



Effects of *Mycoplasma pneumoniae* infection on airway neurokinin-1 receptor expression in BALB/c mice

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ABSTRACT. The aim of this study was to establish a BALB/c mouse model of *Mycoplasma pneumoniae* (MP) infection and to explore the expression of neurokinin-1 receptor (NK1-R) in the trachea and lung tissue and changes in its relative content at different time points (on the 3rd, 7th, 14th, 21st, and 30th days after infection) in MP-infected BALB/c mice. Immunohistochemistry and Western blot analysis were performed to determine NK1-R expression in the trachea and lung tissue and changes in relative content in MP-infected BALB/c mice. After MP infection, the expression of NK1-R on the surfaces of upper tracheal and bronchial epithelial cells, submucosa, and alveolar epithelial cells, as well as around the smooth muscle, was upregulated more significantly in the infection group than in the control group ($P < 0.05$); NK1-R protein expression was enhanced on the 3rd, 7th, 14th, 21st, and 30th days after infection compared with that of the control

group ($P < 0.05$). NK1-R expression in the trachea, bronchus, and lung tissue increased in MP-infected BALB/c mice, which may explain why wheezing occurs after MP infection.

Key words: Mycoplasma pneumonia; BALB/c mouse; NK1-R

INTRODUCTION

Mycoplasma pneumoniae (MP) is a common pathogen that causes atypical pneumonia and bronchitis in children, and its morbidity has gradually increased in recent years.

The relationship between MP and pediatric asthma has attracted the attention of researchers since the 1970s and has remained a focus until the present day. It is generally believed that MP closely correlates with the occurrence, development, and deterioration of pediatric asthma (Biscardi et al., 2004; Ou et al., 2008; Varshney et al., 2009; Guilbert and Denlinger, 2010). However, the relationship between MP infection and the pathogenesis of pediatric asthma remains unclear. Previous studies on asthma have indicated that expressions of neurokinin-1 receptor (NK1-R) in the airway and lung tissue are distinctly elevated in asthmatic rats and guinea pigs (Li et al., 2011; Li and Shang, 2012). MP infects airways and damages ciliated epithelial cells, resulting in exposure of non-myelinated vagal afferent fibers in the mucosa and submucosa, a release of sensory neuropeptide, and an increase of airway hyperresponsiveness. Whether NK1-R expression is enhanced after MP infection remains unclear. Therefore, we explored the expression of NK1-R in the trachea and lung tissue at different time points in MP-infected BALB/c mice.

MATERIAL AND METHODS

Laboratory animals

A total of 100 specific pathogen free (SPF) BALB/c mice, including 50 males and 50 females at 4-6-weeks old with body weight of 14-16 g, were provided by the Laboratory Animal Center of the General Hospital of Shenyang Military Region. All mice were fed a normal diet and each group contained 10 mice. This study was carried out in strict accordance with the recommendations of the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. The animal use protocol was reviewed and approved by the Institutional Animal Care and Use Committee (IACUC) of Shenyang Hospital of China Medical University.

MP culture and colony assay

The MP international standard strain, FH, was provided by the Department of Pathogenic Microorganisms in China Medical University. The main components in the fluid media were bovine brain-heart infusion medium, sterile fetal bovine serum, fresh yeast extract, glucose, penicillin, and phenol red at a pH of 7.6. Bacteria liquid for testing was diluted to 1×10^{-1} – 1×10^{-12} using a serial dilution method; after constant temperature incubation at 37°C for 14 days, the highest dilution when red medium changed to yellow was used as the color change unit. Bacterium concentration was calculated as color changing units (CCU)/mL.

Establishment of MP infection model

A total of 100 BALB/c mice, including 50 males and 50 females at 4 weeks of age, were randomized into an MP infection group (50 mice) and a control group (50 mice). Mice were kept in an SPF animal house and fed separately under standard conditions. The mice were intraperitoneally anesthetized with 0.05 mL 10% chloral hydrate; each BALB/c mouse in the infection group was intranasally vaccinated with 0.02 mL 10^8 CCU/mL MP bacterium fluid with the mouse head tilted at 30° - 45° to ensure that the MP bacterium fluid entered into the lower respiratory tract through natural breath movement. Mice in the control group were intranasally vaccinated with the same amount phosphate-buffered saline (PBS) rather than MP. Five normal mice were used as controls for each time point. Observation of daily activity and body weight was carried out after MP infection; samples were collected from infected and control mice at 4 time points on the 3rd, 7th, 14th, 21st, and 30th days after infection (Hardy et al., 2001; Martin et al., 2001).

Animal grouping and sample collection

BALB/c mice of the same species, body weight (14-16 g), and breeding conditions as well as equal numbers of male and female mice, were randomly divided into the following groups: MP infection of 3 days (MP3), 7 days (MP7), 14 days (MP14), 21 days (MP21), and 30 days (MP30). Respective studies were performed according to different time points and processing factors. For the control group, the same amount of PBS was intranasally vaccinated in mice rather than MP and the samples were collected at times corresponding to sampling of the experimental groups.

The mice were intraperitoneally anesthetized with 0.05 mL 10% chloral hydrate and fixed in supine position before sample collection. The neck and chest were sterilized with conventional iodine and alcohol, and then the chest was opened with sterile equipment [soaked in 0.1% diethylpyrocarbonate (DEPC) for 24 h, followed by autoclaved sterilization]. The entire lung was removed and weighed to calculate the lung wet weight, and then the left lung was collected and stored in 4% paraformaldehyde.

NK1-R detection

Immunohistochemistry was performed on pathological sections to detect NK1-R; the American Universal Imaging Corporation Image Analysis System (Bedford Hills; NY, USA) and MetaMorph (Molecular Devices; Sunnyvale, CA, USA) software were employed to detect the luminous intensity [integrated optical density (OD) total] to determine the relative content of NK1-R. First, 1 or 2 images were randomly selected as standards; the image analysis system was used to identify the shading percentage of red-yellow-blue as a standard when tinctorial-positive cells comprised more than 90%. The OD of NK1-R at different sites on each section was defined as the relative content, and the mean value was calculated and presented as means \pm SD; measurements were repeated for different points, and differences between groups were compared using one-way analysis of variance. $P < 0.05$ was considered to be statistically significant. NK1-R content in the lung tissue was detected using Western

blot analysis and analyzed using the FlourChem V 2.0 gel imaging analysis software (Protein Simple; Santa Clara, CA, USA). The gray-scale value of each protein electrophoresis band was recorded and quantitatively analyzed. The protein content was calculated as the gray-scale value of the sample protein divided by the gray-scale value of β -actin in the same sample.

Statistical analysis

Statistical analysis was carried out using the SPSS 17.0 software (SPSS, Inc.; Chicago, IL, USA). Measurement data are reported as means \pm SD; differences between groups were compared using one-way analysis of variance; $P < 0.05$ was considered to be statistically significant.

RESULTS

NK1-R expression in the trachea of MP-infected BALB/c mice

NK1-R mainly expressed on the surface of upper tracheal epithelial cells, submucosa, epithelial cells surrounding blood vessels, the surface of inflammatory cells, smooth muscle cells, and the surface of glandular cells (Figure 1A). The expression levels of NK1-R on the surface of airway epithelial cells and submucosa were similar to that of the control group on the 3rd (Figure 1B), 21st, and 30th (Figure 1C and D) days after MP infection, but were significantly enhanced on the 7th day (Figure 1E), after which expression declined on the 14th day (Figure 1F) after MP infection.

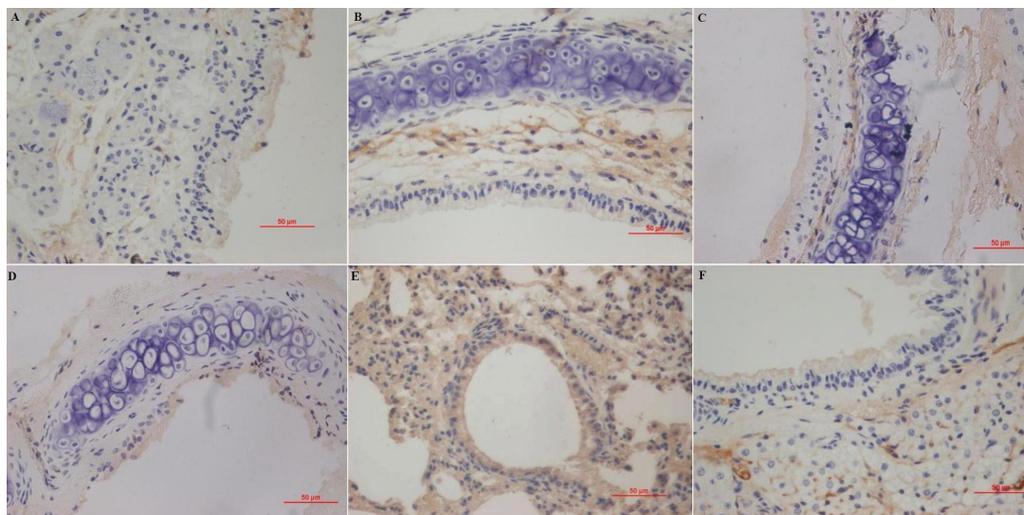


Figure 1. Expression of NK1-R protein in mouse airway from each group 400X. **A.** Control group; **B.** MP3; **C.** MP7; **D.** MP14; **E.** MP21; **F.** MP30.

NK1-R expression in the bronchus and lung tissue of MP-infected BALB/c mice

NK1-R was predominantly expressed on the surface of upper bronchial epithelial cells, submucosa, and the surface of epithelial cells surrounding blood vessels (Figure 2A). The expression level of NK1-R on the surface of upper bronchial epithelial cells, submucosa, and alveolar epithelial cells was similar to that of the control group on the 3rd day (Figure 2B) after MP infection, but distinctly elevated on the 7th, 14th, 21st, and 30th days (Figure 2C, D, E, F) after MP infection.

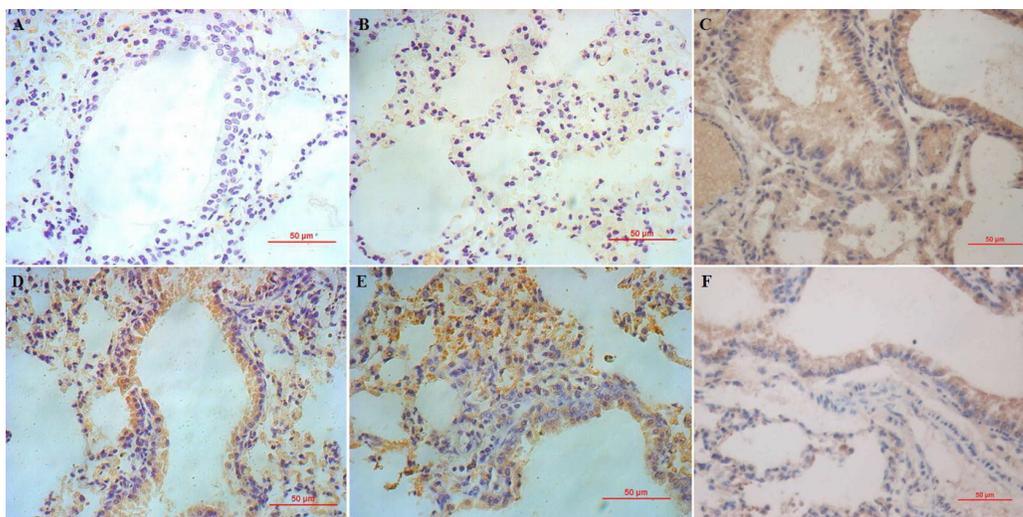


Figure 2. Expression of NK1-R protein in mouse bronchus and lung tissue from each group 400X. **A.** Control group; **B.** MP3; **C.** MP7; **D.** MP14; **E.** MP21; **F.** MP30.

NK1-R expression in the bronchus and lung tissue of MP-infected BALB/c mice represented by OD

After MP infection, NK1-R expression on the surface of upper bronchial epithelial cells, submucosa, and alveolar epithelial cells, as well as near the smooth muscle was up-regulated more significantly in the infection group than in the control group ($P < 0.05$), which slightly declined on the 3rd day and then increased thereafter; NK1-R expression was relatively high on the 7th, 14th, and 21st days, reaching a maximum on the 7th day and then gradually decreasing to the minimum on the 30th day after infection (Figure 3).

Detection of NK1-R protein expression in mouse lung tissue from each group based on Western blot analysis

NK1-R protein expression was enhanced on the 3rd, 7th, 14th, 21st, and 30th days after infection compared with the control group ($P < 0.05$, Figure 4A and B).

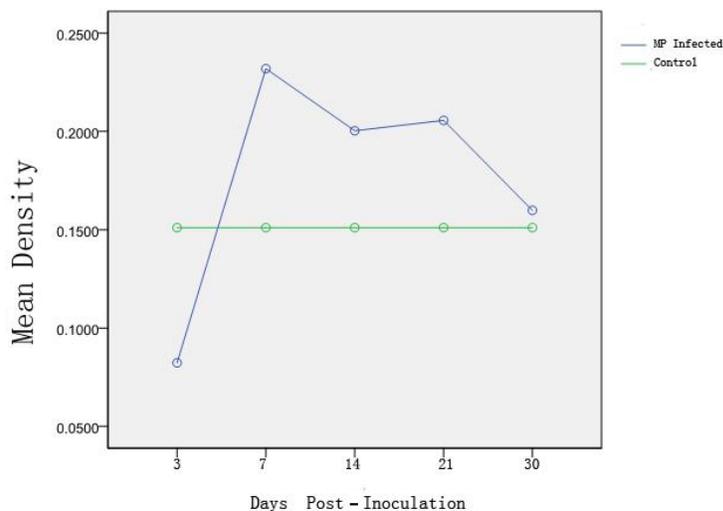


Figure 3. Relative contents of optical density reflecting NK1-R expression in bronchus and lung tissue of *Mycoplasma pneumoniae*-infected BALB/c mice at different time points.

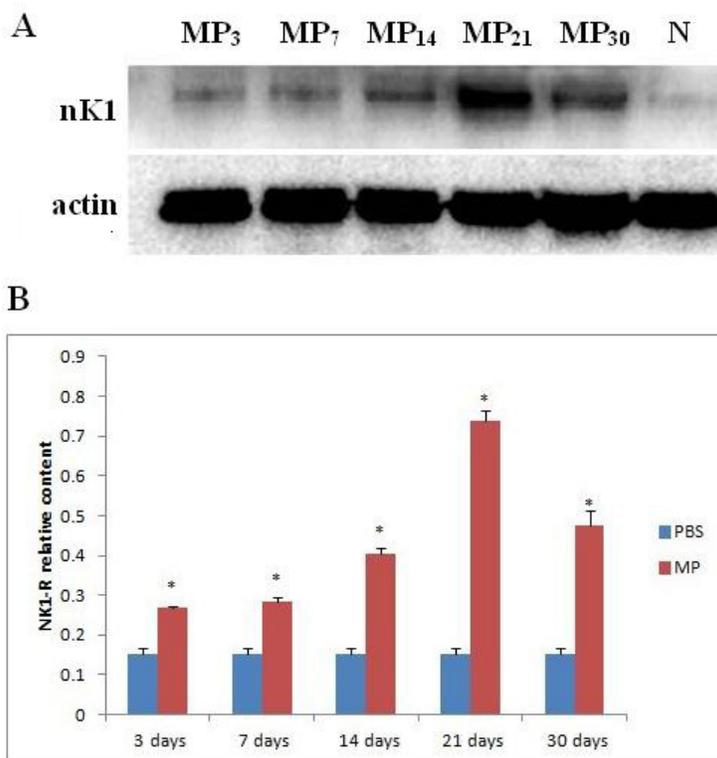


Figure 4. **A.** NK1-R protein expression in mouse lung tissue from each group; **B.** NK1-R protein expression in lung tissue of *Mycoplasma pneumoniae*-infected mice in each group. Lane N = control group. *P < 0.05 was considered to be statistically significant compared with control group.

DISCUSSION

In asthma, modulation of the NK1 receptor appears to influence a variety of pathological symptoms and processes such as inflammation, which induces pronounced and thus readily quantifiable effects in the lungs (e.g., bronchospasm, vasodilatation, vascular leakage, mucus secretion) (Groneberg et al., 2006; Satake and Kawada, 2006; Ramalho et al., 2011).

Asthma is a chronic inflammatory airway disease characterized by variable airway obstruction and bronchial hyperresponsiveness. Several factors affect the development and severity of childhood asthma such as genetic predisposition, atopy, environmental factors, obesity, diet, socioeconomic status, and infectious triggers. MP may play a role in the development of asthma exacerbation during childhood (Annagür et al., 2007).

Acute atypical bacteria are closely correlated with acute asthma attacks, and chronic MP infection plays a particularly important role in persistent asthma (Blasi and Johnston, 2007; Sutherland and Martin, 2007; Korppi, 2010; Maffey et al., 2010). An association between MP and more severe asthma exacerbation has also been observed in a previous study (Cosentini et al., 2008). However, the pathogenesis of MP in asthma remains unclear. A recent study showed that the mechanisms of wheezing attack and asthma exacerbation induced by MP infection primarily include impaired mucociliary clearance, increased mucous production, and eventually, asthma (Bisgaard et al., 2007).

In addition, MP infection stimulates multiple cells and cell components (mast cells, epithelial cells, endothelial cells, macrophages, and smooth muscle cells) to become involved in the pathophysiological process of asthma. These cells release a series of cytokines and adhesion molecules that infiltrate the airway and cause chronic airway inflammation and reconstruction (Jeong et al., 2012; Narita and Tanaka, 2012). Some researchers have hypothesized that MP sensitization and challenge induce collagen deposition in the tracheal and bronchial submucosa of BALB/c mice, resulting in airway reconstruction and causing additional severe bronchial hyperresponsiveness and asthmatic symptoms (Chu et al., 2005). Other studies (Chu et al., 2000) report that infection increases NK1-R expression in the bronchial mucosa in humans.

We observed that NK1-R expression on the surface of upper bronchial epithelial cells, submucosa, and alveolar epithelial cells, as well as near smooth muscle, slightly decreased on the 3rd day after infection and then increased thereafter; the expression level of NK1-R was relatively high on the 7th, 14th, and 21st days, reaching a maximum on the 7th day, and then gradually decreasing to a minimum on the 30th day after MP infection. In addition, NK1-R protein expression was higher on the 3rd, 7th, 14th, 21st, and 30th days after infection compared with levels in the control group; expression of NK1-R protein increased over time to a peak on the 21st day and gradually decreased thereafter. Both immunohistochemistry and Western blot analyses confirmed that NK1-R expression was enhanced after MP infection over time to the peak and then gradually declined. NK1-R is a crucial receptor during asthma attacks and is involved in the chronic airway inflammation induced by increased leakage of blood vessels and tracheal mucosa (Li et al., 2011). MP infection upregulates NK1-R expression in the mouse airway and lung tissue; this explains how the colonization of some pathogenic bacteria make the body more prone to wheezing (Chu et al., 2006; Bisgaard et al., 2007). Infection with MP can precede the onset of asthma, thereby exacerbating asthmatic symptoms, which makes asthma management difficult (Hong, 2012). Moreover, clarithromycin therapy reduced the concentrations of mucosal tumor necrosis factor- α , interleukin-1 β , and interleukin-10 in children with an acute exacerbation of recurrent wheezing (Fonseca-Aten et

al., 2006). Macrolides have both antimicrobial and anti-inflammatory properties. Both mechanisms appear to be important in their clinical efficacy in treating a wide variety of pulmonary disorders, including asthma (Good et al., 2012; Medford, 2012).

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