

Insights into the pathogenesis of *Mycoplasma pneumoniae* (Review)

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Abstract. *Mycoplasma* are the smallest prokaryotic microbes present in nature. These wall-less, malleable organisms can pass through cell filters, and grow and propagate under cell-free conditions *in vitro*. Of the pathogenic *Mycoplasma* *Mycoplasma pneumoniae* has been examined the most. In addition to primary atypical pneumonia and community-acquired pneumonia with predominantly respiratory symptoms, *M. pneumoniae* can also induce autoimmune hemolytic anemia and other diseases in the blood, cardiovascular system, gastrointestinal tract and skin, and can induce pericarditis, myocarditis, nephritis and meningitis. The pathogenesis of *M. pneumoniae* infection is complex and remains to be fully elucidated. The present review aimed to summarize several direct damage mechanisms, including adhesion damage, destruction of membrane fusion, nutrition depletion, invasive damage, toxic damage, inflammatory damage and immune damage. Further investigations are required for determining the detailed pathogenesis of *M. pneumoniae*.

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1. Introduction

Mycoplasma are the smallest prokaryotic microbes present in nature. These wall-less, malleable organisms can pass through cell filters, and can grow and propagate under cell-free conditions *in vitro* (1). *Mycoplasma* contain a 600-1,350 kbp

genome and 23-35% GC. They reproduce predominantly via typical binary fission and have a tendency to form 'fried egg' colonies in solid culture media. At present, seven species of *Mycoplasma* have been found to be pathogenic to humans, including *M. pneumoniae*, *M. urealyticum*, *M. genitalium*, *M. hominis*, *M. fermentation*, *M. penetrans* and *M. pirum* (1). *M. pneumoniae*, which was initially separated, cultivated and named by Chanock and Hayflick in 1962, has been examined the most (2). In addition to primary atypical pneumonia and community-acquired pneumonia, which induce predominantly respiratory symptoms, *M. pneumoniae* can also induce autoimmune hemolytic anemia and other diseases in the blood, cardiovascular system, gastrointestinal tract, and skin, and can induce pericarditis, myocarditis, nephritis and meningitis (3-5).

M. pneumoniae infections are distributed globally with local prevalence. As reported, its infection rate is increasing annually, however, the specific pathogenic mechanism remains to be fully elucidated (2). The pathogenesis of *M. pneumoniae* infection is complex as it involves several mechanisms, including adhesion damage, membrane fusion damage, nutrition depletion, invasive damage, toxic damage, immune damage and inflammatory damage (Fig. 1). However, the specific mechanism underlying its effects remains to be elucidated.

2. Direct damage mechanisms

Adhesion damage. The adhesion of *M. pneumoniae* onto the respiratory epithelia is a precondition dictating the propagation and pathogenesis of *M. pneumoniae* (6). In addition to pseudo-stratified columnar ciliated epithelia, *M. pneumoniae* can also adhere to red blood cells, HeLa cells, fibroblasts, macrophages and tracheal organ cultures *in vitro*, and can adhere to the surfaces of glass or plastics (7). *M. pneumoniae* is asymmetric under electron microscopy (8). The cell membranes at one end can extend outside to form a proline-rich top structure, also termed the apical organ, and specifically adhere onto the neuraminic acid receptors on the membranes of target cells.

Adhesion is an intricate process, as the adhesion structure consists of an interactive adhesion network-like system and adhesion auxiliary proteins. Specifically, the 170 kDa P1 protein functions as a key ligand during adhesion (9). Pulse-tracking tag experiments have shown that 1 h following contact of *M. pneumoniae* with the target cells, the P1 precursor proteins, which are scattered in the cell membranes,

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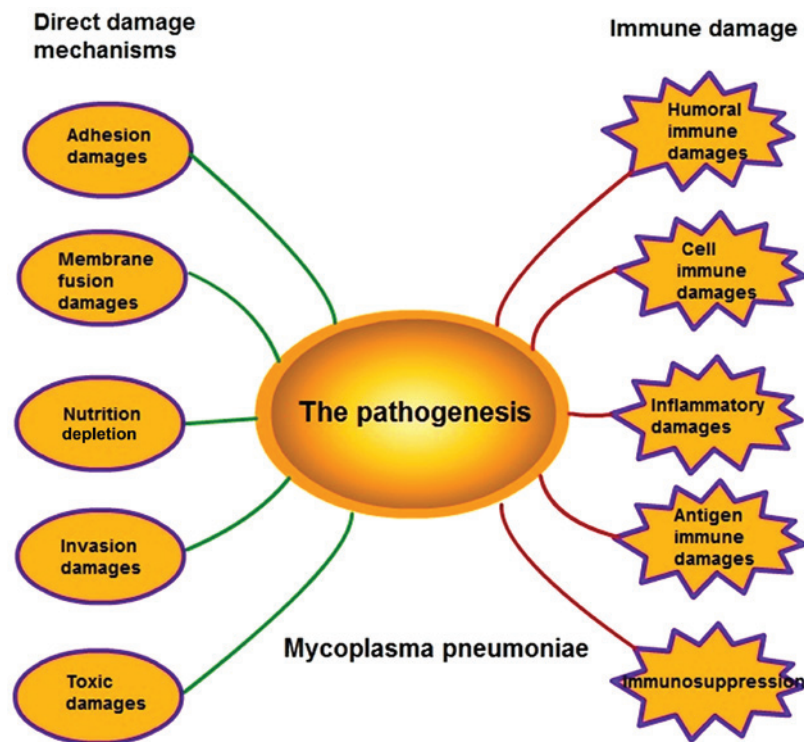


Figure 1. Pathogenesis of *M. pneumoniae*. The pathogenesis of *M. pneumoniae* comprises five direct damage mechanisms, including adhesion damage, membrane fusion damage, nutrition depletion, invasive damage, toxic damage, and five types of immune damage, including humoral immune damage, cell immune damage, inflammatory damage, antigen immune damage and immunosuppression.

rapidly shift to the apical organs, and the leading peptide on their amino terminal is hydrolyzed to mature P1 proteins (10). Due to its sole dependence on the key P1 protein, *M. pneumoniae* is unable to adhere to host cells, however, it can adhere with the assistance of several collaborative auxiliary proteins, including P30 adhesion factor-related protein A (72 KDa), B (85 KDa) and C (37 KDa), HMW 1-5 polypeptides, P40, P90 and P65; these components jointly constitute a characteristic high-electron-density 'adhesion protein complex' (Fig. 2) (11). This complex stabilizes the integrity of the *M. pneumoniae* apical structure by forming a cytoskeleton, anchoring the P1 protein into the cytoskeleton of the adhesive organs, and allowing the P1 proteins on the adhesion cell organs to adhere.

Marking experiments have shown that, in mutant strains with loss of adhesion auxiliary proteins, the P1 protein is chronically dispersed as a precursor in the cell membranes, however, it cannot aggregate to the apical organs or convert into mature P1 protein (9). Electron microscopy has demonstrated that the adhesion of a *M. pneumoniae* variant is concentrated in the adherend in the following order: HMW1, HMW3, P1, P30, P90, P40 and P65, which indicates that these proteins have formed an interrelated adhesion network (12). Specifically, HMW1, HMW2 and HMW3 function as stable adherends and allow other adhesions to locate onto the adherend, and, they are involved in the adhesion onto the respiratory tract epithelia (13). As reported, the *M. pneumoniae* mutant strains, HMW1 and HMW2, can prevent the P1 protein from correctly locating onto the apical structure, which leads to irregular cell morphology, loss of toxicity and sliding ability, and loss of adhering function (14). P30 does not directly affect the positioning of the P1 protein onto the apical structure, however, it

interferes with the binding between P1 and its receptor (15). The loss of P30 or enzymatic cleavage of the carboxyl terminal leads to the complete loss of adhesion function in *M. pneumoniae*, reduced sliding ability, and marked changes in morphology and structure (16). For example, a bifurcate structure appears in the apical tip, and numerous nucleoid-like substances appear in the cytoplasm. When transposon Tn4001 from the genes of an adhesion auxiliary protein C-mutant was used to transform *M. pneumoniae*, the mutant strain showed reduced cell adhesion ability. Following the loss of the P41 protein, the adherend in the sliding process of *M. pneumoniae* was separated from the cell (17). These adhesion auxiliary proteins and adhesion molecules jointly form adhesion protein complexes. The adsorption ability of host cells is decided by the positioning of adhesion proteins and the interaction between the components of the protein complexes.

M. pneumoniae can also utilize the MPN372 protein to combine with lung surfactant protein A (SP-A), pass through the host barrier and permanently adhere to target cells containing the SP-A receptor, including alveolar macrophages, alveolar epithelial type II cells, and other histiocytes inside and outside the lung (18). The pretreatment of *M. pneumoniae* with low-dose proenzyme reduces the binding between *M. pneumoniae* and SP-A by 80-90%, however, pretreatment with mannose does not inhibit the binding between *M. pneumoniae* and SP-A, indicating that *M. pneumoniae* protein components are involved in this process (19).

Membrane fusion damage. The cell membranes of the *Mycoplasma* genus are more durable, compared with those of other prokaryotes, and the cytoskeletal protein network-like

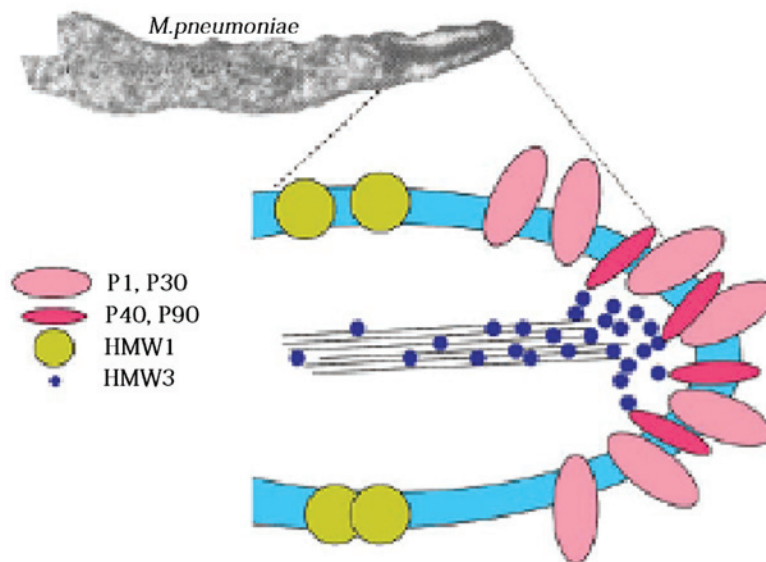


Figure 2. Structure of the *Microplasma pneumoniae* adhesion protein. The adhesion protein of *M. pneumoniae* includes key proteins P1 and P30, adhesion factor-associated proteins, P40, P90, HMW 1 and HMW 3. These components jointly constitute a characteristic high-electron-density 'adhesion protein complex'. This complex stabilizes the integrity of the *M. pneumoniae* apical organ structure by forming a cytoskeleton, anchoring the protein P1 to the cytoskeleton of the adhesive organs, and allowing the P1 proteins accumulating in the adhesion cell organs to adhere.

structure functions as a cell wall in terms of maintaining cell integrity. Following *M. pneumoniae* infection, the lipid bilayer of cell membranes is susceptible to biomembrane fusion, and its structure involves the transcription of specific genes, cytoskeletal changes and changes in the nucleolus (20). Membrane fusion can also cause changes in receptor-identifying sites in the cell membranes, affecting the signal delivery between cells and the production of cellular factors (21).

Nutrition depletion. The small-genome *M. pneumoniae* does not possess the ability to self-synthesize amino acids, fatty acids, cofactors or vitamins. Instead, following permanent adherence via the adherend to the respiratory tract epithelia, *M. pneumoniae* spreads microtubules and inserts them into host cells, enabling oxygen consumption, use of glucose, absorption of cholesterol, ingestion of amino acids and consumption of nutrients in host cells, causing injury to the host cells (22,23).

Invasion damage. *M. pneumoniae* is usually regarded as an extracellular parasite, however, certain studies have shown it can also invade and damage cells. Studies have shown that *M. pneumoniae* can invade A549 lung cancer cells, evidenced by its detection in the cytoplasm and nucleus, and the invasive ability depends on the duration and temperature of infection (24). In cell culture *in vitro*, *M. pneumoniae* has been shown to invade non-phagocytes, survive for >6 months and synthesize DNA inside cells (7). When the clinically isolated RYC15989 strain was utilized to infect human Hep-G2 cells and rat N2A cells, intracellular *Mycoplasma* were observed under laser confocal microscopy, and the intracellular invasion damaging ability of *M. pneumoniae* was also confirmed (25). In addition, during invasion, certain enzymes inside *M. pneumoniae*, including hydrolase, nuclease and phosphoprotein phosphatase shift to the host cells. Nuclease degrades DNA in host cells, whereas phosphoprotein phosphatase interferes with the activity of serine/threonine and tyrosine protein kinase (26,27).

Toxic damage. Adhesion provides conditions for *M. pneumoniae* to induce regional cytotoxic effects, and *M. pneumoniae* can directly induce damage via adhesion, auxiliary proteins, capsular and invasive enzymes. *M. pneumoniae* also exerts its toxin-like effects through its metabolites, exotoxin and exotoxin-like toxic substances, lipids, lipopolysaccharides and membrane lipoprotein (28). Following the adherence of *M. pneumoniae* onto the surface of bronchial cells, with the cytoskeleton rearrangement, *M. pneumoniae* penetrates through the bronchial mucous membranes and releases nuclease and H_2O_2 , which result in swelling, necrosis and a binding of bronchial epithelial cells, slower microvilli movement, structural deformation, and the termination of swinging, thereby inducing the infiltration of lymphocytes, plasma cells and monocytes (22,29). With the lack of superoxide dismutase and catalase in *M. pneumoniae*, the H_2O_2 and superoxide groups synthesized by *M. pneumoniae*, and the endogenous toxic oxygen molecules produced by the host cells, increase the intracellular oxygen pressure in the epithelium, which leads to oxidative stress and subsequent cell death (Fig. 3) (30). The major virulence factors affecting the pathogenesis of *M. pneumoniae* include the accumulation of H_2O_2 inside host cells and the effects of superoxides on the ultrastructure of host cells (31). The ions of *M. pneumoniae*-produced superoxides inhibit the activity and degradation of catalases in the host cells, so that the host cells become more sensitive to the toxic oxygen, resulting in mitochondrial swelling, vascular degeneration, cilia destruction and weakened cilia movement in the epithelium (32,33). *M. pneumoniae* infection leads to the denaturation of red blood cell hemoglobin, loss of reduced glutathione and cytolysis (34,35).

M. pneumoniae is considered to be incapable of secreting cytotoxin. The N-terminal of the *M. pneumoniae*-associated pathogenic factor, MPN372, contains ADP-ribose transferase activity and its structure is similar to the S1 subunit of pertussis toxin, which induces extensive vascular degeneration

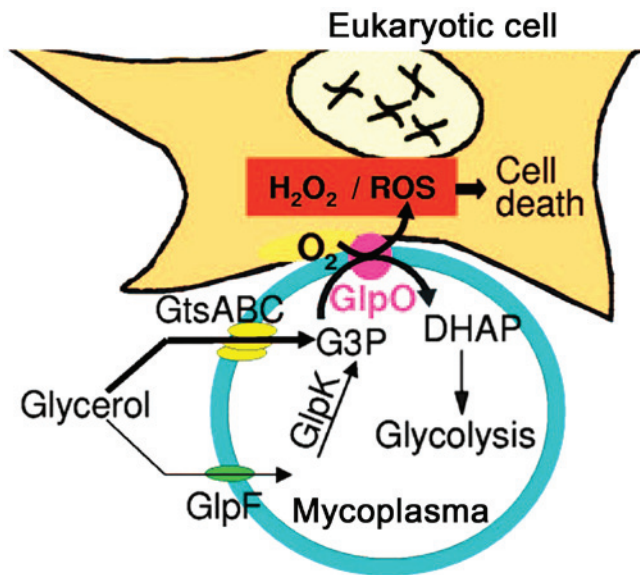


Figure 3. Toxic injury of *Mycoplasma pneumoniae* in glycerol. Following adherence of *M. pneumoniae* onto the surface of eukaryotic cells, cytoskeletal rearrangement occurs, and H_2O_2 and ROS are synthesized and released by *M. pneumoniae* in glycerol, which lead to oxidative stress and subsequent cell death. ROS, reactive oxygen species.

and can cause the death of mammalian cells, thereby inducing chinchough-like clinical symptoms; these are termed community acquired respiratory distress syndrome (CARDS) toxin (36). *M. pneumoniae* CARDS toxin is internalized via clathrin-mediated endocytosis (37), and the CARDS toxin induces pulmonary eosinophilic and lymphocytic inflammation (38,39). Cellular vacuoles induced by *M. pneumoniae* CARDS toxin originate from Rab9-associated compartments (40).

3. Immune damage

Clinical epidemiological findings show that the symptoms of *M. pneumoniae* infection are not observed at infancy, and that the pathogenic peak occurs in children >10 years old (41). In patients with reduced immune function, *M. pneumoniae* infection does not induce notable pathological changes in the lung. Experiments in thymus-excised animals have shown that *M. pneumoniae* infection does not readily induce pneumonia (42). Animal experiments have shown that the histopathologic response occurs 10-14 days following primary *M. pneumoniae* infection, but within 3 days following secondary infection, indicating that the body responds via immune cell accumulation following *M. pneumoniae* infection, but produces a more marked immune response to a second infection (43). These findings indicate that the host immune response is important during the onset of *M. pneumoniae*-induced pneumonia.

Humoral immune damage. The glycolipid antigen on the cell membranes of *M. pneumoniae* induces humoral immunity, and the antibody response is fundamental during the response against *M. pneumoniae* infection. At an early stage of *M. pneumoniae* infection, the body resists *Mycoplasma* settlement predominantly via a non-specific defense mechanism

by secreting inhibitors, alexin and phagocytes (6). Animal experiments have shown that, following infection of the body with *M. pneumoniae*, the levels of complement components C1, C2, C3 and C4 in the bronchial secretions are significantly improved (44). After 2 weeks, the level of alexin begins to decline, whereas the antibody level increases, which indicates the non-specific protective effect of alexin at an early stage of *M. pneumoniae* infection (45). In children infected with *M. pneumoniae*, the contents of C1q, C3, C4 and B in the serum increase to varying degrees in the acute phase and recovery phase, indicating that the alexin classical and bypass activation pathways are involved during *M. pneumoniae* infection (46). With the lack of alexin, the surface of neutrophils have been shown to adhere with and engulf *M. pneumoniae* under electron microscopy, and the *M. pneumoniae* in their phagocytosis vesicles remains active (47). The specific sIgA produced during *M. pneumoniae* infection can protect against infection of respiratory mucous membranes, and its action is key in indigenous resistance (48). It was previously reported, that, 28 days following *M. pneumoniae* infection in pigs, the numbers of B cells in the alveolar lavage fluid and lung parenchyma increased 25-fold (49). In addition, in the acute phase and recovery phase of mycoplasma pneumonia, the contents of IgG, IgM, IgA and immune complex in the serum increase significantly, particularly in severely affected patients (50). Following *M. pneumoniae* infection, the IgM level has been shown to markedly increase in normal children, which usually occurred 7-14 days following infection, peaked in weeks 3-4 and persisted for months (51). *M. pneumoniae* infection can cause an increase in the level of total IgE in the serum, whereas delayed-type and anaphylactic-type allergic reactions induce asthma as an immediate reaction and delayed-phase reaction or a dual-phase reaction, which induce the IgE-mediated airway inflammation and airway hyper-reactivity (52). However, there is no direct evidence that *M. pneumoniae* is the direct cause of asthma. These previous studies indicate that various specific and nonspecific immunoglobulins and complement components are involved during *M. pneumoniae* infection, which assist with the recovery and immunity.

Cell immune damage. Cellular immunity is required by the protein antigens on the cell membranes of *M. pneumoniae*. Following inoculation with *M. pneumoniae* antibody in patients infected with *M. pneumoniae*, a tuberculin-like, delayed-type allergic reaction occurs to differing degrees; the reaction is more severe in severely-affected patients, however, this reaction can be inhibited by anti-thymocyte serum (53). Tuberculin tests in patients infected with *M. pneumoniae* show that the reaction intensity directly affects the degree of lung damage, indicating that cellular immunity is vital during the pathogenesis of *M. pneumoniae* (32). In patients with *M. pneumoniae*, the CD^{4+} T cell count is decreased, the CD^{8+} T cell count is markedly increased and the ratio of CD^{4+}/CD^{8+} is reduced, and these changes are more marked in severely affected patients (53). In adults with *M. pneumoniae* infection, the peripheral blood CD^{4+} T count is decreased, however, the ratio of T-lymphocytes to CD^{4+}/CD^{8+} cells in the bronchoalveolar lavage fluid increases, possibly due to abundant CD^{4+} T cells being involved in the inflammatory reaction (54).

During *M. pneumoniae* infection, the Th1/Th2 ratio is unbalanced, although which type of cell is dominant remains controversial. It was previously reported that, following *M. pneumoniae* infection, Th1-dominated rats exhibit aggregation of peribronchial lymphocytes, whereas Th2-dominated rats exhibit hyperplasia of alveolus mesenchymal cells, which indicate that the imbalance in auxiliary T lymphocyte subgroups is associated with the type of lung damage (53). The mechanism underlying *M. pneumoniae*-induced asthma may be correlated with the enhanced secretion of Th2 cell factors (55).

Inflammatory damage. Inflammatory factors are important during the *M. pneumoniae*-induced inflammatory reaction. Polymerase chain reaction analysis has shown that, following primary *M. pneumoniae* infection in BALB/C rats, the mRNA expression levels of tumor necrosis factor- α (TNF- α), interleukin (IL)-1 β and IL-6 in the lungs were markedly increased, whereas the mRNA expression levels of IL-2 and its receptor were not increased (56). Following the second infection, the mRNA expression levels of TNF- α and IL-6 increased 10-fold, whereas the mRNA expression of IL-2 decreased rapidly within 24 h, and that of IL-10 increased markedly (56). It has been reported that *M. pneumoniae* can induce the production of IL-1 β , which is extensively involved in several types of damage, including tissue destruction and edema formation (57). Following *M. pneumoniae* infection, the serum level of IL-8 increases markedly, whereas white blood cells locate to the site of inflammation and infiltrate, accumulate and release active substances in the affected tissues, causing damage (58). The serum level of TNF- α following *M. pneumoniae* infection in the respiratory tract is significantly increased, and is positively correlated with the severity of illness (59). Serum levels of soluble IL-2 receptor (sIL-2R) can be an important indicator. In children with *M. pneumoniae* infection, the increase in the level of sIL-2R can reactivate the mononuclear cells in the circulation, and is involved in T lymphocyte dysfunction (60). In children with *M. pneumoniae* infection, the level of soluble intercellular adhesion molecule-1 is also markedly increased, which induces the increased bronchial reaction (61). *M. pneumoniae* antigens induce a potent immune reaction and enhance the Th17 cell response *in vivo* and *in-vitro*, with Treg and IL-10 being associated with the suppression of the production of IL-17A (62). The cytoadherence of *M. pneumoniae* induces inflammatory responses through TLR4 and autophagy (6). *M. pneumoniae* infection has been shown to increase inflammatory factors in a rat model of atherosclerosis and aggravate the state of atherosclerosis (4).

Antigen immune damage

Antigenic variation. The *M. pneumoniae* membrane protein is associated with invasiveness, and its variation directly affects the toxicity of *M. pneumoniae*. The molecular weight of the *M. pneumoniae* membrane V-1 antigen can change and is associated with virulence. No toxicity or pathogenesis occurs when its molecular weight is 100-200 kDa, however, toxicity and pathogenesis are observed when its molecular weight is 30 kDa. The gene mutation in V-1 antigen occurs at a 17-amino-acid repetitive sequence at the C terminal of the 94.2 kDa antigen, whereas the 27.4 kDa antigen contains two adjacent, but discontinuous, nine-amino-acid repetitive sequences, and

variation is induced by site-specific DNA inversion (63). In addition, the adsorption of RBCs by *M. pneumoniae* can alter the antigenicity of RBC membranes and induce autoantibody against RBC membrane I antigen, namely the cold agglutinin of RBCs, which induces autoimmune hemolytic anemia (64).

Immune evasion. The *Mycoplasma*-induced viscous polysaccharide capsule, as with other bacteria, is readily formed inside the host, however, it disappears rapidly *in vitro*, indicating phagocytosis by the host cells. *M. pneumoniae* readily induces variation in surface membrane antigens, in order to evade attacks from the host immune system. *M. pneumoniae* may tightly adsorb onto the surface of the host cells, depending on the specific adhesion structure, to avoid phagocytosis prior to exact antibody adjustment (65). The polymorphism of *M. pneumoniae* adhesion antigens also weakens the effects of specific antibodies (66). The glycerophosphatide on *M. pneumoniae* cell membranes shares certain antigenic components with the host cells, and thus can also evade the host's immune surveillance. The invasion of intracellular parasitism assists in enabling *M. pneumoniae* to evade the host's immune clearance and drug effect. Therefore subjecting the patients to chronically infected persons or asymptomatic carriers. Thus, the various immune evasion mechanisms of *M. pneumoniae* constitute the predominant factor explaining why *M. pneumoniae* can survive chronically inside the host.

Cross-reacting antigen. The *M. pneumoniae* membrane antigen is in antigen mimic of the RBC-membrane I antigen, and shares certain antigenic components with *Streptococcus pneumoniae* 23 or 32 and *M. genitalium* (67). As with several plants and bacteria, the membrane glycolipids of *M. pneumoniae* share a common antigen in the brain and lung tissues, which induce cross reaction. The carboxyl end of the P1 and P30 proteins in the adhesive organs of *M. pneumoniae* show high levels of homology to the cytoskeletal proteins, fibrinogen, keratin and troponin in eukaryotes (68). Thus, during infection, autoantibodies in the brain, lung, RBC-membrane, lymphocytes and myocardial cells commonly occur, which form immune complexes and magnify the autoimmune response, leading to multisystem immune damage.

Superantigen. *M. pneumoniae* membranes are full of *Mycoplasma* lipid-associated membrane proteins. At least three types of functional protein have been identified, including *M. pneumoniae* N602 (b subfamily of F0F1-ATPase), *M. pneumoniae* N162 and *M. pneumoniae* N611 (69). Specifically, the inflammatory capacity of *M. pneumoniae* N602 is higher (~100-fold), compared with that of *M. pneumoniae* N161 and *M. pneumoniae* N162, indicating that *M. pneumoniae* N602 is a potential superantigen component (70,71).

Immunosuppression. *M. pneumoniae* infection can induce immunosuppression in the body and cause maladjustment of T cell subgroups. Experiments have revealed that *M. pneumoniae* infection causes severe destruction of B cells and T cells (72). At 13-18 weeks in patients infected with *M. pneumoniae*, the serum level of IgG declines (73). Certain children infected with *M. pneumoniae* suffer from hypoglobulinemia, decreased chemoattraction in neutrophils, lower reactivity to

phytohemagglutinin phytolectin and reduced resistance against combined infections with other pathogens, including *S. pneumoniae* (72). These changes indicate that *M. pneumoniae* infection may induce immunosuppression.

4. Conclusion and perspective

As summarized in the present review, it has been demonstrated over several years that the pathogenesis of *M. pneumoniae* infection is complex; the natural synergy between the various factors involved is summarized in Fig. 1. There is no one factor alone, which is involved. As increased efforts have focussed on investigating *M. pneumoniae* gene structures and functions, and in sequencing, the various pathogenic factors of *M. pneumoniae* membrane proteins, invasive proteins and adhesive proteins can be investigated at the molecular level. This development not only assists with the treatment and prevention of *M. pneumoniae* infection, but is also meaningful for the development of *Mycoplasma* vaccines.

Acknowledgements

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