



Itaconate ameliorates methicillin-resistant *Staphylococcus aureus*-induced acute lung injury through the Nrf2/ARE pathway

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Background: Methicillin-resistant *Staphylococcus aureus* (MRSA) are a critical predisposing factor of sepsis in the clinic. As a product of human energy metabolism and immune response, itaconate can effectively reduce inflammation in the body. This research employed 4-octyl itaconate (4-OI) to illustrate that itaconate exerted anti-inflammatory effects to protect the body from acute lung injury (ALI) induced by MRSA.

Methods: HE staining and immunohistochemistry are used to evaluate the MRSA-induced ALI in mice. WB and qPCR were used to verify the effect of 4-OI on inflammation and oxidative stress caused by MRSA. Molecular docking was used to verify the binding sites of 4-OI and Keap1.

Results: We demonstrated that 4-OI treatment increased the survival ratio, attenuated the pathological damage, inhibited neutrophil infiltration, and reduced lung bacterial burden in the mouse MRSA pneumonia model. 4-OI decreased the expression of inflammatory factors by stimulating the Nrf2 *in vivo* and *in vitro*. Furthermore, 4-OI exerted its effect by promoting nuclear transport of Nrf2 *in vitro*. The results of molecular docking indicated that 4-OI bound to the pocket of Keap1 and exerted a stable interaction. Both Nrf2 inhibitors (ML385) and Nrf2^{-/-} mice abolished the protective effect of 4-OI on MRSA-induced inflammation both *in vitro* and *in vivo*.

Conclusions: 4-OI prevents lung damage caused by MRSA bacteremia via activating Nrf2/ARE pathway.

Keywords: Methicillin-resistant *Staphylococcus aureus* (MRSA); sepsis; itaconate; acute lung injury (ALI); NF-E2-related factor 2

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Introduction

Methicillin-resistant *Staphylococcus aureus* (MRSA), a critical pathogen causing hospital-acquired infections, is a major global health problem (1,2). MRSA infection is a life-threatening disorder because of the bacteria's strong capacity for antibiotic resistance. Patients with MRSA

sepsis often suffer from acute lung injury (ALI), hypoxemia, and respiratory distress (3,4). Due to the continuous emergence of drug-resistant strains, MRSA-induced ALI is associated with high mortality and morbidity worldwide (5). Therefore, new drugs to treat patients with MRSA-induced ALI are urgently needed.

Itaconate, a product of human energy metabolism and immune responses, has strong anti-inflammatory and antioxidant effects. Itaconate can also exert antibacterial effects by inhibiting isocitrate lyase (6). The anti-inflammatory activity of itaconate has attracted increasing attention, with several studies reporting that itaconate dampens the inflammatory response in lipopolysaccharide (LPS)-treated macrophages and mice. Itaconate also inhibits the expression of cytokines and protects mice from ischemia-reperfusion injury (7). As derivatives of itaconate that can penetrate biological membranes, dimethyl itaconate (DI) and 4-octyl itaconate (4-OI) were found to decrease proinflammatory cytokine production in macrophages and protect mice against LPS-induced lethality (8). Additionally, DI has been shown to exert therapeutic effects against fungal infectious diseases, including imiquimod-induced psoriasis-like skin inflammation (9) and *Aspergillus fumigatus*-induced fungal keratitis in mice (10). However, the protective effect of itaconate against Gram-positive bacteria (e.g., MRSA)-induced ALI has not been studied.

Increasing evidence has indicated that nuclear factor-erythroid 2 (NF-E2)-related factor 2 (*Nrf2*) and its downstream gene, heme oxygenase-1 (*HO-1*) and NAD(P)H:quinone oxidoreductase 1 (*NQO1*), play important roles in the anti-inflammatory activity of many natural compounds. As a cell protective enzyme, HO-1 mainly catalyzes the catabolism of heme into ferrous iron, carbon monoxide and biliverdin. The degradation of the heme helps prevent its pro-oxidation effect. Biliverdin and its reduced bilirubin have effective ROS scavenging activity to resist oxidative stress. NQO1 can catalyze the reduction of quinone to hydroquinone, prevent quinone from forming semihydroquinone through a one-electron reduction reaction, and ultimately reduce the ROS generated by the oxidation of semihydroquinone. NQO1 can also prevent environmental stressors from causing oxidative damage to DNA and protect endogenous play an important role in antioxidants. For example, falcariindiol protects against LPS-induced acute intestinal inflammation through the Nrf2/HO-1 pathway (11). *Angelica sinensis* also activates Nrf2/HO-1 and inhibits interleukin 1 beta (IL-1 β) and tumor necrosis factor alpha (TNF- α) in LPS-induced RAW264.7 cells (12). Similarly, 4-OI promotes the nuclear localization of Nrf2 and thereby decreases IL-1 β expression in LPS-treated macrophages and mice (9). However, it is unknown whether the anti-inflammatory effect of itaconate on MRSA-ALI is dependent on the Nrf2/ARE pathway.

This study was conducted to evaluate the molecular mechanisms by which 4-OI alleviates MRSA-ALI. The hypothesis that itaconate attenuates MRSA-ALI by activating the Nrf2/ARE pathway was tested using an MRSA-ALI mouse model, Nrf2-knockout (*Nrf2*^{-/-}) mice, and Nrf2 inhibitors (ML385).

We present the following article in accordance with the ARRIVE reporting checklist (available at <http://dx.doi.org/10.21037/atm-21-1448>).

Methods

Reagents and antibodies

Lipoteichoic acid (LTA) from *S. aureus* (CAS#56411-57-5) and dihydroethidium were obtained from Sigma-Aldrich (St. Louis, MO, USA). The following drugs were purchased from MedChemExpress (Monmouth Junction, NJ, USA) to treat cells or animals: ML385 (Cat# HY-100523), 4-OI (Cat# HY-112675), and corn oil (Cat#HY-Y1888). Antibodies against GAPDH, Keap1, Nrf2, HO-1, and nitric oxide synthase (iNOS), and Cell Counting Kit-8 (CCK-8) were described in our previous report (13).

Bacterial strains and growth conditions

MRSA strain ATCC43300 was cultured and harvested as previously described (14). Briefly, this strain was grown to mid-log phase (OD 600 nm) in Luria-Bertani (LB) with shaking at a rate of 200 rpm in a constant temperature incubator at 37 °C. MRSA bacterial suspension was centrifuged, and the bacteria were resuspended in sterile phosphate-buffered saline (PBS). The bacterial suspension was diluted with sterile PBS and then plated onto LB agar to determine the bacterial concentration; the plated agar was stored in 50% glycerol at -80 °C.

Animals

C57BL/6 wild-type (WT) mice were obtained from Jiangnan University (Wuxi, Jiangsu, China). Male Nrf2-knockout (*Nrf2*^{-/-}) mice (8 weeks old, 18–25 g) were acquired from the Model Animal Research Center (Nanjing, China). Experiments were performed under a project license (No.: JN.No20191230c0400901) granted by ethics committee board of Jiangnan University, in compliance with China national or institutional guidelines for the care and use of animals.

MRSA-induced ALI mice model

WT mice were grouped as follows (n=5): (I) intraperitoneal injection with saline solution (control), (II) intraperitoneal injection with 3×10^8 colony-forming unit/mouse MRSA bacteria (MRSA), and (III) intraperitoneal injection of MRSA and 4-OI (MRSA+4-OI). 4-OI was dissolved in the vehicle (dimethyl sulfoxide: corn oil =1:9). At 1 h after intraperitoneal injection of 4-OI (25 mg/kg) in mice, MRSA was intraperitoneally (i.p.) injected to construct an ALI model (13).

Nrf2^{-/-} male mice were grouped as follows (n=5): Nrf2^{-/-} control, Nrf2^{-/-} MRSA, and Nrf2^{-/-} MRSA + 4-OI. The 4-OI administration and MRSA bacteria were the same as those in WT mice. The mice in all groups were euthanized after 6 hours of treatment to obtain the experimental samples.

Bronchoalveolar lavage fluid (BALF)

The lung tissue of the mice was lavaged with sterile precooled PBS through the trachea. The BALF solution was centrifuged to obtain the supernatant. The protein concentration in the supernatant was determined using a bicinchoninic acid (BCA) kit (Beyotime, Shanghai, China). The total number of cells was determined by flow cytometry (BD C6, BD Biosciences, Franklin Lakes, NJ, USA).

Acquisition and treatment of bone marrow-derived macrophages

Bone marrow flushed from the femurs and tibias of WT C57BL/6 male mice were plated in Dulbecco's Modified Eagle's Medium (Hyclone, Logan, UT, USA) supplemented with 10% fetal bovine serum (Gibco, Grand Island, NY, USA), 1% penicillin-streptomycin (Gibco), and 10 ng/mL macrophage colony-stimulating factor (PeproTech, Rocky Hill, NJ, USA) for 6 days. Bone marrow-derived macrophages (BMDMs) were treated with 4-OI (62.5 μ M) or ML385 (20 μ M) for 4 hours and then challenged with LTA (20 μ g/mL) for 4 hours.

Content of myeloperoxidase (MPO) and malondialdehyde (MDA)

Lung tissue was homogenized, and MPO and MDA were detected according to the manufacturer's instructions.

Histopathology

Lung tissues were fixed in 4% paraformaldehyde and embedded in paraffin. Hematoxylin and eosin (HE) staining was performed to observe lung tissue damage in the mice. The histological ALI scoring system, as described by Matute-Bello, was used (15).

Immunohistochemistry

Lung tissues were cut into 3- μ m sections from the embedded tissue, deparaffinized, and hydrated, and sodium citrate solution was used for antigen retrieval (Solarbio, Beijing, China). The sections were treated with specific serum proteins and immunostained with anti-MPO antibodies at a dilution of 1:100 (ab62141, Abcam, Cambridge, UK). After washing the sections with PBS, the sections were incubated with biotinylated immunoglobulin G (IgG) at 37 °C for 30 minutes. Finally, the sections were incubated in horseradish peroxidase-streptavidin for 30 minutes at room temperature, and the color reaction was developed using diaminobenzidine. The relative content of MPO in the lung tissue sections was determined by Image Pro Plus, as reported previously (13).

Determination of the bacterial load in lung tissue

To determine the bacterial load in the lung tissue, the lung tissues of mice were homogenized and diluted with sterile saline (1:10⁴, 10⁵, 10⁶). Diluted lung tissue homogenate (100 μ L) was inoculated into LB agar plates and incubated overnight at 37 °C. The bacterial load in the lung tissue was determined by counting the number of bacteria on the LB agar plates at different concentrations (colony-forming unit/g).

Survival rates

To determine the effect of 4-OI on the survival time of MRSA-infected mice, WT and Nrf2^{-/-} mice were divided into groups and treated as described above (n=7–10/group). The survival time was recorded every day. Surviving mice in all groups were euthanized after 7 days.

Cell culture and treatment

RAW264.7 cells were purchased from the American Type

Table 1 Primer sequences of qPCR

mRNA name	Sequence (5'-3') forward	Sequence (5'-3') backward
HO-1	CAAGCCGAGAATGCTGAGTTCATG	GCAAGGGATGATTCCTGCCAG
Nrf2	TTCAGCCAGCCCAGCACATC	CGTAGCCGAAGAAACCTCATTGTC
NQO-1	TTCTGTGGCTTCCAGGTCTT	AGGCTGCTTGGAGCAAATA
TNF- α	CATGAGCACAGAAAGCATGATCCG	AGCAGGAATGAGAAGAGGCTGAG
IL-1 β	GCAACTGTTCTGAACTCAACT	ATCTTTTGGGGTCCGTCAACT
IL-6	ACAACCACGGCCTTCCCTACTT	CACGATTCCCAGAGAACATGTG
GADPH	AGGTCGGTGTGAACGGATTG	TGTAGACCATGTAGTTGAGGTCA

Culture Collection (Manassas, VA, USA) and treated with 4-OI (62.5, 125, and 250 μ M) or ML385 (20 μ M) for 4 hours, which was followed by challenge with LTA (20 μ g/mL) for 4 hours. RAW264.7 cells were cultured in RPMI-1640 medium (Hyclone, Logan, UT, USA) supplemented with 10% fetal bovine serum (Gibco, Grand Island, NY, USA) and 1% penicillin streptomycin (Gibco). RAW264.7 cells were grown in an incubator containing 5% CO₂ at 37 °C.

Determination of reactive oxygen species (ROS)

RAW264.7 cells were seeded at a density of 1×10^6 /well in 96-well plates. The cells were divided into the following groups: control, LTA, and LTA+4-OI (62.5 μ M). RAW264.7 cells were treated with 4-OI (62.5 μ M) for 4 hours and then challenged with LTA (20 μ g/mL) for 4 hours. Dihydroethidium (10 μ M) was added to a 96-well plate containing RAW264.7 cells, which were incubated in the dark for 30 minutes. After washing the cells with PBS, fluorescence images were obtained using a Nikon TE-2000 microscope (Tokyo, Japan).

Real-time quantitative polymerase chain reaction

The lung tissue or cells were fully dispersed in TRIzol reagent (CWBIO, Beijing, China), and RNA was obtained according to the manufacturer's instructions. PrimeScript RT kit (Takara, Shiga, Japan) and SYBR Mix (Yeasen, Shanghai, China) were used for the reverse transcriptase reaction and quantitative polymerase chain reaction (qPCR), respectively. The primer sequences are listed in *Table 1*.

Western blot analysis

The methods of extracting, separating, and transferring

proteins from the lung tissues and cells have been previously described (13). Cytoplasmic and nuclear proteins were extracted using nuclear and cytoplasmic extraction reagents (Thermo Fisher Scientific, Waltham, MA, USA). The bands were visualized using an electrochemiluminescence system (Millipore, Billerica, MA, USA).

Molecular modeling

The three-dimensional structure of 4-OI was obtained from PubChem (<https://pubchem.ncbi.nlm.nih.gov/#query=3133-16-2>). Keap1 (PDB ID: 1U6D) was downloaded from the RCSB protein database (<https://www.rcsb.org/>) and imported into AutoDock 4.2. Keap1 was cleaned and hydrogenated, and water was removed using Pymol software. Finally, a flexible program was selected for docking to evaluate the interaction sites and forces.

Statistical analysis

All data are expressed as the mean \pm SD. Differences between the 2 groups of data were compared by *t*-test. Cumulative survival rates were determined using the log-rank test. Differences were considered statistically significant a P value <0.05. GraphPad Prism 9.0 software (GraphPad, Inc., La Jolla, CA, USA) was used to draw the histograms and survival curves.

Results

4-OI attenuated MRSA-induced ALI in mice

HE staining results showed that the alveolar wall was deformed, and numerous inflammatory cells had infiltrated the alveolar space after MRSA infection. The lung injury

score showed that 4-OI attenuated the aforementioned pathological changes in lung tissue caused by MRSA (Figure 1A,B). MRSA infection caused inflammatory cells to migrate into the alveoli, whereas 4-OI reduced the infiltration of inflammatory cells into the lung tissue ($P=0.0179$; Figure 1C). The protein content in the BALF, an indicator of lung vascular permeability, was significantly higher in the MRSA group than in the control group, whereas 4-OI decreased the total protein level in the BALF (Figure 1D). To further evaluate the effect of 4-OI on MRSA-induced lung neutrophil infiltration, we tested the activity and content of MPO in the lung tissue. The results showed that MPO activity was lower in the MRSA + 4-OI group than in the MRSA group ($P=0.01429$; Figure 1E). The results of MPO immunohistochemistry also showed that 4-OI treatment significantly reduced the infiltration of pulmonary neutrophils in the MRSA+4-OI group ($P=0.0181$; Figure 1F,G). Additionally, 4-OI treatment effectively reduced the number of bacteria in the lung organs ($P=0.0452$) (Figure 1H). Importantly, 4-OI treatment significantly increased the survival rate of mice (Figure 1I).

4-OI reduced inflammation and oxidative stress induced by MRSA in mice

Real-time qPCR results showed that MRSA infection significantly increased the messenger RNA (mRNA) expression of *TNF- α* , *IL-1 β* , and *IL-6* in the lung tissue, while 4-OI treatment reduced the expression of these genes (Figure 2A,B,C). MDA content is an important indicator of the degree of tissue cell membrane peroxidation. We determined the concentration of MDA in the lung tissues and found that 4-OI significantly reduced the MDA content that had been increased by MRSA infection (Figure 2D). Notably, 4-OI upregulated the mRNA expression of Nrf2 and its downstream genes (*NQO1* and *HO-1*) in MRSA-infected mice (Figure 2E,F,G).

4-OI inhibited LTA-induced inflammatory response and oxidative stress injury in RAW264.7 cells

The CCK-8 assay results showed that 4-OI at concentrations of 62.5, 125, and 250 μM did not affect the viability of RAW264.7 cells (Figure 3A). As iNOS is an important inflammatory mediator, we evaluated the effect of 4-OI on the iNOS content after LTA challenge in RAW264.7 cells. Western blotting showed that 4-OI (62.5, 125, and 250 μM) significantly inhibited iNOS expression

in RAW264.7 cells stimulated by LTA ($P=0.0244$, 0.0474, 0.0434; Figure 3B,C). In RAW264.7 cells, LTA significantly increased the mRNA expression of *TNF- α* , *IL-1 β* , and *IL-6*; various doses of 4-OI (62.5, 125, and 250 μM) reduced the expression of these proinflammatory cytokines (Figure 3D,E,F). Dihydroethidium probes were used to detect the level of ROS in cells and results showed that 4-OI (62.5 μM) decreased the cellular ROS content (Figure 3G). These results indicate that 4-OI reduced the expression of proinflammatory cytokines and oxidative stress damage induced by LTA in RAW264.7 cells.

4-OI activated Nrf2/ARE pathway in LTA-stimulated macrophages

Nrf2 mRNA level in RAW264.7 cells decreased after LTA stimulation, whereas 4-OI (62.5, 125, and 250 μM) significantly increased Nrf2 expression. High expression of *Nrf2* caused by 4-OI treatment increased the mRNA expression of downstream *NQO1* and *HO-1* (Figure 4A,B,C). Western blotting showed that 4-OI significantly increased the protein expression of *Nrf2* and *HO-1* in a dose-dependent manner (Figure 4D,E,F).

Under oxidative stress conditions, the interaction between Keap1 and Nrf2 in the cytoplasm is disrupted, allowing Nrf2 to dissociate into the nucleus. Therefore, we tested the expression of the Keap1 protein in the cells and found that 4-OI reduced Keap1 expression (Figure 4G). Furthermore, treatment with 4-OI increased the nuclear translocation of Nrf2 compared to that in the control and LTA stimulation groups (Figure 4H,I). Molecular docking analysis also showed that 4-OI inhibited the formation of hydrogen bonds with TYR-426 and ARG-442 in Keap1 (Figure 4J,K).

ML385 abolishes the anti-inflammatory effect of 4-OI in LTA-induced macrophages

ML385 interacts directly with the Nrf2 protein by binding to the Neh1-binding region of Nrf2, preventing binding of the Nrf2-MAFG complex to the anti-oxidant response element (ARE) sequence and reducing transcriptional activity. ML385 abolished the effect of 4-OI on the promotion of the expression of *Nrf2* and *HO-1* (Figure 5A,B,C). Additionally, ML385 upregulated the expression of proinflammatory cytokines (*TNF- α* , *IL-1 β* , and *IL-6*) in the ML385+4-OI group (Figure 5D,E,F). ML385 also abolished the effect of 4-OI on the promotion of the expression of *Nrf2* and *HO-1* and down-regulation of pro-inflammatory

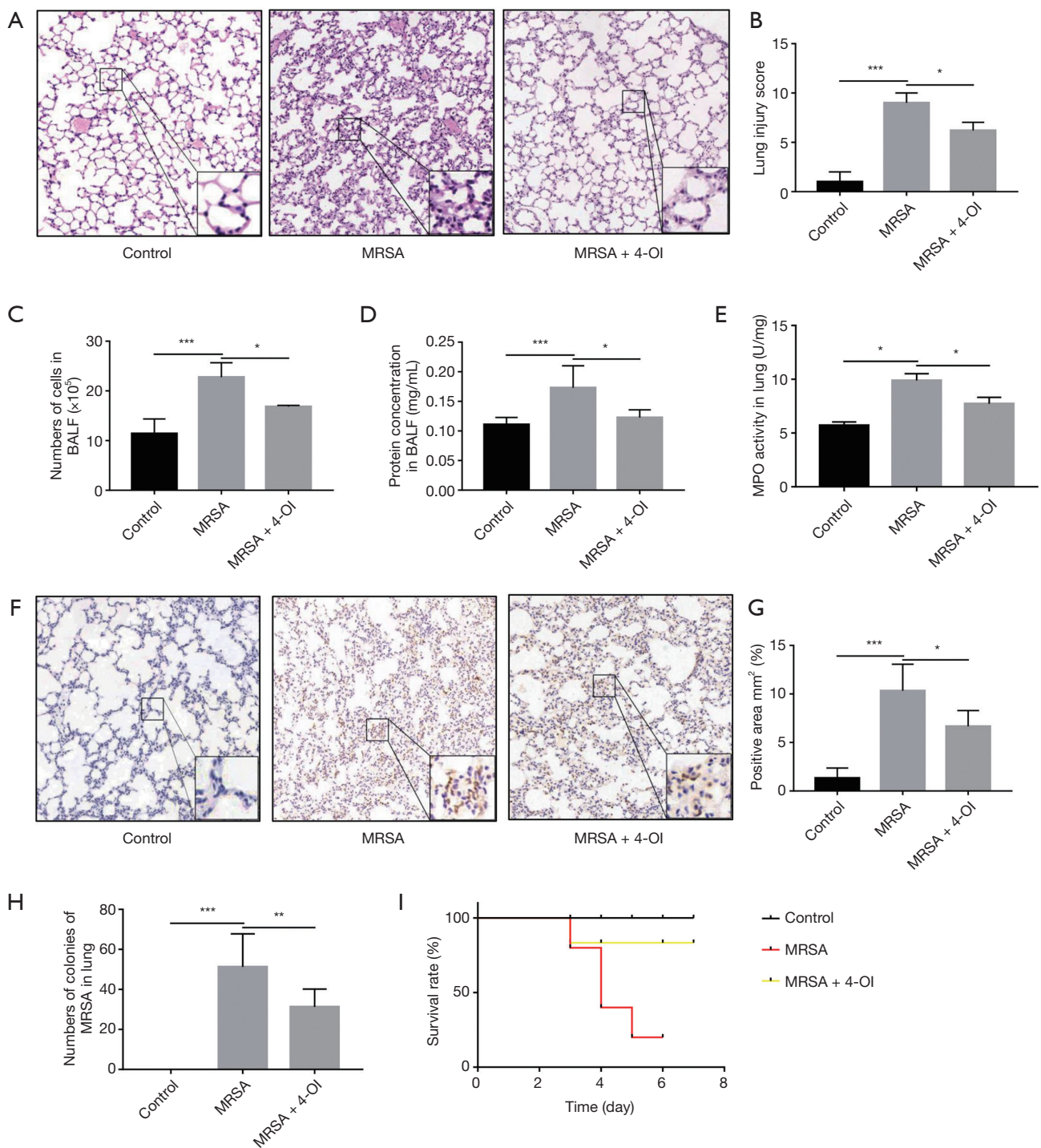


Figure 1 4-OI alleviated MRSA-induced injury symptoms in the ALI mouse model. (A) HE staining (magnification, $\times 200$, bar = $50 \mu\text{m}$; $n=5$). (B) Mouse lung injury score. (C) Total number of cells in BALF ($n=5$). (D) Protein contents in BALF ($n=5$). (E) MPO activities in the lungs. (F) MPO immunohistochemistry (magnification, $\times 200$, bar = $50 \mu\text{m}$; $n=5$). (G) Semiquantitative analysis of MPO immunohistochemistry. (H) MRSA bacterial burden in lung tissue. (I) Cumulative survival curves of MRSA-induced sepsis mice ($n=7-10/\text{group}$). Data are represented as the mean \pm standard deviation (*, $P<0.05$, **, $P<0.01$, and ***, $P<0.001$). MRSA, methicillin-resistant *Staphylococcus aureus*; ALI, acute lung injury; BALF, bronchoalveolar lavage fluid; MPO, myeloperoxidase.

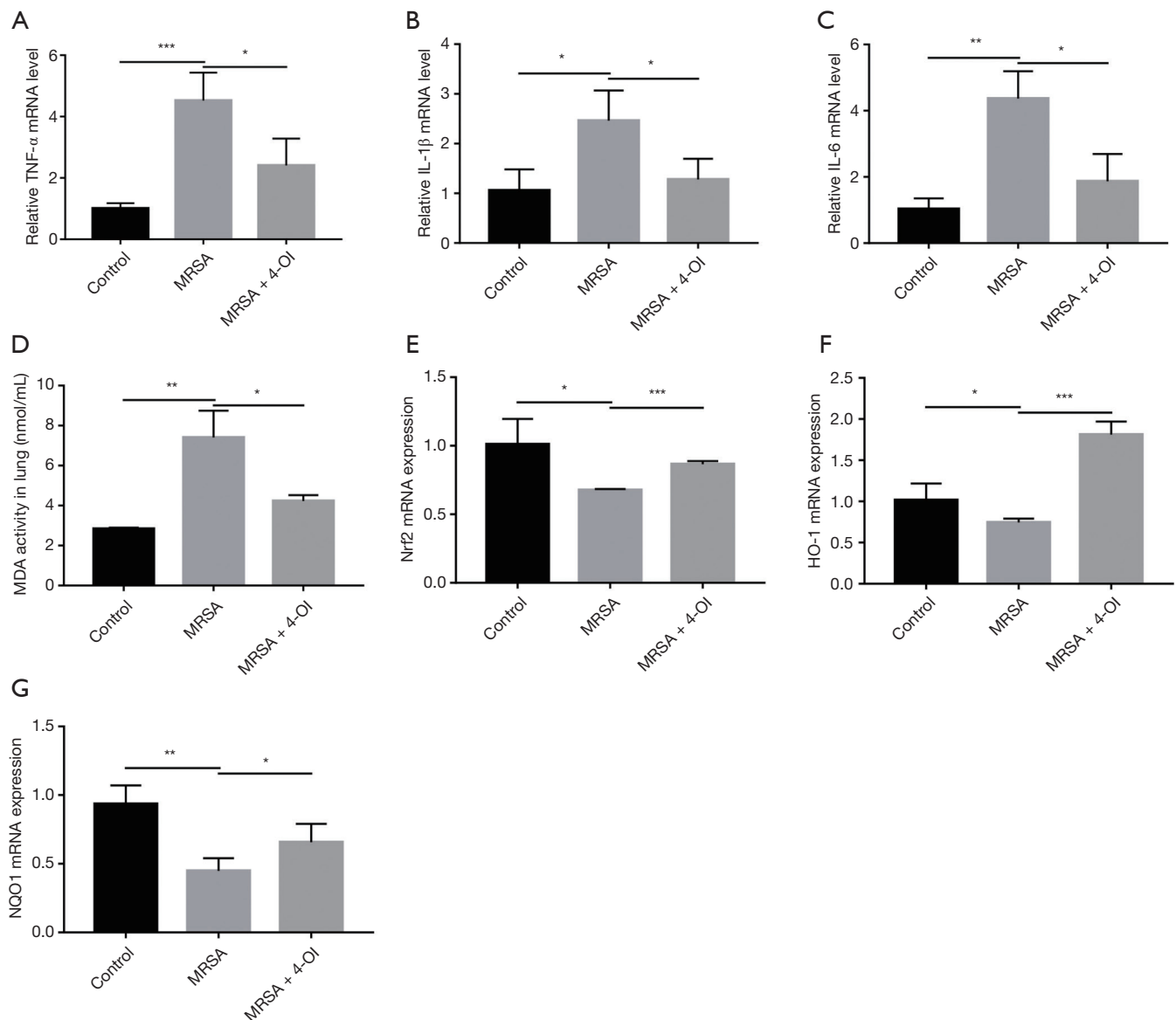


Figure 2 4-OI reduced inflammation and oxidative stress induced by MRSA *in vivo*. (A) TNF- α mRNA levels. (B) mRNA levels of IL-1 β . (C) mRNA levels of IL-6. (D) MDA content in lung tissue. (E) mRNA levels of Nrf2. (F) mRNA levels of HO-1. (G) mRNA levels of NQO1. Data represent the mean \pm standard deviation. Mice in each group =5 (*, $P < 0.05$; **, $P < 0.01$; and ***, $P < 0.001$). 4-OI, 4-octyl itaconate; MRSA, methicillin-resistant *Staphylococcus aureus*; TNF- α , tumor necrosis factor alpha; MDA, malondialdehyde.

factors in BMDMs (Figure 5G,H,I,J,K,L). Taken together, these results show that Nrf2 contributes to the anti-inflammatory effect of 4-OI in LTA-induced macrophages.

Knockout of Nrf2 gene eliminated the protective effect of 4-OI on MRSA-induced ALI in mice

To confirm the role of Nrf2 in the protective effect of

4-OI against MRSA-induced ALI in mice, we used Nrf2^{-/-} mice for further validation. 4-OI treatment did not reverse the changes in the alveolar structure and infiltration of inflammatory cells in Nrf2^{-/-} mice (Figure 6A,B). Additionally, 4-OI did not reduce the protein concentration (Figure 6C) or total cell number (Figure 6D) in the BALF or the MPO content of the lung tissue (Figure 6E,F,G). Furthermore, genetic ablation of Nrf2 increased the

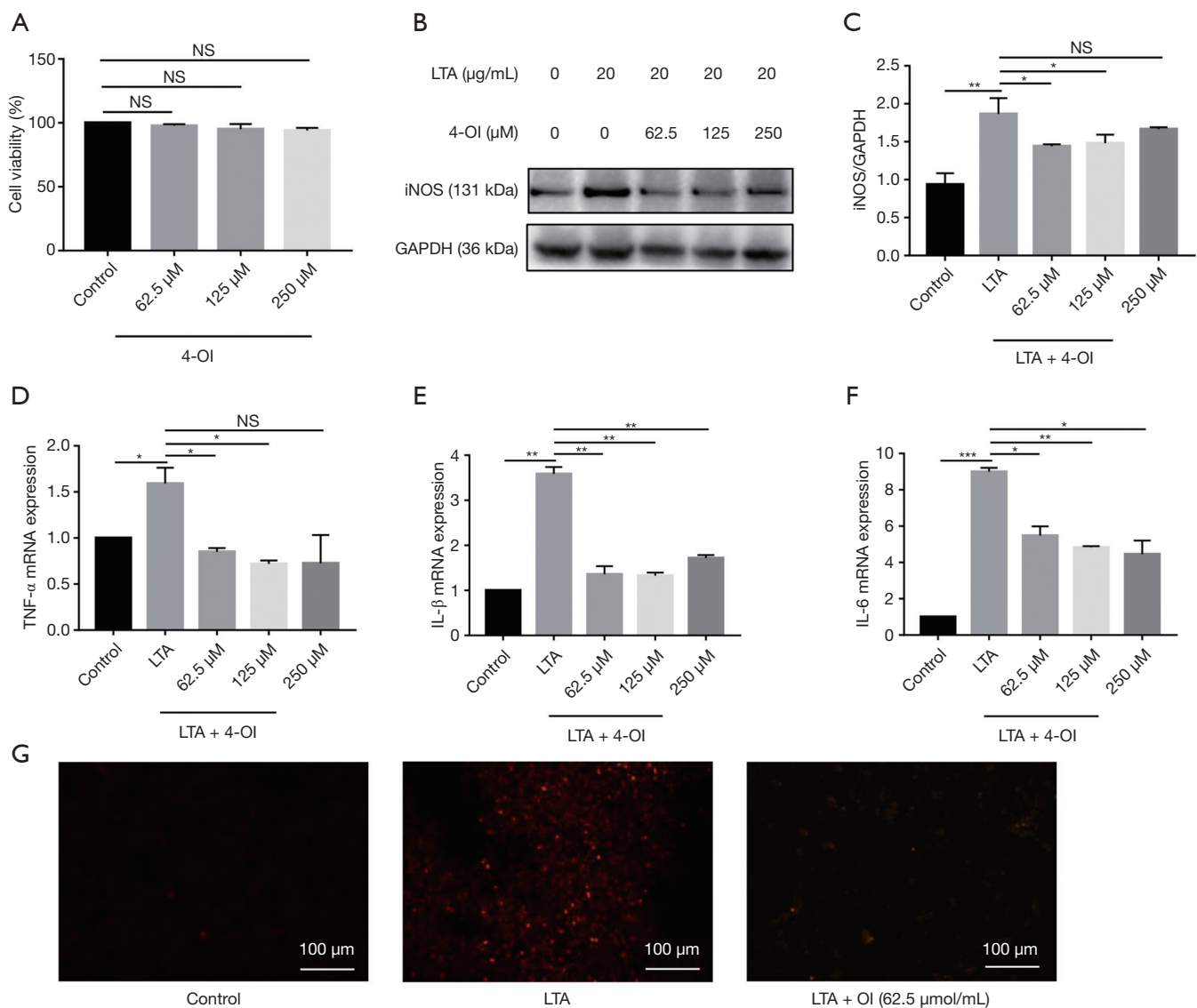


Figure 3 4-OI reduced inflammation and oxidative stress in LTA-induced RAW264.7 cells. (A) CCK-8 assay detected cell viability. (B,C) Western blotting of iNOS and statistical analysis. (D) TNF- α mRNA levels. (E) mRNA levels of IL-1 β . (F) mRNA levels of IL-6. (G) Dihydroethidium staining to detect ROS levels. Data represent the mean \pm standard deviation of 3 independent experiments (*, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$). NS, no significant difference; 4-OI, 4-octyl itaconate; LTA, lipoteichoic acid; TNF- α , tumor necrosis factor alpha.

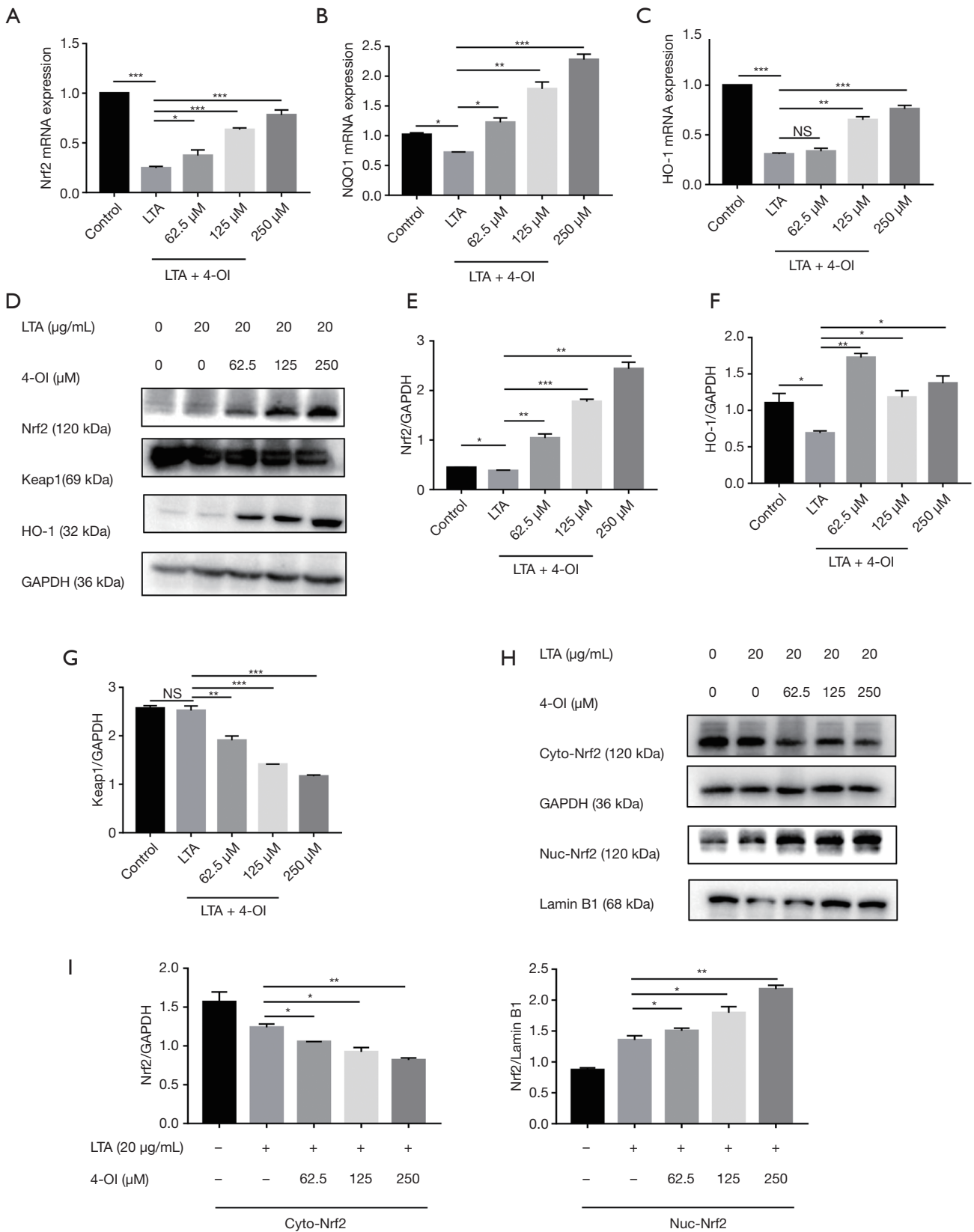
bacterial load in the lungs, and 4-OI could not reverse this condition (Figure 6H). Importantly, 4-OI intervention did not improve the survival rate of MRSA-induced septic Nrf2^{-/-} mice (Figure 6I).

Discussion

In this study, we showed that 4-OI can alleviate MRSA-

induced ALI by reducing the expression of proinflammatory cytokines and oxidative stress genes. The results of molecular docking and western blotting also showed that 4-OI decreased Keap1 levels and activated the Nrf2/ARE pathway to exert anti-inflammatory and antioxidant effects. Our study revealed that 4-OI is an effective therapeutic agent for MRSA-induced lung inflammation (Figure 7).

MRSA is the main pathogen causing acute pneumonia (16),



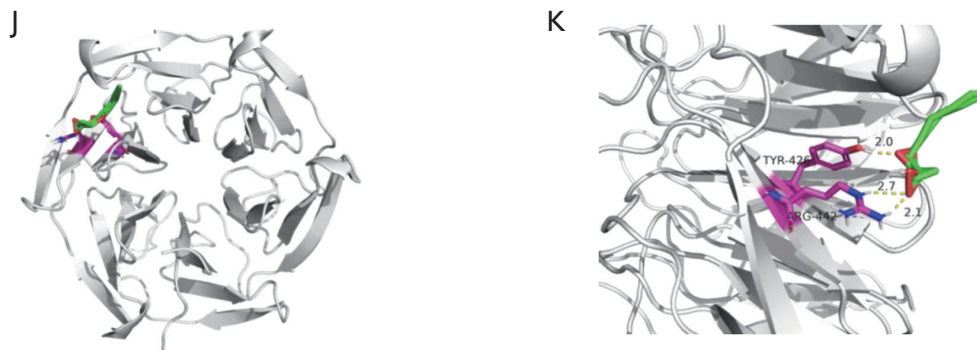


Figure 4 4-OI activated Nrf2 by reducing Keap1 and increasing Nrf2 protein content in LTA-induced macrophages. (A) mRNA levels of Nrf2. (B) mRNA levels of NQO1. (C) mRNA levels of HO-1. (D) Western blotting histograms for Nrf2, Keap1, and HO-1 proteins. (E,F,G) Statistical analysis of Nrf2, HO-1, and Keap1 levels in (D) histograms. (H) Western blot histograms of Nrf2 proteins in the cytoplasm and nucleus. (I) Statistical analysis of Nrf2 levels in (H) histograms. (J) Overall schematic of molecular docking. (K) Binding sites of 4-OI and Keap1. Data represent the mean \pm standard deviation of 3 independent experiments (*, $P < 0.05$; **, $P < 0.01$; and ***, $P < 0.001$). NS, no significant difference; 4-OI, 4-octyl itaconate; LTA, lipoteichoic acid.

chronic obstructive pulmonary disease, fibrosis (17) and other respiratory diseases in hospitals. The renal toxicity of vancomycin makes it difficult to effectively treat respiratory diseases caused by MRSA. As bacterial resistance continues to increase, the mortality rate of MRSA bacteremia has reached more than 20% (18).

Inflammation is an organism's defense response to irritants and pathogens (19). During the inflammatory response caused by Gram-positive bacteria, a large amount of inflammatory cell infiltration aggravates tissue damage by causing the release of inflammatory cytokines (20). Neutrophils are the first immune cells recruited to the injury site. Anti-inflammatory treatment is considered a promising method for treating the early stages of sepsis (21,22). Previous studies have shown that itaconate, a product of human energy metabolism and the immune response, plays an indispensable role in inflammation. However, in the pathological process of MRSA-ALI, the continuous inflammatory response exceeds the regulatory ability of itaconate. However, 4-OI can enter cells where it is hydrolyzed into itaconate. Recent studies have demonstrated that 4-OI can exert anti-inflammatory effects and reduce oxidative stress effects in various cell types (23-25). As an important cell affected by 4-OI, macrophages play an important role in the process of oxidative stress. In this study, we also found that MRSA can cause a mortality rate of more than 50%. 4-OI can effectively improve the survival rate of MRSA mice. After being stimulated by MRSA, both the content of MPO in lung tissue and the number

of neutrophils localized by MPO increased significantly. 4-OI can significantly reduce the increase of MPO in MRSA mice. It has been reported that 4-OI reduces IL-1 β and IL-6 (not including TNF- α) activity in LPS-treated mouse macrophages and exerts anti-inflammatory effects (8). However, it should be noted that 4-OI reduces TNF- α expression in LTA-treated RAW264.7 and MRSA-treated mice. In addition, by activating Nrf2, 4-OI reduces the production of proinflammatory cytokines in human macrophages and peripheral blood mononuclear cells derived from patients with systemic lupus erythematosus (1). In conclusion, our results confirm that 4-OI treatment can reduce the mortality of MRSA-treated mice. 4-OI treatment can also reduce the inflammatory response and oxidative stress damage of MRSA-treated mice and LTA-treated macrophages.

The transcription factor Nrf2 plays a key role in regulating inflammation and oxidative stress (26-28). Activation of the Nrf2/ARE pathway increases the production of downstream enzymes, such as HO-1 and NQO1, thereby protecting the body from oxidative stress. In the resting state, Nrf2 binds to Keap1 in the cytoplasm, and Keap1 controls the expression of Nrf2 (26). When stimulated by the external environment, Nrf2 can eliminate the binding of Keap1 and enter the nucleus to function as a transcription factor (29,30). Itaconic acid esters increase the degradation of cysteine residues on Keap1, leading to further activation of Nrf2 (26). A recent study found that 4-OI can alleviate H₂O₂-induced ROS generation and cell

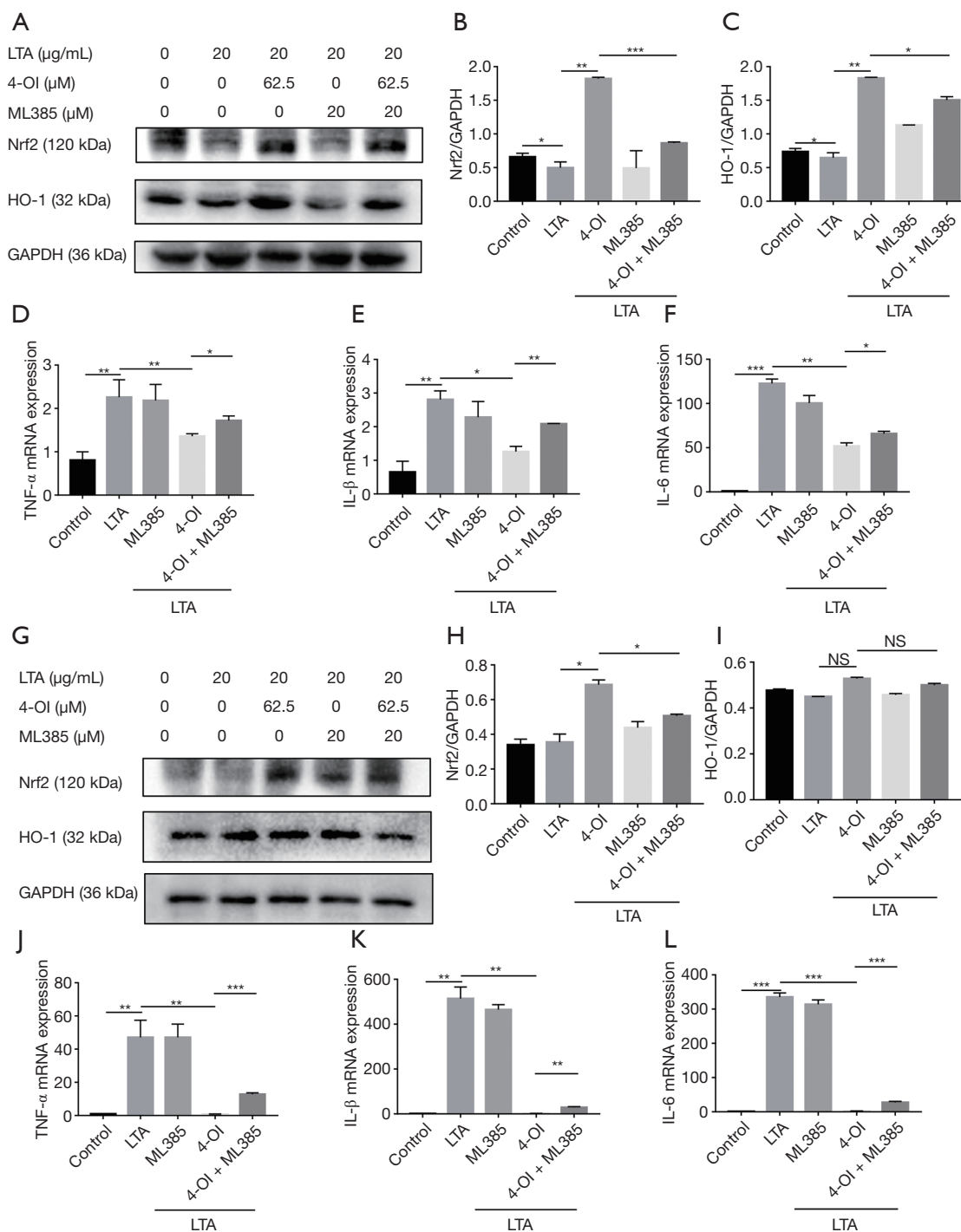


Figure 5 Nrf2 inhibitors abolished the anti-inflammatory effect of 4-OI in LTA-induced macrophages. (A) Western blotting of Nrf2 and HO-1 levels in RAW264.7 cells. (B) Statistical analysis of Nrf2 level in (A) histograms. (C) Statistical analysis of HO-1 level in (A) histograms. (D,E,F) mRNA levels of TNF- α , IL-1 β , and IL-6 were quantified by real-time qPCR in RAW264.7 cells. (G) Western blotting of Nrf2 and HO-1 levels in BMDMs. (H) Statistical analysis of Nrf2 level in (G) histograms. (I) Statistical analysis of HO-1 level in (G) histograms. (J,K,L) mRNA levels of TNF- α , IL-1 β , and IL-6 were quantified by real-time qPCR in BMDMs. Data represent the mean \pm standard deviation of 3 independent experiments (*, $P < 0.05$; **, $P < 0.01$; and ***, $P < 0.001$). NS, no significant difference; 4-OI, 4-octyl itaconate; LTA, lipoteichoic acid; BMDMs, bone marrow-derived macrophages.

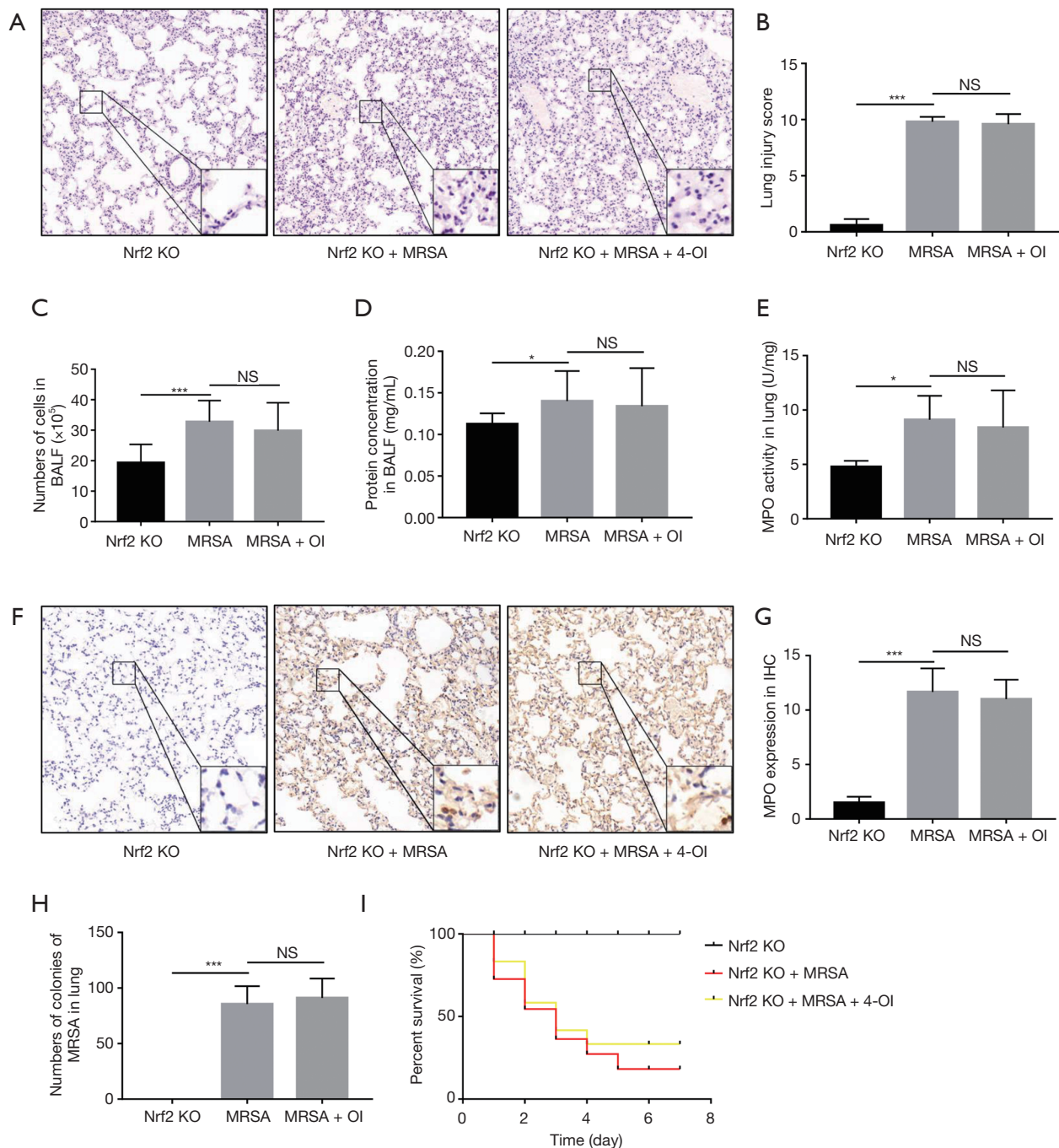


Figure 6 Knockout of Nrf2 gene eliminated the protective effect of 4-OI on MRSA-induced ALI. (A) HE staining (magnification, $\times 200$, bar = 50 μm ; n=5). (B) Mouse lung injury score. (C) Total number of cells in BALF. (D) Protein contents in BALF (n=5). (E) Activity of MPO in lungs (n=3). (F) MPO immunohistochemistry (magnification, $\times 200$, bar = 50 μm ; n=5). (G) Semiquantitative analysis of MPO content by immunohistochemistry. (H) Determination of MRSA bacterial burden in lung tissue. (I) Effect of 4-OI on survival curves of MRSA-induced sepsis in Nrf2^{-/-} mice (n=7–10/group). Data represent the mean \pm standard deviation (*, $P < 0.05$; ***, $P < 0.001$). NS, no significant difference. MRSA, methicillin-resistant *Staphylococcus aureus*; 4-OI, 4-octyl itaconate; ALI, acute lung injury; MPO, myeloperoxidase; BALF, bronchoalveolar lavage fluid.

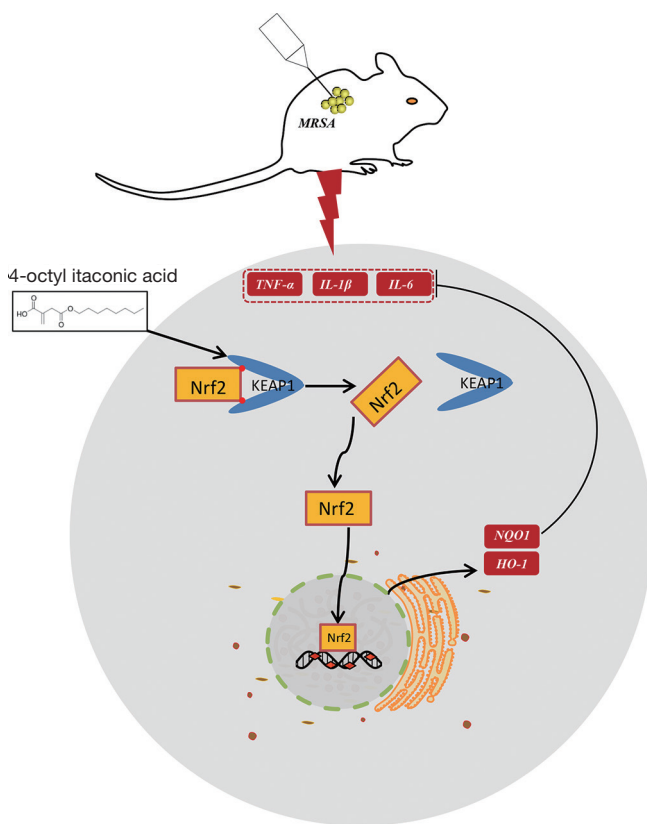


Figure 7 Mechanism of action of 4-octyl itaconate in alleviating MRSA-ALI. 4-OI promotes nuclear translocation of Nrf2 by binding to the amino acid residues of Keap1 (TYR-426 and ARG-442); thereby activating the target genes downstream of Nrf2 (HO-1 and NQO1) to reduce MRSA-induced inflammation and oxidative stress. MRSA, methicillin-resistant *Staphylococcus aureus*; ALI, acute lung injury.

death in SH-SY5Y cells by disrupting the binding of Keap1-NRF2 (31). Many studies have shown that 4-OI activates Nrf2/HO-1 to reduce oxidative stress damage caused by various stimuli. This study is the first to predict the binding sites of 4-OI and Keap1 through molecular docking. 4-OI can activate the Nrf2/HO-1 pathway and promote the nuclear translocation of Nrf2 to resist inflammation and oxidation.

Conclusions

We proved that 4-OI promotes nuclear translocation of Nrf2 by binding to the amino acid residues of Keap1, thereby activating the Nrf2/ARE pathway to protect against MRSA-induced ALI. Thus, 4-OI is a potential therapeutic

anti-inflammatory agent.

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Footnote

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Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. Experiments were performed under a project license (No.: JN.No20191230c0400901) granted by ethics committee board of Jiangnan University, in compliance with China national or institutional guidelines for the care and use of animals.

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