

A Current Review on Mycolic Acid Immunogen of *Corynebacterium pseudotuberculosis*

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Abstract

Caseous lymphadenitis (CLA) is a disease of sheep and goats caused by *Corynebacterium pseudotuberculosis*. Members of *Corynebacterium*, *Mycobacterium*, *Nocardia* and *Rhodococcus* are the CMNR group of bacteria, which are so-called because they possess an outer cell membrane containing mycolic acids (MAs). This outer membrane presumably acts as a permeability barrier that imparts high drug resistance levels to some members of this group. The distinguishing feature of *C. pseudotuberculosis* is that MAs and phospholipase D (PLD) encoded by a β -corynephage in its genome are putative carriers of the toxins. Almost all members of the CMNR group possess a mycolic acid layer or mycomembrane around the cell wall, which is the main virulence determinant and plays a vital role in bacterial survival. Both PLD and MAs are virulence factors that play crucial roles in the pathogenesis of *C. pseudotuberculosis*. Phospholipases are glycopospholipid-hydrolyzing enzymes that facilitate bacterial dissemination in the host and assist the bacteria to evade phagocytosis by depletion of complement and impaired chemotaxis of neutrophils. MAs are waxy coat that protects the bacterium from hydrolytic enzymes within lysosomes and enables bacteria to skip phagocytosis and survive within the host cell. This review presents current research information on the mycolic acids of *C. pseudotuberculosis* and their role in the pathogenesis of the disease which includes host cell responses of acute-phase proteins, pro-inflammatory cytokines, and reproductive hormones changes and cellular tissue changes.

KEYWORDS

Corynebacterium pseudotuberculosis, immunogen Mycolic acid, host cell, responses, acute phase proteins, pro-inflammatory cytokines, reproductive hormones and cellular changes.

BACKGROUND

Corynebacterium pseudotuberculosis is a small, non-spore-forming, curved, gram-positive, intracellular anaerobic rod-shaped bacterium that causes caseous lymphadenitis (CLA) among small ruminants (Collins *et al.*, 1982; Dorella *et al.*, 2009; Jesse *et al.*, 2011; Odhah *et al.*, 2017). The transmission of *Corynebacterium pseudotuberculosis* occurs through direct exposure to the bacterium via physical contact with the infected animal or indirect-

ly through fomites contaminated with organisms released from rupture of abscesses (Stoops, and Thilsted, 1984; Collet *et al.*, 1994; Jesse *et al.*, 2008; Saad *et al.*, 2013).

Corynebacterium diphtheria and *Corynebacterium pseudotuberculosis* belong to the suborder Corynebacterinae, which are pathogenic bacteria species enveloped in a waxy substance known as mycolic acids (MAs) (Daffé and Draper, 1997). The MAs are built in an intricate structural design that renders the bacteria cell membrane impermeable to external assaults (Damien *et al.*,

2004; Jesse et al., 2013a). Mycolic acids are extracted using organic solvents such as glycerol or by esterification of the terminal Penta-arabinofuranosyl units of arabinogalactan (McNeil et al., 1991; Marrakchi et al., 2014; Faeza et al., 2019b). Mutant strains of *Corynebacterium* lack MAs (Portevin et al., 2004; Portevin et al., 2005; Zuber et al., 2008).

The chemical structure of MAs was first elucidated in 1950 (Asselineau and Lederer, 1950). Mycolic acids consist of 3-hydroxyl molecules, a feature that confers its property of being cleaved at extreme temperature, by a reaction similar to a reverse Claisen-type condensation, into a "mero" aldehyde main chain, also called a "meromycolic chain," and a "fatty acid" (Asselineau et al., 1970). Recently, structures of MAs of genera other than that of Mycobacteria were showed to be relatively simple in terms of chemical functions, composing only of a homologous series with various numbers of double bonds, with up to 6 for *Tsukamurella* species (Tomiasu and Yano, 1984; Butler and Guthertz, 2001). In contrast, MAs of Mycobacteria have chains that define MAs' different classes, leading to intricate TLC patterns (Minnikin et al., 1983; Barry et al., 1998; Burkovski, 2013;). Non-polar MAs referred to as α -MAs, contains about 74-80 carbon atoms and two double bonds. A fraction of α -MAs may contain unsaturation, as observed in some strains of the *M. tuberculosis* complex (Watanabe et al., 2001). Polyunsaturated α -MAs represents a significant portion of the MAs in *Mycobacterium fallax* and MAs with 60-62 carbon atoms, known as α MAs, contain one cis double bond. *Segniliparus* was recently shown to have an array of MAs with chain lengths like those of Mycobacteria (Butler et al., 2005; Hong et al., 2012; Laneelle et al., 2013).

Mycolic acids from most mycobacteria possess supplementary oxygen functions located in the distal part of the meromycolic chain, signifying the methoxy and keto MAs wax ester-, epoxy-, and hydroxy-types of MAs (Daffe et al., 1983). The oxygen functions occurring in mycobacterial MAs are typified by the occurrence of an adjacent methyl branch. It is worthy of mentioning that in *M. tuberculosis*, the oxygenated MAs, i.e., hydroxyl MAs and keto, methoxy, contain (84-88) carbon atoms, and are thus four to six carbons longer than the α -MAs from the same strains (Laval et al., 2001; Faeza et al., 2019a). However, there is still a lack of information on the mechanism of action and host cell responses towards Mycolic acid (Carne, 1939; Onon, 1979).

The exotoxin of the bacteria which act as the virulence factor contributes to the penetrability factor for the establishment of the infection in the infected host. In addition to that, mycolic acid and phospholipase D causes necrosis of endothelial cell which contributes to the movement of *C. pseudotuberculosis* through the skin reaching other minute blood vessels supplying the lymph vessels (Egen et al., 1989; Songer, 1997; Komala et al., 2008; Odhah et al., 2017). Furthermore, phospholipase D exerts cytotoxic effects on leukocytes and enhances the lysis of macrophages experimentally challenged goats (Mahmood et al., 2015b). The previous study done using specific anti phospholipase D antibodies showed that exotoxin was essential for the occurrence development of CLA in the host. Vaccination of goats and sheep with inactivated exotoxins prevented the infiltration of bacteria experimentally (Brown et al., 1986). Phospholipase D and MAs serves as determinants of virulence of *C. pseudotuberculosis* because the progeny of PLD-knocked bacteria were unable to produce typical CLA lesion but instead an immunologic reaction (Fontaine and Baird, 2008; Faeza et al., 2019b). These findings suggested that the mutant bacteria may be utilized as a live vaccine specimen for CLA (Hodgson et al., 1992). On the other hand, the virulence status of PLD activity can be demonstrated by the induction of skin necrosis, and direct damage to the "endothelial"

cells. Phospholipase D works "synergistically" with cholesterol oxidase through hydrolysis, and this combined role results in β -hemolysis of small ruminant red blood cells by *Rhodococcus equi* (Bastos et al., 2012). The phosphate ceramide produced during transition is hydrolyzed by phospholipase C of *Rhodococcus equi* (Brown et al., 1987; Pawelczyk and Kremer, 2014). *Corynebacterium pseudotuberculosis* is an intracellular bacterium that multiplies in macrophages and survives by resisting the activities of phagolysosomes due to the cell walls which consist of an external lipid layer. As a result, this facilitates the spread of bacteria into the lymphatic system and the regional lymph nodes (Dorella et al., 2006a; Baird and Fontaine, 2007). After penetration of the causative organism into the host through the nasal cavity, mouth, and ocular mucosa or through the skin, the organism spreads freely or within the macrophages to invade the lymph and reach the internal organs and local lymph nodes (Faeza et al., 2019a; Jesse et al., 2020). The pathogenicity of *C. pseudotuberculosis* depends on its efficiency to affect macrophages, halt phagolysosome activity, and cause tissue necrosis (Batey, 1986a; Jesse et al., 2013b). On the other hand, further spread of the organism from the primary site of entry to the location of CLA lesion depends upon its survival and recurrence within the macrophages facilitated by mycolic acid (Baird and Fontaine, 2007; Jesse et al., 2011). The pyogenic lipid cell layer of the organism is immunogenic and inhibits phagocytosis by elevating its cytotoxicity and destruction of macrophages in abscesses by the release of lysosomal enzymes. It has also been reported that the participating mycolic acid in pathogenicity proves relevant for bacterial survival (Fontaine and Baird, 2008; Stefńska et al., 2010). Due to the paucity of knowledge and the gap of CLA, this review focused on the current findings on mycolic acid from *C. pseudotuberculosis* towards host cell responses such as reproductive hormones, pro-inflammatory cytokines, acute-phase protein and cellular changes.

Corynebacterium pseudotuberculosis

General Characteristics of C. pseudotuberculosis

Corynebacterium pseudotuberculosis has been studied for generations dating back to 1888 when it was first discovered by a French bacteriologist named Edward Nocard who identified it as a peculiar organism isolated from an incidence of lymphatic vessels in a cow. Three years later, Hugo V. P., a bacteriologist from Bulgaria isolated a similar organism from a case of kidney abscess in a ewe and the bacterium was named "Preisz-Nocard" bacillus (Baird and Fontaine, 2007; Papaioannou et al., 2010). In 1948, the bacterium was renamed *C. pseudotuberculosis* which has remained its official designation even though some works of literature still refer to it as *Corynebacterium ovis* (Euzéby, 2003; Euzéby et al., 2004).

C. pseudotuberculosis is a pathogen of socio-economic importance responsible for caseous lymphadenitis (Jesse et al., 2008) with a significant economic loss towards the farmer. Globally, this disease is of serious importance as it is found threatening the production of sheep and goats (Baird and Fontaine, 2003; Latif et al., 2015) with high morbidity resulting in wool loses, milk and meat devaluation hence economic downturn for the farmers (Paton et al., 2003; Paton 2010). In many countries, this disease is still an important problem that is affecting the small ruminant and mutton industry (Ribeiro et al., 2013).

The organism has been isolated from sheep and goats had been done (biovar ovis) and some reports indicate that it can also be isolated from horses and cattle (biovar equi) (Dorella et

al., 2006b; Latif et al., 2015). Many researchers have defined related bacteria from bovine lymphangitis (Yeruham et al., 1996) and ovine renal abscesses (Baird and Fontaine, 2007). So far, the bacterium has been isolated from ulcerative lymphangitis and purulent infections in man and some mammalian species such as goats, horses, and cattle (Corbeil, 2016).

The phylogenic classification of *C. pseudotuberculosis*

Corynebacterium pseudotuberculosis is a member of the genus *Corynebacterium*, which belongs to the family *Corynebacteriaceae*, sub-order *Corynebacterinae* in the order *Actinomycetales*, sub-class *Actinobacteridae* and class *Actinobacteria* (Stackebrandt et al., 1997). This and other genera like the *Mycobacterium*, *Nocardia* and *Rhodococcus* are the entire *Actinomycetes* group (Dorella et al., 2006a; Jesse et al., 2008). The *Corynebacterinae* of the genera dominates the group of CMNR, which share several common features such as cellular organization, consisting of majorly the mycolic acids, peptidoglycans with a reasonable amount of guanine and cytosine genomically. The CMNR group encompasses other species of importance to both the Veterinary and Medical world, which include *M. tuberculosis*, *M. leprae* and *M. bovis*. *Corynebacterium pseudotuberculosis* is further sub-classified into two: the biovar *ovis*, which is generally incriminated in the sheep and goats resulting in an abscess in the viscera and superficial tissues (Biberstein et al., 1971). The second is the biovar *equi* affects equine species and is characterized by ulcerating lymphangitis around the distal extremities, abdominal and thoracic abscesses. The biomolecular technique was used to establish these two biovars (Sutherland et al., 1996; Costa et al., 1998; Connor et al., 2000; Connor et al., 2007).

Biochemical characteristic and virulence factors of *C. pseudotuberculosis*

C. pseudotuberculosis can ferment a variety of oligosaccharides such as galactose, mannose, glucose, and maltose (Henderson, 1979; Jesse et al., 2013a). It also produces and releases different enzymes (Pazatsen et al., 2012). Remarkably, these biochemical properties are indistinguishable among all *C. pseudotuberculosis* strains isolated from many animal species (Sutherland et al., 1996; Abdullah et al., 2013). The oxidation of nitrate to nitrite is a unique biochemical character (Foley et al., 2004). One of the major challenges for *C. pseudotuberculosis* is probably achieving the synthesis of this thick multilayer barrier (Burkovski, 2018). The *C. pseudotuberculosis* envelope is partly responsible for its innate resistance to antibiotics and plays a major role in both its virulence and persistence (Oliveira et al., 2017).

It has been described that *C. pseudotuberculosis* strains collected from some equine and bovines were consistently capable of reducing nitrate. On the other hand, the strains collected from ovine and caprine do not reduce nitrate (Latif et al., 2017). Thus, the nitrate-negative *C. pseudotuberculosis* strains, which are pathogenic in the goats and sheep, were apportioned to biotype I, while nitrate-positive strains, which infect horses and cattle, were apportioned to biotype II (Sutherland et al., 1996; Butler et al., 2005). Serotyping (Barak et al., 1992), molecular analysis (Muckle et al., 1992) and immunoblotting (Perkins et al., 2004) are useful methods to characterize strains of *C. pseudotuberculosis*. The molecular analyses are important in supporting the biochemical findings of two different biovars of *C. pseudotuberculosis* (Khuder et al., 2012).

Several studies on the virulence factors or determinants of most bacterial species had yielded a lot of information but there

is a paucity of data on the virulence determinants of *C. pseudotuberculosis* (Jesse et al., 2020). In recent times, few molecular studies had tried to explain the genetics of *C. pseudotuberculosis* about the identification of two virulent factors including Mycolic acids (MAs) and Phospholipase D (PLD) (Baird and Fontaine, 2007; Mahmood et al., 2015b; Odhah et al., 2019).

Structure of Mycolic acids

Corynebacterium has a complex cell wall structure which is considered as the main virulence determinant (Butler et al., 1986; Collins et al., 1988). This complex cell wall structure is called the mycolic acid or corynomycolic acids (Minnikin et al., 1983). They are characterized by a 2-branched 3-hydroxy fatty acid long-chain and represented the shortest lipid in the group extending twenty and thirty-six carbon atoms in their lengths (Alshamaony et al., 1977; Goodfellow et al., 1982; González et al., 2011). In contrast, Mycobacterial corynomycolic acids extend between sixty and ninety carbon atoms in length (Ionedá and Silva, 1979; Ionedá, 1989). These lipid structures in *C. pseudotuberculosis* are the main virulence determinants and they play a vital role in bacterial survival (Bastos et al., 2012). The lipid nature of this mycolic acid is responsible for its cytotoxic effects (Hard, 1975). It had been indicated that the mycolic acid after purification and inoculation in mice models elevated the clinical signs and resulted in localized congestion, oedema, and hemorrhagic necrosis (Gotoh et al., 1991; Nieto et al., 2009).

In addition, mycolic acids exhibit inflammatory regulatory features, a distinct response that was tolerogenic had been reported in experimental asthmatic conditions, but the chemical constituent is heterogenic (Korf et al., 2006). In vitro studies have shown white blood cell (WBC) degeneration in phagocytosis of these bacteria (Peyron et al., 2008). *Corynebacterium pseudotuberculosis* has the cellular capacity to withstand phagocytic activities of antigen-presenting cells. In pathogenic bacteria, MAs have long been known to induce inflammatory responses and granuloma formations and functions as an adjuvant for T cell responses (Kim et al., 2014; Odhah et al., 2017). A lot of research showed that the virulence factor of *C. pseudotuberculosis* is related to its cell wall lipid contents (Muckle and Gyles, 1983). Based on their studies, the presence of mycolic acid in the cell wall of these bacteria enables it to exert its pathogenic effects and survive as an intracellular bacterium (Dubnau et al., 2000) despite the well-established fact that lysosomes have the lysing ability on any invading organism. However, *M. tuberculosis* survives within the agranulocytes when it is engulfed (Bhatt et al., 2007).

Mycolic acid in *Corynebacterium pseudotuberculosis*

The members of this heterogeneous group share certain characteristics, such as a specific cell wall organization and high guanine and cytosine content on the genome. The cell walls of these genera are composed of a huge polymer complex of peptidoglycan and arabinogalactan and are rich in complex lipid components. The best characterized of these lipids is a long-chain 2-branched 3-hydroxy fatty acid, commonly known as mycolic acid. The mycolic acids of the *Corynebacterium*, also known as corynomycolic acids, are the shortest of the group, being between 20 and 36 carbon atoms in length, whereas those of *Mycobacterium* ranges between 60-90 (Brown and Olander, 1987; Dorella et al., 2006a). The corynomycolic acid is an important virulence factor of *C. pseudotuberculosis* as they contribute to its high tenacity and its ability to persist as a facultative intracellular parasite (Baird, 2003). Two different biovars of *C. pseudotuberculosis* can be dis-

tinguished based on the different nitrate reduction abilities of the two biotypes. Strains isolated from small ruminants tend not to reduce nitrate to nitrite whereas isolates from horses and cattle almost invariably possess a nitrate-reductase and are, therefore, able to reduce nitrate. Thus, Biberstein (1971) suggested the separation into the two biovars *C. pseudotuberculosis* biovar *equi* and *C. pseudotuberculosis* biovar *ovis*. There is a paucity of information in previous literature related to the extracted MAs from *C. pseudotuberculosis* towards the host cell responses such as in clinical signs, haemogram analyses, acute phase proteins (SAA and Hp) responses, antibody (IgM and IgG) titers, reproductive hormones (E2 and P4) concentrations, histopathology cellular changes in vital organs, reproductive organs, bone marrow and its associated lymph nodes.

Structure of Phospholipase D (PLD)

Phospholipase D is an enzyme that belongs to a group of lipases, which can hydrolyze the ester bonds in glycerophospholipids. The letter D indicated the specific ester bond that is targeted or cleaved. In general, phospholipase as an enzyme plays a vital role in the maintenance of cell membrane and signal transduction in eukaryotes (Abra and Quinn, 1975; Abdullah et al., 2015). Barksdale et al. (1981) were the first to describe PLD from *C. pseudotuberculosis*. Later, it was found in almost all isolates of *C. pseudotuberculosis* including biotypes I and II from all species of mammals that were infected with *C. pseudotuberculosis* (Songer et al., 1988). The fact that PLD is a risk factor had been established with some isolates of *C. pseudotuberculosis* that had their PLD genes encoding PLD attenuated as they could not cause abscess formation in the lymph nodes of the infected sheep (McNamara et al., 1994; Latif et al., 2015). PLD hydrolyzes phosphatidylcholine and sphingomyelin found on the cell membranes of mammals, thereby forming choline and ceramide phosphate (Carne, 1939; Onon, 1979). The exotoxins of the bacteria functions as a permeability factor that promotes the systemic dissemination of the pathogen (Jesse et al., 2013a).

In addition, PLD results in the dermal necrosis of cells in the endothelial lining easing *C. pseudotuberculosis* passage into small blood vessels, thereby gaining entrance into the lymphatic vessels (Egen et al., 1989; Songer, 1997). PLD is regarded as a cytotoxic exotoxin for WBC because it destroys the goat macrophages during laboratory trial (Tashjian and Campbell, 1983; Mahmood et al., 2015a). Vaccination of goats with the toxin also prevented the spread of the bacteria after an experimental challenge (Brown et al., 1986; Jesse et al., 2020). PLD mutants are virulence factors of *C. pseudotuberculosis* as they did stimulate an immune response. The bacterial spread by PLD action is explained by dermal necrosis which damaged the endothelial cells. It had also been reported that PLD during hydrolysis works synergistically with an oxidase of cholesterol and the C phospholipase both of which a product of *Rhodococcus equi* to produce in sheep a WBC β -hemolysis. A product ceramide phosphate is produced which gets hydrolyzed by the C phospholipase of *R. equi*, producing ceramide (Jolly, 1965; Brown and Olander, 1987). Dissociation of sphingomyelin into phosphate and choline is done by a sphingomyelin-specific phospholipase (Pepin et al., 1994). It participates in dermal necrosis (Catherine, 1994), macrophage destruction and interferes with neutrophil in goats (Yozwiak and Songer, 1993), synergistic hemolytic action in the presence of *Rhodococcus equi* extracellular factor (Fraser, 1961), inhibits hemolytic action induced by β -lysin produced by *Staphylococcus*. The two latter activities of PLD have been used in laboratories to identify *C. pseudotuberculosis* and thus, diagnose CLA (Zaki, 1976;

Jesse et al., 2016). Phospholipase D hydrolyses sphingomyelin in mammalian cell membranes increasing the vascular permeability especially the endothelial layer, leading to plasma proteins leakage from the blood into the surrounding tissue space and from there into the lymphatic system (Mahmood et al., 2015a). This contributes to the dissemination of *C. pseudotuberculosis* from the primary infection location to other parts of the animal's body (Guimarães et al., 2009; de Sá Guimarães et al., 2011).

Pathogenesis of *C. pseudotuberculosis*

The pathogenesis of CLA in sheep and goats is still poorly understood. To date most research conducted focused on two known virulence factors of *C. pseudotuberculosis*; the phospholipase D and the mycolic acids (Mahmood, et al., 2016; Jesse, et al., 2020). Phospholipases are glycerophospholipid-hydrolyzing enzymes that play an important role in signal transduction and the inflammatory response in eukaryotic cells. Many bacterial pathogens such as *Clostridium perfringens* and *Listeria monocytogenes* produce phospholipases as a part of their invasive strategy (Songer, 1997). *Corynebacterium pseudotuberculosis* produces phospholipase D (PLD), a phosphatidylcholine phosphatidohydrolase that functions as a sphingomyelinase and catalyze the repulsion of sphingomyelin to choline and ceramide (Pepin et al., 1994).

PLD is a virulence factor since *C. pseudotuberculosis* isolates in which the PLD gene was deleted from the genome were incapable of progressing from the site of infection and could not produce the classic lymph node abscesses (Pepin et al., 1994; Jeber et al., 2016). Dermonecrosis, lethality and synergistic lysis of erythrocytes in the presence of *Rhodococcus equi* exotoxin have been reported due to PLD (Pepin et al., 1994; Baird and Fontaine, 2007). Jolly (1965) showed that PLD functions as a permeability factor that increases vascular permeability and causes the leakage of plasma from the vessels through hydrolysis of sphingomyelin in endothelial membranes. This effect may assist in pathogenesis by allowing the bacteria to disseminate from the site of infection to the local lymph nodes (Brown and Olander, 1987; Pepin et al., 1994). This hypothesis was supported by Zaki (1976), who showed that antibodies against the toxin prevented the dissemination of the bacterium from the portal of entry. He concluded that the exotoxin, while not involved in the formation of abscesses, was crucial for the spread of the organism.

Furthermore, PLD may assist *C. pseudotuberculosis* in avoiding phagocytosis early in infection as it was shown to activate complement, and thus deplete it from the surrounding area, as well as impair the chemotaxis of neutrophils (Pepin et al., 1994; Osman et al., 2012b). Since it can replicate within and escape from macrophages, many researchers have suggested that the PLD might also be of significance in the escape from the phagosome and macrophage death (Pepin et al., 1994; McKean et al., 2007; Adza-Rina et al., 2013). The coat of waxy mycolic acids on the cell surface of *C. pseudotuberculosis* plays a major role in pathogenesis where it provides succour to the organism mechanically and biochemically. It has therefore been stipulated that MAs coat also aid the bacterium to extended survival periods which is synonymous with *Actinomycetales* and particularly *Mycobacterium* (Baird and Fontaine, 2007). *Corynebacterium pseudotuberculosis* has environmental resistance and shows a high tenacity in organic material (Augustine and Renshaw, 1986).

However, the cell surface lipid also protects the bacterium from hydrolytic enzymes within lysosomes which enables it to skip phagocytosis and survive within the host cell (Hard, 1972; Brown and Olander, 1987; Williamson, 2001). Pazatsen et al. (2012) first labelled *C. pseudotuberculosis* as a facultative intra-

cellular parasite. He infected mouse macrophages intraperitoneally with *C. pseudotuberculosis* and examined the ultrastructural changes using electron microscopy and observed that five minutes after infection, the bacteria were already engulfed by the macrophages and could be found in large vacuoles surrounded by an electron-dense layer, which was identified as the cell surface lipid of *C. pseudotuberculosis*. Furthermore, the study showed that macrophages that had ingested bacteria underwent rapid degeneration and cell death, releasing bacteria into the extracellular space where the bacteria remained intact. This study concluded that *C. pseudotuberculosis* can survive within the phagolysosomes and causes the death of its host cells due to the toxic effect of its surface lipid (Hard, 1972; Hard, 1975). Tashjian and Campbell (1983) confirmed this hypothesis in an electron microscopy study using caprine mammary macrophages infected with *C. pseudotuberculosis*. They observed that, despite the fusion of macrophage lysosomes with phagosomes containing bacteria, the caprine macrophages underwent progressive degeneration while the bacteria survived (Baird and Fontaine, 2007). The capacity to survive and replicate within phagocytic cells is crucial for the pathogenesis of *C. pseudotuberculosis* and it is thought that the organism migrates within macrophages from the entrance to the site of lesion development (Odhah et al., 2017). There is a consensus amongst researchers that *C. pseudotuberculosis* does not establish a persistent infection at the site of entry but disseminates rapidly, both as free bacteria and within macrophages, to the local lymph nodes (Ayers, 1977; Batey, 1986b; Pepin et al., 1991b). Pepin et al. (1991b) showed that a significant number of neutrophils were first mobilized to the site of infection and began appearing shortly afterwards in the local lymphatic drainage while the number of macrophages in the lesions rose dramatically on the third day after infection. As *C. pseudotuberculosis* can survive phagocytosis, the phagocytic cells serve as vehicles for transporting the bacterium into the local lymph nodes.

Additionally, the PLD exotoxin enhances the transfer of the bacterium via the lymphatics by inducing an inflammatory response and increasing the vascular permeability and lymph flow (Batey, 1986b; Pepin et al., 1991a; Mahmood et al., 2015a). Upon dissemination of the bacterium, micro-abscesses form in the local lymph node, where, after an initial phase of neutrophil dominance, macrophages infiltrate the lesions (Pepin et al., 1991b; Othman et al., 2014a). Even though the macrophages are the main effector cells fighting *C. pseudotuberculosis*, they are not able to eliminate the pathogen. A cycle of phagocytosis, intracellular multiplication and death of the host cells starts, leading to the transformation of the abscesses into epithelioid cell granulomas with a central area of necrosis surrounded by a rim of epithelioid cell and fibrous tissue (Abdullah et al., 2015; Latif et al., 2017). As seen in other intracellular pathogens, the formation of granulomas results in the encapsulation and entrapment of the lesions but also provides the basis for the chronic nature of the disease. Since the bacteria remain viable in the granulomas, the animals are not able to eliminate the infection and often remain life-long carriers. Occasionally, bacteria escape from the abscesses, possibly through emigration of infected phagocytic cells, and spread further through the blood or lymphatic system to other lymph nodes where they are retained in smaller blood vessels and cause the formation of new lesions (Batey, 1986a; Pepin et al., 1991b). Additionally, the cytotoxic properties of the surface lipid directly contribute to abscess formation. The subcutaneous injection of mycolic acids elicits an inflammatory response, resulting in localized swelling and central haemorrhagic necrosis, and induces sterile pyogenic lesions (Hard, 1975; Ayers, 1977; Baird and Fontaine, 2007; Mahmood et al., 2016). Catherine (1994) also

demonstrated a straight relationship existing between cell surface lipid quality of *C. pseudotuberculosis* and its power to induce a chronic abscess formation.

The resistance of host cell to C. pseudotuberculosis

The resistance of host cell to *C. pseudotuberculosis* infection is a complex process that involves the innate and acquired immune response (Zaki, 1976; Batey, 1986a; El-Enbaawy et al., 2005; Moura-Costa et al., 2008; Jesse et al., 2020). The significance of humoral defence had been elucidated by assessing various commercial and experimental vaccine (Eggleton et al., 1991; Stanford et al., 1998; Pinho et al., 2009), bacterial cell-wall fractions (Braga et al., 2007; Brogden et al., 1984), attenuated and killed bacteria (Simmons et al., 1997; Cameron et al., 1998), exotoxins and their subunits (Tashjian and Campbell, 1983) or genetically modified pathogens (Chaplin et al., 1999; De Rose et al., 2002). Significant advances in the field of vaccine production were an effective reduction in the formation of granulomas in experimentally challenged animals during trials. However, for adequate control of infection, macrophagic activation must be improved. A study using mice as a model indicated the potency of levamisole to evoke immunity against *C. pseudotuberculosis*. This finding suggests that cell-mediated immunity might play a significant role in *C. pseudotuberculosis* (Irwin and Knight, 1975). The participation of the complement system in *C. pseudotuberculosis* was studied in experiments demonstrating that type 3 complement receptor (CR3) played a key role during the infection. The findings indicated that treatment of mice with an anti-CR3 mAb resulted in the bacterial proliferation in both liver and spleen with a quite dramatic rise in the amount of mortality recorded 3 days post-infection. Histological investigations revealed that no mononuclear phagocytes gathered at the sites of bacterial proliferation and this confirms the importance of CR3 in the resistance *C. pseudotuberculosis* infection in mice (Lan et al., 1999).

Host Cell Responses to *C. pseudotuberculosis* and MAs

Acute phase protein responses toward C. pseudotuberculosis and its MAs

Acute-phase proteins (APPs) are a specialized group of blood proteins that would change in concentration in animals when subjected to external or internal challenges such as infection, inflammation, surgical trauma, or stress (Abdullah et al., 2013). Therefore, quantification of their concentration can provide diagnostic and prognostic information as biomarkers. The APPs are a composite of negative and positive protein that shows a falling or rising concentration during the response to challenge, respectively. The negative APPs include albumin and transferrin while haptoglobin (Hp), C-reactive protein, serum amyloid A (SAA), fibrinogen and alpha-1-acid glycoprotein are positive APPs. Haptoglobin and SAA are the major APPs in ruminants which can increase over 100-fold on stimulation (Skinner, 1991; Tothova et al., 2014). The acute phase proteins are synthesized by the liver and released following stimulation by cytokines. In ruminants, Hp and SAA are the major APPs that can increase up to 1,000-fold upon stimulation. The acute phase proteins are important in quantifying pathogenesis level in experimental studies and field infection (Abdullah et al., 2013). Odhah et al. (2018) and Faeza et al. (2019a) stated that the acute phase proteins can be used as diagnosis and prog-

nosis markers for various inflammatory diseases, organ malfunction, and infectious diseases in ruminant livestock. Ewes infected with intrauterine bacterial postpartum contamination showed a significant increase in Hp, SAA and fibrinogen concentrations (Jesse et al., 2016; Faeza et al., 2019a). Mahmood et al. (2016) stated that Hp concentration was higher than SAA concentration in goats infected with *C. pseudotuberculosis* and challenged with its exotoxin PLD. In another study, Odhah et al. (2018) showed that a significant increase of Hp concentrations was observed in does challenge with *C. pseudotuberculosis* and its mycolic acid.

Pro-inflammatory cytokine responses toward C. pseudotuberculosis and its MAs

Pro-Inflammatory cytokines are produced by activated macrophages as a proprotein, which is proteolytically processed to its active form by caspase 1. This cytokine is an important mediator of the inflammatory response and is involved in a variety of cellular activities, including cell proliferation, differentiation, and apoptosis (Ingman and Jones, 2008; Jesse et al., 2016). Pro-inflammatory interleukins are polypeptides produced mainly by the WBC, which regulate immunity and inflammatory conditions. They stimulate pathological and physiological function containing non-specific immunity and specific immunity with an overabundance of inflammatory responses (Sirotkin, 2011; Jesse et al., 2017a). Interleukin-1 β , interleukin-6, and tumour necrosis factor- α (TNF- α) are the pro-inflammatory cytokines that showed a significant increase of concentration in goats challenged with *C. pseudotuberculosis* and its MAs (Jesse et al. 2017b; Odhah et al., 2019). Concentrations of IL-1 β and IL-6 increased significantly in non-pregnant female goats challenged with *C. pseudotuberculosis* and MAs via the intradermal route (Jesse et al., 2017a). Another study by Faeza et al. (2019a) stated that bucks challenged with *C. pseudotuberculosis* and its MAs were able to stimulate the responses of IL-1 β and IL-6.

Cellular responses towards C. pseudotuberculosis and its MAs

Lymph node enlargement is the main indicator of intracellular infectious diseases (Jesse et al., 2020) which can be seen in *C. pseudotuberculosis* as it causes pain (Latif et al., 2017). James and Zumla (1999) postulated that granulomas do not define the disease condition but tend to give an idea of the disease because of the characteristics of the pathogenic species. The infections by *C. pseudotuberculosis* cause inflammation of superficial lymph nodes because of systemic infection (Pepin et al., 1991a; Timoney et al., 1988). The common primarily infected superficial lymph nodes in CLA are the retropharyngeal, mandibular, parotid, prescapular, popliteal, mammary, and inguinal lymph nodes (Baird, 2003; Batey et al., 1986b; Roberson et al., 2012). The morphological changes commonly observed are granuloma containing bunches of histocytes with eosinophilic cytoplasm (Carrington et al., 1969; Dorella et al., 2006b; Khuder et al., 2012). Adza Rina et al. (2013) stated that goats challenged with live *C. pseudotuberculosis* through the intranasal route developed abscesses in the mesenteric and supra-mammary lymph nodes compared to oral and intradermal routes.

Kim et al. (2014) reported that treatment with mycolic acid immunogen depleted the T cells in the spleen and suppressed broncho alveolar inflammations, pulmonary inflammations, and asthmatic responses. Khuder et al. (2012) reported histopathological changes such as degeneration, congestion, necrosis, infiltration of polymorphonuclear leukocytes, haemorrhage, oedema and thrombus in the ovaries, uterus, testes, and epididymis

in goats challenged with *C. pseudotuberculosis* and its exotoxin PLD. According to Al-Gaabary et al. (2010), the affected lymph nodes showed lesions of abscessation with a central zone of caseous necrosis surrounded by inflammatory cells consisting of a mixture of neutrophils and macrophages encapsulated by thick fibrous tissues and the liver section showed distorted necrosis in CLA infected sheep. Ibtisam (2008) explained that sheep infected naturally with *Corynebacterium pseudotuberculosis* exhibited abscess and pleural thickening with caseous material at the middle encapsulated by a zone of living and dead neutrophils in the lung whereas in the liver had evidence of congestion with a vacuolar degenerative change of hepatocytes. The heart showed congestion of blood vessels and haemorrhage with severe necrosis of myocardial fibres. The kidneys showed haemorrhage and degeneration of the cells lining the renal tubules. Mice challenged with *C. pseudotuberculosis* infection showed cellular changes in the ovary and uterus such as oedema, congestion, infiltration of neutrophil, macrophages, degeneration, and necrosis. The testis and epididymis exhibited cellular lesions such as oedema, congestion, degeneration, necrosis, and infiltration of polymorphonuclear leukocytes (Jesse et al., 2020). Latif et al. (2017) and Othman et al. (2014b) reported that does challenge with *C. pseudotuberculosis* showed cellular changes such as thrombus formation and necrosis in ovaries, vagina, uterus, and uterine horns. Faeza et al. (2019b) stated that bucks challenged with *C. pseudotuberculosis* and its MAs revealed significant and relatively new findings of cellular changes in testes and epididymis of both challenged groups.

Reproductive hormone responses towards C. pseudotuberculosis and its MAs.

Sex hormones play a vital role in male and female reproductive physiology by coordinating cell differentiation, proliferation, inflammation, apoptosis, homeostasis, and brain functioning (Edwards, 2005; Abdullah et al., 2015; Jesse et al., 2020). Sex hormones are produced by the gonads (testes and ovaries) and transported via the blood pool to act on the peripheral target tissue and the central nervous system (Wilson et al., 1998; Othman et al., 2014c). *Corynebacterium pseudotuberculosis* infection may cause inflammation and lead to the disruption of the hypophyseal-pituitary-gonadal axis (Jesse et al., 2017b). Several studies reported significant histopathological changes from mild to moderate severity in the pituitary gland, ovarian, cervix, uterus, fallopian tube, testes, spermatid and epididymis tissues in goats challenged with wild *C. pseudotuberculosis* and MAs (Khuder et al., 2012; Odhah et al., 2019; Faeza et al., 2019b). A recently published study by Jesse et al. (2020) reported significant cellular changes in the reproductive organs and their associated lymph nodes and changes in the reproductive hormones (estrogen and progesterone) in nonpregnant does experimentally infected with *C. pseudotuberculosis* and its MAs via intradermal routes. Khuder et al. (2012) described a significant decrease in plasma testosterone concentration in mice challenged with *C. pseudotuberculosis* and its exotoxin PLD. Faeza et al. (2019a) reported that male goats challenged with *C. pseudotuberculosis* showed lesions of the irregular shape of testes and shrinkage of seminiferous tubules with interstitial oedema and a significant decrease of testosterone hormone concentration were observed. In another study by Khuder et al. (2012) where male goats challenged with *C. pseudotuberculosis* and its exotoxin PLD showed a significant decrease in scrotal circumference, plasma testosterone concentration and semen quality parameters. A study by Umer et al. (2020) stated that male and female goats challenged with *C.*

pseudotuberculosis showed significant reproductive tissue damages in testes, epididymis, spermatic cord, penis, vulva, vagina, cervix, uterus, fallopian tubes, and ovaries. Abdullah et al. (2015), Khuder et al. (2012) and Jesse et al. (2020) described significant changes in reproductive hormones such as luteinizing hormone (LH), follicle-stimulating hormone (FSH), progesterone, estrogen and testosterone in animals infected with *C. pseudotuberculosis* and its immunogens exotoxin PLD and MAs.

CONCLUSION

In conclusion, this review has summarized the current information and research on mycolic acid from *C. pseudotuberculosis* and its interaction and responses of the host cell in terms of haematology, reproductive hormones, the acute-phase proteins, pro-inflammatory cytokines responses and cellular changes. *C. pseudotuberculosis* can colonise the reproductive tract in males and females and penetrate the physical barrier of the genital tract and cause severe tissue lesions in the vagina and iliac lymph node. Though there remains a dearth of information on the reproductive hormonal influence in female goats. Moreover, the causative agent of CLA reaches an effect on APP, and cytokines through lymphatic ways in infected goats. The review demonstrated that both microbiological and molecular techniques are useful tools in CLA and MAs. Therefore, future studies must be conducted on the *C. pseudotuberculosis* and its immunogens MAs effect on the various parts of the organs system and its physiological changes.

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CONFLICT OF INTEREST

The authors declared that they have no conflict of interest.

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