

Co-Infection of *Mycoplasma hyopneumoniae* and other swine pathogens

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Abstract

Mycoplasma hyopneumoniae (*M. hyopneumoniae*, Mhp) is an etiological agent that causes mycoplasma pneumonia of swine (MPS), this is a chronic respiratory disease present in every major swine-producing country worldwide. It is well elucidated, Mhp infections may downregulate the host immune system and enhance the infection and replication of other pathogens. However, the mechanisms of interaction between Mhp and other pathogens is still missing. Though, additional studies have reported that several cofactors such as bacteria, vaccination failure, stress or crowing and other swine viruses in combination with Mhp, lead to MPS. Aside these cofactors, the co-infection of Mhp with other viruses, such as Porcine Circovirus, Porcine Reproductive and Respiratory Syndrome Virus, Swine Influenza Virus, Pseudorabies Virus, and other bacteria, such as *Pasteurella multocida*, *Actinobacillus pleuropneumoniae* have been widely studied for many years. In this review, we summarized the co-infection of Mhp with other swine pathogens based our studies made, the co-infection with other viruses and bacteria, revealed the interaction mechanism of different pathogens with Mhp in the host

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Running title: *Mycoplasma hyopneumoniae* co-infect with pathogens

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Summary

Mycoplasma hyopneumoniae (*M. hyopneumoniae* , Mhp) is an etiological agent that causes mycoplasma pneumonia of swine (MPS), this is a chronic respiratory disease present in every major swine-producing country worldwide. It is well elucidated, Mhp infections may downregulate the host immune system and enhance the infection and replication of other pathogens. However, the mechanisms of interaction between Mhp and other pathogens is still missing. Though, additional studies have reported that several cofactors

such as bacteria, vaccination failure, stress or crowding and other swine viruses in combination with Mhp, lead to MPS. Aside these cofactors, the co-infection of Mhp with other viruses, such as Porcine Circovirus, Porcine Reproductive and Respiratory Syndrome Virus, Swine Influenza Virus, Pseudorabies Virus, and other bacteria, such as *Pasteurella multocida*, *Actinobacillus pleuropneumoniae* have been widely studied for many years. In this review, we summarized the co-infection of Mhp with other swine pathogens based our studies made, the co-infection with other viruses and bacteria, revealed the interaction mechanism of different pathogens with Mhp in the host.

Keywords

Mycoplasma hyopneumoniae (Mhp); co-infection; swine; virus; bacteria.

Introduction

Mycoplasma hyopneumoniae (*M. hyopneumoniae*, Mhp) forms part of the etiology of a chronic insidious lympho-histiocytic bronchopneumonia, also described as mycoplasmal pneumonia of swine (MPS) or enzootic pneumonia (EP) (DeBey, Jacobson, & Ross, 1992), is a strong impediment to the efficiency and profitability of the global pig industry (Simionatto, Marchioro, Maes, & Dellagostin, 2013). This disease is characterized by high morbidity but low mortality (Maes et al., 2008; Morris, Gardner, Hietala, & Carpenter, 1995), causing cough, asthma, anorexia, and many other symptoms among off-springs, as well as showing decreased daily weight gain and significant lung lesions in the field (Tao, Shu, Chen, Wu, & He, 2019). Mhp was first isolated from the trachea, bronchi, bronchioles and other mucosal surfaces (Blanchard et al., 1992), and it's difficult to be isolated from fields directly because of their slow growth and easy contamination. Despite the continuous improvement of mediums, in *in vitro* culture of Mhp still remains difficult (Calus et al., 2010; Cook, Beddow, Manso-Silvan, Maglennon, & Rycroft, 2016). This limitation, to some extent has stymied the research and the development of improved methods for disease control.

Piglets may be infected with Mhp at the early stage, with prevalence increment during the post-weaning period (Vangroenweghe, Labarque, Piepers, Strutzberg-Minder, & Maes, 2015), the main process involved in the Mhp infection is shown in Fig. 1. Mhp infection causes immunosuppression in pigs, which provides conditions for the secondary infections. Owing to the development of modern swine production, which has intensified the production of swine on a larger-scale, mixed infections by multiple pathogens are becoming more common and serious in swine. Although the exact mechanism of Mhp infection still remains unknown, many studies have reported co-infection with other pathogens for decades, these include Porcine Circovirus (PCV) (Opriessnig & Halbur, 2012), Porcine Reproductive and Respiratory Syndrome Virus (PRRSV), Swine Influenza Virus (SwIV), Pseudorabies Virus (PRV) and other bacterial pathogens, they also serve as possible co-factors for triggering postweaning multisystemic wasting syndrome (PMWS), which have further economic and welfare implications (Ciprian et al., 1988; Krakowka et al., 2000; Maes et al., 2008; Maes, Verdonck, Deluyker, & de Kruif, 1996; Pogradichniy, Yoon, Harms, Sorden, & Daniels, 2002; Wang et al., 2016; Wellenberg, Stockhofe-Zurwieden, Boersma, Jong, & Elbers, 2004). The objective of this review was to summarize the interaction during the co-infection of Mhp with other viruses and bacteria together with other factors leading to infection in recent years, to provide the theoretical basis for studies in the future.

2. Co-Infection of *Mycoplasma hyopneumoniae* with viruses

2.1 Porcine Circovirus 2

Porcine Circovirus 2 (PCV2), is a small non-enveloped DNA virus which contains a single-stranded circular genome of 1.7 kb belonging to the family of *circoviridae* (Hamel, Lin, & Nayar, 1998; Meehan et al., 1998), this virus contributes to the considerable economic losses associated with PCV2-associated disease (PCVAD/PCVD)(Chae, 2005). The most prevalent symptoms of PCVAD/PCVD include porcine dermatitis and nephropathy syndrome (PDNS), which mainly occurs during the growing or finishing stage of pigs (Segales, 2012). According to the different of Cap gene sequence, PCV2 can be divided into several subtypes, PCV 2a-h (Bao et al., 2018; Franzo & Segales, 2018; Yao et al., 2019), the PCV 2a and PCV 2b which are considered to be the most prevalent genotype (Patterson & Opriessnig, 2010; Segales et al., 2008). It has been reported that, PCV2 is crucial but not a sole factor to develop this clinical disease, it must co-infect with other pathogens in order to cause this disease(Tomas, Fernandes, Valero, & Segales, 2008).

Mhp and PCV2 are the major pathogens that cause significant financial loss (Tassis et al., 2017), coincidental infection of Mhp and PCV2 is ubiquitous in clinical conditions and contributes to a range of polymicrobial disease syndromes, such as porcine respiratory disease complex (PRDC) and PMWS (Chae, 2005; Krakowka et al., 2007; Segales, 2012). The co-infection rate of Mhp and PCV2 has been reported to be 35.5% (172/484) in US field cases (Pallares et al., 2002). Fablet and colleagues found that, the detection rate of Mhp DNA in a pig was 33.6% which meant that the PCV2 genome load in sera were 4.310^4 copies/mL (Fablet, Marois-Crehan, et al., 2012). The pigs' lymph nodes were tumefied compared to the normal size, with their lungs having some cranioventral dark purple consolidation, and accompanied by shedding of high levels of PCV2 DNA in semen in coinfection with Mhp and PCV2 cases (Opriessnig, Madson, et al., 2011). Other group also demonstrated, the severity of PCV2-associated lesions in lung and lymphoid tissues of pigs, with a number of antigens within these lesions being significantly higher compared to the control group (Opriessnig et al., 2004).

Both PCV2 and Mhp can target host immune cells and impair host defenses, resulting in a significant increase in the expression of Interferon-gamma (IFN- γ), Interleukin-1beta (IL-1 β), IL-8, Chemokine (C-C motif) ligand 5 (CCL5) and Chemokine (CXC motif) ligand 10 (CXCL10), weak stimulation of IFN- β , IL-6 and IL-10 and downregulated of IL-13 and IFN- α significantly (H. R. Zhang, Lunney, Baker, & Opriessnig, 2011). Furthermore, Mhp enhanced the levels of PCV2 viraemia, whereas PCV2 did not enhance the levels of mycoplasmal nasal shedding in co-infected pigs, compared to singly Mhp or PCV2-infected pigs (Seo, Park, Park, & Chae, 2014). PCV2 replication could be enhanced by subsequent inoculation with Mhp, however, simultaneous PCV2 and Mhp co-inoculation does not potentiate disease in conventional pigs (Sibila, Fort, Nofrarias, Perez de Rozas, et al., 2012; Wang et al., 2016). Other researchers also indicated that, concurrent infection with PCV2 and Mhp did not result in potentiation of clinical signs and lesions attributed to either infection in conventional pigs (Sibila, Fort, Nofrarias, de Rozas, et al., 2012). These results indicated that synergistic effects occur during the process of Mhp and PCV2 sequential infection, and some previous studies also demonstrated that Mhp infection in pigs normally occurs slightly before or around PCV2 infection in the field condition (Chae, 2012; Fachinger, Bischoff, Jedidia, Saalmuller, & Elbers, 2008; Larochele, Magar, & D'Allaire, 2003). Therefore, the study of the interaction between Mhp and PCV2 co-infection based on sequential infection model may be closer to the field condition in pig industry.

2.2 Porcine Reproductive and Respiratory Syndrome Virus

Porcine Reproductive and Respiratory Syndrome Virus (PRRSV), is an enveloped, single-stranded, positive sense RNA virus in the family of *Arteriviridae* (Kavanova et al., 2018) and an etiologic agent that causes porcine reproductive and respiratory syndrome (PRRS), which has been estimated to be the cause of a lost \$600 million in the US swine industry at least every year (Neumann et al., 2005). The typical symptoms of

PRRSV are characterized by blue-ear and failure in pregnant pigs (Lunney et al., 2016), also these infections can induce cell lysis, apoptosis, changes in T-cell subpopulations etc (Labarque, Van Gucht, Nauwynck, Van Reeth, & Pensaert, 2003; Shimizu et al., 1996; Thanawongnuwech, Thacker, & Halbur, 1997).

It has been reported that, the concurrent infection with Mhp and PRRSV is common (Chae, 2016; Fablet, Marois-Crehan, Grasland, Simon, & Rose, 2016b; Pallares et al., 2002). Scott and colleagues have proved their hypothesis that PRRSV and Mhp can be transported via airborne route (Dee, Otake, Oliveira, & Deen, 2009; Otake, Dee, Corzo, Oliveira, & Deen, 2010), these characteristics of pathogens increased the difficulty to control these diseases. PRRSV replicates mainly in porcine alveolar macrophages (PAMs), dendritic cells (DC) in lungs and the upper respiratory tract (Lunney et al., 2016), resulting in viremia as early as 12 h post infection at the early infection stage (Wills, Doster, Galeota, Sur, & Osorio, 2003). For the persistence period, viral replication is primarily localized in lymphoid organs, including tonsil and lymph nodes but not spleen (Allende et al., 2000; Rowland, Lawson, Rossow, & Benfield, 2003; Wills et al., 1997) and gradual decays until the virus becomes extinct in the host (Christopher-Hennings, Nelson, Althouse, & Lunney, 2008; Wills et al., 2003).

The PAMs from pigs infected with PRRSV have significantly downregulated the ability to kill bacteria (Solano, Bautista, Molitor, Segales, & Pijoan, 1998). Although, some researchers have demonstrated that co-infection with Mhp and PRRSV show no potentiating effect (Van Alstine, Stevenson, & Kanitz, 1996). Thacker and colleagues suggested that the Mhp potentiated and prolonged PRRSV-induced pneumonia clinically, macroscopically and microscopically regardless of the inoculation sequence of infection to both pathogens, whereas PRRSV did not aggravate Mhp infection in piglets when inoculated before Mhp under experimental conditions (Thacker, Halbur, Ross, Thanawongnuwech, & Thacker, 1999; Tzika et al., 2015; Van Alstine et al., 1996). Furthermore, Fablet suggested that, the infection by Mhp was associated with PRRSV seropositive status (Fablet, Marois-Crehan, Grasland, Simon, & Rose, 2016a). Moreover, in the concurrent infection with Mhp and PRRSV, many vital functional genes were detected as being differentially expressed (DE) in PAMs (Li et al., 2015), especially, IL-1 β was considerably higher, which is a key component for downstream signal pathways. In addition, the co-infection of Mhp and PRRSV significantly increased the severity and duration of pneumonia in experimentally infected pigs, which was associated with induction of several proinflammatory cytokines (Thacker, Thacker, Kuhn, Hawkins, & Waters, 2000), such as IL-1 α , IL-1 β , IL-6, IL-8, IL-10, IL-12 and Tumor Necrosis Factor-alpha (TNF- α) (Thanawongnuwech, Thacker, Halbur, & Thacker, 2004; Thanawongnuwech & Thacker, 2003; Thanawongnuwech, Young, Thacker, & Thacker, 2001). Other group also found increased levels of both IL-12 and IL-10 in the respiratory tract of pigs experimentally infected with either Mhp and/or PRRSV, IFN- γ and Insulin-like Growth Factor-I (IGF-I) production were lower and delayed in pigs co-infected with PRRSV and Mhp (Roberts & Almond, 2003; Thanawongnuwech & Thacker, 2003). These results suggest that the exacerbation Mhp respiratory disease may be due to viral infection which induces regulatory T cells (Tregs) (LeRoith et al., 2011). In contrast, Fano et al suggested that Mhp did not affect these epidemiological features of PRRSV-associated disease under the conditions of study (Fano, Pijoan, & Dee, 2007).

Other studies also implicated that a single-dose vaccination against Mhp alone decreased the levels of PRRSV viremia and PRRSV-associated pulmonary lesions, whereas single-dose vaccination against PRRSV alone did not decrease nasal shedding of Mhp and Mhp-associated pulmonary lesions in the co-infected pigs (S. J. Park, Seo, Park, & Chae, 2014). Combining vaccines with Mhp and PRRSV did not induce negative interaction which would reduce the efficacy of each individual vaccine (Bourry, Fablet, Simon, & Marois-Crehan, 2015). These results indicated that, the combined vaccination is more efficient than single ones, consequently, further research should focus on the developing of combined vaccine, which can improve the convenient and efficient pig production simultaneously.

2.3 Swine Influenza Virus

Swine influenza virus (SwIV), which belongs to *Orthomyxoviridae* family, causes seasonal epidemic or occasional pandemic outbreaks in pigs worldwide (Taubenberger & Morens, 2008). SwIV can be subdivided into many subtypes, including influenza A, B, and C (Baudon, Peyre, Peiris, & Cowling, 2017; Meiners et al., 2014).

PRDC is an economically enormous problem accompanied by slow growth performance, cough, poor food utilization etc (Baudon et al., 2017). It has been reported that SwIV and Mhp play important roles in PRDC (Deblanc et al., 2016; Deblanc et al., 2012; Fablet, Marois-Crehan, et al., 2012). SwIV can target and replicate in epithelial cells of the upper respiratory tract (Brown, Alexander, Chakraverty, Harris, & Manvell, 1994). Coinfection with SwIV and Mhp was detected in 23 (31%) cases in 74 lungs from 2009 to 2015 retrospective analysis (Rech et al., 2018). Although the interaction between SwIV and Mhp is minimal or even appear independent of each other (Thacker, Thacker, & Janke, 2001), the pre-infection with Mhp was remarkably exacerbated by the clinical symptoms of pigs with H1N1 infection during the first week after virus inoculation (Deblanc et al., 2016). Furthermore, the pig lung lesion caused by inoculation with Mhp and SwIV was more severe than those inoculated with Mhp only (Yazawa et al., 2004). Deblanc and co-workers indicated that clinical signs and macroscopic lung lesions were similar in early time post-H1N1 inoculation compared to pre-Mhp infection or not pig group, and Mhp didn't affect the influenza virus replication and the IFN-induced antiviral responses in the lung, however, there is a higher inflammatory response to H1N1 infection in pre-Mhp infection group. The exact mechanism can be revealed by the massive infiltration of neutrophils and macrophages into the lungs and the increased production of pro-inflammatory cytokines (IL-6, IL-1 β and TNF- α) (Deblanc et al., 2016). Moreover, Deblanc demonstrated that Mhp and H1N1 appeared to act synergistically, as Mhp and H1N2 would compete in pigs that were previously infected with Mhp, resulting in the elimination of Mhp in the lung diaphragmatic lobes by the H1N2 (Deblanc et al., 2012). Furthermore, the miRNAs were differentially expressed (DE) and most of them were downregulated to defend the H1N1 in pulmonary alveolar macrophages during the process of H1N1 infection (Jiang et al., 2015). According to the different level of oxidative stress induced by pre-infection of Mhp, there were different outcomes from the subsequent infection with H1N1 subtype (Deblanc et al., 2013).

These results demonstrated that SwIV did not affect Mhp replication in the co-infected pig, but both Mhp and SwIV decreased or disrupt the function of the mucociliary apparatus and immunosuppression which potentially led to an increased secondary infections from opportunistic organisms under field conditions (DeBey & Ross, 1994).

2.4 Pseudorabies Virus

Pseudorabies virus (Suid herpesvirus 1 or PRV) is a member of the genus *Varicellovirus*, and the family *Herpesviridae* (Peng et al., 2016). PRV is the major causative agent of Aujeszky's disease and contributes to the substantial economic losses in swine production (Freuling, Muller, & Mettenleiter, 2017).

PRV always targets the mucosal epithelium of the pig respiratory and nervous system tissue, causing central nervous system infection and respiratory disease (Pomeranz, Reynolds, & Hengartner, 2005). PRV infection can downregulate the function of macrophage, under conditions such as phagocytosis, killing of phagocytized bacteria, IFN- α production and phagosome fusion were downregulated in PRV infected pigs (Fuentes & Pijoan, 1986; Iglesias, Pijoan, & Molitor, 1992). Shibata and colleagues have indicated that the mean percentage of the lung lesions were 0.1% and 8.3% inoculate with Mhp alone, whereas 9.8% and 17.2% in co-infection with Mhp and PRV in post-inoculation-week (PIW) 2 and 4 respectively (Shibata et al., 1998). Consequently, PRV infection appears to have effect on the severity of experimentally induced acute Mhp in young pigs. Further studies will prove the exact mechanism involved in the interaction of PRV with Mhp, providing reference for the developing of vaccine which can combat both pathogens.

2.5 Co-Infection with Multiple Viruses

Alongside single and dual infections, multiple (including triple or more viruses) infections are also prevalent among pigs (Opriessnig, Gimenez-Lirola, & Halbur, 2011). PRDC is caused by the combination of infectious pathogens, differences in production systems and environmental factors (Hansen et al., 2010). Previous study have demonstrated that PRRSV and Mhp were well known to potentiate PCV2-associated lesions, surprisingly, PRRSV vaccine was found to enhance PCV2 replication and it can be related to the failure of PRRSV vaccine in vaccinated co-infected pigs compared to non-vaccinated co-infected pigs (C. Park, Oh, Seo, Han, & Chae, 2013). Pallares et al have detected that the coinfection rate of PCV, PRRSV and Mhp was 16% (77/484) in 484 cases diagnosed as PMWS (Pallares et al., 2002). A survey also demonstrated that co-infection ratios with three and four pathogens were 17.3% and 7.3% respectively in 110 pneumocystis spp. positive lung samples of Austrian pigs with pneumonia (Weissenbacher-Lang et al., 2016). Vaccination against Mhp significantly protected pigs against multiple viral infections, suggesting that vaccination against Mhp decreases the risk of PMWS and PRDC, and reduces susceptibility of pigs to the other viral pathogens (Chae, 2011). These data indicate that Mhp plays an important role in the co-infection of dual or multiple pathogens *in vivo*.

3. Co-Infection of *Mycoplasma hyopneumoniae* with bacteria

Pasteurella multocida (PMULT), is a capsulated, Gram-negative coccobacillus, this primary pathogen can cause debilitating and fatal porcine pneumonia, especially resulting in pleuritic (Harper, Cox, Adler, & Boyce, 2011; Ross, 2006). Mhp and PMULT are associated with pneumonia at both the pig and herd levels, also the coincidental infections of Mhp and bacteria have also been studied for a long period (Fablet, Marois, et al., 2012). In cases with bacterial pneumonia, PMULT was the most prevalent infectious agent (Mores et al., 2015; Tocqueville, Kempf, Paboeuf, & Marois-Crehan, 2017), while the co-infection with Mhp and PMULT in gilts and sows can render the lungs more susceptible to PMULT colonization and infection (Ciprián, Pijoan, Cruz, Camacho, & Garza, 1988). Though pigs that have recovered from or vaccinated against Mhp infection were resistant to PMULT infection (Amass et al., 1994). In addition, Park and co-workers have elucidated the pathogenic mechanisms through which the Mhp increases the L-fucose composition to enhance adherence of PMULT type A to the bronchial and bronchiolar epithelial cells (C. Park et al., 2016). Moreover, Eileen et al reported that food which contain doxycycline was effective in fattening pigs to some extent and also controlling pneumonia which was due to PMULT and Mhp (Bousquet et al., 1998).

Actinobacillus pleuropneumoniae (APP) is a small, Gram-negative, encapsulated rod with typical coccobacillary morphology and an aetiological agent of porcine pleuropneumonia (Sassu et al., 2018). Concurrent infection with Mhp and APP is common, both of them are responsible for PRDC (Hege, Zimmermann, Scheidegger, & Stark, 2002). Mhp and APP are considered to be the most important primary bacterial respiratory pathogens associated with lung lesions (Fablet, Marois, et al., 2012), co-inoculation with Mhp and APP can induce more severe respiratory disorders in pigs (Haimi-Hakala et al., 2017). The detection rate of 0.1% for APP and 2.6% for Mhp were discovered in 3983 farms across Switzerland (Hege et al., 2002). Marois and colleagues indicated that concurrent infection with Mhp and APP had more severe lesions compared to single does infection in experimental pigs, pigs which were infected with the APP still remained healthy and lung lesions were only observed after the co-infection with Mhp (Marois et al., 2009). Meanwhile, the phagocytic abilities of alveolar macrophage had decreased in pigs co-infected with Mhp and APP (Caruso & Ross, 1990).

Lawsonia intracellularis (LI) are Gram-negative, obligate intracellular bacteria that cause proliferative enteropathy (PE), an economically important disease for the pig industry (Obradovic & Wilson, 2019). Dual infections with LI and Mhp caused the epithelial thickening and post-absorptive metabolic functions altering in pigs that result in reduced nutrient absorption, reductions in growth performance and feed efficiency (Helm, Curry, Schwartz, Lonergan, & Gabler, 2019; Helm, Outhouse, Schwartz, Lonergan, et al., 2018).

Helm et al confirmed that, a dual enteric and respiratory pathogen challenge reduced average daily weight gain (ADG), average daily feed intake (ADFI), Gain:Feed (G:F) and tissue accretion in growing pigs(Helm, Outhouse, Schwartz, Dekkers, et al., 2018).

Moreover, studies have shown that immunosuppression caused by the predisposing of Mhp infection, can downregulate the phagocytic response , causing more serious clinical symptoms through the exposition to other bacterial pathogens, such as *Bordetella bronchiseptica* ,*Haemophilus parasuis* , *Trueperella pyogenes* and streptococci or staphylococci in field outbreaks of MPS(Caruso & Ross, 1990; Maes et al., 2018). It was also, reported that concurrent infections with different Mhp strains have been detected in the same field (Vranckx et al., 2011; Vranckx, Maes, Sacristan Rdel, Pasmans, & Haesebrouck, 2012), resulting in higher severity and prevalence of Mycoplasma-like lung lesions in slaughter pigs (Michiels, Vranckx, et al., 2017). Annelies and colleagues also demonstrated that, pigs that were co-infected with highly virulent strain F7.2C and the low virulent strain F1.12A, showed more severity lung lesions, coughed and larger log copies of Mhp in the bronchoalveolar lavag (Michiels, Arsenakis, et al., 2017). However, simultaneous infection with *M. hyorhinitis* and Mhp did not aggravate the observed lung lesions (Luehrs et al., 2017), the results indicated that different virulent strain can interact with each other, but not between different species of mycoplasma. Further studies is needed to focus on the exact mechanism of how these bacterial pathogens interact with Mhp.

4. Co-Infection of *Mycoplasma hyopneumoniae* with others

Apart from co-infection of Mhp with viruses or bacteria, there are many pathogens which comes from the pig farms diet (Michiels et al., 2018). The presence of mycotoxin in foods and feeds have become a serious global challenge for animals and humans health not only in the developing as well as the developed countries(Stoev, 2013). Fumonisin mycotoxin is a metabolite produced mainly by *Fusarium moniliforme* *Sheld* , which can be subdivided into several subtypes, such as Fumonisin B₁ (FB₁), Fumonisin B₂(FB₂) and Fumonisin B₃(FB₃). The FB₁ is considered to be the main pathogen for inducing porcine pulmonary edema(Harrison, Colvin, Greene, Newman, & Cole, 1990) and pulmonary fibrosis that develops in cases of chronic exposure (Zomborszky-Kovács et al., 2002). Several groups have studied the interaction between Mhp-infection and FB₁-fed pig, their results revealed a strong oedematous changes in the interstitium of lung in addition to deteriorated and extended bronchointerstitial pneumonic process and severe illness requiring euthanasia observed in one pig and evidence of progressive pathology in two pigs between study days 44 and 58 by using computed tomography (CT) and histopathologic review (Posa et al., 2013b; Posa et al., 2016). Therefore, FB₁ aggravated the progression of infection. On the contrast, Michiels and co-workers indicated that the pigs which received feed contaminated with the mycotoxin deoxynivalenol (DON) did not show more severe disease and lesions under experimental pre-Mhp infection compared to the pigs which were fed with non-contaminated feed(Michiels et al., 2018). Furthermore, the opportunistic fungal pathogen Pneumocystis can also be co-infected with Mhp, related interstitial pneumonia in both the Pneumocystis positive lungs and lungs with a mild Mhp infection (Kureljusic, Weissenbacher-Lang, Nedorost, Stixenberger, & Weissenbock, 2016). Consequently, the health level of pigs to some extent can be determine by the kind of feed administered.

Parasitism can also play an important role to negatively impact the pig's ability to respond to respiratory pathogens (Tjornehoj, Eriksen, Aalbaek, & Nansen, 1992). Some adult lungworms that are localized within the terminal bronchioles do elicit bronchitis, bronchoalveolitis, alveolar emphysema and atelectasis(Opriessnig, Gimenez-Lirola, et al., 2011). Furthermore, the pattern of gene expression in the lungs and draining lymph nodes indicated a local Th2-skewed response induced by *Ascaris suum* , infection with *A. suum* significantly compromised the effect of Mhp vaccination (Steenhard et al., 2009), which indicated that it is necessary to control parasite to provide a better environment for pig in farms.

5. Conclusions and Future Perspectives

In summary, many studies have demonstrated that Mhp alone is not enough to induce PRDC or PMWS (Krakowka et al., 2000; Posa et al., 2013a; Wellenberg et al., 2004). The major route of Mhp entry is through the mucosal surfaces of respiratory tract followed by adhering to the cilium of epithelial cells (N. Zhang et al., 2019), the cilium can be gobbled up and shed due to a large number proliferation of Mhp (Tao et al., 2019) and also the duration of Mhp infection can last for not less than 254 days (Pieters, Pijoan, Fano, & Dee, 2009). Mhp infection increases susceptibility of the pig to secondary infections, clinical signs of PCV2, PRRSV, SwIV, PRV and bacterial pathogens infections. Meanwhile, the secondary infection and worse habitat condition also provide a better environment for Mhp infection. The main target sites for these swine pathogens are shown in Table. 1. In addition, once infected with Mhp, pigs can excrete the pathogen lasting for 254 days post-infection (Pieters et al., 2009), this may be a high risk for off-springs infection. Moreover, farming system is a major factor influencing infection and sero-positivity for Mhp infection on pig farms, the virulence status of Mhp isolates may also affect the clinical pattern of infection (Vicca et al., 2002). Consequently, more researches are needed to improve our understanding of the interactions between different swine viruses and bacteria during co-infection with Mhp, encompassing how they interact with the host immune response and how they affect the efficacy of vaccination. At the same time, other experimental models should be developed and analyzed, and field trials can also be conducted to establish best practices for controlling this complex disease syndrome. The thrust of this review may provide point ideas in understanding of MPS and the developing of new strategies to control the disease.

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Conflict of interest statement

The authors declare no conflict of interest.

Data availability statement

The datasets supporting the conclusions of this article are available in the Open Science Framework repository.

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Table 1

Main target sites for various swine pathogens in pig.

	Pathogens	Abbreviation	Target
Viruses	Porcine Circovirus Type 2	PCV2	Lymphoid tissues
	Porcine Reproductive and Respiratory Syndrome Virus	PRRSV	Tonsil, upper respiratory tract and lung
	Swine Influenza Virus	SwIV	Epithelial cells of the upper respiratory
	Pseudorabies Virus	PRV	Mucosal epithelium, nervous system tis
Bacteria	<i>Pasteurella multocida</i>	PMULT	Lungs
	<i>Actinobacillus pleuropneumoniae</i>	APP	Tonsils, necrotic lung tissue and the na
	<i>Lawsonia intracellularis</i>	LI	Intestinal epithelial cells

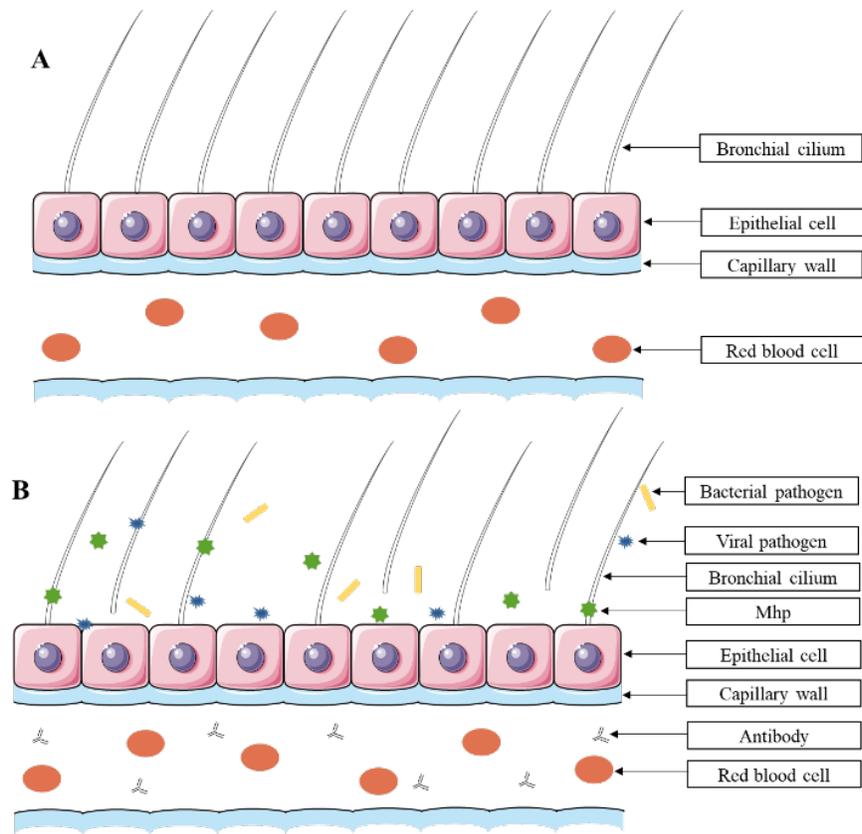
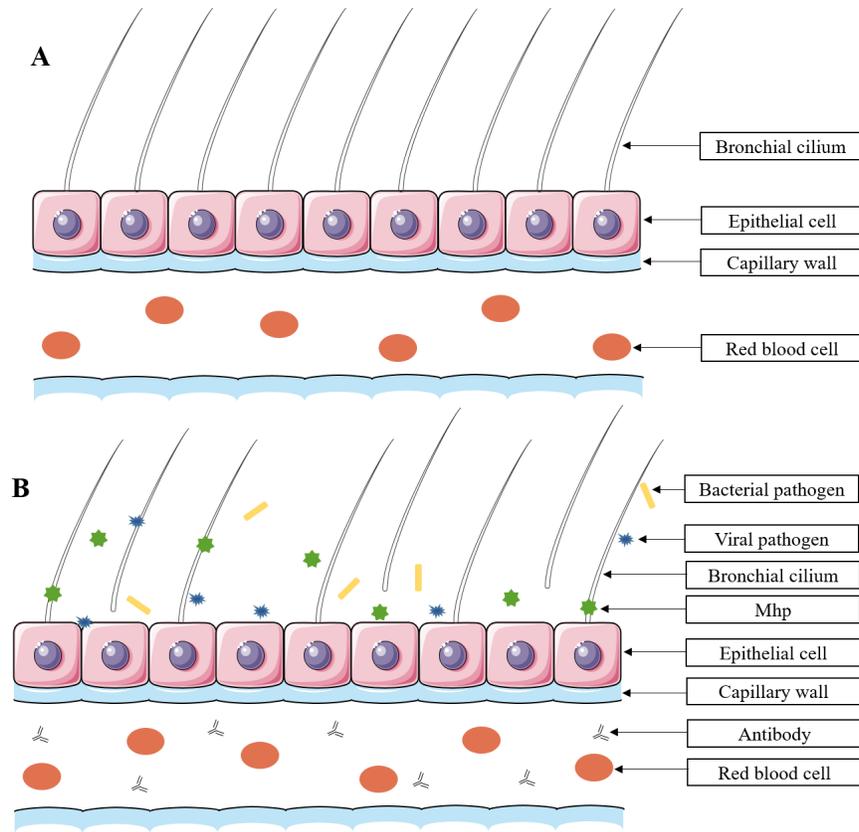


Fig. 1. The main process involved in the Mhp infection. (A) Mhp infection before. (B) Co-infection of Mhp and other swine pathogen.



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Table 1Main target sites for various swine pathogens in pig.docx available at <https://authorea.com/users/322723/articles/451679-co-infection-of-mycoplasma-hyopneumoniae-and-other-swine-pathogens>