the Revert Aid™ First Strand cDNA Synthesis Kit (#K1622; Thermo) with random primers. NLRP3 transcript expression was measured by qRT-PCR with SYBR Green PCR Master Mix (#K0223; Thermo) according to manufacturer instructions with GAPDH as an internal control on an ABI 7300 thermocycler (Applied Biosystems, Foster City, CA, USA). The 2^ΔΔCt relative quantification method was applied. Primer sequences were NLRP3 (5′-AGCCTCAACAAACGCTACAC-3′ and 5′-CATCATCGGGTCAAACAG-3′), GAPDH (5′-AATCC CATCACCATTT-3′ and 5′-AGGCTTGTGTCATACTT-3′).

Measurement of IL-1β, IL-18, and TNF-α

Serum and supernatant IL-1β, IL-18, and TNF-α content was measured by enzyme-linked immunosorbent assay (ELISA; Jiancheng, Nanjing, China).

Mycoplasma Culture

M. pneumoniae strain 29,342 (American Type Culture Collection, Rockville, MD) was grown in broth containing mycoplasma broth base CM403 and selective supplement G SR59 (Oxoid, Hampshire, UK), 0.5% glucose, and 0.002% phenol red at 37°C in the presence of 5% CO2 for 3 days. Viable M. pneumoniae was quantified by colony counting in 10-fold serial dilutions on mycoplasma agar.

Cell Culture and Infection

A549 and 293T cells obtained from the Shanghai Cell Bank, Chinese Academy of Sciences (Shanghai, China) were grown in Dulbecco’s modified Eagle’s medium (DMEM, Hyclone) supplemented with 10% fetal bovine serum (Gibco, Grand Island, NY, USA) at 37°C in the presence of humidified 5% CO2.
Infection was induced by adding 1 mL *M. pneumoniae* to 10 mL cell medium (approximately $1 \times 10^7$ CFU/$10^6$ A549 cells) unless otherwise specified and incubated for 12 h, 24 h, or 48 h. Negative controls were A549 cell cultures with 1 mL bacteria-free mycoplasma broth.

**Preparation of Freeze-Dried MXSG Powder**
The herbal medicines (whole *Ephedra sinica*, *Prunus armeniaca* seeds, *Glycyrrhiza uralensis* roots, and gypsum) were powdered and soaked in purified water for 30 min for 1.5 h. The extracts were then decocted filtered and concentrated at 50°C using a rotary vacuum evaporator. The water extracts were then freeze-dried.

**Lentiviral Overexpression of NLRP3**
The full-length human NLRP3 was cloned into pLVX-puro (Clontech, Palo Alto, CA, USA), then pLVX-NLRP3 (oeNLRP3) and pLVX-puro (vector control) were propagated in 293T cells along with packaging plasmids psPAX2 and pMD2.G.

**Pyroptosis Measurement by PI Staining**
The treated cells were collected and resuspended in staining buffer, then incubated with FLICA 660-YVAD-FMK (FLICA 660 Caspase-1 Assay, Catalog number: 9122) and propidium iodide (PI, Thermo Fisher) for 25 min at 4°C. The ratios of activates caspase-1 and PI-positive cells were analyzed by flow cytometry (BD Biosciences).

**Protein Extraction and Western Blotting**
A549 cells were lysed in ice-cold RIPA lysis buffer (Beyotime Biotechnology), and total protein was quantified by BCA protein assay (Thermo Fisher) according to standard protocols. Protein extracts (25 μg) were resolved by 10% or 12% SDS polyacrylamide gel electrophoresis. Data are expressed as mean ± SD (n = 30), **P < 0.01, vs healthy controls.

![Figure 1](https://www.dovepress.com/)

**Figure 1** Serum levels of IL-1β (A), IL-18 (B), and TNF-α (C), and mRNA expression of NLRP3 (D) in PBMCs of children with *M. pneumoniae* infection. Data are expressed as mean ± SD (n = 30), **P < 0.01, vs healthy controls.
electrophoresis and electroblotted onto nitrocellulose membranes (Millipore). After blocking with 5% nonfat milk, the membranes were probed with primary antibodies of NLRP3 (Abcam, cat. Ab214185, Shanghai, China), IL-1β (Abcam, cat. Ab2105, Shanghai, China), pro-IL-1β (NOVUS, cat. NB600-633, Saint Louis, MO, USA), Caspase 1 (Abcam, cat. Ab74279, Shanghai, China), pro-caspase 1 (Abcam, cat. Ab179515, Shanghai, China), GSDMD-M (NOVUS, cat. NBP2-80427, Saint Louis, MO, USA), and GAPDH (Proteintech, cat. 60004-1-1G, Shanghai, China), followed by incubation with horseradish peroxidase-conjugated secondary antibody (1:2000, Beyotime Biotechnology, Haimen, China). Results were visualized using ECL chromogenic substrate (Bio-Rad Laboratories).

**Statistical Analysis**

Each experiment was conducted at least three times. Statistical analysis between more than two groups and between two groups was performed using ANOVA and Student’s t-test, respectively. A P-value <0.05 was considered significant. Data are presented as means ± standard deviations.

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**Figure 2** *M. pneumoniae* (MP) infection promoted pyroptosis in A549 cells. Controls were cultures with 1 mL bacteria-free mycoplasma broth. (A) Flow cytometry (activated caspase-1 and PI staining); ELISA of IL-1β (B), IL-18 (C), TNF-α (D); (E) qRT-PCR and Western blot analysis of NLRP3 expression in A549; (F) Western blot analysis of IL-1β, pro-IL-1β, caspase-1, pro-caspase 1 (F), and GSDMD-N (G). Data are expressed as mean ± SD (n = 3). *p < 0.05, **p < 0.01, ***p<0.001, vs Control; #p < 0.05, ##p < 0.01, ###p<0.001 vs 12 h; $p < 0.05, $$p < 0.01, $$$p<0.001 vs 24 h.
Results

*M. pneumoniae* Infection Increased Serum Levels of IL-1β, IL-18, and TNF-α, and mRNA Expression of NLRP3 in PBMCs

Serum levels of IL-1, IL-18, and TNF-α, and mRNA expression of NLRP3 in PBMCs were significantly higher in patients than in healthy controls (Figure 1, *P* < 0.01).

*M. pneumoniae* Infection Promoted Pyroptosis in A549 Cells

Flow cytometry showed that *M. pneumoniae* infection-induced incubation-time-dependent pyroptosis in A549 cells and induced expression of pro-inflammatory cytokines IL-1β, IL-18, and TNF-α (Figure 2A–D; *P* < 0.05). *M. pneumoniae* infection also up-regulated expression of NLRP3, pro-IL-1β, Caspase-1, pro-Caspase-1, and GSDMD-N (Figure 2E–G; *P* < 0.05).

All measured indices showed a significant difference by 24 h infection (*P* < 0.05), which was used as the standard infection duration for all subsequent studies.

Maxing Shigan Decoction Mitigated *M. pneumoniae* Infection-Induced Pyroptosis in A549 Cells

Compared to the control, 24 h *M. pneumoniae* infection (vehicle group) induced pyroptosis in A549 cells (Figure 3A, *P* < 0.01). Treatment with NLRP3 inhibitor INF39 (*P* < 0.01) or MXSG (0.1, *P* < 0.05; 0.2 and 0.4 mg/mL, *P* < 0.01) provided dose-dependent suppression of pyroptosis induced by *M. pneumoniae* infection versus the vehicle control. The results are expressed as mean ± SD (n = 3), **p** < 0.01 vs Control (mycoplasma broth without bacteria); *p* < 0.05, ***p*** < 0.01 vs Vehicle (DMEM with 0.1% DMSO); *p* < 0.05, **p** < 0.01 vs 0.1 mg/mL; *P* < 0.05, **p** < 0.01 vs 0.2 mg/mL.

Figure 3 Maxing shigan decoction (MXSG) mitigated *M. pneumoniae* (MP) infection-induced pyroptosis in A549 cells. Cells were treated with *M. pneumoniae* (10⁷ CFU/10⁶ cells) and MXSG (0.1, 0.2, 0.4 mg/mL) for 24 h. Bacteria-free mycoplasma broth was used as the control, and DMEM with 0.1% DMSO was added to cell cultures as the vehicle control. (A) Flow cytometry analysis (activated caspase-1 and PI staining); ELISA of IL-1β (B), IL-18 (C), TNF-α (D); (E) qRT-PCR and Western blot analysis of NLRP3 expression in A549; (F) Western blot analysis of IL-1β, pro-IL-1β, caspase-1, pro-caspase-1 (F), and GSDMD-N (G). Data are expressed as mean ± SD (n = 3), **p** < 0.01 vs Control (mycoplasma broth without bacteria); *p* < 0.05, ***p*** < 0.01 vs Vehicle (DMEM with 0.1% DMSO); *p* < 0.05, **p** < 0.01 vs 0.1 mg/mL; *P* < 0.05, **p** < 0.01 vs 0.2 mg/mL.
vehicle control. MXSG (0.1, 0.2, and 0.4 mg/mL, \( P < 0.05 \)) also suppressed expression of pro-inflammatory cytokines IL-1β, IL-18, and TNF-α and down-regulated NLRP3, pro-IL-1β, Caspase-1, Pro-Caspase-1, and GSDMD-N in M. pneumoniae-infected A549 cells in a dose-dependent manner (\( P < 0.05; \) Figure 3B–G). MXSG significantly reduced all measured infection-induced indices at 0.2 mg/mL, which was selected as the standard dosage for subsequent experiments.

**Maxing Shigan Decoction Mitigated M. pneumoniae Infection-Induced Pyroptosis in A549 Cells by Downregulating NLRP3**

We then explored whether NLRP3 mediated the anti-pyroptotic effects of MXSG in A549 cells. A549 cells overexpressing NLRP3 were exposed to M. pneumonia and 0.2 mg/mL MXSG. The results showed that NLRP3 overexpression significantly suppressed the inhibitory effects of MXSG on NLRP3 expression, pyroptosis, pro-inflammatory cytokines IL-18, TNF-α, and IL-1β, as well as pro-IL-1β, Caspase-1, pro-Caspase-1, and GSDMD-N (Figure 4).

**Maxing Shigan Decoction Mitigated NLRP3 Overexpression-Induced Pyroptosis in A549 Cells**

NLRP3 overexpressing A549 cells were used to determine the anti-pyroptotic effects of MXSG (Figure 5A). NLRP3 overexpression-induced significant pyroptosis in A549 cells compared to the vector group (\( P < 0.01 \)); however, MXSG counteracted this effect (Figure 5B). Our results also indicated that MXSG (\( P < 0.05 \)) suppressed the induction of pro-inflammatory cytokines such as IL-18, TNF-α, and IL-1β induced by NLRP3 overexpression (Figure 5C–F). MXSG also down-regulated NLRP3 (Figure 5A), pro-IL-1β, Caspase-1, pro-Caspase-1 (Figure 5F), and GSDMD-N (Figure 5G) in NLRP3-overexpressing A549 (\( P < 0.05 \)).

**Discussion**

*M. pneumoniae* is one of the most common causes of community-acquired pneumonia and is closely related to the development of asthma. It is widely believed that an outbreak of *M. pneumoniae* would bring about a public health emergency.\(^{1,17}\) We explored the curative effect and
mechanisms of the classic TCM formula Maxing shigan decoction on *M. pneumoniae* infection.

Pyroptosis, closely related to the inflammatory response, is a recently identified pathway of programmed cell death.\(^{18-20}\) Unlike apoptosis, which is physiological suicide, pyroptosis is pathological suicide commonly caused by a viral or bacterial infection. Pyroptosis results in rupture of the cellular membrane and leakage of cytoplasm, triggering an inflammatory response.\(^{21,22}\) Pyroptosis induces pro-inflammatory cytokines such as IL-1β, TNF-α, and IL-18, exacerbating the inflammatory response.\(^{23,24}\) NLRP3 has been associated with the development of pyroptosis.\(^{25,26}\) Here, we found increased serum levels of IL-1β, IL-18, TNF-α, and mRNA expression of NLRP3 in PBMCs of children with *M. pneumoniae* infection compared to healthy children. In the future, we will determine whether pyroptosis occurs in PBMCs of children with *M. pneumoniae* infection. We found that *M. pneumoniae* infection induces time-dependent pyroptosis in A549 lung cancer cells as well as serum IL-1β, TNF-α, and IL-18, and NLRP3 expression. The NLRP3 inflammasome is an important member of the NLR family and comprises NLRP3, ASC, and pro-caspase 1.\(^{27}\) Pyroptosis is mediated by activation of the NLRP3 inflammasome and Caspase-1, as well as Gasdermin D (GSDMD-N).\(^{28-30}\) Importantly, our results suggest that *M. pneumoniae* infection causes time-dependent upregulation of NLRP3, Caspase-1,

Figure 5 Maxing shigan decoction (MXSG) mitigated NLRP3 overexpression-induced pyroptosis in A549 cells. Cells were treated with MXSG (0.2 mg/mL) for 24 h. Bacteria-free broth and DMEM with 0.1% DMSO were used as controls. (A) Western blot analysis of NLRP3 expression; (B) Flow cytometry (PI staining); ELISA analysis of IL-18 (C), TNF-α (D), and IL-1β (E); (F) Western blot analysis of IL-1β, pro-IL-1β, caspase-1, pro-caspase 1 (F), and GSDMD-N (G). Data are expressed as mean ± SD (n = 3), Data are expressed as mean ± SD (n = 3), **P < 0.01 vs Vector; ##P < 0.01 vs oeNLRP3 + Vehicle.
pro-caspase-1, and GSDMD-N in A549 cells. We thus speculated that *M. pneumoniae* induces pyroptosis in these cultures via regulation of the NLRP3 inflammasome. Thus, we determined the effects of the NLRP3 inhibitor INF39 on *M. pneumoniae* infection-induced pyroptosis in A549 cells. Indeed, INF39 abrogated infection-induced pyroptosis and NLRP3 overexpression induced pyroptosis in A549 cells.

TCM provides a broad spectrum of therapies derived from natural active compounds that mitigate disease. Indeed, we found that the effects of Maxing shigan decoction on *M. pneumoniae* infection-induced pyroptosis were similar to that of INF39, inhibiting the production of pro-inflammatory cytokines. This effect was abrogated by NLRP3 overexpression. MXSG treatment also down-regulated the expression of NLRP3, Pro-IL-1β, Caspase-1, pro-caspase-1, and GSDMD-N in *M. pneumoniae*-infected and NLRP3-overexpressing A549 cells.

In conclusion, our findings suggest Maxing shigan decoction mitigates *M. pneumoniae*-induced pyroptosis in A549 cells via the NLRP3 inflammasome, supporting the further pharmaceutical development of Maxing shigan decoction.

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**Disclosure**

The authors declare that they have no competing interests.

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